Biological Activities of Polymers

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FOREWORD

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PREFACE

Biologically active polymers serve as the basis of life and have existed since the beginning of the first forms of life in an aqueous medium. The major macromolecular entities of all living creatures, both animal and plant, are biologically active polymers: proteins, nucleic acids, and polysaccharides. The recognition that some synthetic polymers also exhibit biological activity is of a more recent vintage. The term "biologically active polymers" or "bioactive polymers" encompasses synthetic and natural polymers (native or modified) capable of eliciting a physiological response when applied to or introduced into a living system. The utilization of the biological activities of these natural and/or synthetic polymers is a major and growing area of scientific endeavor (molecular engineering, polymeric drugs, etc.) and is the basis for this book.

The chapters of this volume were chosen to illustrate the broad spectrum of research currently being undertaken related to the biological activities of macromolecules. The initial chapter introduces the basic framework relating the biological activity of polymers to the test conditions, etc., and emphasizes the need to design tests for each specific purpose (i.e., match the tests with the potential end use of the bioactive polymer). The remainder of the book is divided into three sections, each focusing on the current state of the art in particular areas of biological activity. The first section considers a broad range of nonmedical applications, including treatment of wood, control of uranyl ion toxicity, and control of schistosomiasis-carrying snails. Also discussed are polymers acting as herbicides, pesticides, mildew- and rot-resisting agents, mollusicides, algicides, antifungal agents, and antibacterial agents.

The final two sections focus on some medically related applications of bioactive polymers, including targeting and controlled release of drugs. Types of polymer supports and structure–activity relationships are also well represented. The section on drug-related activity also contains chapters on polymer activity as applied to the treatment of specific diseases, and on the emergence of thermal analysis techniques to describe biological activities of polymers more rapidly. The last section is based on applications of biologically active polymers in the broad area of cancer research. Antitumor and antiviral activities are featured along with discussions involving the advantages of polymeric anticancer drugs over the low molecular weight analogous drugs.

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Perspectives in Bioactive Polymers

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Bioactive polymers are macromolecules that can exhibit biological activity in use areas such as medication, weed control, insect control, etc. Natural bioactive polymers are essential to life and include the proteins, nucleic acids and poly- saccharides. Synthetic bioactive polymers are a more recent development but hundreds of possible examples have been reported with potential biological activity. In this brief, introductory review, the history, philosophy, mode of activity and the advan- tages of bioactive polymers are discussed emphasizing synthetic polymers.

Biologically active polymers are of fundamental importance in the life process itself. We need only note that molecules such as the proteins, nucleic acids, enzymes and many complex sugars are macromolecular in nature to realize how dependent life is upon polymer molecules. In addition, many of the polypeptide hormones are derived from a larger polypeptide or are synthesized by means of other macromolecules. Nearly all our foods and most of our bodies are composed of polymers and most of these are bioactive. We can note with certainty that macromolecules are essential to life as we know it. Beyond this, the interaction of macromolecules plays an important role in many areas of life. Examples of this could include such divergent effects as the pain-killing activity of the enkephalins or the toxic nature of snake venom which is a poly- peptide in most cases. In subsequent chapters of this book, other authors will consider a variety of polymeric systems with various types of actual or potential activity with humans, animals and/or plants. In this chapter, we briefly explore the overall realm of biologically active polymers.

History

A complete, detailed history of biologically active polymers is beyond the present scope, but we can note that such a complete history would be very difficult to unravel since the threads of this story are entwined not only in the area of polymer science but also in fields such as biochemistry, pharmacology, molecular biology and medicine. We will restrict this brief survey to synthetic bio-active polymers. Even here, however, it is difficult to draw forth an accurate picture since the early chemical literature abounds with examples of monomers and polymers that could possibly exhibit biological activity. Some of this literature has been reviewed, at least in part, elsewhere (1,2). Some early examples of potentially biologically active polymers include 2-vinylthiophene in 1941 (3), 2-vinylfuran in 1946 (4), 2-vinylpyrrole in 1954 (5), and N-vinyl-pyrrolidone in 1952 (6). Since its synthesis, N-vinylpyrrolidone has been widely used as a bood plasma extender (although this use has been discontinued in the USA) and the aqueous iodine: PVP complex has been marketed as an antiseptic for many years.

The work began by Overberger in the 1950's was aimed at de-veloping biologically active, synthetic polymers and involved the homopolymers and copolymers of the vinyl derivatives of pyrimidine and/or triazine compounds (7,8). Although the polymers had high molecular weights (up to 100,000), no biological activity was observed. In more recent work, Overberger and others have observed significant enzyme-like activity for some synthetic polymers and copolymers of vinyl imidazole derivaties (9,10). This pseudo-enzyme polymer area has been reviewed recently (11,12) and could prove to be of future importance in chemotherapy or in chemical manufacturing.

The major thrust in chemotherapeutic and other biologically active synthetic polymers began in the 1960's. While much of this work involved placing a known bioactive agent on or in a polymer chain, some biologically active polymers were developed which had no active, low molecular weight analogs. Hundreds of examples of such systems have been published by several dozen different investigating groups. We shall only examine a few specific examples and cite only some selected references. (Additional, related references can be found in these papers.)

Donaruma and coworkers have studied copolymers of various sulfa drugs with formaldehyde or dimethylolurea. These copolymers often exhibited bacteriostatic activity and in some cases this activity level was higher than with the corresponding, monomeric, sulfa drug. The polymers often showed lower toxicity levels and, in general, the dimethylolurea derivatives showed higher bacterio-static activity than did the formaldehyde derivatives. In earlier work, Donaruma and coworkers also prepared some antimalarial, polymeric sulfomes and some bioactive tropone derivatives (1,13).

Ringsdorf and coworkers have prepared many monomers, polymers and copolymers that are potentially chemotherapeutic (14,15,16). These included derivatives of cyclophosphamide which could be a potential anticancer compound (15). In addition to homopolymers, they prepared water soluble copolymers with N-vinylpyrrolidone, vinyl amine oxides and other monomers. Other work by this group has included polymers containing steroids, sulfa drugs, chlorambucil and adamantylamine units. In his review article, Ringsdorf discusses the use of some advanced techniques such as a homing device (or directing units), solubilizing units and various spacer units along with the pharmacologically active unit. The purpose of the spacer unit is to move the bioactive unit out from the polymer backbone and to make it more readily available for chemotherapeutic activity. In some cases, the use of spacer units has greatly enhanced the activity of the polymeric drug (14).

Carraher, Pittman, McCormick, Subramanian, Sheats and others have synthesized a wide variety of metal and phosphorus containing polymers which exhibit biological activities (17-22). Tin polyamines, polyesters and polyethers all exhibit broad spectrum antibacterial and antifungal activities. Tin containing polysaccharides derived from a number of sources also exhibit good antifungal and antibacterial activities. Extensive studies have shown that tin containing vinyl polymers exhibit sustained antifouling properties and such materials are currently utilized in the control of barnacles on ship bottoms and shore installations.

Takemoto and coworkers have prepared a wide variety of monomeric derivatives of a number of nucleic acid bases in order to prepare simple analogs of RNA and DNA (23,24). Some of these synthetic polymers do show base-base interactions that are similar to those observed in natural DNA and RNA materials. Pitha has observed anticancer and antiviral activity for some vinyl derivative polymers of uracil or adenine (25).

Gebelein and coworkers have prepared monomers and polymers based on the antineoplastic agents 5-fluorouracil and 6-mercaptopurine (26). These polymers would be potentially useful in treating various types of cancer or leukemia and some derivatives have shown biological activity. Previous work by this group has included the synthesis of polyisoprene derivatives containing sulfa drug, carbamate, urea, oxazolidone and oxazole units (27,28).

Probably the most striking example of a biologically active polymer that does not have an active, low molecular weight analog is the 1:2 divinyl ether: maleic anhydride cyclic, alternating copolymer which is called 'pyran polymer' or DIVEMA. This polymer was first prepared by Butler in 1960 (29) and has been shown to exhibit a remarkable range of biological activity including antitumoral, antiviral, antibacterial, and antifungal activities (30). Another

system which exhibits biological action as a polymer but not as a
monomer are the polymers of 2-vinylpyridine-1-oxide which show
antisilicosis activity (3).   Several groups are attempting to
develop synthetic polymers that would be capable of reversible
oxygen binding and these polymers woulc lead to the development of
artificial blood systmes (32,33).   In addition to these bioactive
polymer systems, much work has been done on the controlled release
of biologically active materials from polymeric matrices or by
cleavage from a polymeric backbone.   These systems have been
reviewed in some recent books (34,35).

Philosophy of Bioactive Polymers

       The elucidation of the biological behavior of compounds can be
approached from many points of view. The first of two extremes is
to treat the test species with various amounts, etc. of the test
material and to observe what occurs either on a gross sense (death)
or on a more selective level (cell, organ, tissue location and
effect).   The other extreme attempts to understand and investigate
on a molecular basis using model systems, in situ tests, etc. The
former allows a quick assay of general biological activity of the
test compound while the latter gives more detailed, but more
limited in scope, information about the biological activity of the
test compound under more controlled, but typically artificial
conditions and extrapolation to real systems is necessary.   Both
approaches have their advantages and both should be utilized in
appropriate situations.

       Desirable objectives in the design of chemotherapy drugs
include the synthesis of materials which have (a) good specificity,
(b) good activity, (c) prolonged activity, and (d) a wide con-
centration range of biological activity.   Thus variability and
specificity are key ingredients.

       Polymeric drugs allow both variability and specificity to be
built into their structure. The placement and chemical environment
of the biologically active agent can be varied within the polymer
along with introducing segments which can vary the macromolecule's
solubility and uptake in select organs and tissues. The molecular
shape and size can also be varied to achieve desirable flow and
isolation (cell walls acting as barriers, etc.).   A similar
polymeric design can be envisioned for polymeric herbicides, pesti-
cides, etc.

Complexity of Bioactive Polymers

       Natural systems are exceedingly complex and interrelated. As
scientists, we become entangled in this web from every side as we
attempt to increase our knowledge of thse varied systems which
include such diverse areas and entities as heart disease, cancer,

nutrition, vitamins, enzymes and nucleic acids woven together into an elaborate network. Slowly we observe that these effects are manifested on a molecular as well as a cellular level, and we observe further that entire systems of organs are frequently directly involved in these interactions. At times it appears nearly impossible to make any simple generalizations in this complex field of biologically active polymers.

Many biological studies are attempted utilizing poorly charac- terized materials, inadequately understood procedures, etc. In our laboratories we have discovered examples where biological analyses were carried out using supposedly pure small molecules purchased from appropriate chemical and biological supply companies which on further analysis were shown to be mixtures of several compounds. Analyses (such as amino acid analyses) are conducted by the bio- logist on a "black box" premise and we as chemists often view the world of biological assays also on a "black box" basis. Much of this "black box" treatment is due to an attempt to cope with a complex entity-life in all its varied forms and levels. Even so we must continue the process of lighting candles in these "black boxes" where possible.

This complexity has given birth to and continues to involve new disciplines such as anatomy, biological chemistry, biophysics, bioengineering, microbiology, immunology, virology, physiology, cardiology, pharmacology, toxicology, dermatology, endocrinology, epidemiology, gastroenterology, infectious diseases, neonatology, nephrology, neurology, otolaryngology, pathology, etc. and directly contributes to the interdisciplinary nature of evaluation of the biological activities of compounds.

The use and understanding of macromolecules as biologically active agents compounds this complexity. Polymer purity, molecular weight, molecular weight distribution, configuration, conforma- tion, nature of end groups all may affect the biological activity of the macromolecules. Time takes on a greater meaning for macro- molecules: aging, changes in the chemical and physical nature, flow dynamics, solubility, ability to penetrate cellular barriers, etc. all may change with time.

An added problem concerns the languages utilized by the various group disciplines in describing test procedures and report- ing biological results. Each of the biologically related disci- plines, government, industrial and clinical laboratories have evolved different reporting procedures typically significant for the mode and purpose of the employed tests. To this is often added additional languages according to the particular disease or tested species. While such diversity is natural, and often legitimate to reflect specific test procedures, organisms, etc., it is necessary that such variances do not impede progress, and interaction between different disciplines is essential in order to bridge this gap.

Interdisciplinary Nature

To realize fully the potential of the biological activity of any compound there must be a marriage between chemistry and biology. While the discipline of biochemistry bridges this gap with respect to many topics, it generally does an inadequate job with respect to drugs, and in particular with respect to polymeric biologically active agents. This void is partially being overcome with the emergency of several interdisciplinary doctorate programs. An even more severe problem exists in situations requiring clinical assays where only qualified MD's are permitted to direct (human) clinical programs.

Typically the chemical and biological areas are bridged through combining the efforts of two or more persons working as a team. In all cases it is best for the team members to attempt to learn the language, strategy of one another and for all to contribute to the planning and conducting of the continuing project on an equal basis.

Modes of Bioactivity

A bioactive polymer could interact directly or indirectly with another molecule or an active site to elicit a biological response. (An indirect mode of action would involve a chemical and/or physical interaction with some other chemical species which would then make the polymer biologically active.) The bioactive moiety could be: (a) the entire polymer chain, (b) segments within the polymer chain, (c) oligiomeric degradation products of the polymer chain, or (d) a low molecular weight (monomeric) fragment derived from the polymer. Modes (c) and (d) would be termed as activity via controlled release of the bioactive agent or species while modes (a) and (b) would be a more direct biological activity of the polymer itself. All these modes (a-d) could occur via direct or indirect interactions.

An additional mode utilizes the polymer as a matrix material for the embedding of the biologically active agent. These systems are called embedded or encapsulated systems. In these systems the biologically active agent is embedded within the polymer matrix and the agent is slowly released by a diffusion process or through the decomposition of the polymer within or about the target species. The polymer embedded agent can be either a reservoir system, in which the agent is coated or surrounded by the polymer, or it can be a monolithic system where the drug is distributed fairly uniformly throughout the polymer matrix. Much research has been done on these systems and embedded drugs show great promise in the treatment of numerous medical disorders such as glaucoma and heroin addiction,

in birth control, and in the control of bacteria, fungus and insects. The topic of embedded drugs will only be lightly dealt with within the following chapters.

Advantages

Many (potential) advantages exist in the use of biologically active macromolecules compared to the usually low molecular weight agents. These include: (a) sustained activity, (b) greater specificity of action (c) lower toxicity, (d) reduced undesirable side effects, and (e) more direct use of the polymer associated active agent. Part of these advantages stem from the ability to incorporate multiple features into macromolecular chains more readily than into a low molecular weight compound and from the ability to restrict the contact which the polymeric material has towards a test organism more readily than with a low molecular weight compound. Conventional drug delivery involving higher test species, mainly animals, involves ingestion of the drug orally into the digestive tract or injection into the circulatory system. The drug is then transported to the diseased site. Unfortunately, significant portions of the drug are typically excreted prior to this and, in addition, a large amount is transported to other tissues and organs which are not diseased resulting in unwanted side effects. Thus the therapeutic index (ratio of toxic dose to the therapeutic dose) of the drug is typically lowered and the duration of activity low. Polymeric drugs can also be delivered through injection and oral administration and through implantation. Polymeric drugs with limited solubility can be implanted or injected at or near the diseased site and would tend to diffuse only slowly. This reduces the undesired side effects and permits a lengthened duration of activity. Alternatively, a soluble polymeric drug can be taken orally or injected into the circulatory system. Two distinct possibilities exist. First, the polymeric drug could diffuse through the body but due to its size, its passage into certain tissues and organs would be restricted. The second possibility is to tailor the properties of the polymeric drug so it would act as a "guided missile" zeroing in on the desired target of activity (1,11,12,26).

The topic of selective targeting of drugs is currently the weakest link in the overall drug delivery system. Work is involved in the use of coupled drugs (insulin tends to make cell walls more permeable, thus it is being investigated in the delivery of anticancer drugs), embedding in synthetic and natural "homing devices" (lipisome embedded drugs) and in the design of drugs with both the targeting and drug activity built in. This work includes use of polymeric drugs in all three areas. Here we will restrict our comments to the latter approach.

It is reasonable to consider that a polymeric drug can be constructed to contain portions which (a) exhibit the desired pharmacological activity, and (b) direct the polymer to a specific site through solubility, steric, electronic and size tailoring. These special features would enable the polymeric drug to have specific activity, reduced side effects and presumably a higher therapeutic index compared to the analogous low molecular weight drug. Similar features could also be built into other bioactive polymers such as polymeric herbicides, etc.

Finally, the macromolecular nature of the polymeric drug might be an advantage. For instance, one mechanism for the anticancer activity of cis-dichlorodiamminoplatinum II,(C-DDP) is that it binds to the two DNA strands through ligand sites supplied by vacated chloride ions thus rendering the DNA dual strands sterile. For this to occur the "window" is quite small with rigorous steric, reactivity and electronic requirements. For instance, hydrolysis of one of the chloro groups prior to or subsequent to initial complexing of the platinum moiety to one DNA strand would prevent the platinum moiety from fulfilling its task. A polymeric analog would have many more opportunities to render the DNA strand couple inactive.

Conclusions

Bioactive polymers have existed since the creation of life on this earth and exhibit many specific interactions. Synthetic, bio-active polymers are of a much more recent origin but show great potential for highly specific bioactivity in the areas of chemo-therapeutic agents, herbicides, pesticides, etc. Future develop-ments in the field of biologically active polymers will probably result in greater control over medication and pest-control agents which will aid greatly in relieving the stress currently being placed on the environment or on the human body by the presently used agents. These advances will result primarily as a consequence of the macromolecular nature ofthese bioactive polymers.

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Biological Activities of Metal-Containing Polymers

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Preliminary biological assay results (bacterial and fungal inhibitory test results) for a number of metal and organometallic polymers are reported emphasizing both the variability and constancy of biological responses. For dextran modified with various organostannane halides, results are dependent on the nature of the organostannane but independent of the modification procedure. Also reported are biological assays of uranium containing compounds where the toxicity of the uranium is greatly reduced through complexation of the uranyl ion with a variety of Lewis bases.

The use of metal-containing polymers for applications in medicine has focused on utilization of siloxane polymers as biologically inert building materials. The purposes of this paper are to illustrate a) the variety of potentially available metal-containing condensation polymers which should exhibit activity toward biological organisms and b) the diversity of biological responses already found for selected metal-containing condensation polymers.

Experimental

The metal-containing polymers cited in this paper were synthesized utilizing low temperature condensation processes - namely the interfacial and solution processes. Where possible, syntheses were accomplished using systems that produced polymers within a minute or less reaction time. The combination of mild reaction conditions and short reaction time permits the use of a variety of potentially unstable reactants where thermal or solution induced rearrangements are possible (1).

Three general biological analysis techniques were utilized. These are the slant tube, disk and protein assays. Slant culture assays were conducted using saline, saline-sugar and saline-nutrient broth solutions. The paper disk assays were conducted with the test organism suspended in sterile water. Appropriate dilutions are made in Sabourand's dextrose agar and the solutions poured into Petri plates. The paper disks are soaked in solutions containing the tested material (solutions) or are placed in the Petri plates with specified weights of test material (solid) placed on them. Protein analyses of samples suspended by continuous vibration were carried out as follows. Cold perchloric acid was added to each culture tube after seven days. Pellets, consisting of the test organism, were collected by centrifugation and subsequently suspended in sodium hydroxide. After autoclaving, the solubilized protein was determined using bovine serum albumin, fraction V, at the standard.

Dimethylsulfoxide, DMSO, and DMSO-water are typically employed as the solvent since many of the polymers are soluble in DMSO and DMSO is not appreciably toxic to man or to most of the organisms tested.

Results are averages of at least duplicate examinations.

## Results and Discussion

Variety of Available Polymers

The use of metal-containing compounds as medicines is wide spread and ancient including merbromine (mercurochrome;mercury), silver sulfadizine (prophylactic in treatment for severe burns), 4-ureidophenylarsonic acid (therapy of ameblasis, tryparsamide) and antimony dimercaptosuccinate (schistosome). The appearance of metal-containing macromolecules in the human body is extensive including the metals of iron (transferrin and hemoglobin), vanadium (hemovanadin), molybdenum (xanthine oxidase), zinc (carbonic anhydrase) and copper (hepatocuprein).

Objectives in the design of chemotherapy drugs include that the drugs have a) good specificity (differentiation), b) good activity, c) long duration of activity and d) wide concentration range of biological activity. Thus variability and specificity are key objectives. Metal-containing condensation polymers, because of the variety of metal size, chemical environment and electronic structure, offer a wide variability of biological activities and the opportunity for good specificity.

Synthesis of most of the metal-containing condensation polymers can be considered in Lewis acid-base terms as described below.

$$X - A - X + Y - B - Y \rightarrow (A - B)$$
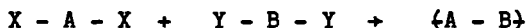
where A has been UO_2, R_3Sb, R_3As, R_3Bi, R_2Pb, R_2Sn, R_2Ge, R_2Si, R_2Ti, R_2Zr, R_2Hf, R_2Mn, $PtCl_2$, MoO_2 and Y-B-Y has been a hydrazine, hydrazide, urea, amine, oxime, amidoxime, diol, dithio, salt of a diacid, sugar, cellulose or if contained on a polymer the Lewis base has contained an alcohol, amine, sulfate, sulfonate, salt of an acid and amidoxime. The Lewis base has also been utilized as a metal carrier with polymers containing the metals Co, Fe, Rh and Ru thus far synthesized. The Lewis base can also contain mixed nucleophiles such as in xanthane dyes, p-aminobenzoic acid (1; Vitamin B_x; antirickettsial), Catalin (2; anticataract), kynurenic acid (3; used in vitamin B deficiency), salicylic acid (4; a nalgesic, antirhenmatic) and Methioprim (5; tumor antagonist).
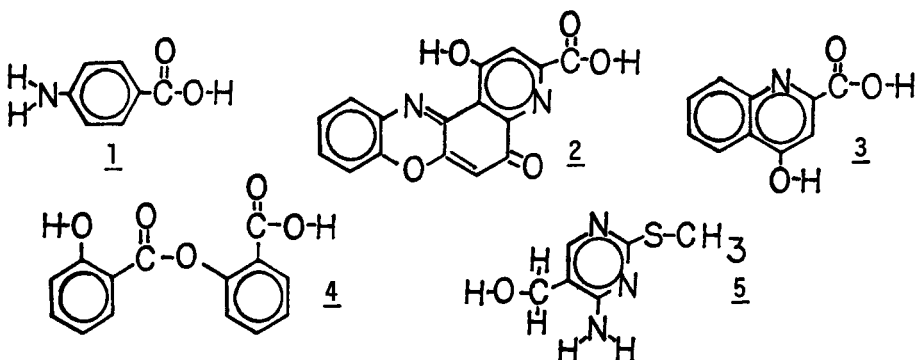

The polar-condensation linkage is advantageous as a site for breakdown and (potential) controlled release of moieties derived from either or both the comonomers.

Many of the employed metal-containing Lewis acid reactants undergo certain reactions for which analogous products are not formed through attempted condensation with acid chlorides utilizing mild reaction conditions. Thus Cp_2TiCl_2 reacts with salts of dicarborylic acid giving titanium-containing polyesters whereas the reaction with organic diacid chlorides does not yield the analogous condensation product under similar reaction conditions (2).

$$R_2TiCl_2 + {}^-O_2CRCO_2{}^- \rightarrow \{Ti-O-C-R-C-O\}$$

$$ClOC-R'-COCl + {}^-O_2CRCO_2{}^- \nrightarrow \{C-R'-C-O-C-R-C-O\}$$
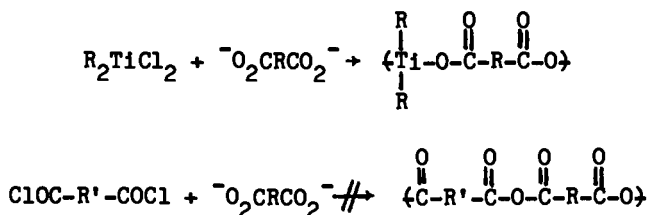
This is an important consideration if a drug as 1-4, which contains carboxyl groups, is to be delivered utilizing a condensation monomer or polymer.

The chemical environment of the metal and/or any desired drug can be varied as to its intended use. Thus the environment about the polar linkage can be unobstructed and the hydrophobic character minimul (6) or through the use of hydrophobic blocking groups the polar linkage can be sterically hindered and highly hydrophobic (7). Thus both intended preferential biological location of the drug (blood, lipid, etc. compatability) and rate of drug release (from fast, to the polymers itself acting as the delivered agent) can, in theory, be built into a polymer.
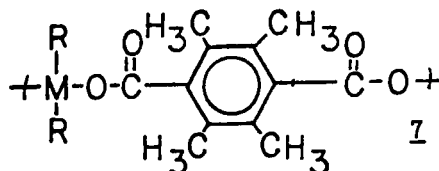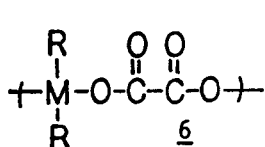
The "delivered-agent" can be either the metal or nonmetal moiety or both moieties and the intended purpose beneficial or detrimental or mixed in nature. Thus 8 was synthesized with the purpose of delivering tin, with the diol portion being coincidental. Polymer 9 was synthesized with the purpose of delivering
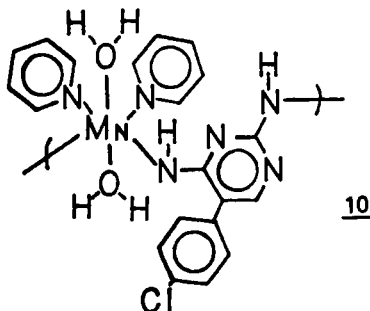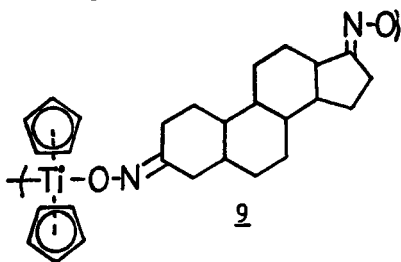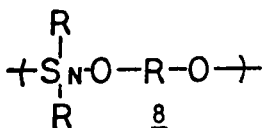
the hormone portion and 10 was synthesized to deliver both the known drug and the essential metal manganese. Regarding the metal, as a general rule Sn, Pb, As, Sb, U can be considered as inhibitory in nature, Mn, Fe as beneficial and Ti, Si etc. as neutral with regard to biological activity.

Thus there exists the potential for "tailormaking" metal-containing biologically active polymers.

Testing Variables

Biological responses are typically dependent on a number of test parameters including state of the tested material (solid or in solution), solvent, organism and test procedure. Since test results are dependent on these variables, it is advantageous to associate testing conditions with an intended application. Much of our research with polymers containing tin, antimony and arsenic is associated with control of mildew and rot for eventual application in topical medications, as thermal insulation and as additives in paints, textiles and paper products.

Figure 1 gives structures of polymers utilized and noted in Table 1. Table 1 contains results as a function of test organism, state and solvent for three tin, two arsenic and four antimony-containing polymers. In the solid state only compounds 1,3 and 8 exhibited good inhibition toward all three organisms. (The organisms are widespread in nature and cause "mildew" and "rot".) Compound 7 inhibited all three test organisms in DMF solution, whereas it was inactive in the solid, whereas compound 3 showed good inhibition in the dry state but only small to no inhibition in HMPA illustrating the variation of biological response with state of the test material. Compound 1 showed good inhibition in DMF but no inhibition when dissolved in HMPA; compound 8 showed good inhibition to the test organisms when dissolved in DMF but no inhibition to the test organisms when dissolved in HMPA illustrating the dependence of biological response on the solvent utilized to deliver the tested compound.

Other instances have been reported relating biological response with state and employed solvent. For instance there exists a direct correlation between the tendency toward inhibition by solid samples of (polyethyleneimine modified through condensation with organostannane halides) and results employing the same solvents dis solved in DMSO (3).

Tin Polydyes

Recently we reported the synthesis of titanium-containing polyetheresters derived from the condensation of Cp_2TiCl_2 with a wide variety of xanthene dyes (4). More recently we effected synthesis of analogous tin containing xanthene dyes through condensation of organostannane dihalides with xanthane salts (5). These products will be called tin polydyes. The xanthene dyes utilized to form the polydyes are typically utilized as cell-coloring agents by

Table 1. Results as a Function of Compound, Test Organisms and Test State

Compound Designation	Amount Tested (mg)	Dry State					DMF	
		Organisms					Organisms	
		As. Flavus	Penicillium sp.	As. Fumagatus	As. Flavus	Penicillium sp.	Penicillium sp.	As. Fumagatus
Figure 1.								
1	0.4	7	6	N	5	S	S	4
2	0.4	N	N	N				
3	1.0	17	25	2				
4	1.0	M	M	M				
5	0.4	S	N	N				
6	0.4	N	S	N				
7	0.4	N	N	N	S	4	4	S
8	0.2	6	I	I	5	5	4	3
9	0.2	N	N	N				
Control	N	N	S	S	S	S	S	N

(continued on next page)

Compound Designation Figure 1.	DMSO Organisms			TEP Organisms		HMPA Organisms	
	As. Flavus	Penicillium sp.	As. Fumagatus	As. Fumagatus	As. Flavus	Penicillium sp.	As. Fumagatus
1	S	S	S	10	N	N	N
2	N	N	N	I	N	*	N
3					S	N	N
4					S	N	N
5	N	N	N		N	*	N
6	N	N	N	10	N	*	N
7	N	N	N	10	N	N	N
8	N	N	N		N	N	N
9	N	N	N		N		
10	N	N	I	N	N	N	

Organisms used – Aspergillus flavus – 240 colony forming units/plate (CFU/plate)
Penicillium species – 500 CFU/plate
Aspergillus fumagatus – 760 CFU/plate

Symbols used – Numbers give the mm of inhibition: I=inhibition, but no clear zone; N=no inhibition; S=slight inhibition, but no clear zone; and * = enhanced growth
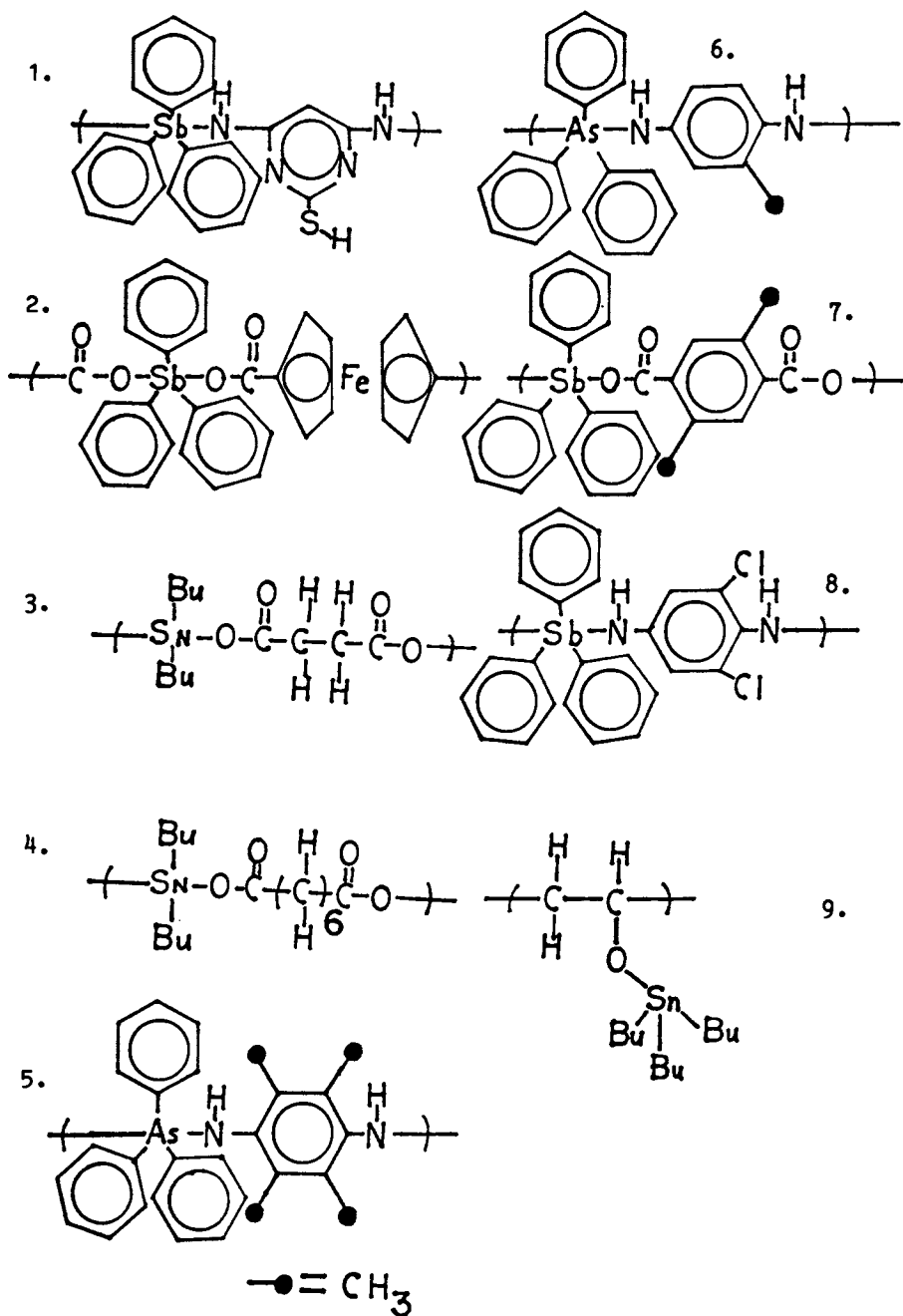
Figure 1.   Structures of compounds utilized in biological testing.

microbiologists, biochemists, etc., and as such polymers containing these dyes may be readily accepted by various organisms, effectively delivering the potentially toxic stannane. Further, mercurochrome, a xanthene dye, unlike the other utilized dyes, is toxic to many organisms because of the presence of mercury.

A number of tin polydyes derived from xanthene dyes were tested against the bacterial species <u>Escherichia</u> <u>coli</u> (C600) and <u>Pseudomonas</u> <u>aeruginosa</u> (7430). Figure 2 shows the structures of tested compounds. Results from testing the compounds as solids appear in Table 2. As expected, the greatest toxicity is from the polydye derived from mercurochrome, probably due to at least the toxicity of the dye portion itself. Most of the compounds exhibited some inhibitory nature consistent with the idea that toxic moieties can, in select situations, be "activated" through coupling with moieties already known to be "biologically acceptable."

Table 2. Inhibition as a Function of Organostannane Polyxanthene

Compound Designation From Figure 2	Organism Tested			
	Pseudomonas Aeruginosa		E. Coli	
	Weight Compound (mg)	Inhibition Zone (mm, diameter)	Weight Compound (mg)	Inhibition Zone (mm)
1	10	0	10	1
2	10	0.8	1	0.7
3	4	1.3	10	1.6
4	6	0	6	0
5	10	2	10	2.5

Polysaccharides Containing Tin

Recent efforts have involved condensation of organostannane halides with polysaccharides derived from various sources (6-8). Two polysaccharide sources are cotton and dextran. The inhibition tendencies of the modified polysaccharide are independent of their source but are dependent on the nature of the substituents on the tin such that inhibition generally decreases as the length of the alkyl chain increases, in agreement with the overall general tendencies of organostannanes (Tables 3 and 4; 6-8). The tests were performed using dry samples more closely approximating the conditions under which the polysaccharide compounds may be used.

To assess whether inhibition is due to an inability of the organisms to metabolize the modified polysaccharide or due to the actual toxicity of the products themselves, studies relating the growth in a liquid medium were undertaken. Trichoderma reesei and Chaetominum globosum readily metabolize polysaccharides including cellulose derived from cotton.
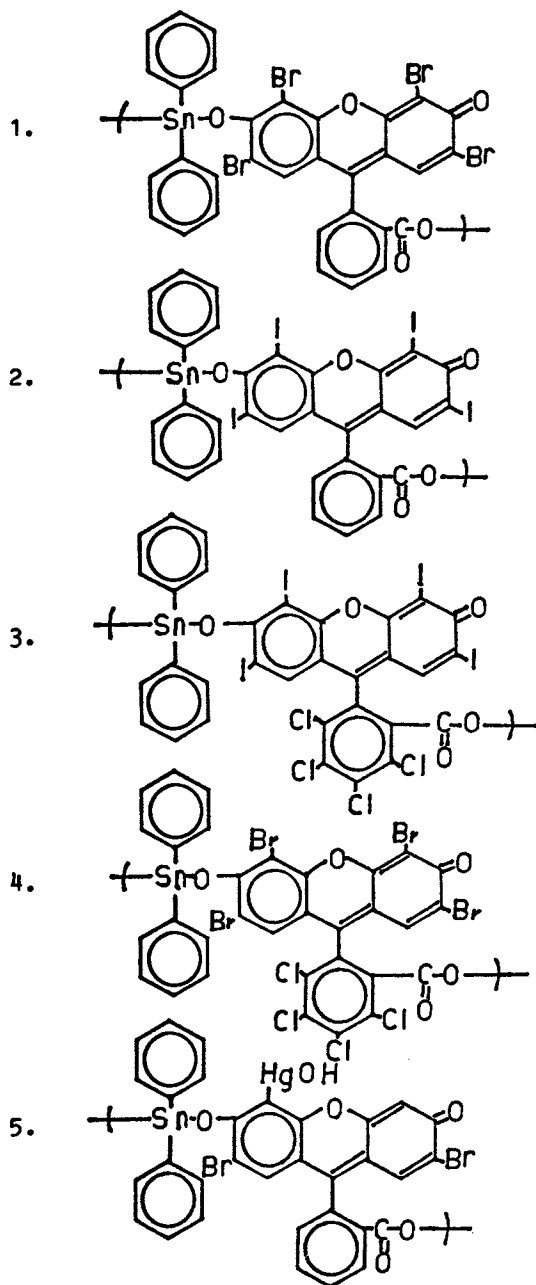
Figure 2.   *Structures of compounds utilized in biological testing.*

Table 3. Inhibition of Organostannane Modified Poly-
          saccharide Compounds.

Organostannane Dichloride	Tin (%)	Ps. Aeruginosa	E. Coli	A. Flavus	A. Niger	A. Fumagatus
Dimethyltin	8	4	4			
Dipropyltin	21			4	4	3
Dibutyltin	37			4	3	2
Dioctyltin	18	0	0	3	0	0
Dilauryltin	15	0	0			
Dibenzyltin	16	0	4			
Triphenyltin	20	0	0	4	3	3

a. 4 = 100%, 3 = 75%, 2 = 50%, 1 = 25%, 0 - 0% inhibition.

Table 4. Growth of Organisms on Organostannane-Containing
          Cellulose Compounds.

Organostannane Dichloride	Tin (%)	Trichoderma Reesei Growth (ug protein/ml)		Chaetomium Globosum Growth (ug protein/ml)	
		With Dextrose	None	With Dextrose	None
Dipropyltin	21	70	70	20	20
Dibutyltin	40	30	30	40	40
Diactyltin	18	280	40	120	50
Cellulose	–	420	250	300	240

Compared to growth of the fungi in solutions not containing
tin modified cellulose, all the compounds inhibit the two fungi
(Table 4).   Growth inhibition in the dextrose containing media
indicates that inhibition is the result of the toxicity of the
modified cellulose compounds on the fungi rather than merely an
inability of the fungi to degrade the test compounds.

The results are indicative of the applicability of such tin-
containing polysaccharides being employed for the retardation of
fungi related rot and mildew. Questions such as duration and mecha-
nism of fungal inhibition by these materials have yet to be
answered.

Uranyl-Containing Compounds

In select cases the purpose for encapsulating the metal or
organometallic in a polymer structure is to diminish biological

toxicity. The uranyl ion, in aqueous solution, is reported to be phototoxic to all tested aquatic fungi and bacteria (9,10). The LD_{50} for human beings is probably below 0.1 mg/kg body weight with uranyl ion exhibiting typical heavy metal toxicity (9-11).

Along with developing procedures for the removal, isolation and concentration of the uranyl ion is the concern for developing less toxic forms of the uranyl ion (2,12). Thus preliminary biological examination was undertaken on several representative bacteria. The test samples included uranyl modified PANa, uranyl nitrate hexahydrate, several uranium polyesters and two uranyl polyoximes.

The uranyl nitrate showed a greater zone of inhibition than that of any of the uranyl polyesters and polyoximes and the uranyl modified PANa (Table 5). In fact, all but one of the uranyl polyesters show no inhibition toward the tested species. Thus complexing the uranyl ion with bidentate salts of carboxylic acids decreases the toxicity of the uranyl ion (at least for the tested species). These tests were conducted on solid samples. Further tests are underway.

Table 5. Results of Preliminary Biological Tests as a Function of Uranyl Compound.

| | | Bacterial Species | |
| | | E.Coli(C600) | Pseudomonas Aeruginosa(7430) |
UO_2^{2+} Reacted with salt of	Sample Amount for Test(g)	Diameter of Inhibition(mm)	Diameter of Inhibition(mm)
PAA	.010	1.3	0.6
UNHH (itself)	.010	2.2	2.1
Terephthalic Acid	.010	NI	NI
Succinic Acid	.010	NI	NI
1,1'-Dicarboxylic Ferrocene Dicarboxylic Acid	.010	NI	NI
2,5-Dichlorotetraphthalic Acid	.010	NI	NI
Adipic Acid	.010	NI	NI
Azelaic Acid	.010	1.1	1.0
Azelaic Acid	.0050	1.0	1.7
p-Benzoquinone Dioxime	.010	NI	NI
Terephthalic Dioxime	.010	0.7	0.7
Terphthalic Dioxime	.0050	0.5	0.5

NI = No Inhibition
Diameter of spot maintained to approximately 0.5mm

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Biocidal Activity of Organotin Polymers in Wood

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The synthesis and preliminary biological assay of organostannane-containing wood is described for the purpose of developing controlled release agents for wood preservation. The organostannane is bonded to the wood using in situ polymerizations. A number of the stannane-containing wood samples show complete resistance to rot and mildew after two years exposure. Further mechanical properties and dimensional stability of the modified wood is also improved.

Trialkyltin compounds have emerged in recent years as broad spectrum toxicants having high toxicity towards marine fouling organisms (1) and wood destroying organisms (2-4) while possessing a tolerable toxicity towards mammals. It has also been shown that trialkyltin compounds pose a minimal hazard to the environment since they eventually degrade to harmless inorganic tin oxides by the action of UV light, microbes, etc. (5,6).

Many controlled release formulations based on tri-n-butyltin groups chemically linked to polymers have been devised in our laboratories and they have performed successfully as antifouling coatings during many months of exposure to marine fouling environment at Miami Beach, Florida (7,8). The effects of chemical structure upon mechanical properties of such polymers (9), as also the copolymerization behavior of organotin vinyl monomers with functional group containing vinyl monomers have been studied (10). Such formulations are effective against Pseudomonas nigrifaciens (marine bacterium), Sarcina lutea (soil bacterium), and Giomeralla cingulata (soil fungus); and the effect of the nature of crosslinks upon leaching rate of the toxin has been determined in our earlier work (11). An investigation of the reactivity of organotin carboxylate polymers has established that the bioactive species released from these polymers in an ocean environment is the trialkyltin chloride (12).

Thus, the aim of our study reported here was to explore ways of adapting the methods of synthesis and application of controlled release organotin antifouling polymers developed in these laboratories to problems of wood preservation (13). The basic approach is to bond the toxic moiety, tributyltin, to a monomer small enough to penetrate wood, and then polymerize this monomer within the wood, i.e., "in situ polymerization" (14). This results in a wood polymer matrix where the accessible void is filled with polymer and, depending upon the monomer, chemically attached to the cellulose structure of wood. Tributyltin compounds, as simple (unbound) additives, are easily leached from wood, and therefore are of limited life expectancy. In the system under study, the treated wood should have a low leach rate because the diffusion of the toxic moiety, bound chemically to the polymer chain, is controlled by the polymer matrix fixed within the wood. This will increase the service life of the wood while ensuring minimal impact on the surrounding environment. Polymer impregnation also has the advantage of increasing the dimensional stability of the wood in water. By decreasing the amount of water which can be absorbed by the wood, the alternating swelling and shrinking of wood in the tidal zone is minimized, and leaching of the preservative is decreased.

Experimental

All treatments to be used in initial field tests were done with standard test coupons provided by the Forest Products Laboratory. These wood specimens are cut from Southern Yellow Pine trees harvested from U.S. Forest Service timber stands and cut into precision 1.5 x 6 x 0.25 inch (3.81 x 15.24 x 0.635 cm) samples with the grain running lengthwise and a ring count of 6 to 10 per inch. By design, each sample therefore has a standard volume of 36.87 cm^3.

All of the samples were treated by one of three different standard procedures for vacuum/pressure impregnation: Full Cell Method, Empty Cell Method (Lowry) or modified Full Cell Method (15). The treatments were applied in a modified pressurized paint tank set up to simulate the procedures used commercially in typical pressure treating cylinders. It was connected to a laboratory vacuum pump in order to pull a vacuum of up to 29 in. Hg (98 kPa), and a compressed air cylinder in order to provide up to 120 psi (827 kPa) air pressure. After treatment, the surface of specimens was wiped free of excess solutions, the samples were air dried overnight, then placed in the circulating air oven and dried to constant weight to remove solvent and moisture before calculating retentions.

The modified Full Cell treatment was employed for impregnating the wood with monomeric materials. This treatment used the treating tank and the vacuum system as a surge tank system. A vacuum-tight glass reaction kettle was used as the treating

chamber. It was equipped with a drain in the bottom, which
doubled as an inlet for treating solution, and a ground glass
jointed addition funnel which has a stopcock connection to the
vacuum tubing that was connected to the surge tank. This was
used to make sure that the treating solution was drawn up to a
level in the addition funnel where there was excess solution in
order to keep the wood totally submerged as it soaked up monomer.
After the solution was drained, each sample was carefully wiped
to remove excess monomer on the surface and cured in an explosion-
proof circulating air oven at the desired temperature. An illu-
strative set of results of in situ polymerization of organotin
monomers in Southern Yellow Pine coupons is given in Table 1.

Six test specimens from each treatment were chosen to be
placed in the field tests at the Key West Naval Station, Key West,
Florida. The test coupons identified with polyethylene tags were
bolted to a fiber glass rack. Each rack contained 2 untreated
samples which were used as controls monitoring the degree of at-
tack over each 6-month period between inspections. The racks are
suspended on nylon line at a depth of ~1 foot (30 cm) below nor-
mal low tide. Consequently, the samples are always submerged but
are very close to the intertidal zone where Limnoria attack is
heaviest, as evidenced by the condition of pilings in that area.

At each semiannual inspection the racks were pulled up, the
surface fouling on the coupons was scraped off to expose the sur-
face of the wood specimen and each small panel was inspected care-
fully and rated by degree of attack on a scale of 0 to 10. A
rating of 10 signifies no attack; 9, trace attack; 7, moderate
attack; 4, heaving attack; and 0, no structural integrity left,
or sample destroyed. Each rack had one control replaced even if
neither had a low rating. The inspections take place in June and
December, with the heaviest attack taking place between June and
December corresponding to the higher water temperatures during
this period. This test method corresponds to ASTM D2481,
Accelerated Evaluation of Wood Preservatives for Marine Services
by Means of Small-Size Specimens.

Results and Discussion

Standard marine creosote, 40% naphthalene creosote, and tri-
butyltin oxide were used to treat test samples by both Full Cell
and Empty Cell Methods. These treatments will serve as perfor-
mance standards in statistical evaluations of the new preserva-
tives.

Two organometallic polymers previously developed, tributyl-
tin methacrylate/methyl methacrylate copolymer (TBTM/MeM) and the
tributyltin ester of methyl vinyl ether/maleic anhydride (TBT-
MVEt-MAn), were selected for use as wood impregnants. Full Cell
and Empty Cell Methods were used to treat wood with TBTM/MeM in
"Rule 66" mineral spirits and in combination with P13 creosote.
Although the polymers seem to be miscible with the creosote, this
combination results in only surface penetration of wood when

impregnated. The tributyltin ester of methyl vinyl ether/maleic anhydride in cyclohexanone was used to treat wood samples by both the Full and Empty Cell Methods. Since it was found to be immiscible with creosote, this combination treatment was not used.

TABLE 1.   IMPREGNATION OF SOUTHERN YELLOW PINE WOOD WITH
           TRIBUTYLTIN METHACRYLATE-GLYCIDYL METHACRYLATE
           MONOMERS AND "IN SITU" COPOLYMERIZATION OF THE
           SAME

Sample No.	Polymer in Wood, %[a]	Sn in Wood, %[b]
Weight % of Monomer in Reaction Solution = 10%		
7-1a	9.79	1.956
7-1b	10.02	1.997
7-1c	9.95	1.985
7-1d	10.83	2.143
Avg of Samples	10	2.0
Weight % of Monomers in Reaction Solution = 25%		
7-2a	25.64	4.475
7-2b	25.01	4.387
7-2c	26.19	4.551
7-2d	26.22	4.556
Avg of Samples	26	4.5
Weight % of Monomers in Reaction Solution = 40%		
7-3a	37.57	5.989
7-3b	38.79	6.129
7-3c	38.20	6.062
7-3d	37.41	5.970
Avg of Samples	38	6.0

a: $\dfrac{\text{[wt of treated dry wood]} - \text{[wt of untreated dry wood]}}{\text{[wt of untreated dry wood]}} \times 100.$

b: $\dfrac{\text{[wt fraction of Sn in polymer] [wt of polymer in treated dry wood]} \times 100}{\text{[wt of treated dry wood]}}$

The length of the polymer chain of these two organometallic polymers makes them rather viscous for use as wood impregnants, so it was decided to try impregnating with monomeric materials and polymerizing "in situ". Methyl methacrylate, tributyltin

methacrylate, 50/50 weight percent tributyltin methacrylate/
methyl methacrylate with and without a crosslinking agent, and
tributyltin methacrylate diluted with mineral spirits were used
as impregnating monomers. Benzoyl peroxide was used as the ini-
tiator, and all of the monomers were cured in the wood. This
method resulted in high retentions of polymer.

The in situ copolymerization of glycidyl methacrylate/
tributyltin methacrylate in Southern Yellow Pine (Table 1) is
comparable to that in Grand Fir wood specimens reported earlier
(13,14). With Grand Fir, it was found that the ultimate flexural
strength, flexural modulus of elasticity, impact strength, and
tensile strength of the treated wood sample were all equivalent
to or better than untreated samples. Exhaustive solvent extrac-
tion indicated a high percentage of the polymer grafted to the
wood. The dimensional stability of the wood was improved by
treatment, as indicated by the antishrink efficiency after a 48-
hour soak in distilled water. Scanning electron microscopy indi-
cated that, in a glycidyl methacrylate treated sample, the poly-
mer's macrodistribution is uniform throughout the sample. Scan-
ning of a tributyltin methacrylate/glycidyl methacrylate sample
and a tributyltin methacrylate/maleic anhydride sample showed
that in both of these specimens the polymer was uniformly distri-
buted at the surface, but decreased toward the center. This may
be a function of the solvent since acetone is a good wood swel-
ling solvent and benzene is not. Electron probe analysis for tin
clearly indicated substantial amounts of tin atoms in the cell
walls and in the filled cells.

Table 2 presents some exposure data from the Key West Test
Site. It is noteworthy that these samples have an unusually
large surface-to-volume ratio when compared to standard pilings
and therefore represent an extremely severe test condition. This
is important in minimizing the length of time a field trial must
be monitored to indicate failure or success of a treatment.

After 2 years of exposure of the first ten sets of samples,
none of the organotin treatments has shown any attack. Untreated
controls have failed completely, demonstrating sample inoculum
present for attack. After 18 months of exposure some attack was
noted in the marine creosote treated controls. After 24 months,
the 23-lb/ft^3 marine creosote had an average rating of 8.3, the
15.85-lb/ft^3 marine creosote had an average rating of 8.4, and the
special marine creosote (40% naphthalene) was rated at an average
of 9.7. In 36 months, they have been attacked and damaged
severely whereas the organotin polymer impregnated systems do not
suffer any deterioration.

The results of the field test confirm the evidence obtained
in laboratory tests of wood decay (13,14). In these tests it was
found that Grand Fir wood specimens treated with organotin poly-
mers were resistant to attack by brown-rot, white-rot and soft-
rot fungae, and a marine bacterium (Coniophoro puteana,Polyporous
versicolor, Quaetomium globosum and Pseudomonas nigrifaciens
respectively).

TABLE 2.   FIELD TEST DATA

Sample No.	Treatment	Rating		
		18 months	24 months	36 months
A1	P-13 Marine Grade	10	10	10
A2	Creosote,(23 lb/ft^3)	10	4	-
A3		10	10	4
A4		10	9	7
A5		9	7	-
A6		10	10	4
A13	TBTM/MEM Copolymer,	10	10	10
A14	(13.91 lb/ft^3)	10	10	10
A15		10	10	10
A16		10	10	10
A17		10	10	10
A18		10	10	10
A31	Tributyl Tin Ester of	10	10	10
A32	Methyl Vinyl Ether/	10	10	10
A33	Maleic Anhydride,	10	10	10
A34	(3.28 lb/ft^3)	10	10	10
A35		10	10	10
A36		10	10	10
A37	P-13 Marine Grade	9	7	-
A38	Creosote, (15.85 lb/ft^3)	10	10	-
A39		10	10	10
A40		10	10	10
A41		10	10	-
A42		10	10	0
A43	Special Marine Creosote,	10	9	7
A44	40% Naphthalene,	10	9	4
A45	(27.17 lb/ft^3)	9	10	4
A46		10	10	10
A47		10	10	10
A48		10	10	9
		12 months	18 months	30 months
B44	TBTM/MeM "In situ"	10	10	10
B45	Polymerization	10	10	10
B46	(22.8 lb/ft^3)	10	10	10
B77		10	10	10
B78		10	10	10
B79		10	10	10
B57	TBTM-GMA "In Situ"	10	10	10
B58	Polymerization, 10%	10	10	10
B59	Polymer in Wood	10	10	10
B60		10	10	10

Conclusions

Protection of wood against microbiological decay can be accomplished by impregnation with trailkyltin polymers. Mechanical properties and dimensional stability may also be improved simultaneously.

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Polymer-Bound Fungicides for Paints

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In order to prepare coatings which contained chemically
bonded fungicides, four vinyl-substituted fungicides were pre-
pared and both copolymerization and terpolymerization studies
were carried out with these monomers. The fungicides were
pentachlorophenol, 3,4',5-tribromosalicylanilide, 8-hydroxyquino-
line, and 2-benzyl-4-chlorophenol. Three monomers of each were
prepared. The acrylates and the chain-extended 2-fungicidalethyl
acrylates were prepared of each fungicide in addition to the
vinyl ether of each fungicide. The acrylates and chain-extended
(2-fungicidalethyl) acrylates were copolymerized with acrylic
monomers and stable terpolymer latexes were prepared from (1)
pentachlorophenyl acrylate (2) 3,4',5-tribromosalicylanilide
acrylate and (3) 2-pentachlorophenylethyl acrylate with the
comonomers methyl methacrylate and n-butyl acrylate. Terpolymer
latices of vinyl acetate, 2-ethylhexyl acrylate, and 3,4',5-
tribromosalicylanilide acrylate were made. Copolymers and latexes
containing pentachlorophenyl and 3,4',5-tribromosalicylanilide
acrylate were tested against <u>Auerobasidium</u> <u>pullulans</u> in agar dish
accelerated growth studies and were found to prevent growth.
Several polymers did not exhibit a zone of inhibition around the
treated sample as did the free fungicides which could migrate
into the agar medium. However, when the polymers were first
irradiated by UV light, a zone of inhibition was found indicating
photohydrolysis occurred to free the fungicide. This represents
a novel mechanism for the controlled release of fungicide.

The long term protection of paint films against mildew
defacement is, as yet, an unachieved goal. The problem has
received increased attention from the Paint Research Institute
since 1971.[1] A variety of fungicides

have been used as additives to paints, but several problems are
encountered when fungicides are simply dispersed into the paint.
Since films are thin and have high surface areas, water leaching,
evaporation, or migration of the fungicide occur. When the
fungicide concentration drops below a critical level, mildew may
begin to grow.[2] If a fungicide were chemically bound to the
polymer backbone of the film, a number of advantages might be
realized.[3] Leaching of the fungicide should be retarded.
Vaporization might be eliminated. These differences could lead
to longer mildew resistant lifetimes of the film. Solubility
problems would be eliminated. Biocides which have high water
solubilities or high vapor pressures now could be candidates for
use in paints whereas they could not have been considered
previously. Finally, the toxicity of the paint (to humans) might
be reduced. However, polymer-bound biocides may have reduced
effectiveness and this point needs to be experimentally tested.

In designing mildewcidal coating polymers, the fungicides
can be bound to the polymer in a permanent fashion or in a way
that the fungicide may be slowly released. Chemical anchoring
of biocides in paints was the subject of a detailed PRI-supported
literature search,[4] but little research has appeared on this
topic outside of the area of undersea marine coatings where con-
siderable interest existed.[5] Pittman et.al. first chose to
prepare polymers where the biocidal functions were attached by
ester and amide groups, because it was envisioned that these
could be cleaved in the environment created by developing
colonies of Aureobasidium pullulans, the major paint defacing
organism,[6] and both preinvading and coinvading microorganisms
which accompany these colonies. The broad spectrum biocide,
pentachlorophenol, was converted to its acrylate and copolymeriz-
ed with vinyl acetate or ethyl acrylate to provide polymers where
ester group cleavage would release pentachlorophenol.[7] These
polymers proved to resist the growth of Pseudomonas sp.,
Aspergillus sp., Alternaria sp. and A. pullulans in petri dish
tests.[7] Unlike polymers to which pentachlorophenol was simply
added, films of the biocidal copolymers did not exhibit zones of
growth inhibition adjacent to the films.[3,7]

To further test the merit of the polymer-anchoring concept,
we have synthesized a series of fungicidal monomers where the
active agent is attached by both hydrolyzable (ester) groups and
nonhydrolyzable (ether) groups. Co- and terpolymerization of
these monomers into model paint binderlike polymers was complet-
ed. In addition, some stable paint-like latices were prepared.
This synthetic program was undertaken to provide reasonably
"paint-like" biocidal polymers for both laboratory and outdoor
fence testing of the mildew resistance of coating films. Herein,
we describe a portion of the synthetic work and some test results.

Fungicidal Monomer Synthesis

Acrylate derivatives were prepared of the known fungicides(1)
pentachlorophenol (3) 3,4¦5-tribromosalicylanilide (5) 8-
hydroxyquinoline and (7) 2-benzyl-4-chlorophenol using acryloyl
chloride and amine bases. These are outlined in Scheme I.
These acrylates, (1), (3), (5) and (7) were then used to prepare
acrylic paint binder polymers. Since each of these fungicidal
acrylates are phenolic esters, they are readily hydrolyzed. The
various phenoxides function as good leaving groups. Resonance
stabilization of charge into the phenyl ring stabilizes the
phenoxy anion making it a better leaving group in ester hydrol-
ysis. Another group of chain-extended acrylates was prepared by
introducing an ethyl group between the phenol and ester group.
The resulting chain-extended acrylates will not hydrolyze as
rapidly because they are alkyl esters. The alkoxide anion
leaving group is less stable than a phenoxide and therefore
is a poorer leaving group. Thus, in addition to the acrylates,
the chain-extended acrylates, (2), (4), (6) and (8), of
each of the four fungicides was prepared. The acrylate
and chain-extended acrylate syntheses are outlined in
Scheme I.
Fatty The acrylates (1), (3), (5), and (7) were each prepared
using acryloyl chloride under normal Shotten-Baumann conditions.
The syntheses of chain-extended acrylates (2), (6), and (8) were
accomplished by reacting the respective fungicidal phenoxide
anion with 2-chloroethanol to generate the corresponding 2-
phenoxyethanol. Treatment of these 2-substituted ethanol
derivatives with acryloyl chloride and triethyl amine gave the
chain-extended acrylates. This route was unsuccessful, however,
starting with 3,4¦5-tribromosalicylanilide because the displac-
ment of chloride from 2-chloroethanol failed. The oxygen of
3,4¦5-tribromosalicylanilide is severly hindered by the ortho
bromo and amide functions which sterically retard the S_N-2
displacement. The same reaction on 2-iodoethanol proceeded in
poor yield (25%) giving the 2-substituted ethanol which was then
esterified with acryloyl chloride to give (4). The direct
displacement of tosylate from 2-tosylethyl acrylate resulted in
only an 8% yield of chain-extended acrylate, (4). The structure
of monomers (1)-(8) were confirmed by infrared and nuclear
magnetic resonance spectroscopy.
Vinyl ethers of the fungicidal monomers were also prepared
since an ether linkage connecting a fungicide to a polymer would
be highly resistant to hydrolysis. Thus, vinyl ether polymers
were a synthetic goal in order to determine if the polymers,
themselves, could be biocidal since release of the biocide could
not occur. Three general synthetic routes were examined. First,
the 2-bromoethyl ethers were made of the fungicidal phenols from
1,2-dibromoethane followed by base-catalyzed dehydrohalogenation.
Secondly, based on literature precedent,[8] the phenol was treated
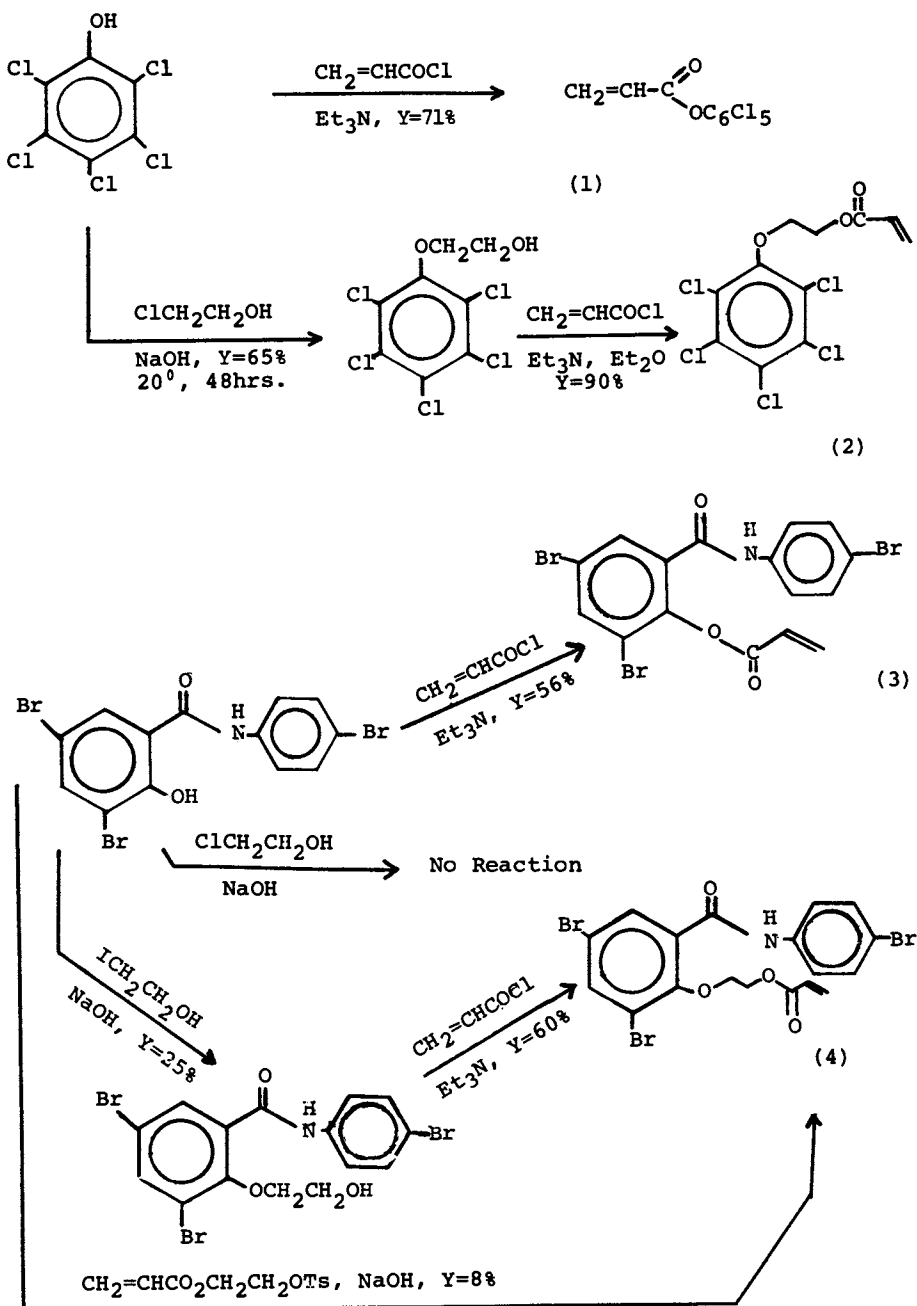
Scheme I.   (Part 1)
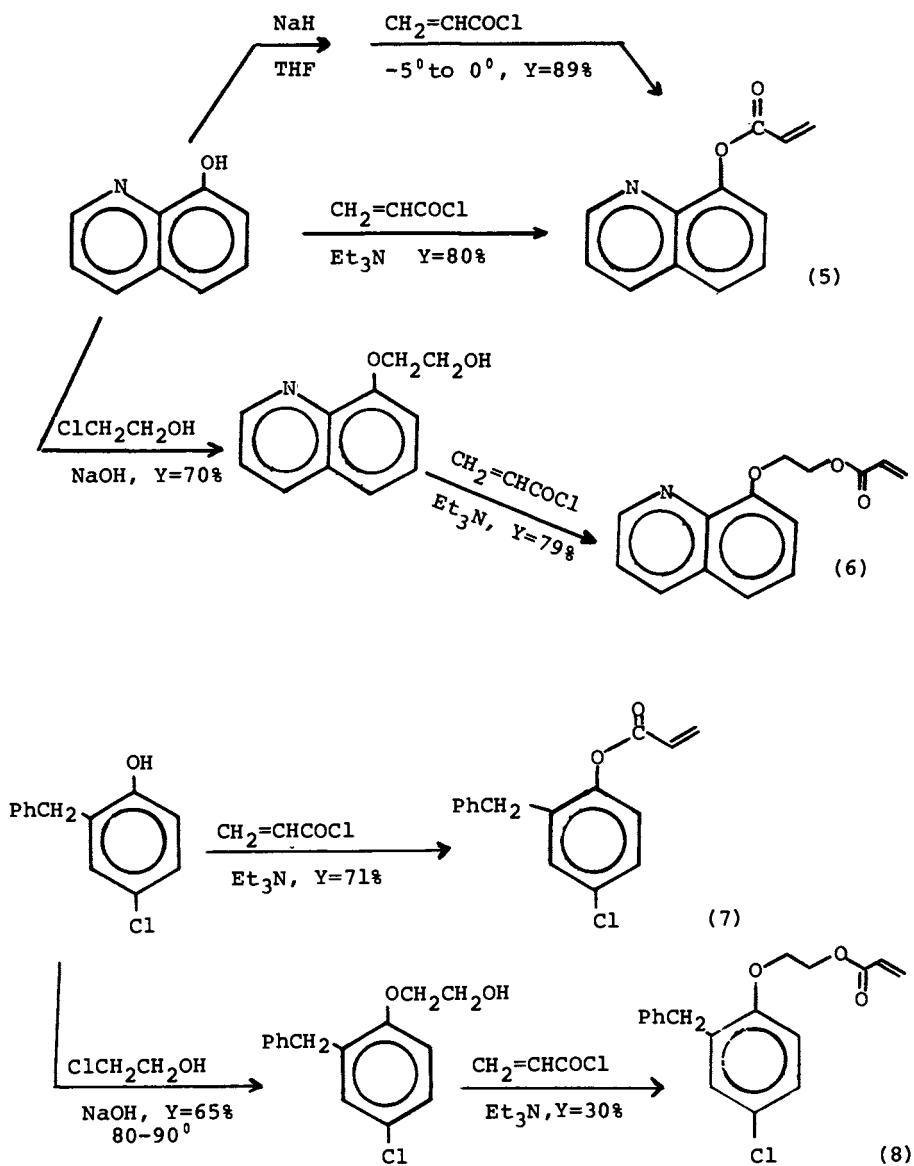
Scheme I. (Part 2)

with vinyl acetate in the presence of mercuric acetate to give
the vinyl ether in an addition-elimination sequence. A third
route was based on 1,2-vinylbis(triphenylphosphonium)dibromide.
Reaction of the phenol with this salt in chloroform and triethyl-
amine, followed by treatment with aqueous NaOH, gave the vinyl
ethers of pentachlorophenol and 2-benzyl-4-chlorophenol. This
method and the vinyl acetate route were not successful in giving
the vinyl ether of the highly hindered phenol, tribromosalicyl-
anilide. 8-hydroxyquinoline was converted to its vinyl ether by
treatment with acetylene in base at elevated temperatures and
pressures. The routes employed are summarized in Scheme II.

Polymerizations

        The fungicidal acrylates, (1), (3), (5), and (7), were
solution homopolymerized using radical initiation (AIBN) and
copolymerized with hexyl methacrylate in emulsion reactions
(sodium lauryl sulfate, $K_2S_2O_8$, 60^0). Similar studies were
performed on the chain-extended acrylates, (2), (4), (6), and
(8). Emulsion copolymers of (7) and (8) and homopolymers of
(7) were of low molecular weight, persumably due to chain-
transfer caused by the benzylic hydrogens. The fungicidal
monomers give homopolymers with high T_g values. Coatings with
fungicidal monomers therefore must have one comonomer which would
lower T_g. Thus, n-butyl acrylate, n-hexyl acrylate, or 2-
ethylhexyl acrylate were employed in copolymers and terpolymers
giving latices which serve as binders in test coating systems.
        Pentachlorophenyl acrylate, (1), was terpolymerized with
methyl methacrylate (MMA) and n-butyl acrylate (nBA) (Scheme III)
to give a latex containing 53% solids and a pH of 4.7 which was
adjusted to 6.8 by adding aqueous NaOH. The latex was stable up
to pH = 10. A small aliquot was coagulated and the resulting
polymer purified. Its intrinsic viscosity was $3.1^{d\ell}$/g and
analysis indicated 2 mole percent (1), 58% MMA and 40% nBA.
Similar terpolymer latices were prepared from acrylates (2) and
(3) (Scheme III). Another terpolymer latex made from (3), vinyl
acetate, and 2-ethylhexyl acrylate contained 54% solids. These
latices and their compositions are summarized in the Table 1 and
a sample experimental procedure is given in the experimental
section.
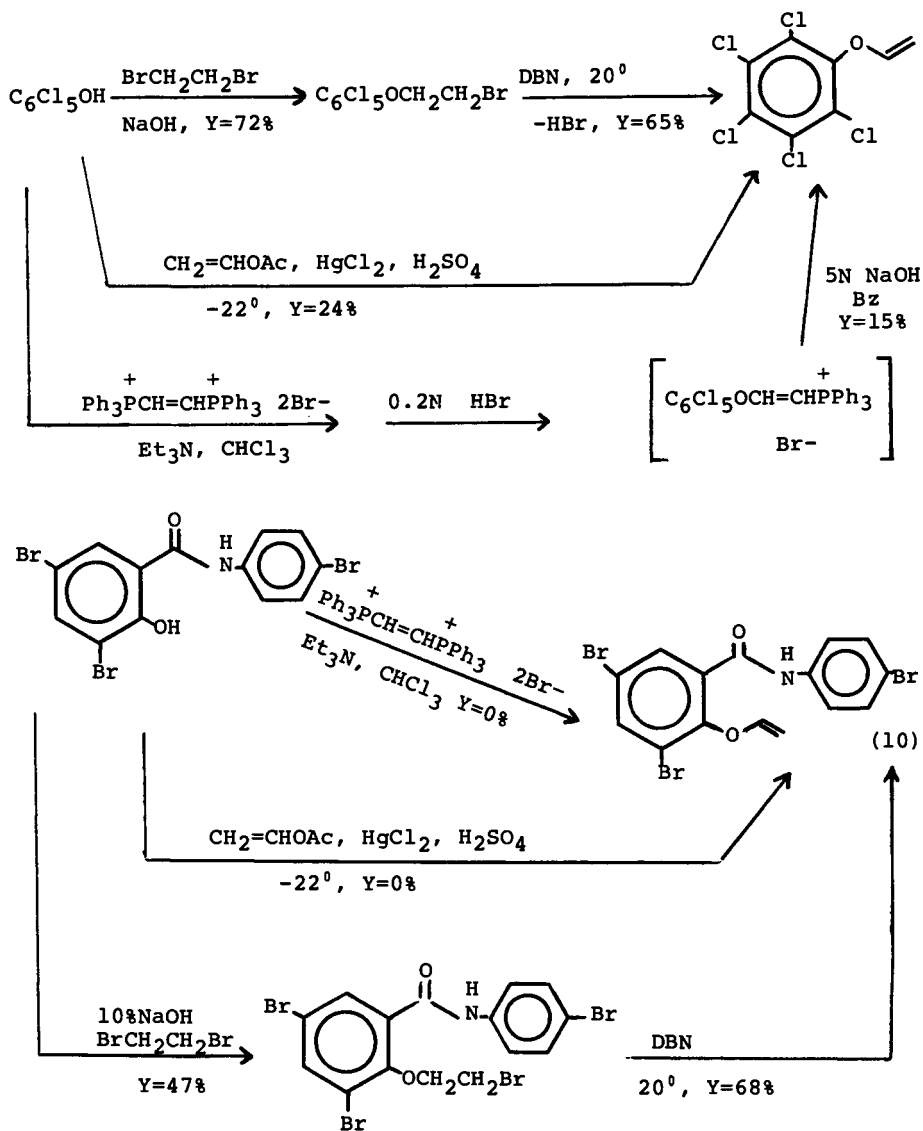
Scheme II. (Part 1)
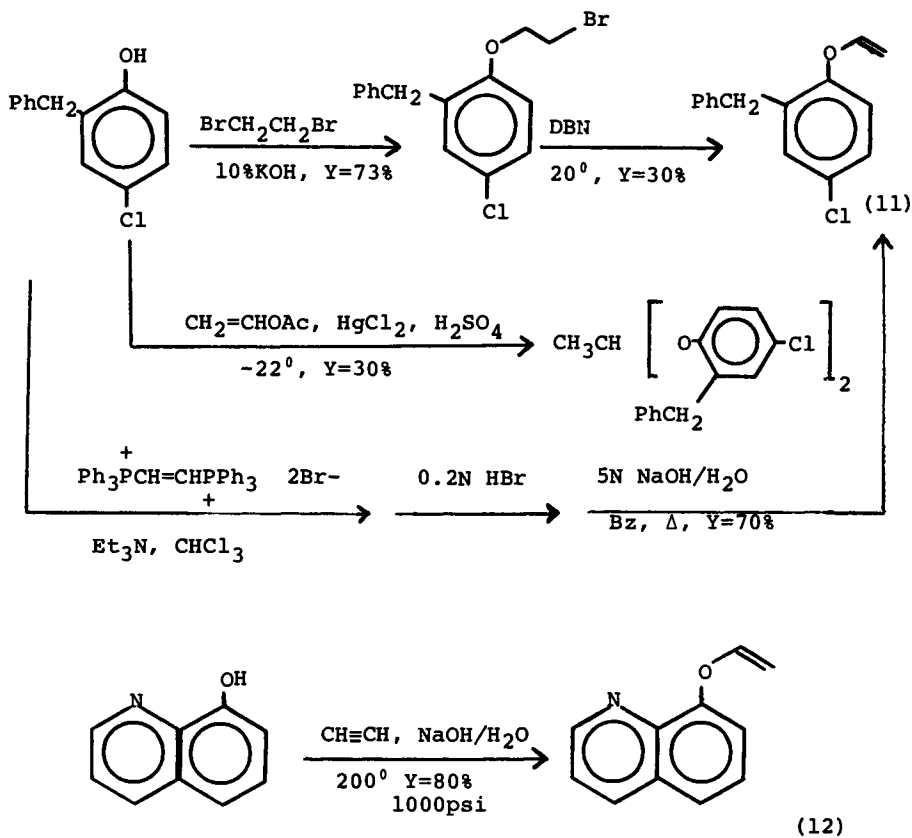
Scheme II.   (Part 2)

Table 1. Composition of Stable Latices of Model Paint Binders
Containing Chemically Bound Fungicides.[a]

Number	Monomers (mole % in Polymer) [b]	Solids (% wt. of latex)	$\|\eta\|$ [c] dl/g
1	1(2), MMA(58), nBA(40)	49.7	3.1
2	1(5), MMA(64), nBA(32)	40.0	2.3
3	2(11.4), MMA(6), nBA(82.6)	54.2	3.5
4	3(2), MMA(47), nBA(51)	48.3	3.1
5	3(5.8), MMA(43), nBA(51.2)	40.0	2.8
6	3(1.4), VA(87.4), EHA(11.2)	54.0	1.3

[a]These latices were prepared as described in the text. MMA =
methyl methacrylate, nBA = n-butyl acrylate, VA = vinyl acetate,
EHA = 2-ethylhexyl acrylate.
[b]The numbers in brackets refer to the mole % of the monomers in
the binder polymer isolated from the lattices and purified.
These were determined by elemental analysis.
[c]The intrinsic viscosities are those of the binder polymer after
coagulation, isolation, and purification by two reprecipitation
steps.

A variety of petri dish and coated wood accelerated growth
tests have been performed with both copolymers and latexes. The
results may be generalized as follows. Copolymers of the fung-
icidal monomers (1) and (3) prevent the growth of mildew without
producing zones of inhibition which reach out into the nutrient
medium. This was also true of coatings prepared from latex
samples 1, 4, and 5 (Table I). This may be contrasted to the
free fungicides, pentachlorophenol and 3,4,5-tribromosalicyl-
anilide, which do give zones of inhibition when simply blended
(or dissolved) into the coating polymer.

The concentration of fungicide in the coating polymer often
must be higher when the fungicide is polymer-bound to prevent
mildew growth. This is illustrated for pentachlorophenyl
acrylate, (1), and its homopolymer ($\bar{M}n$ = 10,000) in Table II.
When tested against Escherchia Coli, Staphylococcus Aspergillus
niger, Penicillium citrinum, and Saccharomyces cerevisiae in
agar dish tests, to determine the minimum inhibitory concentra-
tion (MIC), the homopolymer was less active than monomer (1)
when the organisms were E. coli and Saccharomyces cerevisiae.
The polymer was of comparable activity for the other organisms in
the test. When tested after UV irradiation, coatings prepared
from latex numbers 1, and 4 did exhibit zones of inhibition in
accelearated growth tests against Auerobasidium pullulans. This
illustrated that photohydrolysis was occurring which caused
saponification of the ester group, thereby freeing the fungicidal
phenol.
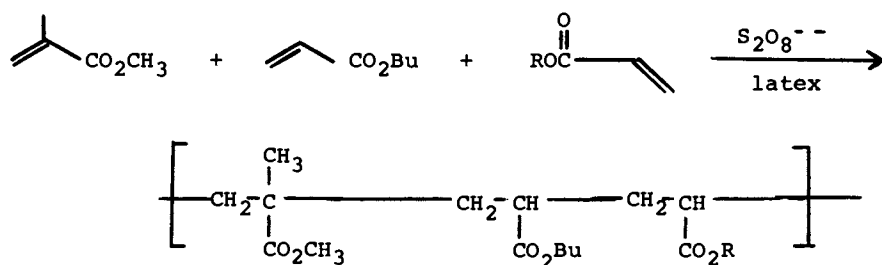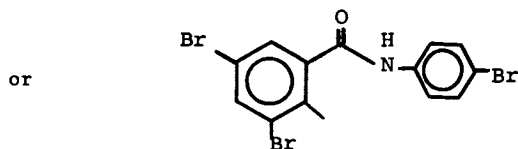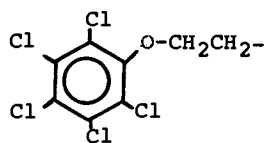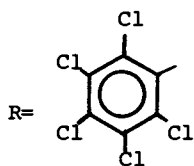
Scheme III.

Several copolymers (1)/HMA, (3)/HMA for example) were formulated into test alkyl paint formulations which were painted onto cedar panels for outside evaluation. Latices and copolymers are currently undergoing accelerated growth testing (Zabel, Syracuse University) and they are being formulated into a standard paint (Singer, Troy Chemical Co.) for testing on outdoor fences (Matthews, duPont). Attempts are underway to incorporate vinyl ether into suitable binder polymers. Results of the biological testing will be published later.

Homopolymers and both ethylacrylate and vinylacetate copolymers of 8-hydroxyquinolyl acrylate, (5), gave large zones of inhibition in agar dish tests due to rapid self-hydrolysis from the polymers. Thus, the polymer systems are not suitable for slow release. Tests are underway on polymers of the extended chain analog, (6).

Table II. Minimum Inhibitory Concentrations of Pentachlorophenyl Acrylate, (1), and its Homopolymer Toward Micro-ogranisms Using an Agar Dilution Method. (- means no growth; + means growth).

Compound		MIC					
		1000	500	250	100	50	10
(1)	E. Coli	-	-	-	-	+	+
	Staphylococcus aureus	-	-	-	-	+	+
	Asperigillus niger	-	-	-	+	+	+
	Pencilillium citrinum	-	-	-	+	+	+
	Saccharomyces cerevisiae	-	-	-	-	+	+
homo polymer of (1) Mn=10,000	E. Coli	+	+	+	+	+	+
	Staphylococcus aureus	-	-	-	-	+	+
	aspergillus niger	-	-	-	+	+	+
	Penicillium citrinum	-	-	-	+	+	+
	Saccharomyces cerevisiae	-	-	-	+	+	+

Experimental Section

The synthesis of monomers (1) and (5) have previously appeared. Thus, they will not be further described.

2-Pentachlorophenoxyethanol. 26.6 g (0.1 mole) of sub-limed pentachlorophenol was dissolved in water (150 ml) containing NaOH (6 g , 0.15 mole) by warming. After cooling this clear solution to room temp., 2-chloroethanol (12 g , 0.15 mole) 2-chloroethanol in 25 mls of water was added slowly (over 3 hrs) with mechanical stirring. After 10 hrs of stirring, the solution was heated to 80°C for 2 hrs. A copious white precipitate appeared which after cooling was filtered, dissolved in methylene chloride, and washed three times with NaOH (0.5 N) followed by water. The solvent was removed under reduced pressure giving a

white crystalline product which was crystallized from methanol/
water or diethyl ether/pet. ether 20/80 (60%) yield 18 g mp
91°-92° C, = 30.95, H = 1.61 found: 31.03, H = 1.63. IR and nmr
spectra agreed with the structure.

2-Pentachlorophenoxethyl Acrylate, (2). 2-Pentachloro-
phenoxyethanol (16.5 g 0.0532 mole) was dissolved in dry methy-
lene chloride (100 mls) and triethyl amine (6.5 g) was added.
The solution was cooled in ice bath to (0-5°) to which freshly
distilled acrylolyl chloride (5.6 g 0.061 mole) in 20 ml dry
methylene chloride was added under dry conditions. The solution
was allowed to thaw overnight and worked up by addition of satu-
rated solution . of sodium carbonate and then by extraction in
methylene chloride. On removal of solvent an oily residue
remained. This was crystallized from methanol/diethylether
(90/10) as light yellow fluffy crystals. M.P. 69-70°C yield 17
gms (90%).

Acryloyloxy-3-4,5-Tribromosalicylanilide, (3). 3,4,5-
Tribromosalicyclicylanilide (30.0 g, 66.7 mmol) and triethylamine
(7.1 g, 70.3 mmol) were dissolved in benzene (100 ml) and cooled
to 5°C. Acryloyl chloride (6.3 g, 69.6 mmol) was added dropwise
with stirring and triethylamine hydrochloride precipitated. The
mixture was warmed to 20°C for 3 hrs., filtered from the tri-
ethylamine hydrochloride, washed twice with water, and dried over
anhydrous $MgSO_4$. Removal of the benzene in vacuo gave a white
solid which recrystallized from a 4:1 dioxane/water mixture to
give 18.9 g (53.4%) of _3_ mp 149.5 - 151°C. Analysis: Calcd.;
C 38.10; H 1.98; N 2.78; Found: C 38.21; H 1.98; N 2.78; IR
$(KBr; cm^{-1})$; 3300 (NH), 1740 (CO, ester), 1660, 1630 (CO, amide),
1530, 1390, 860. nmr ($CDCl_3$; ppm): 7.4 - 8.2 (m, aryl and N_H, 7
H), 6.1 (m, 3H, vinyl). The multiplet at 6.1 - 6.7 exhibits an
ABC pattern.

2-(2-Hydroxyethoxy)-3,4,5-Tribromosalicylanilide. Method A:
A solution of 3,4,5-tribromosalicylanilide (4.0 g, 8.8 mmol) in
10% aqueous NaOH 96 ml) was heated to refluxing and 2-chloro-
ethanol (12. g, 15.0 mmol) was added dropwise. After 11 hrs.
the mixture was cooled and extracted with ether and the solution
dried with anhydrous $MgSO_4$. Removal of the ether in vacuo gave
0.9 g of a brown solid which proved to be 3,4,5-tribromosalicyl-
anilide. The aqueous layer was acidified and extracted with
ether. Removal of the ether in vacuo gave back unreacted
3,4,5-tribromosalicylanilide quantitatively.
Method B: 3,4,5-Tribromosalicylanilide (36.7 g, 81.6 mmol)
was dissolved in a solution of ethanol (50 ml), dimethylform-
amide (200 ml) and water (50 ml) containing KOH (5.9 g, 105.2
mmol). The solution was added to 2-iodethanol (15.1 g, 87.8
mmol) and the reaction was refluxed for 3 days, filtered and the
filtrate and precipitate were extracted with ether. The combined

extracts were dried with anhydrous MgSO4 and cooled to obtain
white crystals of the title compound. Yield: 8.8 g (21.5%).
mp 161.8 - 163.0°C. Analysis: Calcd.: C 36.5; H 2.40; N 2.80;
Found: C 36.52; H 2.27; N 2.86. IR (KBr; cm^{-1}); 3500 (OH),
3300 (NH), 2900 (aliph. CH), 1650 and 1580 (CO), 1530, 1240
(COC), 1080 (COC). nmr (CDCl3-DMSO-d$_6$ ppm): 7.8 (4H, amide
ring), 7.5 (q, 2H, acid ring), 10.3 (broad S, 1H, N$\underline{H}$), 4.1 (m,
2H, OC$\underline{H}_2$), 3.8 (m, 2H, BrC$\underline{H}_2$), 4.8 broad S, O$\underline{H}$).

   2-(2-Acryloyloxethoxy)-3,4,5-Tribromosalicylanilide, (4).
Method A: The triethylamine salt of 3,4,5-tribromosalicylanilide
(7.3 g, 13.2 mmol) was dissolved in dimethylformamide (10 ml)
and added to a solution of 2-(p-toluenesulfonyl)ethyl acrylate
(5.5 g, 20.4 mmol) in benzene (20 ml). The mixture was heated
to 93°C for 1 hr. and then cooled to 20°C for 16 hrs. The
reaction was filtered and water (30 ml) added to the filtrate.
The filtrate was extracted with ether and the extract dried with
anhydrous MgSO4. Removal of the ether in vacuo gave a brown
solid and a brown liquid (unreacted 2-(p-toluenesulfonyl)ethyl
acrylate). The solid was recrystallized from chloroform to
obtain 0.4 g (5.6%) of (4). mp 153-154°C. Analysis: Calcd.: C
39.42; H 2.55; N 2.55; Found: C 39.53; H 2.61; N 2.52. IR (KBr;
cm^{-1}): 3500 (NH), 1740 (CO ester), 1650 and 1580 and (CO amide),
1250, 1120, 1070. nmr (DMSO-d$_6$; ppm): 8.0 (d, 1H, acid ring),
7.7 (d, 1H, acid ring), 7.5 (d, 4H, anilide ring), 7.4 (s, N$\underline{H}$),
5.8 (m, 3H, vinylic, ABC pattern), 4.3 (m, 4H, the ethylenic
hydrogens show an A$_2$B$_2$ pattern).
   Method B: 3,4,5-Tribromosalicylanilide (13.4 g, 29.7 mmol)
was dissolved in a mixture of dimethylformamide (20 ml) and
ethanol (50 ml) and the solution was added to 9.9% aqueous NaOH
(11.9 g, 29.7 mmol of NaOH). The resulting solution was heated
to refluxing and 2-bromoethyl acrylate (5.3 g, 29.7 mmol) was
added dropwise. After 8 hrs. the reaction was cooled and (4)
crystallized out of the solution. The (4) was recrystallized
from chloroform to give 1.2 g (8.1%).

   2-(8-Quinolinyloxy)ethanol. 8-Hydroxyquinoline (14.5 g, 100
mmol) was dissolved by warming in 200 ml of water containing
NaOH (4 gm). After cooling to room temperature, 2-chloroethanol
(12 g, 150 mmol) was added to the solution over 5 hrs. The
mixture was stirred for 49 hrs. at 20°C and a precipitate was
filtered. The precipitate was water washed, dried in vacuo and
recrystallized from benzene/ether/pet.ether 50/20/30. Yield
14.5 g (70%). mp 84-85°C (lit. 83-84°). A pure sample
was prepared by chromatography on an alumina column using chloro-
form as the eluent. Analysis (calculated for C$_{11}$H$_{11}$NO$_2$·H$_2$O):
Calcd.: C 63.77; H 6.29; N 6.76; Found: C 63.49; H 6.26; N
6.66. IR (KBr; cm^{-1}): 3400 ()H), 1310 (CO alcohol), 1260 (COC),
1110 (CO alcohol), 1070 (COC). nmr (CDCl3; ppm): 8.8 (m, 1H,
aryl), 7.9 (m, 1H, aryl), 7.2 (m, 3H, aryl), 6.9 (q, 1H, aryl),

5.1 (broad S, O$\underline{H}$), 4.1 (m, 4H, ethylenic showing A_2B_2 pattern).
The hydrogens of the nitrogen ring system show an AMX pattern
where H^1 is a doublet of doublets at 8.8, H^3 is a quartet at 6.9,
and H^2 is a doublet of doublets at 7.9.

    2.17 2-(8-Quinolinyloxy)ethyl Acrylate, (6). 2-(quinolinyl-
oxy)ethanol (9.7 g, 51.3 mmol) and triethylamine (5.2 g, 51.3
mmol) were dissolved in benzene (117 ml) and cooled to 5°C.
Acrylolyl chloride (7 g, 63.0 mmol) was added dropwise and, after
the addition was complete, the mixture was warmed to 20°C for
12 hrs. The benzene solution was washed twice with 5% aqueous
NaHCO$_3$ (100 ml) and three times with water (100 ml). The benzene
solution was dried with anhydrous MgSO$_4$ and the benzene removed
in vacuo to give 9.9 g of a brown oil which recrystallized from
carbon tetrachloride to obtain 9.2 g (73.8%) of mp 96.5-97.0°C.
Analysis: Calcd.: C 69.14; H 5.35; N 5.76; Found: C 68.95;
H 5.44; N 5.71. IR (KBr; cm^{-1}): 3050 (aryl CH), 2850 (aliph.
CH), 1750 (C=O), 1630 (vinyl C=C), 1600 (aryl C=C), 1200 (ArOC),
1100 (COC), 1040 (ArOC), 980, 930. nmr (CDCl$_3$, ppm): 8.8 (m,
1H, aryl), 8.0 (m, 1H, aryl), 7.2 (m, 4H, aryl), 6.0 (m, 3H,
vinyl: ABC pattern), 4.5 (m, 4H, ethylenic, A_2B_2 pattern).

    o-Benzyl-p-chlorophenyl Acrylate, (7). o-Benzyl-p-
chlorophenol(29.7 g, 126 mmol) and triethylamine (12.7 g, 126
mmol) were dissolved in benzene (100 ml) and cooled to 5°C.
Acryloyl chloride (11.4 g, 126 mmol) was added dropwise with
vigorous stirring over a period of 45 min. and then the reaction
was warmed to 20°C for 3 hrs. The reaction mixture was washed
once with 10% aqueous NaHCO$_3$ and twice with water, dried with
anhydrous MgSO$_4$. Removal of the benzene in vacuo gave a brown
oil which was distilled in vacuo (bp $_{0.7}$ 210-212°C) to obtain
24.6 g (71.2%) of (7). The distilled (7) crystallized upon
cooling. mp 41-42°C. Analysis: Calcd.: C 70.46; H 4.77;
Found: C 70.50; H 3.93. IR (neat; cm^{-1}); 1750 (CO), 1480, 1400,
1220, 1150. nmr(CDCl$_3$; ppm): 7.1 (m, 8H, aryl), 6.0 (m, 3H,
vinyl, ABC pattern), 3.8 (s, 2H, benzyl).

    2-(o-Benzyl-p-chlorophenoxy)ethanol. 2-Chloroethanol (11.4
g, 141.6 mmol) was added to a solution of o-benzyl-p-chlorophenol
(20.2 g, 91.5 mmol) in 9.9% aqueous NaOH (52 ml, 5.1 g NaOH,
127.5 mmol). The reaction mixture was refluxed for 15 hr., cooled
and extracted with diethyl ehter. The extract was dried over
anhydrous MgSO$_4$ and the ether was removed in vacuo to obtain a
brown oil. The oil was dissolved in boiling hexane and white
crystals formed upon cooling. Yield: 11.3 g (43.3%). Analysis:
Calcd.: C 68.57, H 5.71; Found: C 68.32, H 5.81. IR (KBr;
cm^{-1}); 3300 (OH), 1600 (C=C), 1480, 1250 (ArOC), 1100 (COC),
1060 (ArOC). nmr (CDCl$_3$; ppm): 7.1 (s, 6H, aryl), 6.5 - 6.7 (m,
2H, aryl), 3.8 (m, 7H, O$\underline{H}$), ethylenic and benzylic hydrogens).
Upon addition of D$_2$O to the nmr sample the integration became 8:6
8:6 aryl/ethlenic and benzylic.

2-(o-Benzyl-p-chlorophenoxy)ethyl Acrylate, (8). 2-(o-Benzyl-p-chlorophenoxy)ethanol (20.4 g, 77.7 mmol) and triethylamine (8.6 g, 85.5 mmol) were dissolved in benzene (100 ml) and cooled to 5°C. Acryloyl chloride (7.5 g, 85.5 mmol) was added dropwise. Triethylamine hydrochloride precipitated immediately upon addition of the acryloyl chloride. The mixture was warmed to 20°C for 4 hrs. and then washed once with 5% aqueous NaHCO$_3$ (100 ml) and twice with water (100 ml). The organic layer was dried with anhydrous MgSO$_4$ and the benzene was removed in vacuo to give a yellow oil which crystallized upon cooling. Recrystallization from acetone yielded 7.3 g (29.7%) of (8). mp 74–75°C. Analysis: Calcd.: C 68.25; H 5.37; Found: C 67.88; H 4.93. IR (KBr; cm^{-1}); 1720 (C=O), 1640 (C=C), 1250, 1070, 1050, 990. nmr (CDCl$_3$; ppm): 7.2–7.6 (m, 6H, aryl), 6.8–5.8 (m, 5H, aryl and vinyl), 4.5 (m, 2H, OCH$_2$), 4.1 (m, 2H, BrCH$_2$), 3.9 (S, 2H, benzylic). The ethylenic hydrogens at 4.1 and 4.5 showed an A$_2$B$_2$ pattern.

2-Pentachlorophenoxyethyl Bromide. A solution of pentachlorophenol (50.1 g, 188.7 mmol) in ethanol (50 ml) was added to 10% aqueous NaOH (85.5 ml) and the resulting mixture was added slowly to refluxing 1,2-dibromoethane (176.6 g, 939.4 mmol). After 12 hrs. water (100 ml) was added to the reaction and the organic layer separated and dried over anhydrous MgSO$_4$ for 8 hrs. Removal of excess 1,2-dibromoethane in vacuo yielded a light brown solid. Recrystallization from 95% ethanol gave 55.5 g (71.9%) mp 84–85°C (lit. 81–82°). Analysis: Calcd.: C 25.70; H 1.34; Found: C 25.9, H 1.06. IR (KBr; cm^{-1}): 1450, 1400, 1300, 1220 (ArOC), 1140 (COC), 1060 (ArOC). nmr (CDCl$_3$; ppm): 4.3 (t, 2H, OCH$_2$), 3.7 (t, 2H, BrCH$_2$); J = 6 Hz.

Vinyl Pentachlorophenyl Ether, (9). Method A: 2-Pentachlorophenoxyethyl bromide (0.4 g, 1.1 mmol) was dissolved in tetrahydrofuran (3 ml) and 1,5-diazabicyclo-5,4,0-undec-5-ene (DBU) (0.16 g, 1.1 mmol) was added dropwise to the solution. The mixture was refluxed for 30 minutes and water (100 ml) was added, and the solution was extracted with chloroform. The chloroform extract was dried with anhydrous MgSO$_4$ and the chloroform removed in vacuo to yield 0.2 g (64.5%) of (9).
Method B: Pentachlorophenol (9.7 g, 36.5 mmol) was dissolved in vinyl acetate (18.8 g, 218.6 mmol) and the solution was cooled to −30°C. Mercuric acetate (0.9 g, 2.8 mmol) and 2 drops of conc. H$_2$SO$_4$ were added to the solution. Reaction was allowed to proceed for 3 hrs. at −30° and then for 12 hrs. at −5°C. The excess vinyl acetate was removed by passing air over the mixture and the solids were chromatographed on grade III alumina using a 4:1 hexane/benzene eluent. The material eluting first was recrystallized from chloroform to obtain 2.7 g (25.0%) of (9).

mp 84–85°C (lit. 82–84°C).[9] Analysis: Calcd.: C 32.82; H
1.03; Found: C 33.15; H 1.00. IR (KBr; cm^{-1}): 1620 (C=C),
1390, 1150, 1120, 980, 950, 910, 850. nmr (CDCl$_3$–DMSO-d$_6$; ppm):
6.5 (Q; 1 J$_{AX}$ = 6 Hx; J$_{BX}$ = 3 Hz), 4.3 (quartet of doublets, 2H,
J$_{AB}$ = 2 Hz).

Method C: Pentachlorophenol (5.1 g, 18.8 mmol), 1,2–vinyl-
bis(triphenylphosphonium)dibromide (13.3 g, 18.8 mmol) and
triethylamine (2.1 g, 20.9 mmol) were dissolved in chloroform
(50 ml) and refluxed for 15 hrs. The chloroform solution was
washed with 0.2 N HBr (50 ml) and then with water (50 ml) three
times, dried over anhydrous MgSO$_4$ and filtered. The chloroform
was removed in vacuo to leave a dark brown oil. The oil was
relfuxed in a 60:40 benzene/5N aqueous NaOH mixture (100 ml) for
5 hrs., the benzene layer separated and the NaOH solution was
extracted with benzene. The combined benzene layers were dried
with anhydrous Na$_2$SO$_4$. Removal of the benzene in vacuo yielded
a dark brown oil which was chromatographed on grade III alumina
using a 4:1 hexane/benzene eluent to give 0.8 g (14.5%) of (9).

   2-(2-Bromoethoxy)-3,4;5-Tribromosalicylanilide. The sodium
salt of 3,4;5-tribromosalicylanilide (8.5 g 18.0 mmol) was
dissolved in a 75:25 dimethylformamide/water mixture (40 ml) and
the solution was added dropwise to 1,2-dibromoethane (45.8 g,
243.5 mmol). The mixture was heated to 90°C for 30 min.,
washed with water (100 ml) three times, and the organic layer
was dried with anhydrous MgSO$_4$. Upon removal of the 1,2-
dibromoethane in vacuo, a brown solid was obtained which re-
crystallized from an acetone – ethanol mixture to give 4.7 g
(46.9%) of 2-(-bromoethoxy)-3,4;5-tribromosalicylanilide. mp
173–174.5°C. Analysis: Calcd.: C 32.52; H 1.98; N 2.51;
Found: C 32.52; H 1.78 N 2.39. IR (KBr; cm^{-1}); 3300 (NH), 1670
and 1600 (C=O), 1550, 1270 (ArOC), 1090 (COC), 1020 (COC). nmr
(DMSO-d$_6$, ppm): 9.2 (broad s, NH; 1), 8.1 (d, acid ring, 1),
7.8 (d, acid ring, 1), 7.5 (d, amide ring, 4H), 4.3 (t, OCH$_2$, 2),
3.7 (5, BrCH$_2$, 2).

   Vinyl 3,4;5-Tribromosalicylanilide Ether, (10). Method A:
2-(2-bromoethoxy)-3,4;5-tribromosalicylanilide (6.9 g, 12.4
mmol) was dissolved in tetrahydrofuran (30 ml) and the solution
heated to 40°C. DBU (2.0 g, 13.3 mmol) was added dropwise and a
white precipitate began to form. After 20 min., the mixture
was cooled to 20°C and stirred for 13 hrs. Water (50 ml) was
added and the reaction was filtered. The solids from the react-
ion was filtered. The solids from the reaction were dried in
vacuo and recrystallized from an acetone-ethanol mixture to give
4.0 g (67.8%) of (10), mp 203–204°C. Analysis: Calcd.: C
37.82; H 2.10; N 2.94; Found: C 37.61; H 1.92; N 2.87. IR
(KBr; cm^{-1}): 300 (NH), 1650 (C=O), 1580, 1270, 1250, 1230, 1160,
1040. nmr (DMSO-d$_6$; ppm): 8.0 (s, 1H), 7.5 (m, 1H), 7.2 (m,
2H), 4.5 (m, 1H), 3.9 (m, 2H).

Method B: 2-(2-Bromoethoxy)-3,45-tribromosalicylanilide
(3.3 g, 5.93 mmol) was dissolved in dimethylformamide (10 ml)
and added dropwise to a solution of NaOH (0.8 g, 20 mmol) in
95% ethanol (20 ml). After 10 min., a yellow precipitate formed.
After 1 hr., the mixture was filtered and extracted with ether.
The extract was dried with anhydrous MgSO₄ and the ether removed
in vacuo to obtain 1.3 g of 2-(-bromoethyoxy)-3,4¦5-tribromo-
salicylanide. The yellow precipitant was recrystallized from an
acetone-ethanol mixture to give 0.5 g (17.7%) of (10). The
water/ethanol dimethyformaide mixture was acidified and a white
precipitate formed. The precipitate was filtered out and dried
to obtain 0.8 g of 3,4¦5-tribromosalicylanilide.

2-(O-Benzyl-p-chlorophenoxy)ethyl Bromide. A solution of
o-benzyl-p-chlorophenol (25.0 g, 114.4 mmol) in 10% aqueous KOH
(loo ml) was added dropwise to refluxing 1,2-dibromoethane
(54.0 g, 287.2 mmol). After 15 hrs. the organic layer was
separated from the reaction, washed three times with 5% aqueous
KOH, three times with water, and dried with anhydrous MgSO₄.
Removal of the excess dibromoethane in vacuo gave a brown oil
which recrystallized from hexane to give 27.1 g(73.0%) of the
title compound. Analysis: Calcd.: C 55.30; H 4.30; Found:
C 55.34; H 4.24. ir (KBr; cm⁻¹): 1600 (C=C), 1500, 1250 (ArOC),
1070 (COC), 1050 (ArOC). nmr (CDCl₃; ppm): 7.1 (s, 7H, aryl),
6.4 (m, 1H, aryl), 4.0 (t, 2H, OCH₂), 3.8 (t, 2H, BrCH₂), 3.3
(s, 2H, benzyl).

Vinyl o-Benzyl-p-chlorophenyl Ether, (11). Method A: o-
Benzyl-p-chlorophenol (5.0 g, 22.9 mmol) and 1,2-vinyl bis-
(triphenylphosphonium)dibromide (16.3 g, 23.0 mmol) were mixed
and dried in vacuo. The mixture was dissolved in chloroform
(50 ml) and triethylamine (2.6 g, 25.7 mmol) in chloroform (8 ml)
was added dropwise at 20°C. After complete addition of the
triethylamine, the mixture was refluxed for 16 hrs. The solu-
tion was washed twice with 0.2 N HBr, three times with water and
poured into ether to give a brown oil which was isolated and
refluxed for 5 hrs. with a 60:40 benzene/5N aqueous NaOH mixture
(100 ml). The aqueous layer was separated and extracted with
benzene. The extract was combined with the benzene from the
mixture , washed twice with water, and dried over anhydrous
MgSO₄. Removal of the benzene in vacuo gave a brown oil which
was chromatographed on grade III alumina using a 4:1 hexane/-
benzene eluent. Removal of the eluent gave 3.9 g (69.6%) of (11).
bp 109–112°C. Analysis: Calcd.: C 73.62; H 6.13; Found C
73.60; H 5.13. IR (neat, cm⁻¹): 1640 (vinyl C=C), 1480, 1250
(vinyl COC), 1150, 1030 (COC), 750, 700. nmr (CDCl₃; ppm): 6.9
(m, 8H, aryl), 6.3 (q, 1H, vinyl), 4.3 (m, 2H, vinyl CH₂), 3.7
(s, 2H, benzyl). The quartet at 6.3 and multiplet at 4.3 show
an ABX pattern: J_{AB} = 1.5 hz, J_{BX} = 6.2 hz, J_{AX} = 14.0 hz.

Method B: o-benzyl-p-chlorophenol (1.8 g, 8.6 mmol) was dissolved in vinyl acetate (4.4 g, 51.4 mmol) and cooled to -20°C. Mercuric acetate (0.6 g, 1.8 mmol) and 2 drops of conc. H_2SO_4 were added. After 12 hrs., the mixture was chromatographed on a silica gel column using a 2:1 hexane/benzene eluent. The first material eluting off the column was 1,1-bis-(o-benzyl-p-chlorophenoxy)ethane (the acetal produced from vinyl o-benzyl-p- chlorophenylether and o-benzyl-p-chlorophenol). None of the vinyl ether was found however. nmr of the acetal in $CDCl_3$ showed a multiplet at .9 (aryl), a quartet at 5.7 (CH), a singlet at 3.8 (benzylic) and a doublet at 1.3 (CH$_3$).

Method C: 2-(o-benzyl-p-chlorophenoxy)ethyl bromide (3.3 g, 10.1 mmol) was dissolved in tetrahydrofurna (50 ml) and the solution heated to refluxing. Potassium t-butoxide (1.4 g, 12.1 mmol) in tetrahydrofuran (8 ml) was added slowly to the refluxing solution. After 16 hrs., the mixture was filtered and water (50 ml) was added. The mixture was extracted with ether, the extract dried with anhydrous $MgSO_4$, and the ether removed in vacuo to obtain a light yellow liquid which was distilled in vacuo to get 1.7 g (67.8%) of (11). b.p.$_{0.04}$ 107-108°C.

General Procedure for Synthesis of Stable Lactices. Illustrated for Latex 1 (Table II).

A. Preparation of Kettle Charge. Aerosol 22 (tetrasodium N-(1,2-dicarboxy-ethyl)-N-octadecylsulfonsuccinamate (10.8 g) methanol (9.5 g), sodium bicarbonate (1.8 g) were dissolved in water (176 g) and an ammonium persulfate solution (38 g of a 10% by wt. aqueous solution) was added. The solution was purged with nitrogen for 15 min. and charged to the polymerization vessel and heated to 60°C. B. Preparation of Pre-Emulsified Monomer Charge. Aerosol A-102 (disodium ethoxylated alcohol half ester of sulfo succinic acid) (10.8 g), methanol (9.5 g) and sodium metabisulfite (0.8 g) were dissolved in water (59.1 g). The monomers were slowly added with stirring to this solution and the resulting emulsion was purged with nitrogen for 15 min. The amounts were: monomer 1 (20 g, 62 mmol.), MMA (200 g, 2.2 moles) and nBA (160 g, 1.25 moles). The resulting monomer emulsion was charged to the reaction vessel (A) from a dropping funnel fitted with a stirrer and agitation was maintained in the dropping funnel during the entire addition procedure. C. Preparation of Delayed Charge. A delayed charge of itaconic acid (3.8 g), diammonium phosphate (0.6 g) in water (19.1 g) was purged with nitrogen for 15 min. D. Polymerization. After the reaction kettle (with its initial charge) reached 60°, the preemulsified monomer charge was added at a rate of 1-2% per minute. When approximately 15% of the monomer charged had been added, the addition was stopped until polymerization was indicated by an exotherm. The delayed charge was then added to the kettle and the monomer charge was started again. After all the

monomer had been added, the polymerization was continued for 1 hr. at 60°, cooled and filtered through a 100 mesh nylon screen to remove any coagulated material. A portion of this latex was poured into a 7:1 methanol/saturated aqueous NaCl solution to precipitate the polymer which then was twice reprecipitated from THF into methanol. Analysis: C 61.09, H 8.14, Cl 2.95.

Acknowledgments

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Poly(thiosemicarbazide) Copper(II) Complexes as Potential Algicides and Molluscicides

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Polymeric poly (thiosemicarbazide) copper (II) complexes have been made and are being evaluated as algicides and molluscicides. This endeavor showed considerable promise in the laboratory and slow release data for copper (II) indicate that reusable cartridge type copper (II) systems for schistosomaisis and algae control may be feasible.

A number of poly (thiosemicarbazides) have been prepared which complex copper (II) so strongly that mineral acids and even ethylenediamine tetraacetic acid do not effectively remove the copper ion from the polymers (see Scheme 1)[1]. However, in water, initial studies indicated that over longer periods of time, copper (II) ions were released in such fashion that the polymers might be useful for the slow release of copper (II) to provide algae or schistosomiasis control agents (see Table 1) in natural waters. The effect of algae polluting natural waters is a common sight which we all are familiar with.

Schistosomiasis is a serious, often fatal, disease which attacks humans where aquatic snails are the disease carrier[2]. Such snails live in irrigation ditches, streams, swamps, and other natural waters in the warmer climates. The disease carrying snails can be killed by adding copper (II) to their aquatic habitat, and the classic source of the needed copper (II) is cupric sulfate. However, in this case and for algae the salt rapidly migrates away in moving waters or is exchanged into the surrounding soil being very rapidly lost. Thus, frequent additions are required to provide effectiveness. One way tried to retard loss of copper (II) has been to mill copper sulfate into rubber pellets from which water slowly leaches out copper (II) to provide the needed ions to kill snails. However, in natural waters these pellets float away or sink into the channel soil. Of course, any ion exchange resin could provide a substrate to hold copper (II) so that natural water soluble salts would ex-
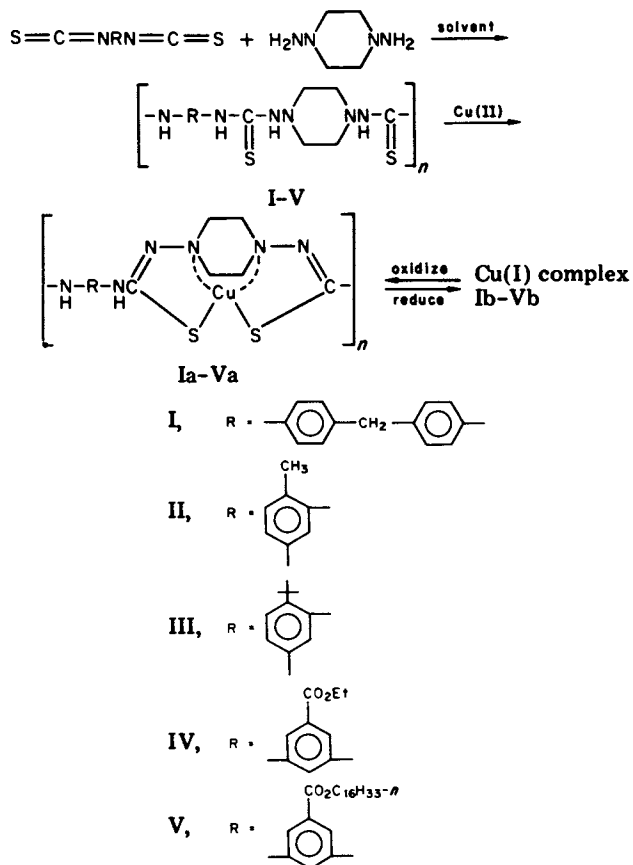
Scheme I.   Preparation of poly(thiosemicarbazides).

Table 1

Release of Copper (II) From Polymers I-V as a Function of Time

% Release Over Time*

Polymer	1 da.	2 da.	3 da.	4 da.	7 da.
I	14.6(11.9)	15.6(20.4)	17.4(21.8)	18.8(23.2)	17.3(27.4)
II	4.3(5.8)	5.8(8.6)	7.2(11.6)	8.7(14.4)	10.1(17.3)
III	11.6(18.9)	14.5(20.3)	14.5(21.7)	14.4(24.6)	14.4(21.4)
IV	11.6(13.1)	13.1(14.5)	11.6(15.9)	13.0(17.4)	11.5(18.9)
V	<0.1(0.6)	<0.1(0.6)	0.3(0.6)	0.3(0.6)	<0.1(.84)

Polymer	9 da.	11 da.	14 da.
I	21.6(28.8)	23.3(31.1)	23.2(33.5)
II	10.1(20.2)	9.8(25.6)	12.9(25.8)
III	19.4(28.8)	19.4(32.3)	24.5(34.8)
IV	11.5(20.2)	19.4(19.4)	18.1(23.2)
V	<0.1(1.1)	<0.1(4.9)	0.5(5.0)

* Figure in parentheses is with agitation of the samples.

change away the copper (II) while the regenerable ion exchanger
was in a fixed accessible site. However, the copper must be
slowly released and not exchanged away from the ion exchanger too
quickly. Thus an ion exchanger which binds copper (II) tightly
is needed. Yet, the binding of the copper (II) must not be so
tight that it will not exchange away from the polymer. The[4]
desired rate of copper (II) release should be 1-2% per day.
Knowing that copper (II) was tightly bound to I-V, the rate of
release from the polymer in water was observed over a period of
time.

     As can be seen from Table 1 data, the preliminary data
indicate that a slow release of copper (II) may be occurring over
a fourteen day period. This release apparently is characterized
by an initial surge followed thereafter by release of smaller
amounts as time goes on. Over a fourteen day period after the
initial surge, the release for polymers I, III, and IV averages
about in the neighborhood of 1% per day.

     It was desired to study the release of copper (II) ions
from I-V in detail particularly in synthetic natural waters which
contain dissolved ligands capable of eluting copper ions from the
polymer and/or precipitating free copper (II).

Experimental

     Copper release experiments were performed by placing a
weighed amount of resin in a 250 ml Erlenmeyer flask to which a
known amount of water of a given chemical composition had already
been added. If pH adjustment was required, it was done im-
mediately after the polymer was added. The flasks were then
sealed on a rotary shaker (Lab-Line) at 125 rpm. After a given
time either the total or dissolved copper concentration in the
water was determined.

     A semidynamic system was used to examine the resin re-
lease properties if introduced to natural water. Resin was placed
in nylon mesh bags and put in a flask as previously described.
After a certain time interval the nylon bags were removed and
placed in a fresh flask of water. This was repeated until no
further release of copper was observed.

     The resin for these tests was loaded using a batch method.
A known weight of resin was placed in a flask containing a known
volume and copper concentration at pH = 3. The flasks were then
placed on the rotary shaker for 48 hours. They were then removed
and the copper solution was filtered through a 0.45μm membrane
filter. The solution was subsequently analyzed for copper con-
centration. The polymers were then stored in a moist state until
used. If rinsing was performed, it was done immediately prior
to resin addition to the release water.

     Computer equilibrium calculations were performed using
MINEQL[5] and an IBM 360-65. The solubility of copper in equili-
brium with $Cu(OH)_2(s)$ and $Cu_2(OH)_2CO_3(s)$ (malachite) was examined.

Composition of Synthetic
Natural Water$\frac{3}{}$

pH$_2$ = 7.80 - 8.15
Ca$_{+2}$ = 67 mg/L as CaCO$_3$
Mg^{+2} = 67 mg/L as CaCO$_3$
Alk = 100 mg/L as CaCO$_3$
Fe^{+3} = 0.1 mg/L
SO4 = 54 mg/L
Cl$^-$ = 52 mg/L

Copper Release Experiments

Both I and IV were initially used in the release ex-
periments. First, the effects of pH were examined. As shown
in Figure 1 A and B, at pH = 3 a quick initial release (approxi-
mately 15 minutes for IV, 6 hours for I) occurred and after this
time no more copper was released. The difference in Figure 1
A and B is that Figure 1 B takes into account resin weight present
in the flask.

IV appears to reach the equilibrium release concentra-
tion earlier. This result is attributed to the following two
reasons. First, under acid conditions IV may be partially eluted
with acid solutions. Second, the smaller particles would have a
greater surface area and tend to have higher rates of transfer.
All the measurements were of dissolved copper unless noted other-
wise.

Since natural waters are obviously not at pH=3, the
polymers were next placed in a 2×10^{-3}M sodium bicarbonate
solution at a pH=6.0 - 6.5. As can be seen in Figure 2, the
same type of quick initial release occurred. The measured copper
values are somewhat lower due to precipitation of copper but
probably also due to the fact that less copper is released at
higher pH values. The solutions were not buffered well at this
pH and bicarbonate condition, and as a result the pH of most
solutions when removed for measurements was around 4.0.

The result for the same type of solution at an initial
pH of 8 were as expected. Once the copper loaded resin was added,
copper was immediately precipitated. The pH of the solution
decreased to 7.5-7.6. IV once again tended to release a fraction
of its copper quickly, while I released its fraction slightly
slower as evidenced by the plateaus in Figure 3 A and B.

As a result of these tests I in the pellet form was
chosen for additional investigation because its release charac-
teristics were slower. The rest of the release experiments were
performed in a synthetic water to better model natural systems.
The first experiment involved placing the pellets in the synthetic
natural water (SNW). As was expected results similar to the
pH = 8 experiment were observed as shown in Figure 4.
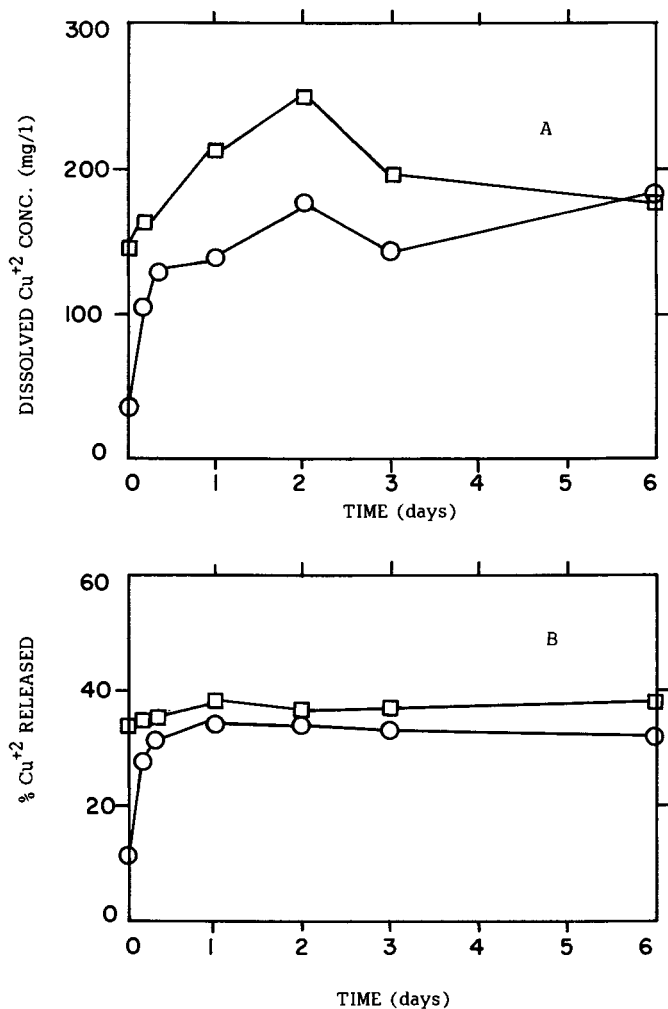
Figure 1. Release of copper by I (○) and IV (□) in deionized water at pH 3. Key: A, dissolved Cu²⁺ concentration (mg/L); and B, percent Cu²⁺ released.
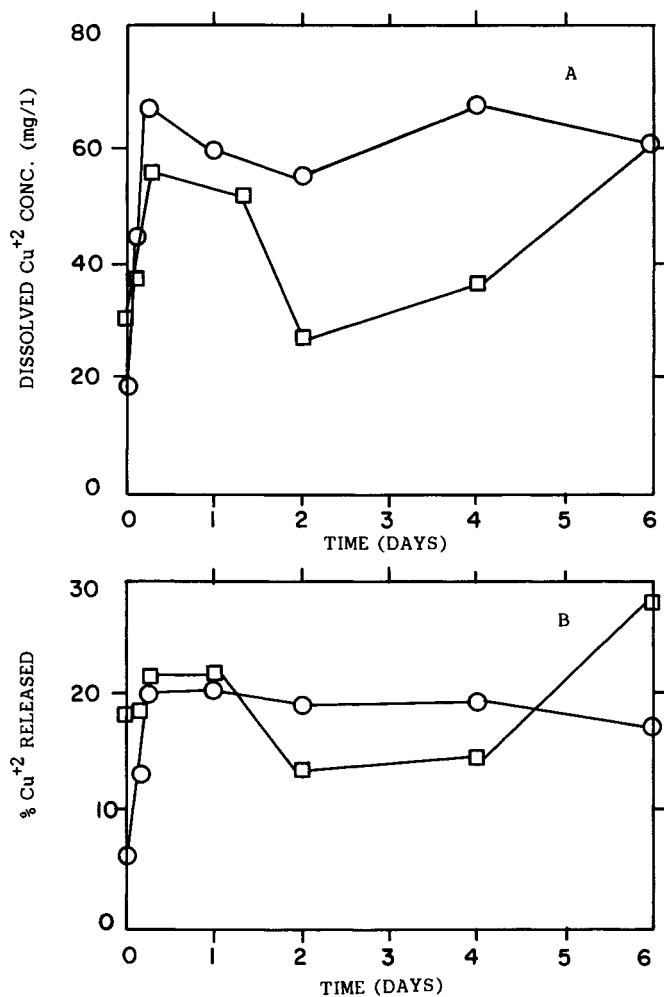
Figure 2.   *Release of copper by I (○) and IV (□) in 2 × 10⁻³ M NaHCO₃ solution at pH 6–6.5. Key: A, dissolved Cu²⁺ concentration (mg/L); and B, percent Cu²⁺ released.*
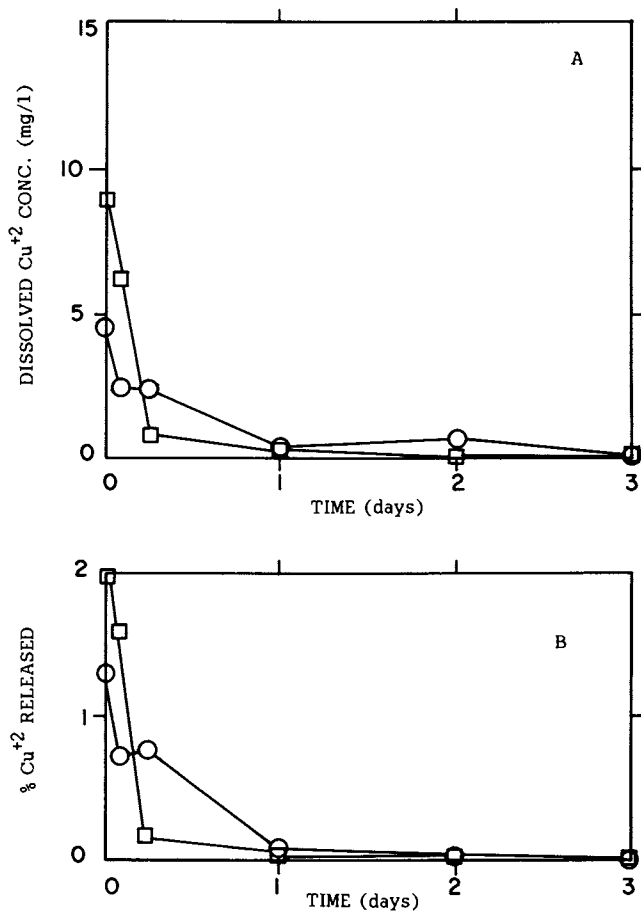
Figure 3.   Release of copper by I (○) and IV (□) in 2 × 10⁻³M. NaCO₃ solution at pH 8. Key: A, dissolved Cu²⁺ concentration (mg/L); and B, percent Cu²⁺ released.
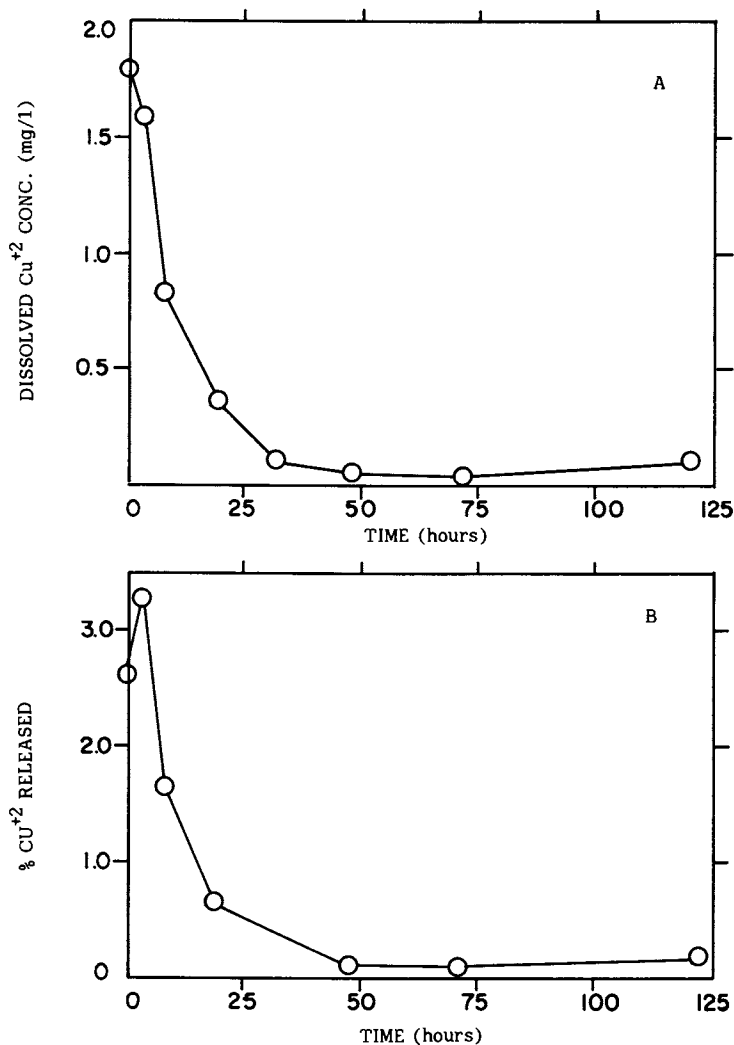
*Figure 4.   Release of copper by I in synthetic natural water at initial pH 7.8.
Key: A, dissolved Cu²⁺ concentration (mg/L); and B, percent Cu²⁺ released.*

        In order to more closely simulate the dilution effects
of a natural water body, a semidynamic system was evaluated
using the SNW. As is evidenced by Figure 5A, in both the rinsed
and unrinsed cases the pellets released approximately 80 percent
of their total copper in the first 15 minutes. After 30 minutes
very little additional copper was released from the resin.
        The effect of particle size was also examined in the
semidynamic system. I (20 x 50 mesh) was exposed under the same
conditions (Figure 5 B) and the rinsed cases behave much the
same; therefore, it is safe to assume that a good portion of the
copper in most of these experiments came from the surface water
associated with the moist resin, with more surface water being
associated with the smaller size particles. This copper most
likely originated from the solutions used to load the polymers.
        A final series of experiments were run to evaluate the
effects of the presence of an organic complexing agent on re-
lease properties. EDTA was chosen as a model organic ligand.
As in evidence by Figure 6 the EDTA complexes as much copper as
it can. The dissolved copper concentrations at the end of the
test for the 10^{-5} M EDTA case are very close to the theoretical
values calculated 0.63 mg/l Cu^{2+} as will be presented below.
When the EDTA concentration exceeded the stoichiometric amount
of copper initially on the resin, a total release of 80 percent
of the initially bound copper was observed. This result is not
unexpected because of the high formation constant for copper –
EDTA complexes. In addition some of the copper in solution was
probably due once again to the surface water.

Computer Equilibrium Calculations

        To better understand the mechanisms involved in the re-
lease studies, computer equilibrium calculations were performed
using Mineql (4). The equilibrium speciation of copper with
hydroxide as a function of pH was first examined. As can be
seen in Figure 7 for a 10^{-3} Cu_T solution; the solid phase takes
over from the free copper ion as the dominant phase at a pH of
6.0. For the much lower copper concentrations typically found
in water the transition pH is shifted higher.
        Next, calculations were performed to demonstrate the pos-
sible speciation occurring in the synthetic natural water used.
In Table 2 a summary of the values calculated can be found, It
can easily be seen that $Cu_2(OH)_2CO_3(s)$ (malachite) is the
dominent species for this particular pH and ion composition, and
accounts for essentially all the copper. The free copper con-
centration is relatively low 3.8 X 10^{-9}M (0.24 ,ug/L). The total
soluble Cu(II) concentration is very close to the value found
in actual experiments (see Figure 4 and Table 2). The $Cu(OH)_2(s)$
concentration is not available due to the method of solution used
in Mineql (4). It performs calculations to determine the dominant
phase and once it selects one the other is ignored.
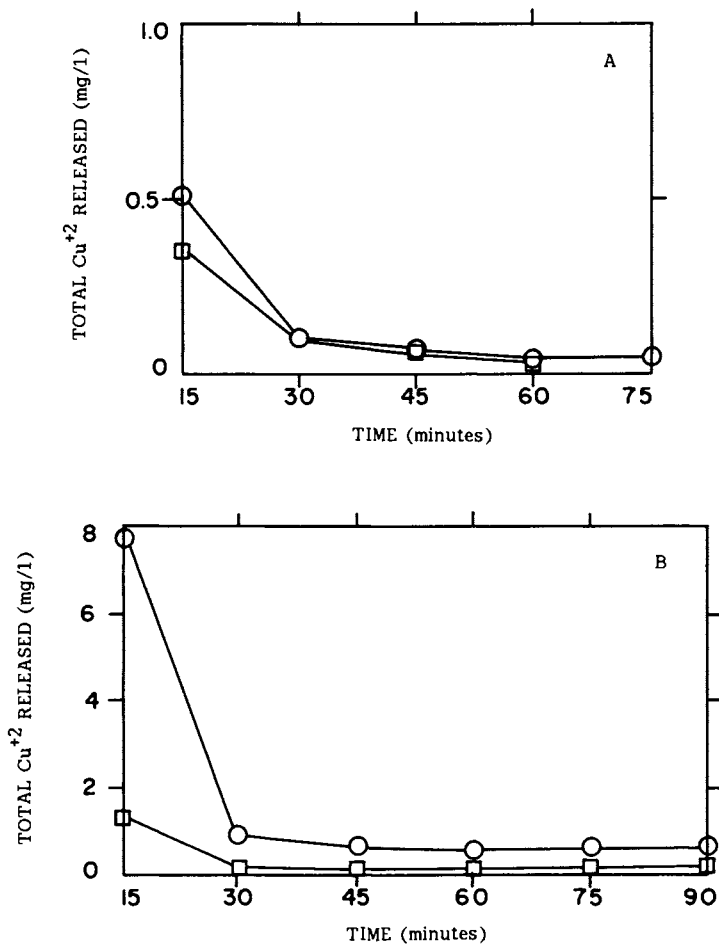
Figure 5.   Release of copper by I (20 × 50-mesh pellets) in a semidynamic system using synthetic natural water.   Key: ○, unrinsed; □, rinsed; A and B, total Cu²⁺ released (mg/L).
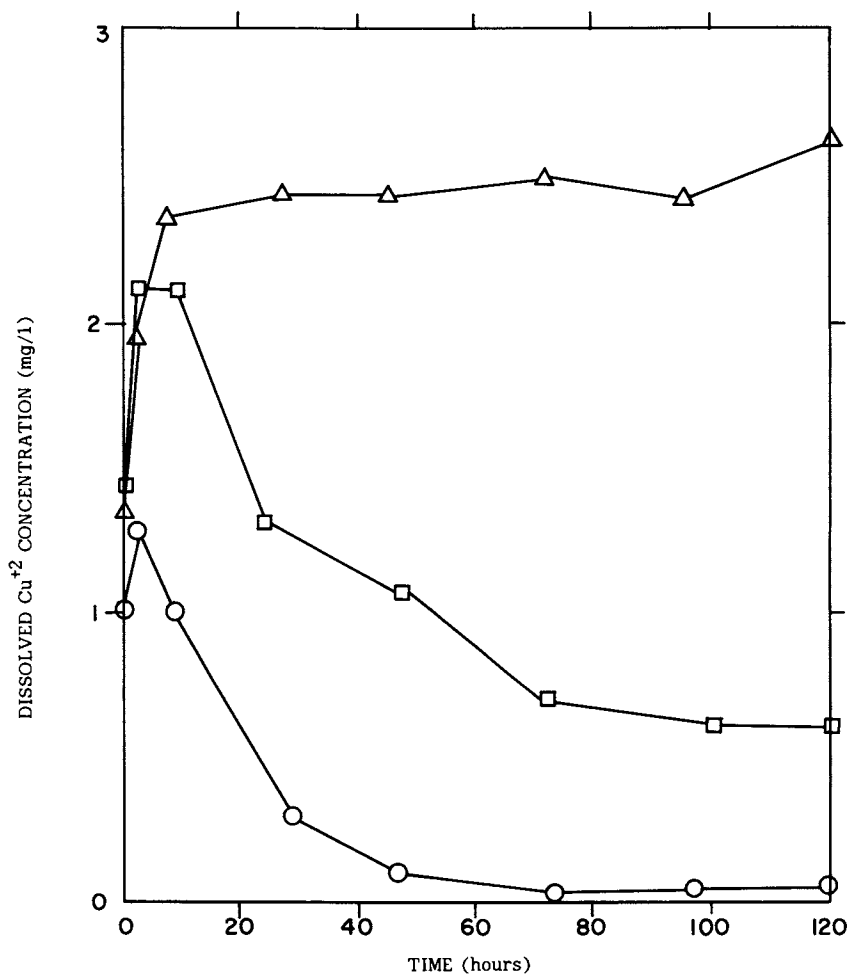
Figure 6.　Effects of EDTA in synthetic natural water on the release of I pellets at average initial pH 8.05.　Key: ○, 10^{-6}M EDTA; □, 10^{-5}M EDTA; and △, 10^{-4}M EDTA.
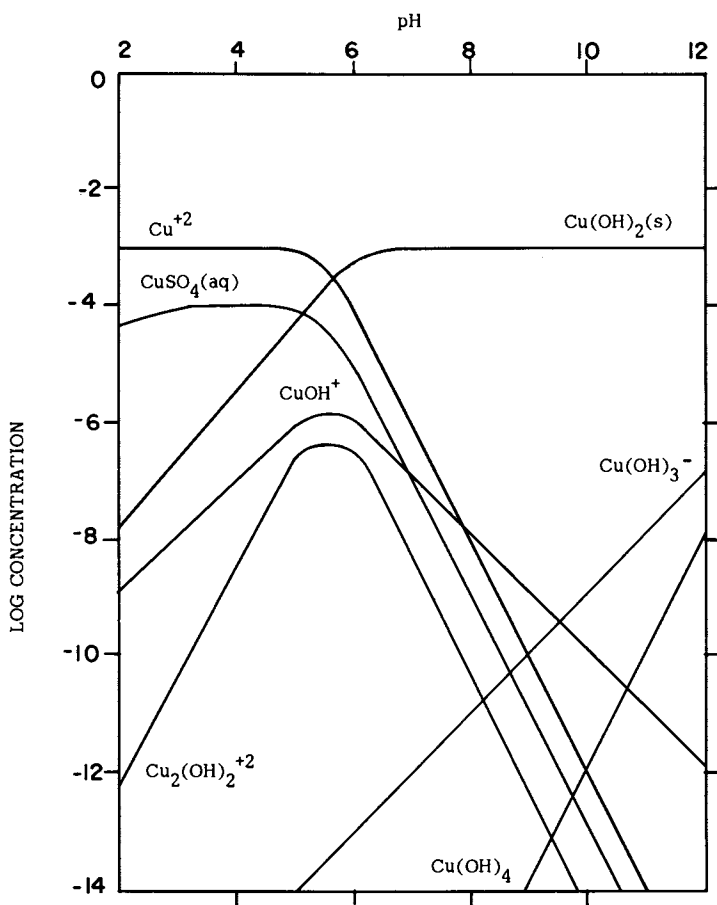
Figure 7. Copper hydroxide equilibrium for a $10^{-3}M$ solution of $CuSO_4$ as a function of pH.

Table 2   Speciation of Copper in Synthetic Water at pH = 8 (Cu_T^{+2} = 2 X 10^{-5}M; C_T = 2 X 10^{-3}M)

Species	-Log Concentration
Cu^{+2}	8.41
$CuCO_3(aq)$	6.80
$Cu(CO_3)_2^=$	8.36
$CuSO_4$	9.97
$CuCL^+$	10.84
$CuCL_2$	13.98
$CuOH^+$	8.40
$Cu_2(OH)_2$	11.21
$Cu_2OH_2CO_3(s)$	5.00
$Cu(OH)_2$ (s)	N.A.

N.A. -   Exact value not available due to the method of computer solution. $Cu_2(OH)_2CO_3(s)$ was the only solid phase considered.

The shift of dominance in solid phase for the synthetic natural water used, occurs between total inorganic carbon concentrations 10^{-4} and 10^{-3} M as shown in Table 3. Therefore, for most of the experiments conducted in the laboratory it is assumed that the solid phase present was malachite. A natural water could easily reduce the free copper concentration to a very low value even in the presence of high total copper. This can easily be seen in Tables 3 and 4.

The presence of organic complexing agents and a variation of pH were also examined to see what effects these might have in synthetic water. EDTA was once again chosen for the model organic ligand and pH was varied from 6 to 9. For lower pH values higher free copper concentrations were found as shown in Figure 8. As EDTA concentration increased the copper that it complexed seemed to originate from the solid phase, thus not affecting the free Cu^{2+} until the solid phase was completely dissolved. At this point the free copper along with some other species such as $CuOH^+$ and $Cu_2(OH)_2^{+2}$ provided the source for the EDTA. Figures similar to Figure 8 can be used to predict free copper concentration changes that might result from pH or organic ligand concentrations.

On the basis of these findings and the examination of the literature it is quite evident the overall picture of copper speciation should be examined in toxicity studies. Organic and inorganic complexation play a very important role in the speciation and toxicity and can not be ignored. Before applying these resins to aquatic organism control more testing in dynamic systems along with toxicity studies should be performed. The resins show high promise as controlled release substrates for copper(II).

Table 3   Comparison of Speciation of Copper in Varying $CO_3^=$ Concentrations at pH = 8 for 1 X 10^{-3} Cu_T (-Log Concentration)

Species	$CO_3^= = 0$	$CO_3^= = 10^{-4}$	$CO_3^= = 10^{-3}$	$CO_3^= = 10^{-2}$
Cu^{+2}	8.06	7.94	8.13	8.66
$CuOH^+$	7.90	7.90	7.98	8.71
$Cu_2(OH)_2^{+2}$	10.24	10.24	10.63	11.76
$Cu(OH)_3^-$	10.84	10.84	11.03	11.56
$Cu(OH)_4^=$	15.82	15.81	15.97	16.35
$CuSO_4$	8.88	8.89	9.14	9.98
$Cu_2HCO_3^+$	-	9.97	9.49	8.89
$CuCO_3$	-	7.57	7.09	6.49
$Cu(CO_3)_2^=$	-	10.46	9.25	7.22
$Cu(OH)_2(s)$	3.00	3.00	N.A.	N.A.
$Cu_2(OH)_2CO_3(s)$	-	N.A.	3.3	3.3
$CuCO_3(s)$	-	N.A.	N.A.	N.A.
$Cu_3(OH)_2(CO_3)_2(s)$	-	N.A.	N.A.	N.A.

N.A. - Not available due to the method of computer solution.

Table 4 The Effect of pH and EDTA on Cu^{+2}, $CuOH^+$, $Cu_2(OH)_2^{2+}$ Concentration in Synthetic Water with $Cu_T = 2 \times 10^{-5} M$

pH = 6

-log EDTA	Cu^{2+}	-log $CuOH^+$	$Cu_2(OH)^{2+}$
10^{-6}	5.21	7.2	8.81
10^{-5}	5.21	7.2	8.81
1.5×10^{-5}	5.38	7.36	9.14
2.0×10^{-5}	6.30	8.29	10.99
5.0×10^{-5}	11.46	13.44	21.30
10^{-4}	11.93	13.92	22.25

pH = 7

-log EDTA	Cu^{2+}	-log $CuOH^+$	$Cu_2(OH)^{2+}$
10^{-6}	6.89	7.87	10.16
10^{-5}	6.89	7.87	10.16
1.5×10^{-5}	6.89	7.87	10.16
2.0×10^{-5}	7.87	8.86	12.13
5.0×10^{-5}	11.46	12.45	19.31
10^{-4}	11.94	12.93	20.27

pH = 8

-log EDTA	Cu^{2+}	-log $CuOH^+$	$Cu_2(OH)^{2+}$
10^{-6}	8.41	8.40	11.21
10^{-5}	8.41	8.40	11.21
1.5×10^{-5}	8.41	8.40	11.21
2.0×10^{-5}	8.81	8.80	12.01
5.0×10^{-5}	11.47	11.46	17.33
10^{-4}	11.95	11.93	18.28

pH = 9

-log EDTA	Cu^{2+}	-log $CuOH^+$	$Cu_2(OH)^{2+}$
10^{-6}	9.92	8.90	12.22
10^{-5}	9.92	8.90	12.22
1.5×10^{-5}	9.92	8.90	12.22
2.0×10^{-5}	9.92	8.90	12.22
5.0×10^{-5}	12.30	11.29	16.99
10^{-4}	12.74	11.73	17.87
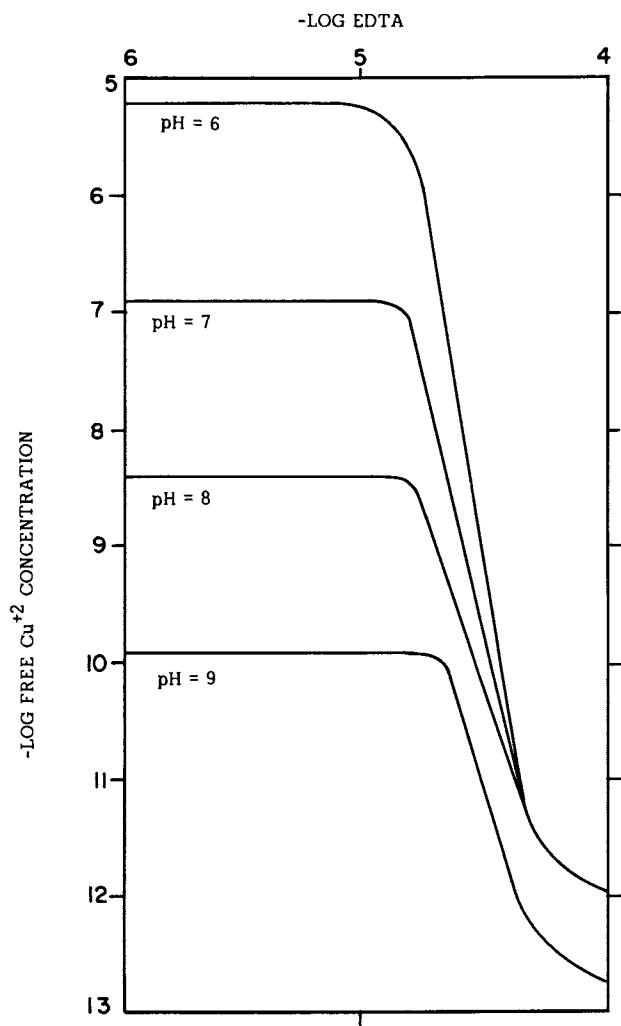
Figure 8. Effect of EDTA and pH on free copper concentration in synthetic water.

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Potential Polymeric Herbicides Derived from Poly(vinyl alcohol): Modification of Polymers[1]

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Isocyanates can be reacted with poly(vinyl alcohol) in homogeneous solutions in dimethylsulfoxide to produce modified polymers that contain between 10 and 100% of the hydroxyl groups replaced by the appropriate carbamate group. In this study, modified polymers were prepared using phenyl, 3-chlorophenyl and 4-chlorophenyl isocyanates and these polymers bear a strong structural similarity to known carbamate herbicides.

Many types of biologically active polymeric systems have been described in the literature including drug, insecticide, herbicide and fungicide systems. These systems could involve biologically active polymers or the controlled release of a low molecular weight species from a polymeric matrix and these systems have been reviewed in several recent reports (1-6). In the controlled release system, the biologically active agent is contained within the polymer in some fashion and could be a 'solution' or a dispersion of this agent in a polymeric matrix or the enclosing or encapsulation of the agent in the polymer (as in a microcapsule). The primary objective of such controlled release systems is to permit the release of a fairly constant level of the biologically active agent for a relatively long period of time. A biologically active polymer, on the other hand, would exhibit its biological activity as a polymer and would not require the release of a low molecular weight species for its action. The potent biological activities of the nucleic acids and the enzymes are examples of polymeric biologically active agents and a number of synthetic polymers (e.g. the divinyl ether- maleic anhydride cyclocopolymer) have shown this type of behavior (7). Many polymeric systems lie between these extremes and sometimes these may exhibit biological activity directly (as a polymer) or serve as a source of controlled release of a low molecular weight species that is attached to the polymer. In the latter case, the system would bear a close

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[1] This is Part IV in a series.

thermokinetic relationship to an ideal monolithic controlled
release system if the bioactive agent were uniformly distributed
along the polymeric backbone. In actual practice, it is not
always obvious whether a polymeric molecule derives its activity
directly or by the release of a low molecular weight species and
this can only be deduced experimentally. If a polymeric system
showed release of a known bioactive agent, it is probable that
this system acts as a controlled release system but this does not
preclude the possibility of this system also functioning as a
polymeric biological agent. On the other hand, if a system does
not show the release of any bioactive agent and still exhibits
biological activity, then this system must function as a polymeric
biologically active system.

     Both the polymeric systems and the controlled release systems
could be valuable in medicinal areas or in pest control. In each
case, prolonged activity of the bioagent would be expected as well
as the maintainance of a low, fairly constant concentration of
this agent. A non-degrading, polymeric bioagent could maintain
its activity for an indefinite period but could create new envi-
ronmental challenges. It is important to take this potential
problem into consideration in designing a polymeric bioagent since
some polymers can survive for a very long time in the natural
environment and could create major pollution problems. Neverthe-
less, a polymeric bioagent has several unique potential advantages
over a controlled release agent. For example, the polymeric
bioagent could be made water soluble fairly readily (e.g. by
copolymerization with a solubilizing monomer) and, in principle,
could be directed to a specific target site (e.g. a diseased
organ or a specific type of weed) by careful, selective copolymer-
ization reactions. Such specific effects cannot be expected for
controlled release systems. These special considerations have
been discussed recently for polymeric drugs (7,8).

     Polymeric bioagents can be prepared by the modification of
existing polymers (9,10) or by the synthesis and polymerization of
new monomeric agents (3,7,8). In the present paper, we are de-
scribing the synthesis of some new, potential herbicide polymers
by the modification of poly(vinyl alcohol) in a continuation of
our previous work in this area (11-13).

Experimental

     Materials Used. The poly(vinyl alcohol) samples used in this
study were two fully hydrolyzed materials which were vacuum dried
several days at 100°C and 10 mmHg to remove any residual water.
These PVA's had molecular weights of 14,000 and 60,000. Dry,
analytical grade dimethylsulfoxide (DMSO) was used as the reaction
solvent. The isocyanates were used as supplied and included
phenylisocyanate (Eastman), 4-chlorophenylisocyanate (Columbia)
and 3-chlorophenylisocyanate (Eastman).

Modified Polymer Preparation. The basic procedure for the preparation of these modified polymers is outlined in Equation (1) and has been described in detail in our earlier publications (11-13). The basic procedure is to dissolve the dried PVA in DMSO and then add, with stirring, a solution of the isocyanate in DMSO. (While the isocyanate could be added directly, this procedure minimizes any problems that could arise by adding a high concentration of the isocyanate in preparing a partially substituted product which could give a highly heterogeneous material.) A small amount of triethylamine is added as a catalyst and the system is stirring several days at room temprature. The modified polymer can be isolated by pouring the system into a large amount of water (preferred) or methanol while stirring rapidly. The precipitate is collected by filtration, washed with non-solvent, air dried and vacuum dried to give the modified polymer. Using this general procedure, modified polymers can be prepared with 10 to 100% substitution of the hydroxyl groups in recovered yields that normally exceed 90%.

In a typical example, 6.6 g. PVA (MW of 14,000) was dissolved in 100 ml. DMSO and a solution of 27.6 g. 3-chlorophenylisocyanate (20% excess) in 100 ml. DMSO was added, with sitrring, followed by 1 ml. triethylamine. After 5 days stirring at room temperature on a magnetic stirrer, the polymer was isolated by pouring into several liters of water. After filtering and drying under vacuum, 26.7 g. modified polymer (90.2% of theory) was isolated which had essentially all the -OH groups substituted by carbamate groups. The isocyanates used in this study included phenyl, 3-chlorophenyl and 4-chlorophenyl isocyanates.

Polymer Properties. Infrared spectra were determined for the modified polymers using a Beckman Acculab 4 Spectrophotometer on films cast on NaCl plates from dimethylacetamide solutions. As the degree of substitution increased, the -OH peak (3300 cm^{-1}) decreased and peaks due to the amide II band (1560 cm^{-1}), -NH and aromatic bands increased.

As the degree of substitution increased, the polymers became more brittle and film formation became more difficult. Polymer solutions, in DMSO or DMAC, became less viscous as the degree of substitution increases even though the polymer molecular weight increases. Below about 20% substitution, these solutions tend to gel in the same manner as unsubstituted PVA1 solutions. (The molecular weight of the repeat unit increases about 450% in PVA1 which is completely substituted with a ClC_6H_4NCO and a polymer with an initial molecular weight of about 14,000 would increase to about 62,800.)

Hydrolysis studies were run by placing a 0.5 g. polymer sample in 50 ml. distilled water and allowing this to stand with occasional stirring. After two months, an aliquot of the supernatent liquid was analyzed spectroscopically in a Cary 14 for the presence of the appropriate aniline derivative. A polymer fully

substituted with $4\text{-}ClC_6H_4NCO$ only releases 0.46% of the 4-chloro-
aniline in this two month period.

Discussion

Poly(vinyl alcohol) readily reacts with a wide variety of
isocyanates in DMSO solution (Equation 1) to produce modified
polymers that contain 10-100% of the original hydroxyl groups
substituted with carbamate groups. The yields of such reactions
are essentially quantitative. Similar results have been reported
from other research groups (14-16). Under these conditions,
poly(vinyl alcohol) has been reacted with 3- and 4-chlorophenyl-
isocyanate (this study), phenylisocyanate (12-14), methyl, ethyl,
isopropyl and 1-naphthyl isocyanates (14) and methoxymethyliso-
cyanate (15,16). The reaction of PVAl with phenyl, tolyl and 4-
chlorophenyl isocyanates in dimethylacetamide solutions has also
been reported (17).

$$-\left(CH_2\underset{OH}{\overset{|}{CH}}\right)_n \quad \xrightarrow[\text{DMSO, Et}_3\text{N}]{x\ R-NCO} \quad -\left(CH_2\underset{OH}{\overset{|}{CH}}\right)_{n-x}-\left(CH_2\underset{\underset{\underset{NHR}{|}}{\overset{|}{\underset{C=O}{O}}}}{\overset{|}{CH}}\right)_x$$
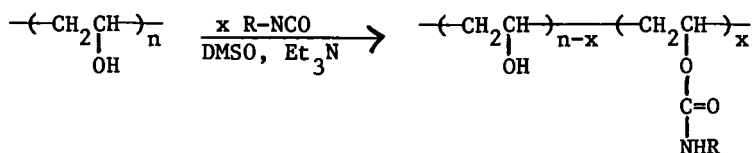
(Equation 1)

The carbamate containing polymers prepared in this study bear
a close structural relationship to some relatively common carba-
mate herbicides. A comparison of these structures is shown in
Figure 1. Carbamate herbicides are known to function by inhibit-
ing cell division and are classed as antimitotic agents. These
carbamate herbicides are normally applied to the soil rather than
the foliage and are best applied as a pre-emergent control materi-
al. Although each herbicide has a specific group of plants to
which it is either harmless or toxic, the following generaliza-
tions can be made. The carbamate herbicides are normally harmless
to the following plants: flax, garlic, marigolds, mustard, onions,
peas, rape, safflower, spinach, soybeans, sugar beets and sun-
flower. On the other hand, the carbamate herbicides are usually
toxic to the following plants: barley, crabgrass, darnel (rye-
grass), fescue, goosegrass, maize, quackgrass (counchgrass) oats,
timothy, watergrass, wild oats and wheat. It is well established
that the carbamate herbicides degrade in the soil. via microbial
action, to produce an aniline derivative as shown in Equation 2.
This hydrolysis occurs fairly rapidly and the original herbicide
structure does not survive long in the natural environment.
Whether the resulting aniline derivative can also be a source of
antimitotic activity does not appear to be established but the
toxicity of aniline derivatives is well known (18-20). The
antimitotic activity is generally believed to occur via the
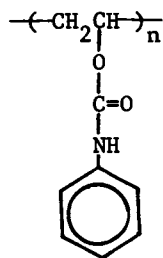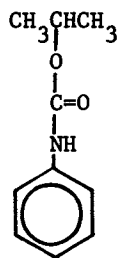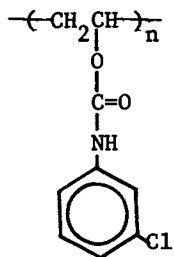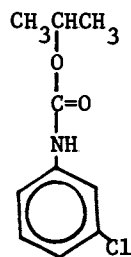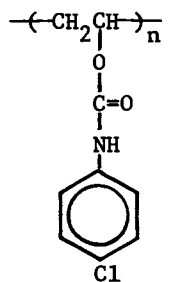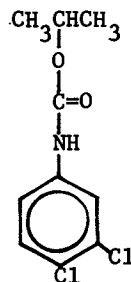
Figure 1.   *Structural relationships between the modified polymers and some carbamate herbicides.*

carbamate molecule as a whole rather than from some degradation
product. In addition, not all carbamate herbicides contain a
N-phenyl group and an aniline derivative is not always a degrada-
tion by-product. (For example, dichlormate is 3,4-dichlorobenzyl
methylcarbamate and would yield methyl amine and 3,4-dichloro-
benzyl alcohol.) A number of insecticide carbamates exist which
are known to function by inhibiting acetylcholinesterase (21,22).
These are usually derivatives of methyl carbamate.

$$CH_3\underset{\underset{\underset{\underset{\underset{C_6H_5}{NH}}{|}}{C=O}}{\overset{|}{\underset{|}{O}}}}{C}HCH_3 \quad \xrightarrow{H_2O} \quad CH_3\underset{\overset{|}{OH}}{C}HCH_3 \quad + \quad C_6H_5NH \quad + \quad CO_2$$

(Equation 2)
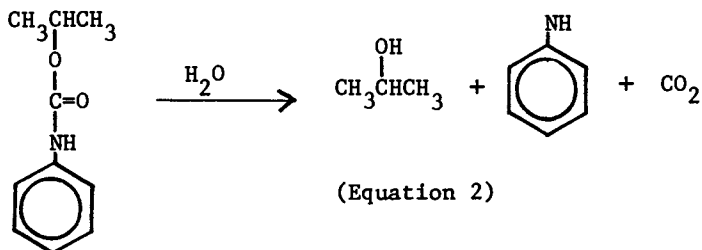
     The controlled release of various types of herbicides have
been studied by many workers and has been summarized elsewhere
(4-6, 23). Less is known, however, about the controlled release
of herbicides for land plants than for aquatic plants. Previous
studies by other research groups have reported that isocyanate
modified polymers, made from chitin, cellulose or poly(vinyl
alcohol), hydrolyze slowly in aqueous media to release the corres-
ponding aniline derivative (17,24,25). No data appears to be
currently available on how effective these agents are as herbi-
cides.
     In the present study, we have prepared a series of polymers
which contain the carbamate units shown in Figure 1. While these
materials do resemble known herbicides closely, this does not
necessarily mean these polymers would also exhibit this type of
activity although there are many instances of polymeric drugs that
are biologically active in the same way as low molecular weight
analogs without these polymeric drugs serving as a controlled re-
lease source of this drug (1,3,7). The systems studied here do
release the expected aniline derivative when exposed to water but
the rate of release is very slow (approximately 0.23% per month)
and these modified polymers can be isolated from the preparation
reaction media (DMSO) by pouring this solution into water. The
polymers themselves are completely insoluble in water, therefore
we would not expect them to be highly mobile under the use con-
ditions that would be encountered in agriculture, etc. This would
be an advantage in maintaining the proper herbicide levels and
also in preventing unwanted migration of this agent to other areas
where it could cause environmental damage. At the present time,
biological activity data are not available for these polymers.
     The solution viscosity behavior of the new 3- and 4-chloro-
phenyl isocyanate derivatives are similar to that reported earlier

for the phenyl isocyanate derivative in that the viscosity de-
creases as the extent of substitution, and overall molecular
weight, increases (12). This effect is believed to be due to
either a decrease in hydrogen bonding interactions as the fraction
of hydroxyl groups decreases or to an increase in chain stiffness
due to the substitution of the bulky carbamate units for these
hydroxyl groups which would reduce chain entanglements and would
lower the viscosity of these solutions. Further studies are in
progress on this viscosity effect and will be reported in greater
detail in a future publication.

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In Vivo Studies on Drug—Polymer Sustained-Release Systems

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The pharmacodynamic performance was studied for three drug–polymer matrix systems and comparisons were made between the in vivo and the in vitro results. The systems studied were: (a) gentamicin in silicone rubber, (b) tetracycline in HEMA/MMA copolymers, and (c) niridazole in silicone rubber. In general, the in vitro models fall short of predicting the actual in vivo results accurately, even in those cases where the controlled release of the agent showed positive, desirable results in the test animals.

Sustained release of biological agents is a concept that has received increased attention over the past decade. The primary goal of sustained drug release is the prolonged delivery of a drug to a particular body compartment or anatomical target site. This goal is accomplished by the application of a therapeutic drug delivery system designed to control both temporal and spatial aspects of drug disposition.

Numerous drug–polymer delivery systems have been proposed with the expressed purpose of releasing a biologically active agent into the surrounding medium at a constant (zero-order) release rate. Attempts to achieve zero-order release rates have included infusion pumps (1), erodable matrices (2), controlled geometries (3), and various membrane–enclosed reservoir devices (4, 5, 6). Unfortunately, many of these systems fall far short of their goal, suffering from prolonged burst effects, device breakdown, and constraints in design, fabrication, or material properties. The result is usually an exponential release pattern with time and a relatively limited period of constant release.

Previous investigations in the field of controlled drug delivery have centered on the design and in vitro testing of the device. Animal experimentation has been largely limited to reports on the temporal pharmacodynamic performance of the drugs released from the implanted, inserted, or surface-applied polymer vehicles. Few studies have attempted to provide details of the kinetic

behavior of these release systems throughout the period of _in vivo_
evaluation. These data are essential to exploring the
pharmacokinetic characteristics of such systems and providing for
the development of predictive models. These models also increase
experimental efficiency by allowing the use of preliminary
measurements and analyses to optimize the further design of the
controlled drug-release systems, conserving time and animal
resources.

     We wish to review here our efforts which have been directed
toward evaluating the _in vivo_ release behavior of various
drug-polymer devices. We have used dogs, rats and mice in these
studies for reasons which will be explained. Our studies, which
are briefly summarized in Table 1, have had two major themes: the
investigation of drug release behavior in a disease model in an
animal, and the investigation of the pharmacokinetics of a
drug-polymer sustained release system in an animal model. Both of
these themes are pertinent to the evaluation of drug-polymer
sustained release systems intended for use in humans or animals.
We intend only a brief review here and refer the reader to the
references for experimental techniques, details and an in-depth
discussion of each system.

Table 1. Drug-Polymer Sustained Release Systems

General Theme or Disease Model	Studies	Animal	Drug	Polymer	References
Prosthetic Valve Endo- carditis	in vitro in vivo pharmaco- kinetics	dog	Genta- micin	Silicone Rubber	7-11
Antibiotic Delivery	in vitro in vivo pharmaco- kinetics	rat	Tetra- cycline	HEMA/MMA	12-15
Schisto- somiasis	in vivo	mouse	Niri- dazole	Silicone Rubber	16

Sustained Release of Gentamicin From Prosthetic Heart Valves

     Although prosthetic heart valves have been extensively and
successfully employed clinically, problems still exist during the
postoperative period concerning thromboembolism, infection and
valvular incompetence. Infectious prosthetic valve endocarditis
(PVE) remains as a serious and potentially devastating consequence
of valve replacement surgery with mechanically fabricated

prosthetic valves. Prophylactic antibiotic usage has reduced the
clinically confirmed incidence of PVE but early mortality due to
such infections remains high, 70% or greater as reported in several
studies.

     An earlier study has indicated the ability of parenteral
prophylactic antibiotic therapy to reduce the incidence of
prosthetic mitral valve failure in dogs associated with valve
related infections. As an alternative to the above approach, we
have investigated the therapeutic efficacy of a system providing
for a sustained release of gentamicin, incorporating the antibiotic
in silicone rubber which is in the sewing rim of a mitral valve
prosthesis implanted in dogs.

     In vitro studies with the uncoated gentamicin loaded valve rim
inserts revealed an initial burst of drug released which peaked on
the first day and slowly fell through day 10. At that time, a near
constant pattern ensued during which 2.5-7.0 mg/day were released
for a period of 2-3 wks followed by lower levels of drug release
for up to 2 mos. Valve rim inserts with the additional drug-free
silicone rubber layer (coated samples) exhibited a different
release pattern with a reduced burst effect and greater persistence
of long-term drug delivery. Peak release with the coated valve rim
insert was delayed to the third or fourth day and after 4 days mean
cumulative gentamicin release was only 59% of that of the uncoated
inserts.

     Serum measurements in dogs receiving the uncoated antibiotic
loaded valve rims containing the ^{3}H-gentamicin exhibited a large
early burst of gentamicin release, similar to that observed in the
in vitro experiments. Peak serum gentamicin levels occurred during
the cardiopulmonary bypass procedure. Serum drug concentrations
rapidly fell over the first 3 to 4 days following valve
implantation and remained at levels of approximately 0.2 μg/ml for
the next 2 to 3 wks. Serum gentamicin concentrations were
measurable for 1 to 2 mos before falling below detectable levels (<
0.1 μg/ml). The initial burst of gentamicin release was absent as
measured in the serum of dogs implanted with prosthetic valves
containing the coated rim inserts. The additional layer of
drug-free silicone over the drug loaded valve insert effectively
reduced peak serum gentamicin concentrations such that levels
remained relatively constant over the observable drug delivery
period.

     Survival data on dogs implanted with mitral valves containing
drug-free silicone rubber rim inserts (controls) and rim inserts
loaded with 40% gentamicin sulfate, by weight, both uncoated and
coated samples, are presented in Table II. A significant increase
in survival is readily apparent for dogs receiving gentamicin
loaded valves (uncoated and coated) compared to the animals
receiving the control valves with the drug-free silicone rubber
inserts. Long-term survival (dogs living 5 mos or greater) was 70%
for the combined uncoated and coated gentamicin loaded valve
implants, compared to only 31% in the control group. Three of the

dogs in the gentamicin loaded valve rim group were alive 22 mos or
more postimplantation.

Table II. Survival Data on Dogs Receiving Control and Gentamicin
                 Loaded Prosthetic Mitral Valves

Group (Total # Animals)	Died Before 1 mo.	Alive After 1 mo.	Percent Survival After 1 mo.
Control Valves 9	6	3	33%
Uncoated Valves 24	4	20	83% (p<0.05)
Coated Valves 13	3	10	77% (p<0.10)

    To investigate the in vivo efficacy of sustained-release
preparations a comprehensive pharmacokinetic model is necessary.
The two basic processes depicted within a given analysis are the
primary diffusive transport of the active agent through the polymer
delivery vehicle into the biological receiving medium and the
subsequent transport, distribution and elimination of the drug.
The latter process has been detailed in numerous pharmacokinetic
models relating the observations on absorption, distribution, and
elimination of orally or parenterally administered conventional
pharmaceutical agents. The former process is unique to the topic
of sustained drug release and is related to the physical and
chemical characteristics of the drug and its polymer delivery
vehicle and the biological response elicited by the implanted
system in situ.
    To couple the in vitro release rate behavior of the gentamicin
releasing prosthetic valve system with observations on
serum-gentamicin levels after implantation of the valve in dogs, an
open one-compartment pharmacokinetic model was added to the release
rate equations describing in vitro drug delivery. Gentamicin is
not metabolized, and its distribution kinetics can be modeled
adequately (within a 15% error) by a one-compartmental system.
Elimination kinetics and the apparent volume of distribution in the
individual dogs were derived from the gentamicin clearance studies.
    Attempts were made to fit the in vivo serum concentration
results to pharmacokinetic models based on revised diffusion-model
equations for the uncoated and coated valve-rim inserts. To couple
the in vitro diffusion equations to an in vivo simulation, a
classical one-compartment model of drug distribution was employed.
The mean apparent volume of distribution, V_D = 5.63 liters, and the
first order eliminating constant, K_e = 0.695/hr, for gentamicin
were determined in the intravenous injection-clearance studies.

Diffusion-model constants were recalculated using the results from in vitro flow-cell studies with suture punctured, Dacron mesh enclosed, valve-rim inserts. In general, all the model results fall short of predicting the actual serum-gentamicin concentrations determined experimentally over the observed valve implantation period.

From the results obtained in the attempts to model the in vitro release of gentamicin from the uncoated and coated valve rim inserts, it would appear that the actual drug-release process is more complex than predicted by relatively simplistic numerical analyses. For the uncoated rim insert system, computer fitting of the data was good over the early period of release but somewhat less adequate during the later time when a constant diffusion coefficient throughout the analysis was employed. The same trend was observed in the analysis of the coated rim insert system under identical conditions.

Release of gentamicin in vivo is an even more complex process. After implantation of the valve in the dog, the rate of release of the drug from the rim-insert surface would be dependent on such factors as: the relative amounts of surface exposed to the blood and the tissue, blood perfusion of the tissue, intracellular sequestration of the drug (both at the valve site and at peripheral organs), and the biological response of the blood and tissue layers adjacent to the valve surface. The fibrin pseudo-intima which forms early over the blood-contacting surface of the valve sewing ring might offer an additional diffusional resistance to the drug delivery device. At the tissue interface, fibrous capsule formation and the pooling of fluids within the surgically damaged tissue, or either one, may produce similar effects, modifying the release process.

In the later serum-gentamicin determinations, the elevation of measured serum-drug concentration over levels predicted by the pharmacokinetic model is most likely due to the inability of a one-compartment distribution model to properly describe the observed tissue sequestration of gentamicin and the later return of the drug from the tissues to the vascular compartment. This process would delay total drug excretion and cause an elevation of the later-time measurements of serum gentamicin concentration. Over the low-serum concentration range observed in the experiments after three to four days of valve implantation, the effects of tissue drug depots on serum gentamicin concentration would become significant.

Future pharmacokinetic models dealing with sustained drug release must take these effects into account through the use of time varying elimination parameters and multi-compartmental simulations. To this end, investigations are being conducted to better measure the tissue handling of gentamicin delivered from intramuscular sustained-release devices with different initial drug loadings and the effect of the cardiopulmonary-bypass procedure on drug-elimination characteristics.

In our bacteriology studies, it is apparent that parenteral antibiotic does provide some protection against the development of PVE when these results are compared to an earlier experiment which used prophylactic systemic antibiotic for a 10 day period following valve replacement. In addition, it appears that the gentamicin loaded valve inserts provide protection against the development of PVE. To better appreciate the pathogenesis of early-onset PVE and the therapeutic efficacy of the gentamicin releasing prosthetic valve, we are developing an experimental protocol which utilizes Staphylococcus aureus bacteremias and direct inoculation in the periprosthetic tissue.

These studies are important in that they demonstrate the absorption of gentamicin into the local tissue environment, periprosthetic cardiac tissue and sewing rim thrombi, following release from the device. In addition, they provide some indication as to the role that the local tissue environment plays in determining the serum and organ levels of gentamicin. The role that the healing phenomena in prosthetic valves may play in deter-mining drug levels in adjacent tissues and the availability of the drug to the systemic circulation is important. We have carried out experiments using a similar gentamicin-silicone rubber release system in dog skeletal muscle and have demonstrated the temporal and spatial variations of gentamicin levels in muscle tissue surrounding rod implants. These studies provide corroborative support for the effect of the fibrous capsule on retarding the permeation and sequestering of gentamicin in the preprosthetic tissue.

Sustained Release of Tetracycline From HEMA/MMA Copolymers

Previous experience with a hydrogel matrix-drug delivery system led to the investigation of a membrane-controlled drug delivery device in which a hydrophilic polymer core is supersaturated with a given pharmaceutical. The core-drug mixture is coated with a drug-free polymer layer which, by its hydrophobic nature (relative to the core material), functions as the rate-limiting membrane in drug diffusion from the device. Membrane-enclosed drug delivery systems maintain a constant activity source of the permeating drug at the interface between the core and the membrane-coating material. The result is a constant drug release rate that is proportional to the concentration gradient established over the diffusion rate-limiting outer membrane.

A membrane-controlled drug delivery device was developed to release tetracycline at zero-order rates. The tetracycline delivery vehicle is a trilaminate disk consisting of core and coating membranes fabricated from a series of 2-hydroxyethyl-methacrylate and methylmethacrylate copolymers. Appropriate adjustment of the monomer composition ratio imparts a hydrophobic nature to the copolymer outer coating membrane (relative to the

core material), which serves as the rate-limiting membrane in drug
diffusion. The trilaminate disks demonstrated a zero-order
tetracycline release over 4 months in vitro. The zero-order
release rate was a function of the general device geometry, coating
membrane thickness, disk surface area, level of core reservoir drug
loading, and membrane coating copolymer composition. Permeability
parameters of tetracycline diffusion through a series of
2-hydroxyethylmethacrylate–methylmethacrylate copolymer membranes
were determined by a flux-lag time method. Equilibrium hydration
values of these membranes also were determined. The ability of
trilaminate 2-hydroxyethylmethacrylate–methylmethacrylate devices
to release tetracycline at constant rates over a prolonged period
offers unique therapeutic and investigational possibilities.

The trilaminate tetracycline-releasing disks were evaluated in
vivo by implantation in the peritoneal cavities of female
Sprague-Dawley rats. The implanted disks were coated with
copolymer membranes of either 2:98 or 22:78 2-hydroxyethyl-
methacrylate–methylmethacrylate composition. In vivo release rates
of tetracycline were constant with time, as measured by the
cumulative drug release from disks recovered over different
implantation periods. Steady-state tetracycline release rates from
the trilaminate devices--158 µg/day for the 22:78 2-hydroxy-
ethylmethacrylate–methylmethacrylate copolymer-coated disks and 123
µg/day for the 2:98 2-hydroxyethylmethacrylate–methylmethacrylate-
coated disks--were in excellent correlation with the steady-state
total daily excretion of the drug (153 and 110 µg/day for the two
systems, respectively) by the rats. Plasma tetracycline con-
centrations reached steady-state levels within 2–3 days of
trilaminate disk implantation and remained constant over the 14-day
implantation period. Steady-state tissue tetracycline concen-
trations averaged 10–12 µg/g in the bone; 3–4 µg/g in the liver,
kidney and GI tract; 2–3 µg/g in the muscle; and <1 µg/g in the fat
tissue. A flow-limited pharmacokinetic model was constructed to
simulate in vivo tetracycline delivery from the trilaminate disks
and the subsequent drug distribution. Model predictions of tissue
and plasma tetracycline concentrations based on the experimentally
determined zero-order release rates were in good agreement with
experimental measurements.

The ability of trilaminate controlled-release disks fabricated
from 2-hydroxyethylmethacrylate–methylmethacrylate copolymers to
deliver tetracycline at a constant release rate when implanted in
rats has been demonstrated. A full range of in vivo zero-order
release kinetics is attainable through variations in the copolymer
composition of the diffusion-controlling coating membrane of the
trilaminate disk. The ability of such a device to deliver a
pharmaceutical at a desired constant release rate over a long
period is extremely valuable both as a therapeutic and an
investigational tool. This system may be extended to the study of
other drugs where long-term constant delivery rates are desired. A
flow-limited pharmacokinetic model is presented that utilizes the

ability of the trilaminate controlled-release device to deliver
tetracycline at constant rates over a prolonged implantation
period. Predictions based on this model are consistent with tissue
tetracycline concentrations measured in experiments with these
implant devices. The physiological pharmacokinetic model can be
utilized to provide reliable predictions of tissue drug levels and
is useful in studies of differential organ toxicities due to
effects caused by the parent drug or its metabolites. The model
also contributes information valuable for the design of proper drug
dosing regimens and for the fabrication and placement of controlled
drug delivery systems designed to deliver drugs to selected
physiological target sites.

The coupling of the principles of controlled release from a
zero-order rate device and long-term physiological pharmacokinetic
modeling is a unique research concept and will be used in
investigational systems where such drug delivery characteristics
help elucidate physiological rate mechanisms.

A controlled tetracycline delivery device, consisting of a
membrane enclosed trilaminate disc fabricated from a series of
2-hydroxyethylmethacrylate–methylmethacrylate copolymers,
demonstrated the ability to deliver tetracycline at zero-order
rates _in vitro_ and _in vivo_ in rats and was applied to study the
pharmacokinetics of tetracycline in the pregnant rat. The
trilaminate discs containing tetracycline were implanted in
pregnant Sprague-Dawley rats on the eighth day of gestation. The
animals were sacrificed on day 19, 20, and 21 of gestation in order
to measure the distribution of the controlled release tetracycline
in the maternal, fetal, and placental tissues. Constant plasma
tetracycline levels were attained in both the maternal and fetal
circulations after 4 to 5 days postimplantation of the trilaminate
discs. Placental transfer of tetracycline appeared rapid and no
partitioning of the drug was observed between the maternal and
fetal plasma. Tetracycline levels did not differ significantly in
the maternal and fetal soft tissues (liver, kidney, G.I. tract,
muscle, placenta) as measured over the last 3 to 4 days of the
animal's gestational period. Highest tetracycline concentrations
were determined in the fetal bone samples. In addition, some
accumulation of the drug occurred in the amniotic fluid. A
flow-limited pharmacokinetic model was constructed to simulate the
distribution of tetracycline, delivered at a constant rate from the
trilaminate device, in the pregnant rats. Predictions of fetal
growth and maternal and fetal tissue tetracycline concentrations
were in good agreement with the experimental measurements. The
ability of these copolymer systems to deliver tetracycline at
zero-order rates over extended periods offers numerous potential
therapeutic and investigational applications, especially where such
drug delivery characteristics are beneficial to the elucidation of
physiological rate mechanisms, as in the pregnant animal.

Sustained Release of Niridazole From Silicone Rubber

Niridazole was incorporated into rubber implants to invest-
igate the potential of a sustained release of the drug in the
therapy of murine Schistosomiasis mansoni. The infected animals
(200 cercariae for 6 weeks) were randomly divided into three
groups: one group received silicone rubber implants containing 50%
by weight niridazole; a second group received blank silicone rubber
implants with no drug; and the third group received no implants.
Mortality 4 weeks later was in excess of 80% for animals with no
implants or with the blank silicone rubber implants. In contrast,
10% mortality was observed in the mice receiving the niridazole-
silicone rubber implants over a 10-week period. The worm burden in
the niridazole-silicone rubber implant group was reduced at 10
weeks postimplantation by 77%.

The dramatic differences in mortality and morbidity between
the niridazole-treated and control mice in our experiments are
probably the result of reduction of worm burden to levels which can
be tolerated by the animals. In addition, the low levels of
niridazole diffusing from the implants may contribute to
amelioration of disease by suppressing granuloma formation. As
delayed hypersensitivity reactions in the host have been shown to
play a key role in the immunopathogenesis of schistosomiasis, a low
level of niridazole may help to alleviate the consequences of
disease.

In this initial investigation, an intraperitoneal implantation
site was chosen for the purposes of delivering the drug to the
appropriate physiological target site, i.e., the mesenteric
circulation. Other polymer-drug systems are under investigation
for delivering niridazole and other antischistosomal drugs over a
range of release rates sufficient to accomplish the combined tasks
of reducing worm burdens and minimizing disease symptomatology.

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Polymeric Delivery Systems for Macromolecules

Approaches for Studying In Vivo Release Kinetics and Designing Constant Rate Systems

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Methods for quantitating in vivo release kinetics
and for designing constant-rate release systems
for macromolecules are described. These systems
are composed of ethylene-vinyl acetate copolymer
and the incorporated drug. Approaches for directly
quantitating in vivo release kinetics have been
established using the polysaccharide, inulin. In
vitro and in vivo release kinetics of inulin from
the copolymer system were nearly identical. An
approach for achieving zero-order release kinetics
was tested by constructing matrices in the form of
a hemisphere with all portions laminated with an
impermeable coating except for a cavity in the
center face. Zero-order release kinetics of bovine
serum albumin were observed for 60 days.

Over the past 20 years research has been conducted in 2 primary areas concerning the use of polymers to improve the ways drugs are delivered to the body. One way has been to attach the drug to a polymer in order to impart a different affinity or specificity to the drug. The second approach has been to incorporate the drug inside a polymeric matrix to cause it to be slowly released into the body. In this paper, two aspects of the second approach are reviewed for polymeric systems for delivering large molecular weight (M.W. > 600) drugs.

Despite increasing advances in the use of controlled release polymeric delivery systems (1-6), it has, in general, proved difficult to develop such vehicles for the long-term administration of macromolecular drugs. This is largely due to the fact that most biocompatible polymers such as silicone rubber are impermeable to molecules over 600 molecular weight (7). Recent studies in our laboratory have demonstrated, however, that solvent casting of a variety of polymeric materials (ethylene-vinyl acetate

copolymer, polyvinylalcohol, poly-2-hydroxyethyl-methacrylate in
the presence of powdered drug permits continuous release
of macromolecules for over 100 days (8). It appears that the
powders cause a series of interconnecting channels to be formed
in the polymeric matrix. These channels are large enough to
permit macromolecular diffusion but tortuous enough to cause a
slow and continuous release (9). We have also examined factors
affecting in vitro release kinetics (10) and have demonstrated
that the macromolecules released from the systems are biochem-
ically active in vitro and biologically active in vivo (11). The
polymers used have also been shown to be biocompatible in sensi-
tive animal tissues (12). In this paper we review two areas of
our most recent research that provide further information on
these unique drug delivery systems. These are (1) the development
of techniques for comparing in vivo and in vitro release kinetics;
and (2) the development of approaches for achieving zero-order
release kinetics.

COMPARISON OF IN VITRO AND IN VIVO RELEASE KINETICS

In earlier studies in which macromolecules were released
from polymers in vivo (11), the substances tested were proteins
or DNA which were metabolized. Thus, a direct comparison of in
vivo and in vitro release was impossible. We therefore chose the
polysaccharide inulin which has a molecular weight of 5200 daltons.
Inulin was chosen as a marker because it is not metabolized, not
excreted by the glomerulus, and is neither reabsorbed or secreted
by the kidney tubules. It is not bound by plasma proteins and is
not toxic (13). Complete recovery of ^{3}H-inulin (14) or C^{14}-inulin
(15) in urine has been observed from intravenous injection into
rats, rabbits, dogs and humans. Thus, inulin provides an
excellent marker for comparing in vitro and in vivo release
kinetics because inulin recovered in urine can be directly com-
pared to inulin collected in vitro.

Methods. Polymer slabs containing inulin were made as
follows: 1) ^{3}H-inulin was diluted with non-radioactive inulin to
a specific activity of 3.49 µCi/mg; 2) 586 mg of this inulin was
suspended in 15 ml of 5% ethylene-vinyl acetate copolymer in
methylene chloride; and 3) this suspension was cast in a square
7 cm x 7 cm x 0.5 cm glass mold, dried and cut into nine 1 cm x
1 cm squares as described previously (10).
     In the in vitro kinetic experiments, 4 polymer squares were
placed in 5 ml of Phosphate Buffered Saline (PBS) at pH 7.4 and 37°C.
Polymer slabs were transferred daily to fresh PBS and the amount
of ^{3}H-inulin released was assayed as follows: a 250 µl sample of
PBS was added to 5 ml of Biofluour scintillation cocktail and
0.5 ml of distilled water. The emulsion was vortexed and counted
in a scintillation counter.
     In the in vivo experiments, the remaining 5 squares were im-
planted subcutaneously in rats. The rats were housed in metabolic
cages and urine was collected daily. A 250 µl urine sample was

placed into 5 ml of Biofluour scintillation cocktail, vortexed
and counted as above. Quenching of urine was corrected for using
the channels ratio method (16).

    Results. Excellent correlation between in vitro and in vivo
release rates was observed (Figure 1). As in internal control,
the inulin polymer squares were removed from the rats at the end
of 450 hrs and rat urine analyzed 4.5 hrs later. As shown in
Figure 1, the recovery rate of ^{3}H-inulin dropped 300 fold. This
control confirmed that ^{3}H-inulin is rapidly cleared from the
blood stream and excreted into the urine. Thus inulin is an ex-
cellent marker for directly assessing release rates from the poly-
mer implant.

APPROACHES TO ACHIEVING ZERO-ORDER RELEASE KINETICS

    Another limitation of the systems which were fabricated in
the shape of a slab in earlier studies was that they released
macromolecules at decreasing rates as a function of time. How-
ever, in many cases, it is desirable to have constant release
kinetics so that the body will maintain a constant level of the
released drug. The reason for the decreases in release rates
observed in previous studies (10), may be explained by considering
a typical implant, in the shape of a slab, as a model (17). Soon
after implantation, the solid drug dissolves from the surface
layer of the implant and then diffuses out of the implant. Since
the drug doesn't have very far to travel to reach the surrounding
media, release is rapid. As time elapses, the surface layer
becomes depleted, so drug from deeper within the implant must
dissolve and diffuse to the surface. The drug now has a longer
distance to travel, so release rates decrease.
    We thought it would be possible to compensate for this
phenomenon if a polymer implant was designed in such a way that
an increased area of drug would be available for release as the
distance from the release surface increased. A variety of shapes
were analyzed from a theoretical standpoint (18); the best results
were obtained with a hemispheric device laminated with an imperme-
able coating, except for a small cavity in the center face (Figure
2). This can be envisioned as a small cantelope cut in half,
where the orange pulp of the melon is the drug-polymer mixture.
The melon half is then coated everywhere except where the seeds
were, so all the drug must be released through the small exposed
section. In this case although the drug must still travel in-
creased distances at a later point in release time, this new shape
compensates by providing more surface area of available drug as
time elapses. To test this hypothesis, procedures were recently
developed to construct hemispheric systems for the release of
macromolecules (19) and these are discussed below.

    Methods. The molds used for fabricating hemisphere shaped
systems for macromolecules were composed of glass and had hemis-
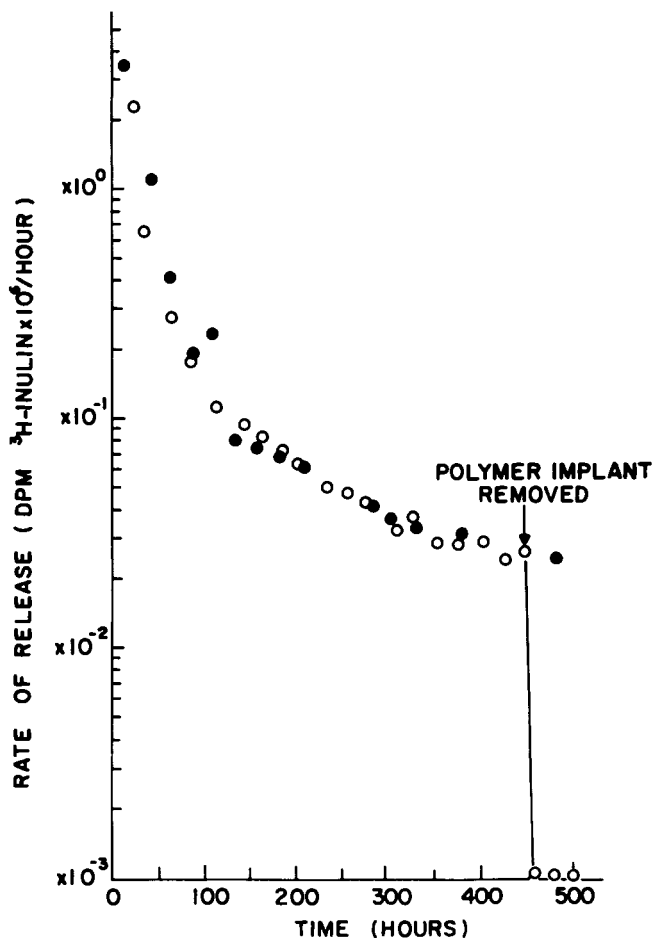
Figure 1. *In vivo vs. in vitro comparison of rates of release of ^{3}H-inulin from ethylene–vinyl acetate copolymer at 44% loading. Key: implants into 5 rats; in vivo, $\bigcirc$, N = 5 polymers; and in vitro, $\bullet$, N = 4 polymers.*
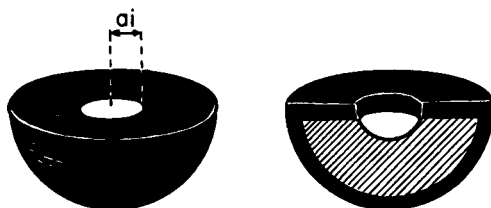
Figure 2. *Schematic of an inwardly releasing hemisphere (a_i = inner radius). The hatched lines represent the dispersed zone of drug in polymer, and the black represents laminated regions through which release cannot occur. Key: top view, t = 0, (left); and side view cross section, t = 0 (right).*

pheric bottoms (13 mm diameter x 11 mm height). These molds were
made from glass test tubes by cutting off the   bottoms. To
prevent these molds from falling over, an embedding platform was
made. This platform was constructed by pouring 15 ml of molten
paraplast into a bacterial petri dish, and then placing empty
glass molds into the paraplast so that they touched bottom.
After allowing the paraplast to harden, the molds were removed
and the indentations left behind in the paraplast could be used
to subsequently support the molds.

To make the hemisphere polymer systems, earlier methods of
preparing polymer-macromolecule slabs (10) were adapted. The
procedure is illustrated in Figure 3 and described in detail as
follows. First, blank glass molds were positioned in the inden-
tations in the paraplast embedding platform. Then, 6 ml of 20%
ethylene-vinyl acetate copolymer (EVA) solution made in methylene
chloride and 514 mg of bovine serum albumin (BSA) powder (particle
size range (10) from 150 to 180 μm) were mixed in a glass vial at
room temperature and vortexed for 1 minute to yield a uniform
suspension. Into each hemispheric glass mold, 0.8 ml of the
protein-polymer dispersion were pipetted with a Pipetman. The
petri dish containing these samples was then transferred onto a
block of dry-ice for 10 minutes. The samples gelled within one
minute. The petri dish containing these samples was then dried
first at -20°C for two days and then at 20°C for another two
days as reported previously (10).

The hemisphere pellets were twice coated with 20% EVA solution
(containing no macromolecules) to form an impermeable barrier.
The coating procedure is described as follows: (i) the tip of a
cylindrical metal stick (1.8 diameter x 30 mm in length) was
inserted into the center of the flat surface of each hemisphere
pellet to a depth of approximately 3 mm. (ii) Using the uninser-
ted portion of the stick as a handle, the hemisphere pellets were
placed directly on the surface of a block of dry ice for 10
minutes. (The round bottoms of the pellets were touching the
dry ice.) (iii) Again using the metal stick as a handle, the
cooled hemisphere was then immersed into 20% EVA solution at 20°C
for 10 seconds, removed, and then placed immediately on the same
dry ice for 10 minutes. (iv) A second layer of coating was done
in the same manner as described above. (v) The hemispheres
were then put in the freezer (-20°C) for two days followed by
further drying at 20°C for two days in a dessicator under a house-
line vacuum (600 millitorr) to remove residual solvent (10).
(vi) Finally, to create the exposed cavity in the face of the
hemisphere pellet, the metal stick was removed by gently encirc-
ling the polymer surface immediately surrounding the stick with
a scalpel-blade.

As controls, hemispheres of pure EVA were prepared uncoated
and completely coated (including the cavity). Similarly,
completely coated hemispheres containing BSA were prepared. All
of these controls were tested for release kinetics.

FABRICATION of HEMISPHERE-SHAPED
POLYMER MATRICES
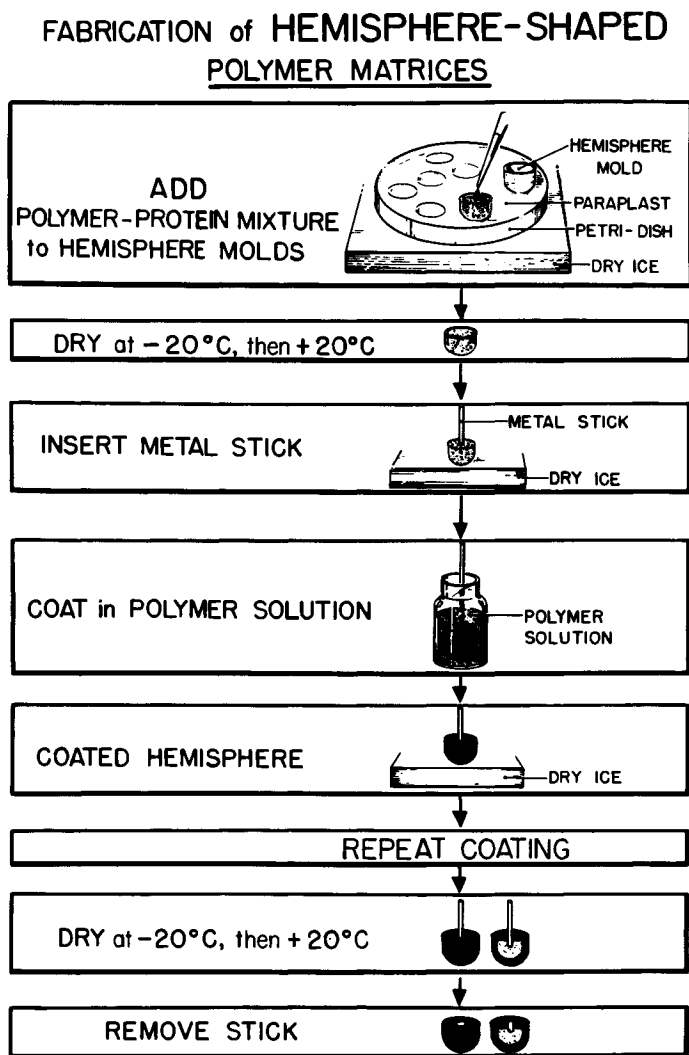
Figure 3. *Flow sheet for preparing hemisphere-shaped devices for the release of macromolecules. In the last two blocks of the flow sheet, both the top view and a cross-sectional view of the same hemisphere device are shown. (See text for detailed procedures.)*

The kinetics of BSA release from the hemisphere shaped systems were followed by methods described previously (10). BSA concentrations were measured spectrophotometrically by determining absorbance at 280 nm or 220 nm (19).

Results. Figure 4 shows the release kinetics of BSA from the hemisphere shaped systems. Each point represents 8 samples. A linear relationship between cumulative percentage release and the time of release was observed for 60 days. The release rate was approximately 0.5 mg of BSA per day. Standard error of the mean of cumulative release at each time point was within 12%. Control hemisphere shaped devices completely coated with 20% EVA solution, did not release any protein. The media collected in the release experiments of the control hemisphere shaped devices prepared from pure EVA solution (without BSA) showed no spectrophotometric absorbance.

DISCUSSION

The first set of experiments provides strong support for the assumption in earlier studies (8,9,10) that in vitro experiments would yield useful estimates for in vivo release kinetics. Further studies must still be conducted with other molecules and for longer times to fully validate the agreement between in vitro and in vivo release kinetics. Nevertheless, these studies demonstrate excellent agreement in the case studied here and establish a potentially important new technique for determining in vitro-in vivo release kinetics from polymeric delivery systems.
    The second set of studies demonstrate that hemispheric matrices can act as constant release systems for high molecular weight compounds. The fabrication procedures for these systems are relatively simple and can be performed without expensive apparatus.
    The release rate of macromolecules or proteins, such as BSA, from ethylene-vinyl acetate copolymer matrix slabs can be varied by as much as 2000-fold by manipulating three fabrication parameters, (i.e., drug loadings, aggregate sizes, and matrix coating) (10). The release rate of macromolecules from a hemisphere shaped device, can conceivably be changed by manipulating the same fabrication parameters and the size of the opening. In addition, we have previously demonstrated approaches for increasing the release rates of macromolecules from polymer slabs when desired by placing steel beads in the slabs and applying an external magnetic field   (20). It is conceivable that a similar approach could be employed with the hemisphere devices to yield a constant release system supplemented by occasional bursts when needed (e.g., insulin delivery in diabetes). The design of such a system is currently an ongoing project in our laboratory.
    Although the hemisphere design resulted in zero-order release kinetics for macromolecules such as BSA, the release mechanism for these systems has not yet been fully established. However,
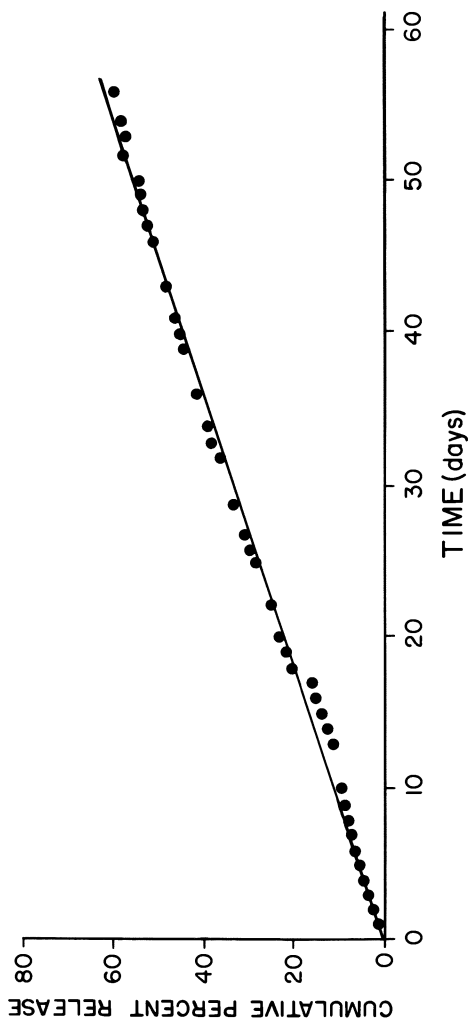
Figure 4.   Cumulative release of bovine serum albumin vs. time.   The matrix was made of ethyl-ene–vinyl acetate copolymer and bovine serum albumin. Standard error of the mean of the cumu-lative release at each time point was within 12%.

evidence continues to indicate that the release of macromolecules
from polymeric matrices occurs via diffusion through interconnec-
ting pores (9). The present study is consistent with such a
mechanism. However, a mathematical model that can be used to
predict release kinetics from macromolecular release systems has
not yet been developed.
  Although only BSA was used as a model macromolecule in the
second study, initial results have also indicated that zero-
order release rates were obtained for over one month for lysozyme
and β-lactoglobulin when they were incorporated into hemispheric
matrices. The hemisphere systems have not yet been tested in
vivo. However, future studies will explore the inulin model used
in the first section of this paper.
  The major problems encountered in the fabrication procedures
in the present study are the techniques of both coating the
matrix and opening the cavity uniformly. The development of a
technique for opening the hole on the flat face of the hemisphere
is very critical to achieve reproducibility. Besides the methods
practiced in these studies, improved techniques to create holes
on the flat face of the hemisphere shaped device should be ex-
plored. Techniques such as computer aided drilling may prove
useful in future studies.
  The delivery systems for macromolecules have been used in a
number of biological and potential medical applications. These
include:

● vehicles for bioassays of informational macromolecules (11);
● tools in biological studies of neovascularization (21);
● sources of chemical gradients in chemotactic studies (22);
● histochemical tools in brain research (23);
● insulin delivery systems (24);
● improved vehicles for immunization (25,26);
● interferon delivery systems (27); and
● delivery systems for anti-cancer agents (28).

  In summary, the studies reported in this review provide 2
important demonstrations (1) that in vitro release kinetics of
macromolecules such as inulin from ethylene-vinyl acetate copoly-
mer matrices are identical to their in vivo release kinetics,
and (2) that zero-order release for macromolecules can be achieved
for over 60 days using a hemisphere design. Further experimenta-
tion in these areas should provide information that will be useful
in the eventual design of controlled release systems for insulin
and other important bioactive macromolecules.

Acknowledgments

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Design of Polymeric Iron Chelators for Treating Iron Overload in Cooley's Anemia

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Iron chelating polymers based on hydroxamic acid and catechol groups have been synthesized for use in iron chelation therapy for treating iron overload in Cooley's Anemia. A mouse screen for in vivo activity of iron chelators has shown that poly(N-methacryloyl-β-alanine hydroxamic acid) (P-11) is effective and about equal to the standard drug desferrioxamine. The 11-atom spacing between hydroxamic acid groups is about optimum for intramolecular iron chelation and good solubility of the iron complex. A similar polymer, poly(N-methacryloylhydroxamic acid) (P-3), which has only 3 atoms between hydroxamic acids, is also active, but also rather toxic, a result that can be attributed to the tendency of P-3 to form insoluble intermolecular crosslinked complexes. A polymer bearing catechol groups is weekly active in removing iron in the mouse screen. Improvement may be possible with better control of the spacing of the catechol groups.

Cooley's Anemia is a genetic disorder relatively rare in the United States, but found more often among peoples of the Mediterranean area and within a band extending through India and southeast Asia (1). In addition to severe anemia, the disease causes bone malformation and general deterioration and distruction of the vital organs leading eventually to death.

The disease is characterized by an inability of the body to synthesize adequate amounts of the β chain of hemoglobin. The usual and really the only treatment is to administer blood transfusions on a regular basis throughout the lifetime of the individual. However, such a continual whole blood transfusion raises the iron level to the point where iron deposits form in the

liver, spleen, bone, heart, and other organs. Heart failure in
the teens and early 20's is usually the final outcome (2).

In order to retard or prevent this accumulation of iron, a
powerful iron chelator, desferrioxamine-B, is administered either
by injection or by intravenous infusion. This chelator is capa-
ble of forming a very stable complex with iron and then carrying
the iron out of the body with the urine or stool. Iron chelation
therapy greatly improves the well-being of the patient and
extends the lifetime of the patient appreciably (3).

Desferrioxamine-B (DFB) (I) is produced in a fermenta-
tion process by Ciba Pharmaceutical Co. and is supplied as the
methane sulfonate salt under the name Desferal. Although there
are many excellent iron chelators known, DFB is the only one that
not only works in humans, but is available in sufficient amounts
to supply the need for iron chelation therapy.

The iron chelating ability of DFB is due to the hydroxamic
acid, a functional group that possesses a natural affinity for
iron(III). Also inherent in this structure is a powerful chelate
effect due to the 9 atom spacing between hydroxamic acid groups,
which permits three neighboring hydroxamic acids to fit the
octahedral coordination sphere of the iron(III) without severe
steric strain.

The chelate effect of DFB causes the formation constant to
be greater than that of the monomeric acetohydroxamic acid by two
orders of magnitude, Table 1. Other naturally occurring iron
chelators, called siderochromes, include various modifications of
DFB, ferrichrome (II), rhodotorulic acid (III) and enter-
obactin (IV), all of which possess high formation constants
for iron, due in part to powerful chelate effects, Table 1.

Table 1.
Formation Constants of Iron Chelators

Chelator	log K	Reference
Acetohydroxamic Acid	28.2	(4)
Desferrioxamine-B	30.6	(4)
Desferrioxamine-E	32.5	(4)
Desferrichrome	29.1	(4)
Rhodotorulic Acid (RA)	62.3[a]	(5)
Enterobactin	52	(6)

$^a Fe_2(RA)_3$

Although DFB therapy is effective in removing large quanti-
ties of iron rapidly, there are some drawbacks in its use. One
of these is the short plasma residence time, about 30 minutes,
which causes a significant reduction in the efficiency of DFB to
remove iron. To counteract this rapid plasma clearance, the
chelator is often administered by means of a portable pump that
is worn by the patient and which continuously administers a
controlled amount of the drug (7). On the other hand, the pump
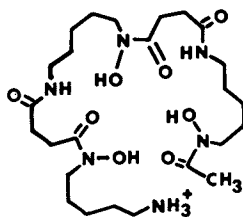
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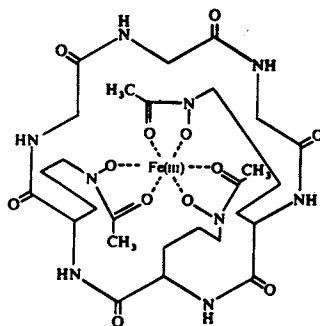
II
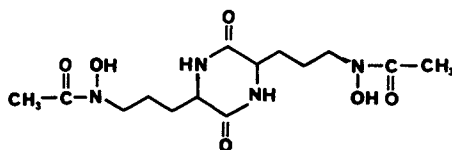
III
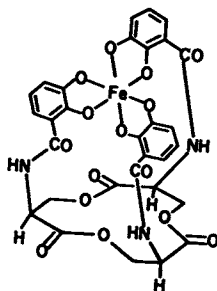
IV

is often psychologically unacceptable, and frequent injections throughout the day are painful and not very practical.

For these reasons a program designed to synthesize new iron chelators is being supported by the Cooley's Anemia Foundation of New York and the National Institute of Arthritis, Metabolism and Digestive Diseases of the National Institutes of Health. The object is to devise structures that not only remove iron from the body but also possess properties that provide a better mode of administration. A definite advantage would be achieved if the new drug could be given orally, the most convenient route. Also, if the plasma survival time could be extended, the efficiency of drug utilization would be increased, and thus the frequency of treatment and quantity of drug required would be reduced.

Advances have been made in the area of designing and synthesizing chemical structures for binding iron tightly and specifically, and possessing the appropriate solubility characteristics and chemical stability for drug use. On the other hand, there is no assurance that any such compound, no matter how carefully designed, will actually work in vivo. Although the compound must, of course, have a high affinity for iron, there is no simple relation between this property and the ability of the compound to remove excess iron from living systems (8). Adding to the complexity of the problem is the idea that there are several iron pools open to attack by an iron chelator, such as transferrin, ferritin and iron in transit. However, knowledge as to which of these pools is available to an iron chelator such as DFB is not understood. Also, there is evidence that the rat and mouse screens, used to provide a preliminary evaluation of the drug, do not always provide a measure of the behavior of the drug in humans. For example, rhodotorulic acid, which appeared promising on the basis of animal tests, nevertheless produced painful reactions when used in humans (9). Clearly there is much to be learned about iron transport, the manner in which iron chelators really work, and the effect of structure on adverse physiological reactions.

Design of Iron Chelators

The design of new iron chelators has been directed largely toward mimicking the naturally occurring siderochromes such as desferrioxamine and enterobactin. Three functional groups are under serious investigation, hydroxamic acids, catechols, and phenols. These functional groups have been incorporated into a variety of structures in order to produce compounds having exceptionally high stability constants for iron, close to, or higher than that of desferrioxamine. Unfortunately, a high iron binding constant does not assure high activity in vivo (8), and thus many compounds will have to be prepared and screened biologically in order to find a satisfactory balance of chemical and physiological properties.

Several powerful iron chelators, some recently synthesized specifically for the Cooley's Anemia program, are being tested to ascertain their potential as drugs for use in iron chelation therapy for treating iron overload in Cooley's Anemia.   Some of the more promising ones include:

Ethylenediamine-N,N'-bis(2-hydroxyphenylacetic acid) (EDHPA) (V) (8)

Derivatives of N,N'-bis(2-hydroxybenzyl)ethylenediamine-N,-N'diacetic acid (HBED) (VI) (8),

1,3,5 -N,N',N"-tris(2,3-dihydroxybenzoyl)triaminomethylbenzene (VII) (10,11),

1,5,9-N,N'N"-tris(2,3-dihydroxybenzoyl)triazacyclotridecane (VIII) (10)

poly(N-methacryloyl-β-alaninehydroxamic acid) (P-11) (IX) (12).

All of these compounds have exceptionally high binding constants for iron.

Polymeric Iron Chelators

The structure of the hydroxamic acid polymer P-11 was designed to maximize the formation constant for iron chelation. The formation constants of a series of such polymers is directly related to the magnitude of the chelate effect, which is, in turn, a function of the side chain length, Table II.   A maximum chelate effect appears at a spacing of 11 atoms.   A shorter spacing, 9 atoms, (P-9) is insufficient to permit three neighboring hydroxamic acids to fit about a single iron without strain, hence the lower value of log K.   With increasing side chain length, P-13 and P-15, the chelate effect decreases with the increasing degrees of freedom and a grandual decrease in log K results (12).

Table II.
Formation Constants of Iron Complexes of
Hydroxamic Acid Polymers (12)

Polymer[a]	n[b]	log K_{Fe}
P-9	1	28.6
P-11	2	29.7
P-13	3	29.3
P-15	-[c]	29.0

[a]Code number indicates number of atoms
     separating hydroxamic acid groups.
[b]Numbers refer to structure IX.
[c]Spacer is glycylglycine.
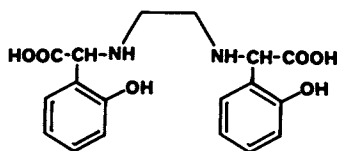
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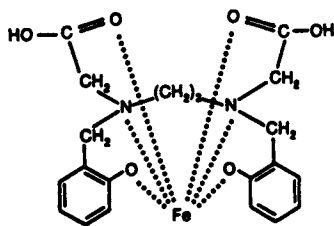
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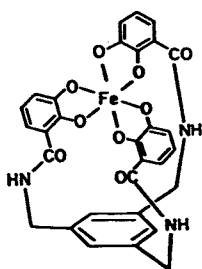
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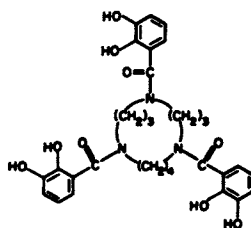
VIII
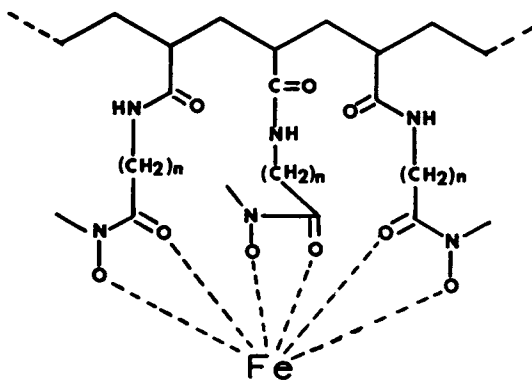
IX

Another hydroxamic acid polymer, poly(N-methylacryloyl-hydroxamic acid) (P-3) (**X**) in which the hydroxamic acid units are separated by only three atoms was similarly prepared. Because of the short side chains and closely spaced hydroxamic acid groups, P-3 tends to cross link and precipitate on complexation with iron. In cases such as this, the groups are too closely spaced to give soluble intramolecular complexes with iron.
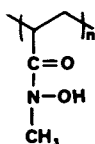
X

Hydroxamic Acid Polymer P-11 has been prepared in two molecular weight varieties, $[\eta]_{DMF-H_2O}$ 2.24 (HP-11) and 0.87 dl/g (P-11). The formation constant reported in Table II is for the lower molecular weight sample. Although the formation constant of the higher molecular weight material has not been determined, it is expected that there would not be much difference between the two.

A polymer (P-DHB) (**XI**) based on catechol, the active functional group of enterobactin, was recently synthesized by the reaction of polyvinyl amine with the ethyl ester of 2,3-dihydroxybenzoic acid (DHB). Only about one third of the amine groups was found to be substituted with DHB units. The formation constant of the iron(III) complex (log K = 40) is the same as that reported for the simple dimethyl amide of DHB and so there does not appear to be any appreciable chelate effect.
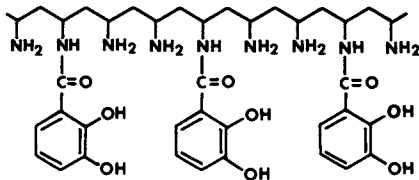
XI

Bioassay of Polymeric Iron Chelators

The iron chelator screen consists essentially of treating iron overloaded mice with the iron chelator by daily I.P. injections over a period of 7 days. Urine and stools are collected daily. At the end of the week, the mice are sacrificed and the livers and spleens are collected. The samples are then reduced to a suitable form and analyzed for iron by atomic absorption. An active chelator should produce increased iron levels in the urine and stool and decreased iron in the liver and spleen. Polymers P-11 (both molecular weight forms) P-3 and P-DHB were subjected to the mouse screen. The results are shown in Table III and compared with the drug DFB, which is used as a standard for comparison.

Overall, P-11 and HP-11 are judged to be about equal in potency to the standard DFB and except for lower fecal output occasionally observed at higher dose levels, both appear to be completely non-toxic. In the case of P-3, it is suggested that the extreme toxicity and high mortality is due to the tendency of P-3 to cross-link and precipitate on iron chelation. Such an event occurring in the blood stream would have obvious deleterious effects. Since the structural features of P-3 are similar to P-11 it is difficult to account for the toxicity on any basis other than the spacing of the groups. Regardless of the exact cause of the observed toxicity, it is clear that changes in spacing of the chelating groups profoundly affect both chemical and physiological activity.

P-DHB shows some weak activity in the mouse screen and gives some hope to the idea that a better placement of the DHB groups may result in a more positive biological outlook.

Potential Advantage of Polymeric Iron Chelators

Although the synthetic work on polymeric iron chelators has been directed primarily toward achieving strong chelate effects by appropriate spacing of functional groups, there is another means by which polymers might be of some distinct advantage over smaller molecules. And that concerns the residence time of the chelator in the plasma. As mentioned above, DFB is almost completely cleared from the plasma in 30 minutes, an effect which lowers the efficiency of DFB considerably. Although we do not know the direct cause of the rapid clearance of DFB, two possibilities have been suggested. Either the uncomplexed form is being degraded by metabolic processes, or the chelator is simply being lost by normal diffusion out through the membranes. Although degradation is a likely path, nevertheless, there is evidence to show that the iron complex of DFB is eliminated intact. If the loss of DFB is by simple diffusion, a high molecular weight analog of DFB, polymer P-11 for example, might diffuse out less rapidly, increasing the active lifetime in the

Table III.

Results of Mouse Bioassay of Polymeric Iron Chelators[a]

Treatment		Iron Content, % change			
Drug	Dose mg/kg	Spleen	Liver	Feces	Urine
		– – – – First Series – – – – –			
None[b]	–	[+190]	[+206]	[-10]	[+33]
DFB	125	0	-15	+28	+181
	250	+3	-25	+76	+238
	500	+7	-35	+81	+450
P-11[c]	100	-17	-24	+44	+138
	500	-41	-49	-3	+331
		– – – – – Second Series – – – – –			
None[b]	–	[+594]	[+347]	[+9]	[+30]
DFB	250	-5	-22	+10	+498
HP-11[d]	500	-35	-12	-12	+631
P-3[e]	375	-46	-6	-80	+73
P-DHB[f]					

[a]Percent changes are based on the values of transfused control mice.

[b]Values in brackets compare tranfused controls with nontransfused controls.

[c]Decreased fecal output was observed at the 500 mg/kg dose level, otherwise no toxic signs were observed.

[d]No toxic signs observed.

[e]Severe toxic reactions, 6/10 mice died during the 7 day treatment.

[f]Weakly active as indicated by a 60% elevation of iron in the urine compared to control.

plasma and thus increasing the efficiency of iron removal by the hydroxamic acid complexes.

A dynamic factor involving changes in chain conformation on complexation might be involved. The uncomplexed polymer is highly solvated and would exist in an expanded state as a random coil. On complexation with iron, the polymer shrinks to occupy a smaller volume. When we consider that passage through membranes depends not only on the molecular weight, but also on the molecular volume, it is not unreasonable to suppose that a polymer of some select molecular weight might easily be passed by a membrane while in the complexed state, but be retarded or blocked by the membrane when in the form of a larger uncomplexed random coil. Such a polymer would circulate in the plasma until sufficiently complexed with iron to be small enough to diffuse out through the membranes.

Evidence that such polymers are capable of this dynamic behavior are the following observations. When an aqueous solution of a hydroxamic acid-acrylamide copolymer (containing 1-2 mole percent hydroxamic acid units) is treated with iron, the viscosity of the solution changes. At high polymer concentrations the viscosity increases dramatically, indicating an increase in molecular weight through the formation of inter-molecular crosslinks involving iron-hydroxamic acid complexes. On the other hand, at low polymer concentrations, addition of iron causes a decrease in viscosity, clearly indicating a reduction in the molecular volume of the copolymer caused by intramolecular crosslink (13).

Iron complexation also affects the passage of the polymer through membranes. A low molecular weight variety of P-11 was dissolved in water and divided into 4 portions. Two of the portions were treated with standardized iron solution in an amount sufficient to form the 3:1 HA:Fe complex, and then placed in Spectrapor membrane tubes, one of 6000 and one of 12,000 mol. wt. cut off. The two other portions were diluted to the same extent with distilled water and similarly placed in two membrane tubes. After 24 hours the water surrounding the iron complexed samples had become red-brown, indicating passage of the polymer complex through the membrane. Samples of the water surrounding the membrane tubes containing uncomplexed polymers were treated with a few drops of the iron solution to form the colored complex of any polymer that had passed through the membrane. Some color was produced, but the color was not as intense as that from the iron complexed pair. Thus, the rate of diffusion of the polymer-iron complex through the membrane was greater than the rate of diffusion of the pure uncomplexed polymer. In principle, these results support the idea that a polymeric iron chelator could be designed to possess a long plasma residence time to collect iron and a relatively short residence time for the complex itself. On the other hand, since the actual processes and mechanisms of iron chelation therapy are largely unknown, whether or not such a drug

really could be made to work in this fashion is difficult to say. But clearly further work is needed to clarify these points.

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Structural and Functional Interrelationships of Anterior Pituitary Hormones

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The hormones of the anterior pituitary gland fall
into three groups on the basis of structural
similarities. Corticotropin (ACTH), the melano-
tropins (α-MSH and β-MSH), the lipotropins (β-LPH
and γ-LPH) and endorphin (β-EP) comprise the first
group of closely related flexible polypeptides.
All the hormones of this group are derived from a
single macromolecular precursor named proopiomelano-
cortin. Growth hormone (GH) and prolactin (PRL)
are compact globular proteins with characteristic
three-dimensional structure. These two hormones
exhibit considerable sequence homology. The gonado-
tropins, lutropin (LH) and follitropin (FSH), as
well as thyrotropin (TSH) are closely related
glycoproteins forming the third group. These hormones
are composed of two nonidentical subunits of compar-
able size. The α-subunit is common to the three
glycoprotein hormones. Structure-function
relationships of the pituitary hormones have been
studied by the synthesis of analogs and by selective
chemical modification.

The hormones of the anterior pituitary gland fall into three
groups on the basis of structural similarities (1). The structural
features of these hormones include all levels of organization of
protein structure. Corticotropin (ACTH), the melanotropins
(α- and β-MSH), lipotropin (LPH) and endorphin (EP) comprise the
first group of flexible polypeptides in which information is
organized in the linear sequence of amino acid residues. Growth
hormone (GH) and prolactin (PRL) exhibit considerable structural
homology and are compact globular proteins possessing character-
istic three dimensional structure. The unique feature of the
hormones of the third group is their quaternary structure. The
three members of this group, namely, lutropin (LH), follitropin
(FSH) and thyrotropin (TSH) are closely related glycoproteins

composed of two non-identical subunits of comparable size. In the
case of the hormones of groups 2 and 3, information is contained
in the distinct conformation of the protein. Disruption of the
three dimensional structure results in loss of biological activity.

Peptide Hormones

Structure-function relationships of the hormones in the first
group have been investigated in detail by assessing the biological
activities of numerous analogs of these hormones which have been
synthesized. The rapid progress in the field of peptide synthesis
in the late fifties and early sixties made it feasible to
synthesize peptides ranging in size from 10 to 40 residues. The
introduction of several new protecting groups and improvements in
separation procedures have vastly increased the utility of solid-
phase techniques for peptide synthesis. As a result, the total
synthesis of all the hormones of the first group have been
accomplished (2-6).

ACTH and MSH

ACTH is a single chain polypeptide composed of 39 amino acid
residues (Figure 1). The amino acid sequence of α-MSH is identi-
cal to the sequence of the first 13 residues in ACTH. In α-MSH
the amino and carboxyl terminals are blocked with an acetyl and
amide group, respectively (Figure 1). The heptapeptide sequence
Met-Glu-His-Phe-Arg-Trp-Gly present in ACTH and α-MSH is also
contained in the β-MSH molecules isolated from the pituitary
glands of several species (7). The primary physiological function
of ACTH is the acute regulation of glucocorticoid production by
the adrenal cortex and the long term maintenance of the adrenal
cortex. The major recognized function of the melanotropins is
their ability to stimulate the dispersion of melanin in amphibian
melanophores. ACTH is also able to induce melanin dispersion.
In addition, both ACTH and MSH stimulate lipolysis in rabbit
adipose tissue. That the melanophore stimulating activity
associated with purified ACTH is intrinsic to the hormone and not
due to contamination with the melanotropins, was conclusively
demonstrated by the synthesis of a nonadecapeptide corresponding
to the first 19 residues of ACTH (8). This synthetic peptide
stimulated steroidogenesis in the adrenal gland and also elicited
melanin dispersion.
Studies with synthetic analogs of ACTH and MSH provided an
understanding of the manner in which information is organized
along the linear sequence of amino acid residues (9, 10, 11).
Inspite of the high degree of structural similarity between ACTH
and α-MSH, the biological peoperties of the two hormones are
highly specific. α-MSH has less than 0.04% of the potency of
ACTH in stimulating glucocorticoid production in the adrenal
gland. Conversely, ACTH is only 1% as potent as α-MSH in

α-MSH: Ac-Ser-Tyr-Ser-Met-Glu-His-Phe-Arg-Trp-Gly-Lys-Pro-Val-NH$_2$
       1   5   10   13

β-MSH: Asp-Glu-Gly-Pro-Tyr-Lys-Met-Glu-His-Phe-Arg-Trp-Gly-
          1   5   10

          Ser-Pro-Pro-Lys-Asp
               15   18

ACTH: Ser-Tyr-Ser-Met-Glu-His-Phe-Arg-Trp-Gly-Lys-Pro-Val-
          1   5   10

          Gly-Lys-Lys-Arg-Arg-Pro-Val-Lys-Val-Tyr-Pro-Asn-Gly-
               15   20   25

          Ala-Glu-Asp-Glu-Ser-Ala-Glu-Ala-Phe-Pro-Leu-Glu-Phe
                      30   35   39

Figure 1.   Structure of corticotropin and the melanotropins.

eliciting melanin dispersion in the melanophores of nonmammalian animals. The minimal structural features required for eliciting either response appear to be the same. The pentapeptide sequence His-Phe-Arg-Trp-Gly has been shown to stimulate both adreno-cortical cells (12) and melanophores (13) to the same maximal degree as the intact hormones, although nearly million-fold higher concentration of the pentapeptide was required. It appears that the same pentapeptide segment of both ACTH and α-MSH is involved in functional interaction of the hormones with their respective target cells. Specificity of interaction is conferred by the amino acid residues attached to either side of the pentapeptide segment. The decapeptide ACTH[1-10] which includes the His-Phe-Arg-Trp-Gly segment is fully active as a steroidogenic agent at high concentrations whereas ACTH[11-19NH$_2$] is inactive even at very high concentrations. The nonadecapeptide ACTH[1-19NH$_2$] is equipotent to the intact hormone. These results imply that link-ing the functional ACTH[1-10] peptide to the ACTH[11-19NH$_2$] sequence serves to lower the concentration of the peptide required for full activity by several orders of magnitude. The linear sequence corresponding to residues 11-19 of ACTH serves as an attachment site. The interaction of this region of the hormone with the molecular surface of the receptor increases the local concentration of the functional segment of the peptide in the microenvironment of the receptor. α-MSH which lacks the strongly basic amino acid sequence Lys-Lys-Arg-Arg (residues 15-18 of ACTH) is unable to attach efficiently to the adrenocortical receptor at physiological concentrations and therefore does not stimulate the adrenocortical cell even though it contains the information necessary for activating the receptor. The structure of α-MSH is designed for productive interaction with the melanophore. Des-acetyl α-MSH is only 25% as potent as α-MSH. The lack of the acetyl group at the amino terminal and the presence of an extended peptide chain beyond residue 13, hinder the interaction of ACTH with melanophores and, therefore, at physiological concentrations, only α-MSH is able to activate melanin dispersion in the melano-phores.

Further studies have shown that although the same amino acid sequence is involved in the activation of receptors on adreno-cortical cells and melanophores, there are significant differences in the structural features of these receptors even in the region involved in interaction with the active pentapeptide sequence (14). Selective chemical modification of the single tryptophan residue in ACTH by reaction with o-nitrophenylsulfenyl chloride caused profound alterations in the biological properties of the hormone (15-19). The analog, o-nitrophenylsulfenyl-Trp[9]-ACTH (NPS-ACTH) was found to be inactive as a lipolytic agent on rat adipocytes (15) and only 1/100 as potent as ACTH in stimulating steroido-genesis in rat adrenocortical cells (18). NPS-ACTH was found to be a potent anagonist of ACTH-induced lipolysis (15) as well as adenylate cyclase stimulation in both rat adipocytes (17) and

rat adrenocortical cells (18, 19). On the other hand, NPS-ACTH was 3.4 times as potent as ACTH in stimulating amphibian melanophores (16) and 4.4 times as potent as ACTH in stimulating lipolysis in rabbit adipocytes (14). Similar selective modifications of the Trp residue in α-MSH converted the hormone into a more potent melanin dispersing agent (16) and lipolytic agent in rabbit adipocytes (14). These results suggest that the complementary region of the specific ACTH receptor involved in interacting with the His-Phe-Arg-Trp-Gly segment must be structurally different from the complementary region of the MSH receptor involved in the interaction with the same peptide segment. Furthermore, the receptors on rat adipocytes and rat adrenocortical cells appear to be very similar and highly specific for ACTH. The receptors on amphibian melanophores and rabbit adipocytes resemble each other closely and are specific for MSH. Comparison of the steroidogenic, lipolytic and melanophore stimulating activities of a number of analogs of ACTH and MSH confirmed the above findings (11).

LPH and EP

Lipotropin was discovered fortuitously during the investigation of a simplified isolation procedure for ACTH (20). Two peptides exhibiting lipolytic activity in rabbit adipose tissue were isolated from ovine pituitaries and were designated β- and γ-LPH. Ovine β-LPH (21) is a linear polypeptide composed of 91 amino acid residues (Figure 2). The 58-residue γ-LPH is identical to the sequence region 1-58 of β-LPH. Residues 41-58 in LPH correspond to the sequence of β-MSH. The biological activities of β- and γ-LPH are similar to those of β-MSH and can be accounted for by the presence of the β-MSH sequence within the LPH molecule. The physiological role of LPH has remained obscure. Recent evidence strongly suggests that the biological function of LPH may be that of a prohormone for the opioid peptide β-endorphin.

The discovery of endogenous opiate peptides has been one of the most exciting developments in the peptide hormone field. In December 1975, Hughes, Kosterlitz and collaborators (22) reported the isolation, structure and synthesis of two similar pentapeptides from porcine brain with potent opiate agonist activity. They recognized that the entire peptide sequence of one of these peptides, namely, methionine enkephalin, is present in residues 61-65 of the lipotropin molecule (Figure 2). In their search for LPH from camel pituitaries, Li and Chung (23) isolated a 31-amino acid residue peptide, which corresponded to residues 61-91 of LPH. This peptide was found to be more potent than the enkephalins in its analgesic properties as well as in its affinity for brain opiate receptor, and was named endorphin (EP). β-EP has been isolated from porcine (24), ovine (25), bovine (26), rat (27) and human (28) pituitaries. The structure of β-EP appears to be highly conserved. Both camel (29) and human (30) β-EP

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H-Glu-Leu-Thr-Gly-Glu-Arg-Leu-Glu-Gln-Ala-Arg-Gly-Pro-Glu-Ala-
                    5                       10                      15

  Gln-Ala-Glu-Ser-Ala-Ala-Ala-Arg-Ala-Glu-Leu-Glu-Tyr-Gly-Leu-
                   20                      25                      30

  Val-Ala-Glu-Ala-Glu-Ala-Ala-Glu-Lys-Lys-Asp-Ser-Gly-Pro-Tyr-
                   35                      40                      45

  Lys-Met-Glu-His-Phe-Arg-Trp-Gly-Ser-Pro-Pro-Lys-Asp-Lys-Arg-
                   50                      55                      60

  Tyr-Gly-Gly-Phe-Met-Thr-Ser-Glu-Lys-Ser-Gln-Thr-Pro-Leu-Val-
                   65                      70                      75

  Thr-Leu-Phe-Lys-Asn-Ala-Ile-Ile-Lys-Asn-Ala-His-Lys-Lys-Gly-
                   80                      85                      90

  Gln-OH
```

Figure 2.   Structure of LPH.   Endorphin corresponds to residues 61-91.

have been synthesized, in addition to a large number of analogs (31-35). Analysis of the structure-function relationships using the guinea pig ileum assay showed that the potency increases as the peptide chain length is increased from that of enkephalin to endorphin. The complete amino acid sequence of β-EP was found to be necessary for eliciting full analgesic activity in vivo. Blocking of the amino terminal of β-EP by acetylation results in loss of analgesic activity. This explains why β-LPH which contains the entire β-EP sequence has no analgesic activity. In addition to its analgesic activity, EP exhibits a remarkable array of biological properties modulating behaviour. Endorphin appears to be the most potent peptide in all biological tests. Structure-function studies using diverse biological assays may lead to the separation of these biological activities. Recent clinical studies indicate that synthetic EP exhibits beneficial effects in patients with severe pain, narcotic abstinence, schizophrenic behaviour and depression. Endorphin appears to interact with a variety of receptors in the brain. As a result of the enormous interest in brain peptides, it has become apparent that the other pituitary hormones such as ACTH and α-MSH are also present in discrete regions of the brain and that the hormones of this group may be involved in complex interactions in different areas of the brain, thus effecting modulation of behaviour.

Concurrent with the discovery of EP, it was recognized that all the hormones of the first group are formed biosynthetically by posttranslational processing of a high molecular weight precursor (36, 37, 38). The precursor, proopiomelanocortin, has a molecular weight of 31,000. The complete amino acid sequence of proopiomelanocortin has been deduced from the nucleotide sequence of the complimentary DNA (39). Lipotropin forms the carboxyl terminal and is preceded by the ACTH sequence. ACTH and LPH are separated by the paired basic residue sequence Lys-Arg. The amino terminal region also contains a sequence of residues common to both ACTH and LPH, namely His-Phe-Arg-Trp. In common with other protein precursors, the biologically active residues in proopiomelanocortin are flanked by paired basic residues at which proteolytic processing takes place.

Protein Hormones

The biological activities of the hormones of groups II and III are dependent on the unique conformations of these molecules. In this case, information is not confined to any linear segment of the polypeptide chain but is derived from the noncovalent interactions of distant segments of the polypeptide chain. Owing to the large size of these hormones structure-function studies have depended largely on selective chemical modification of one or more amino acid side chains. Studies with human growth hormone (hGH) provide a good illustration of this approach.

Growth Hormone (GH)

Human GH is a single chain polypeptide composed of 191 amino acid residues cross-linked by two disulfide bridges between cysteine residues at 58 and 165, and 182, and 189, respectively (Figure 3). Reduction of both disulfide bridges and alkylation with iodoacetamide yields the tetra-S-carbamidomethylated hGH which retains full biological activity in animals and humans, and its secondary and tertiary structure are indistinguishable from the structure of the intact hormone (40).

It was discovered recently that limited proteolytic digestion of hGH with human plasmin did not alter the biological properties of the hormone (41, 42). Li and Gráf (43) found that plasmin caused the excision of a hexapeptide corresponding to residues 135-140. The fragments 1-134 and 141-191 are held together by the disulfide bridge between cysteine 53 and 165 in plasmin-digested hGH. Reduction and alkylation of plasmin-digested hGH severs the covalent link between the two peptide fragments and they can then be separated from each other by gel filtration (43). Whereas plasmin-digested hGH is as potent as the intact hormone, the separated fragments have very little biological activity. However, recombination of the two fragments results in the formation of a noncovalent complex which is indistinguishable from the unmodified hormone in terms of biological activity as well as conformation as shown by circular dichroic spectra (44).

The dissociation of tetra-S-carbamidomethylated plasmin-digested hGH has been investigated and found to be a slow, exergonic equilibrium reaction (45). The major products of the dissociation have been shown to be the monomeric form of hGH(141-191) and a trimeric form of hGH(1-134). The carboxyl terminal fragment appears to be a random coil, whereas the amino terminal fragment, hGH(1-134) retains a considerable degree of secondary and tertiary structure. These results suggest that the noncovalent interaction between the fragments is essential for maintenance of the biologically active conformation of the hormone. Since hGH(1-134) by itself has about 14% activity and the activity of hGH(141-191) is negligible, it is likely that hGH(1-134) contains much of the information for favorable interaction with the GH receptor and hGH(141-191) probably stabilizes the biologically active conformation of hGH(1-134).

An important consequence of these findings is that it is now possible to investigate structure-function relationships of hGH through semi-synthesis. Li et al. (46) showed that combination of the synthetic peptide corresponding to hGH(141-191) with the natural hGH(1-134) resulted in the full restoration of biological activity and immunoreactivity. Synthetic analogs containing norluecine in place of methionine and alanine in place of cysteine in hGH(141-191) as well as in hGH(145-191) all complemented with natural hGH(1-134) to generate full growth promoting activity

H-Phe-Pro-Thr-Ile-Pro-Leu-Ser-Arg-Leu-Phe————————————
   1   5   10

————————————————————————————————————Cys————————————
                                                       53

——Asp-Gly-Ser-Pro-Arg-Thr-Gly-Gln-Ile-Phe-Lys-Gln———Cys——
     130   134   140   165

————Cys-Arg-Ser-Val-Glu-Gly-Ser-Cys-Gly-Phe-OH
       182   189
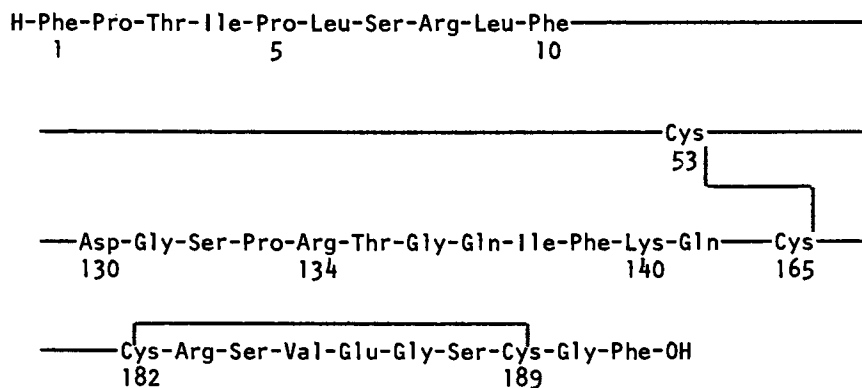
Figure 3.   Structure of human growth hormone.

(47, 48). The importance of various amino acid residues in the
fragment hGH(141-191) can be examined systematically by the
semisynthetic approach.

Lactogenic Hormones

The primary protein hormone of the pituitary gland involved
in regulation of lactation in mammals is prolactin (PRL). In
addition to this, in humans there are two other hormones with
lactogenic activity. Both hGH and human chorionic somatomammo-
tropin (hCS) have identical lactogenic activities in the standard
pigeon crop sac assay but their potencies are considerably lower
than that of PRL. In the rabbit and mouse mammary gland assays,
however, the lactogenic potencies of the three hormones are
comparable. Comparison of the amino acid sequences of ovine PRL,
hGH and hCS shows that there is considerable structural homology
(49). Among all the known growth hormones, hGH alone has both
growth-promoting and lactogenic activities.
PRL has also been selectively cleaved by limited proteolysis
with fibrinolysin (50). This results in the cleavage of the
peptide chain between residues 53 and 54. The two fragments
PRL(1-53) and PRL(54-199) are devoid of both lactogenic activity
and immunoreactivity. Recombination of the fragments resulted in
full restoration of immunoreactivity but not biological activity.
The physicochemical properties of the recombinant molecule were
very similar but not identical to those of the native hormone.
These results emphasize the enormous importance of the integrity
of the conformation of these hormones for their biological
functions.
Human CS, the placental hormone, is very similar to hGH in
its primary structure (96% homology) yet has very low growth-
promoting activity. Plasmin treatment of hCS results in the
removal of a heptapeptide from the hormone (51). Reduction and
alkylation of plasmin treated hCS gave two fragments corresponding
to residues 1-133 and 141-191 of hCS. The fragments were devoid
of any biological or immunochemical activity. Complementation
of the carboxyl terminal fragment, hCS(141-191) with hGH(1-134)
yielded a product with 50% growth promoting activity and nearly
full immunoreactivity (52). The other hybrid, made up of hCS-
(1-133) and hGH(141-191) had very low growth promoting activity
(<10%). The immunological and receptor-binding properties of
the two hybrids have been analyzed in detail (53). These studies
have shown that the determinants for binding hGH to hepatic GH
receptors as well as lactogenic receptors are present in the
hGH(1-134) fragment of the hormone.

Glycoprotein Hormones

The gonadotropins LH and FSH, as well as TSH are all composed
of two non-identical, non-covalently complexed, glycopeptide

subunits. One of the subunits, designated α, is common to all three hormones. The β subunit of a glycoprotein hormone, therefore, determines the hormonal specificity. Individual subunits normally have little or no intrinsic biological activity (54). There is extensive structural homology (40-45%) between the β subunits. The glycoprotein hormones are characterized by a carbohydrate content of 15-25% and a high cystine content. All α subunits possess five disulfide bridges and β subunits possess six. LH has a high proline content, FSH is rich in acidic amino acids and TSH has a high tyrosine content. The placental hormones, human chorionic gonadotropin (hCH) and equine chorionic gonadotropin (eCG or PMSG) are structurally very similar to the pituitary glycoprotein hormones.

Selective chemical modification and recombination of the α and β subunits of the glycoprotein hormones to give intra and interspecies hybrids are the two major experimental approaches employed in elucidating the structural and functional interrelationships of this group of hormones. The alteration of an average of one histidine residue out of six in ovine LH by Rose Bengal-sensitized photooxidation resulted in 90% loss of biological activity (55). Based on a variety of chemical modifications of LH including nitration of tyrosine residues and acylation of the ε-amino groups of lysine residues, Ward (56) has proposed a model for the location of various functional groups involved in intersubunit contact and in the interaction with the receptor. In the α subunit the tyrosine in position 92 or 93, or both are considered essential for biological activity. Ward concludes that the α subunit accounts for the majority of the receptor-site binding, but the β subunit specifies the α subunit conformation and possibly one-third of the receptor-site binding specificity (56).

The role of the carbohydrate moiety in the actions of the glycoprotein hormones has been investigated by specific removal of carbohydrate by enzymatic (57) and chemical (58) methods. Chemical deglycosylation of ovine LH by treatment with anhydrous HF for 75 min at 0° resulted in the removal of nearly two-thirds of the carbohydrate moiety (58). Deglycosylated LH (DG-LH) was completely different from the native hormone in its physicochemical and biological properties. DG-LH could bind to LH receptors but did not stimulate cAMP production. The steroidogenic response of DG-LH was significantly attenuated compared to LH. Furthermore, DG-LH was found to be an effective antagonist of LH both in vitro and in vivo. These results suggest that the carbohydrate groups of the glycoprotein hormones are important for biological activity and that alteration of the carbohydrate may lead to antagonists which may prove useful agents for contraception.

Acknowledgment

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Covalent Binding of Trypsin to Hydrogels

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Covalent binding of trypsin to hydrogels, i.e.,
derivatives of poly(2-hydroxyethyl methacrylate),
PHEMA has been effected. Reaction of p-nitro-
benzenesulfonyl isocyanate with PHEMA or copolymers
containing HEMA and varying amounts of styrene
followed by reduction of the nitro group with
sodium dithionite produced p-aminobenzenesulfonyl
carbamates of hydrogels (p-ABSC-HEMA). The precise
composition of the hydrogel carbamates was
determined by nonaqueous titration of the carbamate
moiety with tetrabutylammonium hydroxide in
isopropanol/dimethylformamide.

Treatment of p-ABSC-HEMA with glutaraldehyde
produced enzyme supports capable of binding up to
55 wt % trypsin. Incorporation of hydrophobic
styrene units into the support reduced the capacity
to 2-4 wt % but enhanced the specific activity of
the trypsin. The esterase activity of bound
trypsin, assayed with TAME, was found to range
from 11% to 45% of that exhibited by the free
trypsin. Active-site titration of a PHEMA-
trypsin conjugate with p-nitrophenyl-p'-
guanadinobenzoate HCl indicated the active species
to be 31% of the total amount of protein bound.
Thus, 45% of the active sites present in free
trypsin survived the immobilization process.

In recent years, a growing interest in hydrogels as solid
supports for enzyme, hormones and pharmacons has become evident.
Poly(hydroxyethyl) methacrylate), PHEMA, has been utilized to
entrap trypsin and glucose oxidase (1) but the radical systems

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employed deactivated a significant fraction of the enzymes.
Higher retention of enzymatic activity was observed when entrap-
ment was effected using low temperature, radiation induced poly-
merization of enzyme-monomer mixtures in a glassy state (2).
Covalent binding of trypsin to crosslinked acrylamide-HEMA co-
polymers has been achieved by activating the support with cyanogen
bromide; the immobilized enzyme exhibited 35% of the initial free
enzyme activity (3). Hydrogels containing PHEMA and varying other
components proved to be non-thrombogenic and stabilizing supports
for a number of enzymes (4). A combination of entrapment and
covalent binding has been particularly effective; copolymerization
of acrylamide and acryloxysuccinimide to an active prepolymer
followed by crosslinking with diamines in the presence of enzymes
produced active enzyme matrixes from more than sixty enzymes (5).
Copolymers of HEMA with p-acetaminophenylethoxy methacrylate or
p-nitrophenol esters of ω-carboxymethacrylamide have been
employed successfully to covalently bind enzymes (6).

　　　Although the cyanogen bromide technique for activating
hydroxyl containing polymeric supports has gained widespread
acceptance (6), problems with removal of residual cyanide ions (7)
has prompted a search for alternate activation techniques.
Recently, the application of 2-(3-aminophenyl)-1,3-dioxolane as a
synthon for diazotized or isothiocyanate functionalized polyvinyl
alcohol has been described (8). We have elected to use p-nitro-
benzenesulfonyl isocyanate as a synthon with general utility for
the introduction of active binding sites on hydroxyl containing
polymers. Using hydrogels, we have demonstrated the utility of
this binding agent by immobilizing trypsin.

Experimental

　　　Reagents. Hydroxyethyl methacrylate (HEMA) was extracted
with hexane to remove bis-esters and distilled in vacuo, b.p. 69°/
0.1 mm Hg. p-Nitrobenzenesulfonyl isocyanate (b.p. 160°/4 mm Hg)
was synthesized in 63% yield by phosgenation of p-nitrobenzene-
sulfonamide in the presence of butyl isocyanate (9). Homo- and
copolymers of HEMA were prepared by solution polymerization in
DMF in the presence of benzoyl peroxide and the results are
summarized in Table I. The hydrolytic stability of the hydrogels
was estimated by slurrying 0.5-1.0 g of polymer in 5.0 mL 0.35 N
KOH in 10 mL culture tubes equipped with teflon lined screw caps.
The samples were heated at 100° for up to 24 hr in an aluminum
constant temperature block. Immediately upon removal from the
heating block, the samples were cooled and acidified with 6N HCl.
The precipitated gel was washed and soaked in distilled water for
30 min before the carboxylic acid content was estimated by
titration with standard NaOH.

　　　Preparation of Hydrogel p-Nitrophenyl Carbamates. A 20 mL
solution of hydrogel (2-3.0 g) in pyridine was treated with 3 g

TABLE I.   SYNTHESIS OF HYDROGELS

Hydrogel System	Monomer Feed[a] HEMA g	Styrene g	Monomer Ratio in Copolymer[b] HEMA:STY	Precipitant	Conv. %	Mol. Wt.[c] 10^4	Solvents
Poly(2-Hydroxyethyl Methacrylate) PHEMA (1a)	18			Benzene	79	3.1	DMF Pyridine
Copoly(2-Hydroxyethyl Methacrylate-Styrene), CP(HEMA-STY) (2a)	13	10	5 : 4	Water	80	2.7	THF, DMF Pyridine
Copoly(2-Hydroxyethyl Methacrylate-Styrene), CP(HEMA-STY) (3a)	5	19	1 : 5	Water	78	2.6	Benzene CHCl3, THF DMF, Pyridine
Poly(Hydroxypropyl Acrylate) PHPA (4a)	HPA, 30 g			Benzene	70	2.4	DMF Pyridine
Poly(2-Hydroxyethyl Acrylate) PHEA (5a)	HEA, 30 g			Benzene	60	2.2	DMF Pyridine

[a] Monomer:solvent (DMF) – 1:5 v/v, initiator = BPO (0.3 wt % of solvent), reaction time = 14 hours, temperature = 70°C. [b] Based on oxygen content in the copolymer. [c] GPC (polystyrene standard)

of p-nitrophenyl isocyanate. The solution was heated at 70° for 6 hr before precipitating the polymer adduct by adding the reaction solution to 200 mL of water. The adduct was redissolved in THF, reprecipitated in isopropanol and dried in vacuo at 60° overnight before the extent of carbamation was estimated by elemental analysis (Table II).

Preparation of Hydrogel Arylsulfonyl Carbamates. To a solution of hydrogel (1-4.0 g) in dry pyridine (20-40 mL) was added either p-toluenesulfonyl isocyanate (pTSI), 3.0 g, or p-nitrobenzenesulfonyl isocyanate (p-NBSI), 0.5-3.0 g and the mixture was heated at 70° under nitrogen for 4 hr. The adduct was isolated by precipitation in benzene or ethanol. Reprecipitation from acetone or DMF yielded the desired products, IR (film) 3170 cm^{-1} (-NH), 1530 cm^{-1} (-NO$_2$), 1340, 1180 cm^{-1} (-SO$_2$). The precise degree of substitution was determined by non-aqueous titration (Table II).

Reduction of Hydrogel p-Nitrobenzenesulfonyl Carbamates. A solution of p-NBS carbamate, 1.0 g, in 20 mL of THF was blended with 15 mL of 10% aqueous Na$_2$S$_2$O$_4$. The two phase system was stirred vigorously and heated to gentle reflux for 4-6 hr. The reduction was monitored by observing the disappearance of nitro absorbance at 1530 cm^{-1}. Evaporation of the THF phase, followed by exhaustive washing with distilled water yielded the reduced adduct. Loss of product during the isolation procedure due to partial solubility in water was normally observed. The extent of reduction is reported in Table II.

Non-Aqueous Potentiometric Titration of Hydrogel Arylsulfonyl Carbamates. An approximately 0.1 N solution of tetrabutyl-ammonium hydroxide (TBAH) in isopropanol was prepared by passing 100 mL of a saturated isopropanol solution of tetrabutylammonium iodide through an Amberlite IRA-400 column which has been converted to the hydroxide form. The column was washed with 400 mL of isopropanol and the eluates were combined and standardized by titrating benzoic acid dissolved in DMF.
    A sample of the hydrogel carbamate (50-80 mg) was dissolved (or slurried) in 3 mL DMF. The solution was titrated in a microtitration vessel equipped with a 5.5 mm combination electrode. The end points and half neutralization potentials (HNP) were identified by the inflections in the EMF (MV) titration curve. A 0.3 thymol blue isopropanol solution also served as an effective end point indicator (yellow to blue).

Glutaraldehyde Immobilization of Trypsin. A sample of hydrogel-p-aminobenzenesulfonyl carbamate (0.1-0.2 g) was suspended in 5.0 mL pH 7 phosphate buffer solution in a 10 mL vial equipped with a stirring bar and Teflon lined screw cap. To the suspension was added an excess of 50% glutaraldehyde (0.5 mL), and the

mixture was stirred at room temperature for 1.5 hours. After
centrifuging, the supernatant was removed by decantation, the
residue was washed with five 7.0 mL portions of pH 7 buffer solu-
tion, and the washings were discarded. To the washed residue was
added 8.0 mL pH 7 buffer and an excess of trypsin (20-50 mg). The
mixture was stirred at 5°C overnight (17-20 hours). The residue
was then washed with aliquots of pH 7 buffer until the wash
exhibited no activity toward TAME. The immobilized enzyme was
stored in pH 8 borate buffer (10-30 mL) at 4°C.

   Titrimetric Analysis of Trypsin Activity. Into the micro-
titration vessel were introduced 0.6 mL borate pH 8 buffer, 0.2
mL 0.01 M $CaCl_2$ solution, and various increments of TAME, i.e.,
0.1 mL, 0.2 mL, 0.4 mL, 0.7 mL, and 1.4 mL of a freshly prepared
0.08 M TAME solution in pH 8 buffer. At zero time, an aliquot of
0.1 mL of the immobilized trypsin stock suspension or 0.1 mL of a
known free trypsin solution was introduced, and 0.1 N NaOH solu-
tion was added dropwise to maintain a constant pH reading. The
time and volume of base added were recorded.

Results and Discussion

   Arylsulfonyl isocyanates are very reactive acylating agents
which form adducts with alcohols, amines, polysubstituted double
bond and activated aromatic rings (9). The arylsulfonyl carbam-
ates produced upon addition to alcohols exhibit an acidity
comparable to that of benzoic acid. Reaction of cellulose with
p-toluenesulfonyl isocyanate, which produces the corresponding
carbamate without pretreatment, has been utilized to produce a
cation exchange resin (10). In neutral and alkaline media, the
ionized arylsulfonyl carbamates enhance the hydrophilicity of
hydrophobic copolymers and would provide counterions for trace
metals frequently associated with enzymes. Thus we elected to
design a coupling agent based upon arylsulfonyl isocyanate
chemistry to take advantage of the diversified chemistry exhibited
by these reagents. p-Nitrobenzenesulfonyl isocyanate appeared to
be the best candidate for evaluation.
   Although hydrogels exhibit excellent compatability with
enzymes, the gels generally exhibit poor mechanical properties.
Tear resistance is very low; data on PHEMA indicates a tensile
strength of 50 psi, an ultimate elongation of 70% and an elastic
molulus of 140 psi (11). Since copolymerization of HEMA is
readily effected (12), we decided to strengthen the hydrogel and
moderate the hydrophilicity of the support by incorporation of
styrene. Soluble copolymers were prepared to facilitate charac-
terization but the corresponding compositions could be prepared
in a crosslinked form by incorporating divinylbenzene. The
efficacy in enzyme binding by the copolymer supports was compared
with that of PHEMA using trypsin as a standard. The low

hydrolytic stability of hydroxyalkyl acrylates forced us to limit
our studies to methacrylate derivatives (Figure 1).

   Hydrogel Modification. Treatment of either pyridine or
tetrahydrofuran solutions of the hydrogels with p-nitrobenzene-
sulfonyl isocyanate (PNBSI) at 70° afforded the corresponding
arylsulfonyl carbamates in 40-100% yield. The extent of substitu-
tion increased when styrene copolymers were treated indicating
that the hydrophobic character of the support favors modification.
The reaction of PHEMA with p-toluenesulfonyl isocyanate (PTSI) and
p-nitrophenyl isocyanate (PNPI), a common enzyme coupling agent,
was conducted under similar conditions and the ease of carbamation
decreased in the order: PNBSI > PTSI > PNPI. As expected, the
carbamation enhanced the hydrophilicity of the gels; the adducts
were more swollen in water and tended to dissolve in base. The
stability of the hydrogel adducts in acid or base appeared to
parallel that of the original gels. For example, when PHEMA-PNBS
carbamate was subjected to 2 N HCl or 2 N NaOH for 12 hours at
room temperature, the extent of substitutent loss was found to be
4% and 17%, respectively, based upon loss of nitrogen content.
   To activate the hydrogel adducts for enzyme immobilization,
it was necessary to reduce the nitro function to the corresponding
amine. From the many techniques available to effect this modifi-
cation, we selected sodium dithionite as the mildest procedure
with the least potential problems with residual reagent. Since
the hydrogels were quite hydrophilic it was possible to use a two
phase solvent system (THF/H_2O) to effect the reduction. The
arylsulfonyl carbamates were reduced selectively under these
conditions; no reduction of p-nitrophenyl carbamate was observed.

TABLE II.   MODIFICATION OF HYDROGELS

Hydro-gel	Deriva-tive PTS(b) Carbamate % substn.	PNBS(c) Carbamate % substn.	PABS(d) Carbamate % substn.	PNP(e) Carbamate % substn.	PAP(f) Carbamate % substn.
PHEMA (1a)	1b, 28	1c, 43	1d, 65	1e, 18	∿0
CP(HEMA STY) (2a)	2b, 30	2c, 60	2d, 84	2e, 25	∿0
CP(HEMA-STY) (3a)	3b, 40	3c, 100	3d, 100	3e, 34	∿0
PHPA (4a)	4b, 60	4c, 66	4d, 70	—	—
PHEA (5a)	5b, 37	5c, 51	5d, 65	—	—

b) p-Toluenesulfonyl; c) p-Nitrobenzenesulfonyl; d)p-Aminobenzene-
sulfonyl carbamate; e) p-Nitrophenyl; f) p-Aminophenyl
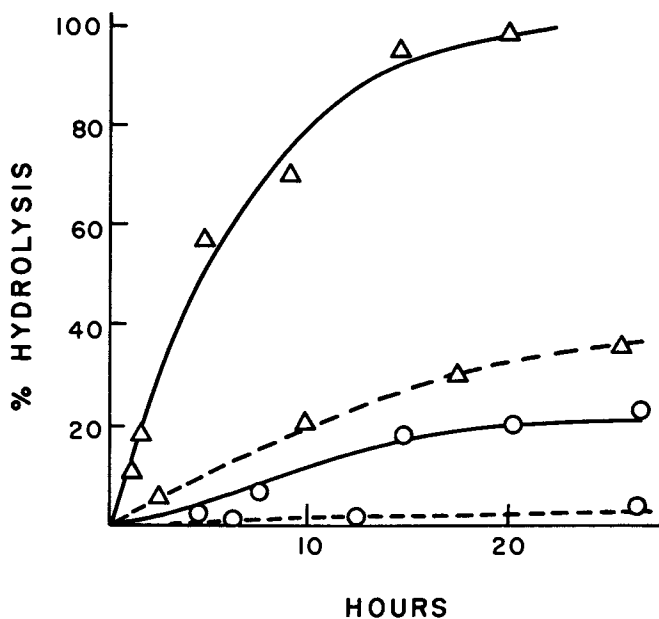
Figure 1.   Hydrolysis of poly(2-hydroxyethyl methacrylate) (○) and poly(hydroxy-propyl acrylate) (△) in 0.4 M KOH. Conditions: ——— *100°; and* – – – *25°.*

A major problem encountered in polymer modification is the precise analysis of the substituents introduced. Multiple modifications compound this problem and introduce further sources of error. However, we observed that the relative acidity of the arylsulfonyl carbamates was dependent upon the substituents and that the various end points could be resolved using nonaqueous titration techniques. Using dimethylformamide as a solvent, and tetrabutylammonium hydroxide (TBAH) in isopropanol as a titrant, the potentiometric titrations can be conducted with normal glass and calomel electrodes. End points were recognized at the potentiometric breaks or with the aid of indicators and the relative strengths of the acid could be expressed as half neutralization potentials. The hydrogel derivatives exhibit half neutralization potentials within $\pm$ 200 mv, which corresponds roughly to pKa values between 5 and 9 in aqueous media (Scheme I). Combining elemental analysis with nonaqueous titration enables us to determine the precise composition of a hydrogel derivative.

Scheme I.   HALF NEUTRALIZATION POTENTIALS IN DMF

EMF(mv)   Hydrogel Derivative

```
+200  ┼
      |  ---- Trichloroacetic Acid
      |                           O
+100  ┼                           ‖
      |  ---- R-CH₂CH₂-O-C-NHSO₂〈O〉NO₂
      |                           O
  0   ┼                           ‖
      |  ---- R-CH₂CH₂-O-C-NHSO₂〈O〉CH₃
-100  ┼
      |  ---- R-CH₂CH₂-O-C-NHSO₂〈O〉NH₂
      |                    ‖
      |                    O
-200  ┤  ` `Benzoic acid
      |  ------ R-H              CH₃
-300  ┤                    ┌         ┐
      |            R =     ┤CH₂-C-CH₂CH├
                           │    │       │N
                           │   C=O 〈O〉 │
                           └    │       ┘
                                O
```

R-CH₂CH₂-O-$\overset{O}{\overset{\|}{C}}$-NHSO₂〈O〉NO₂

R-CH₂CH₂-O-$\overset{O}{\overset{\|}{C}}$-NHSO₂〈O〉CH₃

R-CH₂CH₂-O-$\underset{O}{\underset{\|}{C}}$-NHSO₂〈O〉NH₂
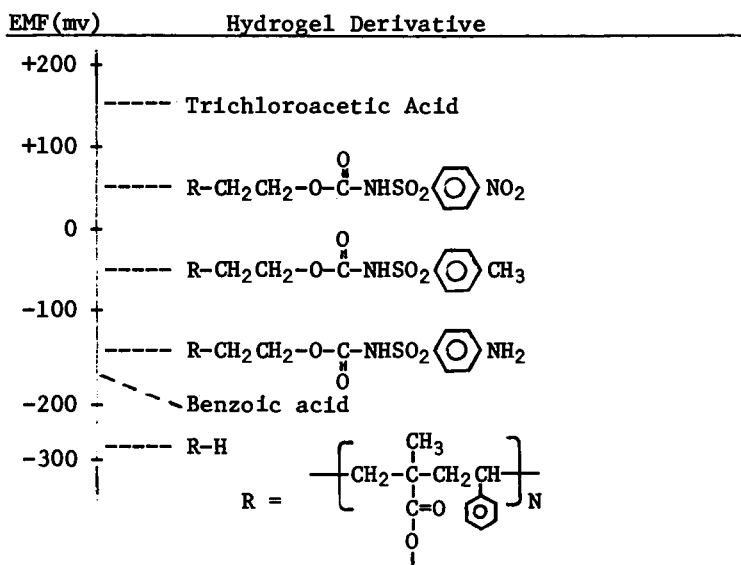
Trypsin Immobilization. The acidity of the arylsulfonyl carbamates limited the methods for immobilizing trypsin. Although treatment of a p-aminobenzenesulfonyl carbamate adduct with isoamyl nitrite generated a diazonium salt, no enzyme coupling could be detected. Thus, we were forced to couple trypsin to the hydrogel adducts with glutaraldehyde (Scheme II). Control runs demonstrated that simple adsorption was not responsible for the observed activity. The esterase activity was ascertained utilizing α-N-p-toluenesulfonyl-L-arginine methyl ester (TAME) as a

substrate and monitoring extent of reaction by titration. The
total enzyme bound was estimated by complete hydrolysis of the
enzyme conjugate and determination of selected amino acids by gas
chromatography (13). The amino acids were double derivatized;
first esterification with n-propanol and then treatment with
heptafluorobutyric anhydride to acylate the amino groups. The
quantitative GC analysis was simplified by using a difunctional
standard, DL-α-ε-diaminopimelic acid. Figure 2 illustrates a
typical chromatogram obtained from hydrolysis of enzyme conjugate
3. For comparison, the chromatogram of the free trypsin, used as
the reference standard, is reproduced in Figure 3. By selecting
four or five well resolved peaks and ratioing them to the
corresponding peak areas of pure trypsin, one obtains a very
reliable estimate of the protein bound (Table III).

Scheme II. TRYPSIN IMMOBILIZATION

CVB-Trypsin
Activity Retained
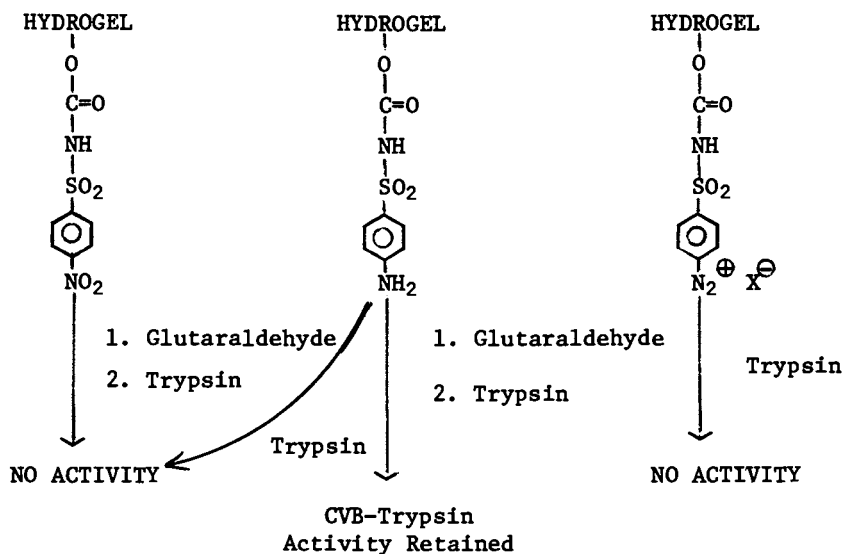
Active-site Titration. In investigating immobilized enzymes,
it is important to be aware of another highly significant consid-
eration, i.e., the relative amounts of active, partially active,
and inactive enzyme molecules per total bound enzyme. In
principle, this information can be obtained with active-site
directed titration reagents. For example, Fritz et al. (14) used
p-nitrophenyl-p'-guanidinobenzoate (NPGB) to determine the
"active" trypsin molecules in a trypsin cellulose conjugate. NPGB
proved to be an excellent active-site titrant for tyrpsin; Ford
and co-workers (15) used it on trypsin covalently linked to porous
glass particles, employing a recirculating reactor system capable
of providing accurate results at active enzyme concentration of
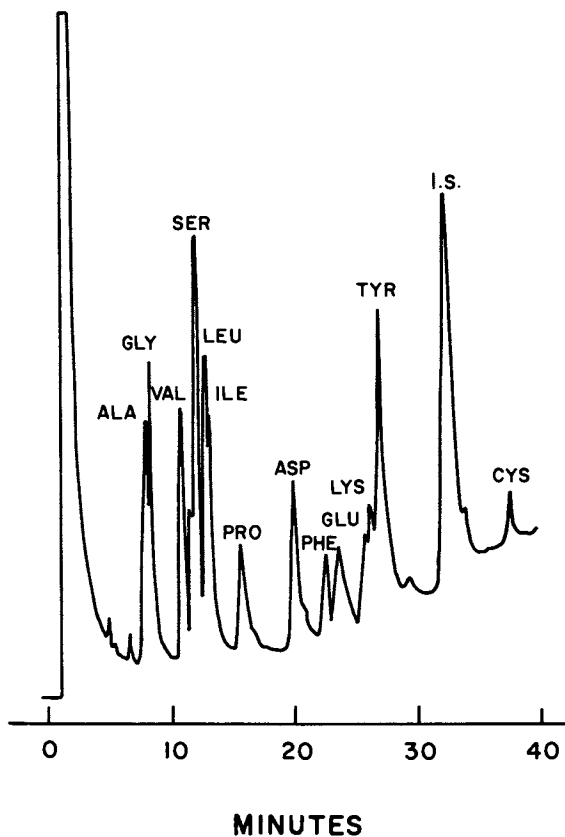
Figure 2. Gas chromatogram of the N-HFB n-propyl derivatives of the hydrolysate of trypsin CVB hydrogel sulfonyl carbamate (3f). Conditions: 12 ft. × 1/4 in. glass column packed with 3% OV-1 on Chromasorb-W; initial temp was 100° for 4 min, programed to 250° at 4°/min.
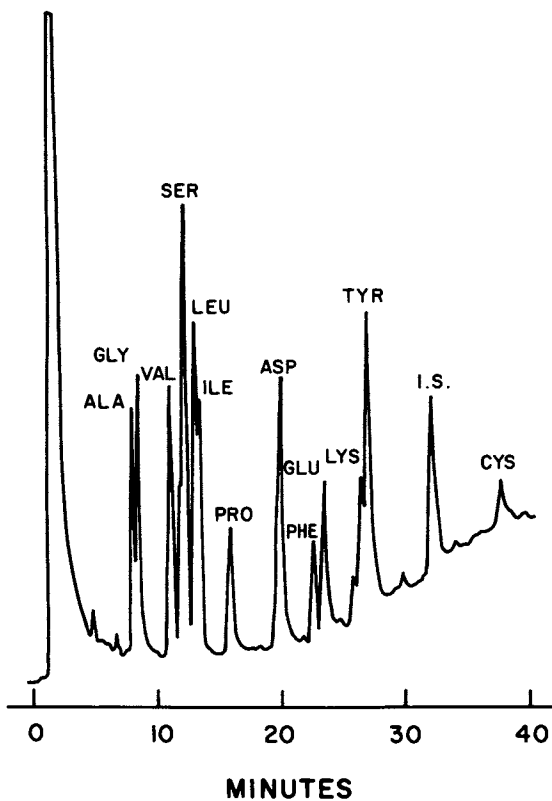
Figure 3.   Gas chromatogram of the N-HFB n-propyl derivatives of the hydrolysate of free trypsin. Conditions: 12 ft. × 1/4 in. glass column packed with 3% OV-1 on Chromasorb-W; initial temp. was 100° for 4 min, programed to 250° at 4°/min.

TABLE III.   SUMMARY OF ACTIVITY ANALYSIS FOR TRYPSIN CVB
             HYDROGEL SULFONYL CARBAMATES

Analysis \ Trypsin-cvb- Hydrogel- PABSC	PHEMA		CP(HEMA-STY)		
			(1:1)		(1:5)
	1f	1g	2f	2g	3f
Percent Carbamation	43.0	7.9	60.0	3.5	100
$\underline{\text{Mg Trypsin}}$ mL stock suspn.	0.33	0.10	0.30	0.06	0.26
Enzyme bound wt percent	55.0	1.0	2.6	0.4	4.4
Km(app)x10^{3}M	6.8	18.0	8.2	13.0	16.0
V'$_{max}$ $\frac{\mu \text{ mole}}{\text{Min-Mg}}$ a	44.0	60.0	30.0	125	47.0
Percent Sp. Activity	17.0	21.0	11.0	45.0	38.0

a) Free Trypsin: V'_{max} = 280 μ mole/Min-Mg

                  $Km = 4.8 \cdot 10^{-3}$ M

1×10^{-6} moles trypsin per liter fluid volume. To demonstrate
the feasibility of using the Ford method to determine the
active-site of our immobilized enzyme systems, trypsin CVB-PHEMA-
PABS-carbamate was treated in a circulation reactor with NPGB and
the titration is illustrated in Figure 4. The amount of p-nitro-
phenol produced by the burst is equal to the amount of the active
immobilized trypsin which, for this particular system, turns out
to be 31% of the total bound enzyme. Active-site titrations of
soluble trypsin were performed according to Chase and Shaw (16),
and the active molecules for free trypsin was found to be 70% of
the total protein involved. Consequently, the retention of
active molecules for the immobilized enzyme was calculated 45%.
The specific activity is 17% (Table III) for the same system so
the efficiency of the system, based on the actually available
active sites, was 38%. Thus, 62% of the initially active trypsin
bound has lost its activity upon binding.
     The use of kinetic assay methods requires several assump-
tions regarding the kinetic parameters of the immobilized enzyme
and the influence of diffusion on the results. As has been
discussed earlier, for many substrate-immobilized enzyme combina-
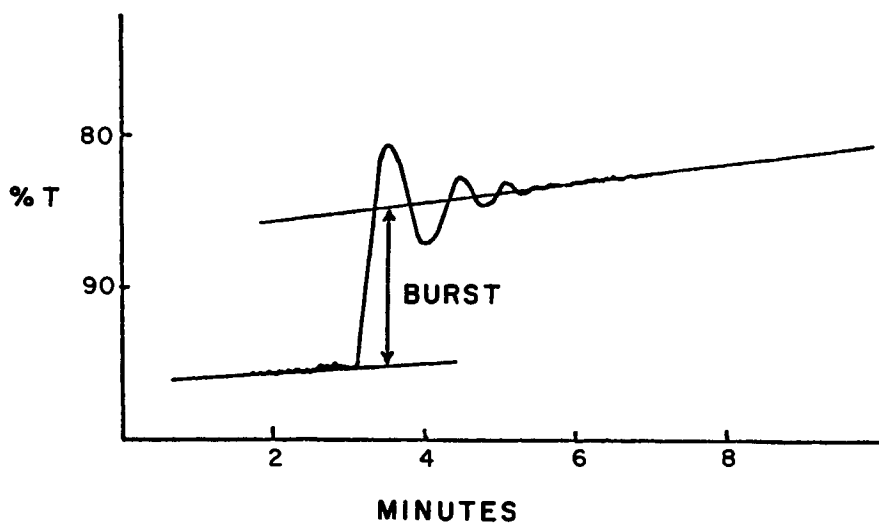
Figure 4.   *Active site titration recorder trace of 17.0 mg trypsin CVB hydrogel sulfonyl carbamate (1f) being titrated by 19.75 mL of 0.12 mM NPGB in pH 8.3 veronal buffer; burst, 0.060 A.*

tions the rapid rate of reaction provides sizable concentration
gradients both outside the immobilizing matrix and within its
pores. These concentration gradients greatly retard the rate of
reaction, even when the substrate concentration within the pores
is much greater than the immobilized enzyme Km (17). This
diffusional effect serves to decrease the apparent active enzyme
concentration. In order to obtain valid active enzyme concentra-
tions from a kinetic assay, the K_{cat} for the immobilized enzyme
must be known. However, the effects of the micro-environment and
the immobilization process itself cause the kinetic parameters of
the immobilized enzyme to differ frequently from those of the
soluble enzyme. This shift in K_{cat} of the immobilized enzyme as
result of changes in the micro-environmental pH (pH optima shift)
has been well documented (18).

    Influence of Support Structure upon Trypsin Activity. The
apparent Michaelis constant, Km(app), is much greater for
immobilized enzymes than Km of free enzyme. In our systems Km
(app) increases as the hydrophobic component increases (Table III),
apparently substantiating the theory that Km(app) of an immobil-
ized enzyme will be decreased by attachment to a hydrophilic
polymer. Another trend shown in Table III is the increase of Km
values with the decrease of the degree of carbamation. For
example, in PHEMA, the value of Km is 6.8×10^{-3} M when its
carbamation is 43%, but it jumps to 18×10^{-3} M with a much lower
degree of substitution. The change is understandable when one
realizes that with a higher percentage of carbamation the degree
of ionization due to the acidic carbamate groups is greater and,
at the same time, the enzyme to carrier ratio increases. The
former factor enhances the hydrophilicity of the system, and the
latter reduces the ratio of components that are foreign to the
enzyme; both contribute to a smaller diffusion restriction and,
as a result, a smaller Km(app) for the system with a higher
degree of carbamation.
    Table III also shows that trypsin-cvb-hydrogel derivatives
retain in the range of 11-45% the original enzyme activity, which
is about average for the covalently bound trypsin on polymers
containing hydrogel. For comparison, Mosbach (3) immobilized
trypsin on a crosslinked copolymer of acrylamide and hydroxyethyl-
methacrylate via CNBr coupling, and analyzed it photometrically
with α-N-benzoyl-Dl-arginine-p-nitroanilide (BAPNA), reporting a
35% retention of activity. O'Driscoll and co-workers (1) however,
reported an equivalent of 1.3% efficiency when trypsin was
physically entrapped in a crosslinked PHEMA gel and assayed with
TAME.
    Some reports in literature have shown that the specific
activity of a covalently bonded enzyme increases as the solubility
of its support increases (19-20). Others have observed that a
rough inverse correlation exists between the activity of an
immobilized enzyme and the amount of protein bound on the support

(21). These relationships were found true to some extent in our
experiments. As seen in Table III, the bound enzyme efficiency
increases when the amount of protein immobilized decreases six-
fold, and it also increases with a substantial decrease of enzyme
content. However, it should be noted that, in both cases, the
increase in efficiency occurs on polymer supports with a
supposedly offsetting effect of decreasing hydrophilicity. Thus,
regardless of the effect of its hydrophilicity, a hydrogel-enzyme
conjugate appears to exhibit a higher degree of activity retention
with a lower enzyme load. An exception is in the case of (2f) and
(3f), in which a relatively small increase of enzyme bound results
in a substantial increase of activity. In addition, this increase
in efficiency is noticeably accompanied by a decrease of hydro-
philicity of the polymer support. The fact that hydrogel systems
with high contents of styrene or low degrees of carbamation seem
most often associated with high values of specific activity leads
us to believe that phase separation in copolymers containing
fractions of highly different hydrophilicity and interactions
between enzyme molecules and hydrogel (or carbamates) are probably
the causes for the discrepancies. Thus, when the styrene content
in a hydrogel system is high or the percent carbamation is low,
phase separation occurs; the hydrophobic section effectively
forces the smaller populations of the hydrophilic carbamate-
trypsin conjugates to be separated, localized, and concentrated.
When this happens the enzyme molecules are in a more natural
environment, and they exhibit better activity efficiency, such as
in the cases of (2g) and (3f). On the other hand, when the
contents of the hydrogel monomers or carbamate substituents in
the systems are high, there is a possibility that some interpene-
tration between the bound enzyme and the hydrogel network might
occur. The enzyme molecules are in a less desirable environment
compared with systems with higher degrees of phase separation,
and their structure might even suffer some disruption. As a
result, the favorable diffusion effect due to the presence of
highly hydrophilic elements might still enhance the activity of
the bound enzyme but not to the extent as expected. This may
explain in large part the relatively low values of specific
activity exhibited by (1f), (1g) and (2f). However, it is
extremely difficult to ascribe precisely the cause and magnitude
of an alternation in the activity of an immobilized enzyme.

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Chitin–Protein Complexes

Ordered Biopolymer Composites

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The skeletal materials of insects and many
other "lower" animals are biopolymer composites,
in which chitin (polysaccharide) fibrils are
separated by a matrix of protein. X-ray fiber
diagrams show that in many of these systems both
the fibrous and matrix components are ordered.
We have developed a model for the three-dimensional
structure of a typical insect chitin-protein com-
plex, that from the ovipositor of the ichneumon
fly *Megarhyssa*. The x-ray pattern shows that
crystalline α – chitin fibrils are arranged
hexagonally with a center to center distance of
$7 \cdot 25$nm. This hexagonal packing is confirmed by
electron microscopy of stained ovipositor sec-
tions. The protein forms a helical sheath around
each chitin fibril, with six protein subunits
repeating in $3 \cdot 06$nm along the fiber axis. Fourier
reconstruction of the electron micrographs also
reveals a sixfold arrangement of subunits in the
protein matrix. The structure provides a basis
for the discussion of fiber-matrix interactions
in these and other structural tissues, such as
collagen and cellulose containing systems.

The polysaccharide chitin forms the component of the
skeletal materials of many lower animals, (1, 2, 3) and serves
in an analogous way to collagen in mammalian connective tissues
and cellulose in plant cell walls. All of these tissues can be
classified as fiber-matrix composites: in the cell walls of
higher plants the cellulose fibrils are surrounded by other
polysaccharides, proteins, and also lignin in some cases; in

mammalian connective tissues collagen fibrils are surrounded by
a gel of proteoglycans and other proteins; and in chitinous
tissues the fibrils are in a matrix of protein. In addition,
the animal tissues are frequently calcified. The mechanical
properties of these skeletal materials necessarily depend on
the structures of the fiber and matrix phases and their mutual
interaction. At present we have extensive knowledge of the phys-
ical structures of all three types of fibers, but information
with regard to the structure of the matrix has been more diffi-
cult to obtain. For the cellulose and collagen containing
tissues, there has been extensive study of the matrix chemistry,
but x-ray studies indicate that the physical structures are
amorphous. The chitinous tissues have received less attention,
but it has been shown that in certain of these systems the
matrix proteins are ordered, so that it is possible to develop
a three-dimensional model for the fiber-matrix structure and
interactions. The present paper describes our studies using
x-ray diffraction and electron microscopy to derive the struc-
ture of one of the more ordered chitin-protein complexes, that
from ovipositor of the ichneumon fly *Megarhyssa*.
     Chitin is a repeating homopolymer of $\beta(1,4)$-linked N-
acetyl-D-glucosamine (NAG) and is obtained after chemical or
enzymic deproteinization of the native tissues. Such specimens
of chitin generally contain some unacetylated residues, (4, 5)
and there has been some speculation that these may be the points
for covalent linkage to the protein. (6-9) Much further work is
necessary in this area, but evidence has been presented for co-
valent chitin-protein linkages at some of the chitin chain ends
in certain tissues. It is interesting to note that the only
reported chitin that is pure poly NAG occurs in the spines of
certain marine diatoms, and exists completely free of pro-
tein. (10) Chitin has been shown to exist naturally in two
polymorphic forms, known as α- and β-chitins. (11) In both
forms the chitin chains have a 2_1 helical conformation, (12, 13,
14) with two NAG residues repeating in approximately 1·03nm
similar to that observed for cellulose. (15, 16) The two forms
of chitin differ in terms of the polarity of adjacent chains,
which are antiparallel in α- and parallel in β-chitin. This
leads to different intermolecular hydrogen bonding networks, (17,
18) with a result that β-chitin is swollen by water and α-chitin
is not, which may explain why α-chitin is by far the more common
form found in nature. The chitin in *Megarhyssa* ovipositor is
α-chitin, and has the structure shown in fig. 1, as was deter-
mined previously in this laboratory. (18)
     Electron micrographs of insect cuticles in which the pro-
tein is stained e.g. with uranyl acetate show a hexagonal array
of (unstained) chitin fibrils in a matrix of protein. Neville
et al (19) have reported microfibrils with diameters of 2·4 -
3·0nm separated by 5·1 - 8·3nm in a survey of insect cuticles
from a wide variety of species. Rudall (20) observed a hexagonal
lattice dimension of a = 6·9nm for *Megarhyssa* ovipositor. He also

a

b
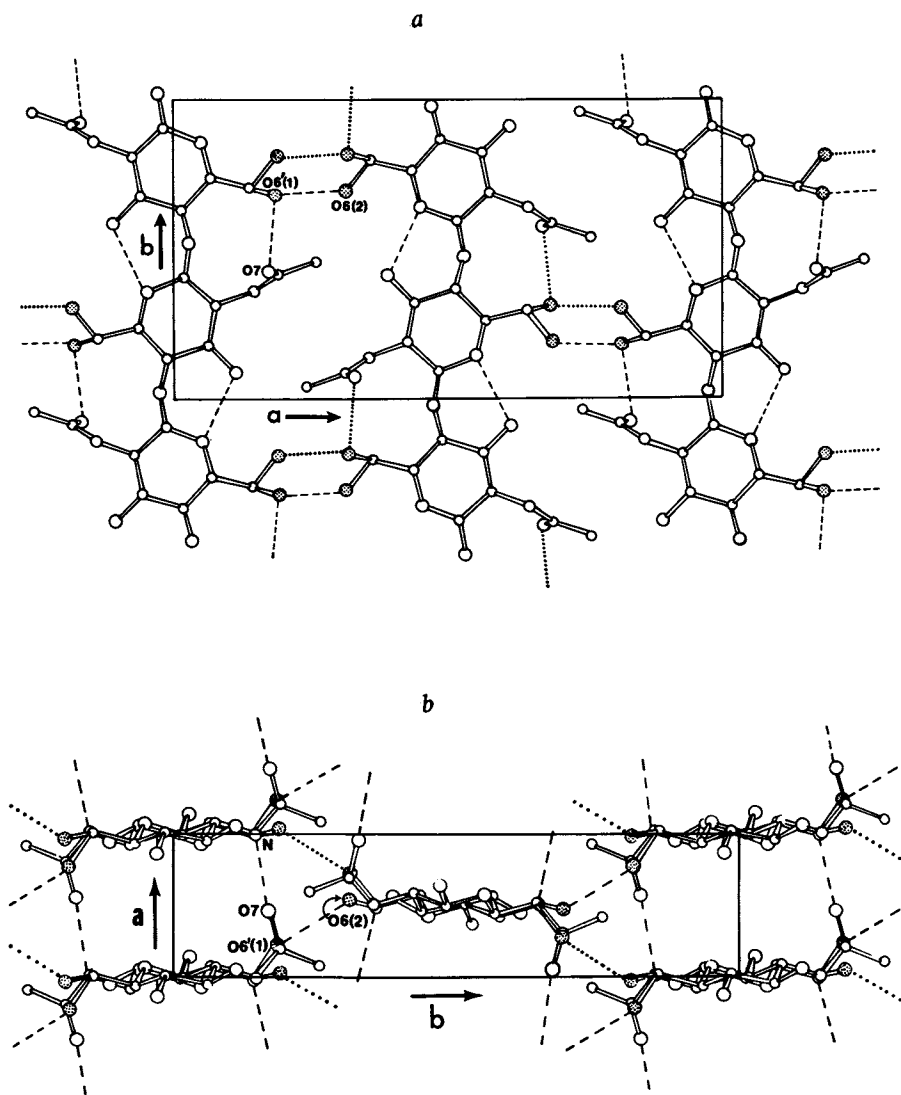
Figure 1. Structure of α-chitin. Key: a, ac projection and b, ab projection. (Reproduced, with permission, from Ref. 18. Copyright 1973, Academic Press.)

showed (20, 11) that this and many other insect cuticles gave
x-ray patterns that indicated that the protein matrix is or-
dered. This led to a classification (21, 22) of the chitin-
protein complexes based on their fiber repeats of 3·1, 4·1, or
6·2nm, which are respectively three, four, and six times the
chitin repeat.

At present relatively little is known about the chemistry
of the matrix protein, although significant differences have
been reported between proteins extracted from different tissues.
Extraction data indicate the presence of numerous protein moi-
eties, (23, 24, 25, 6) but interpretation is difficult because
of extensive cross-linking (tanning) in mature cuticles. Rudall
and Kenchington (21) report that the evidence for ordering of
the protein is eliminated by extraction with 7M urea, suggesting
a non-covalent interaction between protein and chitin.
Rudall (22) also cites Skerrow, who showed evidence for a basic
protein subunit of molecular weight 27,000 extracted in 7M urea
from young honeybee (*Apis mellifera*) cuticle, a tissue giving a
similar x-ray pattern to that for *Megarhyssa* ovipositor.

In the work described below we have derived a structure for
the chitin protein complex of *Megarhyssa* ovipositor, for which
confirmatory evidence has been obtained by Fourier reconstruc-
tion of electron micrographs. Details of this work have been
published separately (see refs. 26 and 27).

Materials and Methods

Mature female ichneumon flies (*Megarhyssa lunator* and *Mega-
rhyssa atrata*) were generously provided by Dr. S. Teraguchi of
this university, Dr. A.G. Nielson of Ohio Agriculture Research
and Development Center, Wooster, Ohio, and Dr. K.M. Rudall, of
Auckland, New Zealand. Fresh and museum specimens proved to be
indistinguishable by x-ray and electron microscope techniques.
The ovipositors were easily separated longitudinally into three
sections (1st and 2nd volvifers) and cut into 1 cm lengths to
serve as x-ray specimens. Deporteinized specimens were prepared
by boiling in 5% KOH solution for 24 hrs.

X-ray patterns were recorded on Kodak No-Screen film using
Ni filtered CuKα radiation from a Rigaku Denki rotating anode
source. Small angle and wide angle patterns were obtained using
a Rigaku Denki and Searle (toroidal) camera respectively. An
E.D.P. scanning optical densitometer was used to measure inten-
sities and to determine the small angle d-spacings and the
positions of the maxima on the protein layer lines.

Specimens for electron microscopy were prepared as thin
cross-sections and stained with uranyl acetate and lead citrate.
A JEOL 100B electron microscope was used to obtain the electron
micrographs. Reconstruction of sections of these micrographs
by Fourier filtration methods was done in collaboration with
Dr. J. Frank of New York State Department of Health, Albany, N.Y.

(as described in refs. 28 and 29). In addition, a number of
individual fibrils were selected where the neighboring protein
appeared to be well ordered. These fibrils were averaged after
optimization of their rotational orientation.

Results and Discussion

     Figures 2 and 3 show the x-ray fiber diagrams of intact
and deproteinized *Megarhyssa* ovipositor. Figure 3 is charac-
teristic of α–chitin and most of the features of this pattern
can also be seen in fig. 2. Thus it can be concluded that the
intact ovipositor contains crystalline α-chitin. The additional
data in fig. 2 are quite extensive and indicate that there is an
ordered superstructure. Firstly, in the small angle region
there are three equatorial reflections at $d = 6 \cdot 29$, $3 \cdot 62$, and
$\sim 3 \cdot 14$nm. These can be indexed by a two-dimensional hexagonal
lattice with dimension $a = 7 \cdot 25$nm. This compares favorably with
$a \simeq 6 \cdot 9$nm reported previously by Rudall, (20) and $a \simeq 7 \cdot 5$nm for
the center to center distance of hexagonally packed chitin
microfibrils seen in our electron micrographs (see below).
Comparison of the squared Fourier transforms of solid cylinders
with the observed small angle intensities (F^2) showed best
agreement for a diameter of $7 \cdot 25$nm. We therefore envisage a
structure comprised of microfibrils, each consisting of a crys-
talline core surrounded by a sheath of protein, which are
closely packed on a hexagonal lattice as shown in fig. 4. The
calculations also indicate that the protein and chitin have ap-
proximately the same electron density, and therefore the radius
of the chitin core is not defined. The complex contains ap-
proximately 85% protein, from which a core diameter of $2 \cdot 8$nm can
be calculated on volume considerations. This however assumes a
homogenous distribution of the chitin and protein in the bulk
structure. From our electron micrographs the fibril diameter is
in the range $3 - 4$nm and it is difficult to be more specific due
to the granular appearance of the stained protein. It has been
suggested by Atkins (30) that certain wide angle equatorial re-
flections due to the chitin may arise from a crystallite four
chains wide, which would correspond to a dimension of $3 \cdot 8$nm.
Thus the diameter of the chitin core is uncertain (within the
above limits) but this is not critical to the analyses that
follow for the arrangement of the protein subunits.
     In the wide angle region of the x-ray pattern for the in-
tact ovipositor there appear to be three layer lines for every
one for the purified chitin: layer lines 3, 6, and 9 in fig. 2
are very similar to layer lines 1, 2, and 3 in fig. 3. The
extra layer lines (1, 2, 4, 5, 7, 9, 10...) and the additional
intensities on layers 3, 6, and 9 must arise from the protein
(and its interaction with the chitin). The layer line repeat
is $3 \cdot 06$nm, and thus we envisage a protein sheath structure with
a repeat approximately three times that of the chitin, i.e. six
NAG residues.
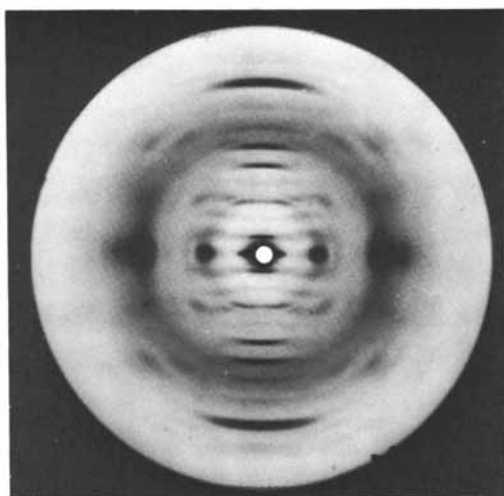
Figure 2.   X-ray fiber diagram of intact ovipositor of **Megarhyssa** **lunator.** *(Re-produced, with permission, from Ref. 26. Copyright 1980, Academic Press.)*
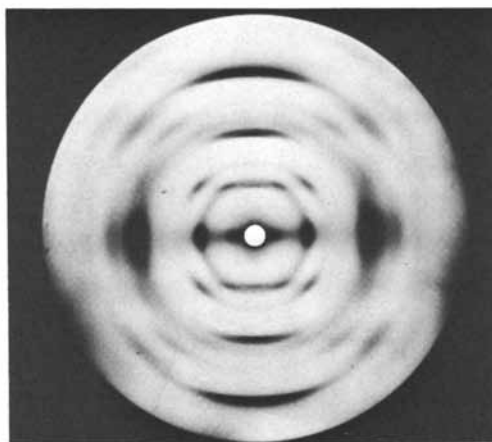
Figure 3.   X-ray fiber diagram of deproteinized ovipositor of **Megarhyssa** **lunator.** *(Reproduced, with permission, from Ref. 26. Copyright 1980, Academic Press.)*
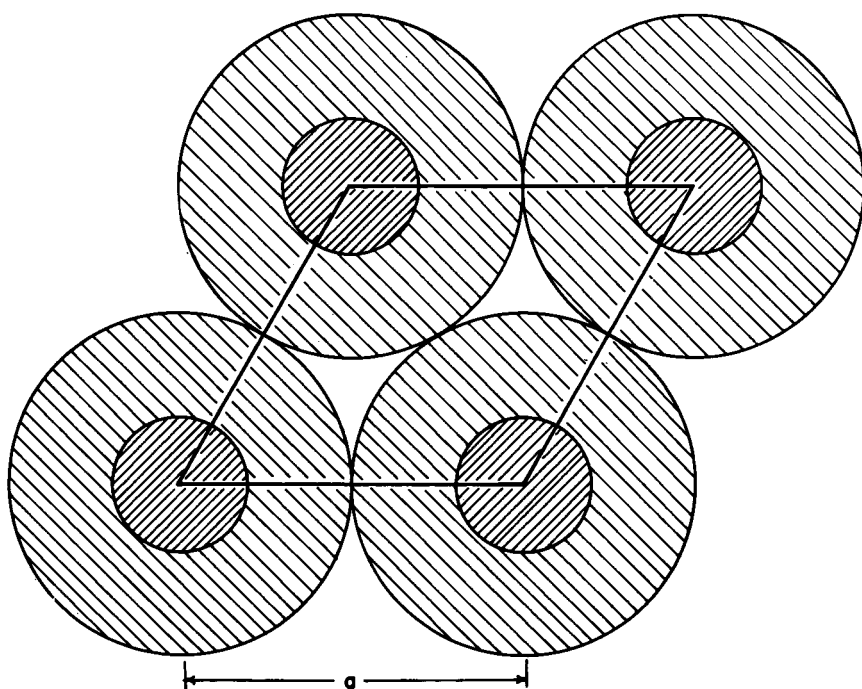
Figure 4.   Hexagonal arrangement of core-sheath microfibrils in the proposed structure of Megarhyssa *chitin–protein complex. (Reproduced, with permission, from Ref. 26. Copyright, 1980, Academic Press.)*

The intensity distribution on the protein layer lines suggests a helical arrangement of protein subunits. The strong intensity on layer line 1 is seen on close inspection to be split, i.e. it is off-meridional; and similarly, the intensity at the meridian is split in photographs of tilted specimens. Compared to layer lines 1 and 5, layer lines 2 and 4 are weaker and have their maxima further from the meridian. Layer line 3 is mainly due to the chitin; the protein contribution is weak and the first maxima is even further from the meridian than those on layers 2 and 4. This intensity distribution is suggestive of a helical structure with six units per turn repeating in 3·06nm. Such a structure would have a meridional reflection on the sixth layer line, as is observed, although of course this coincides with the strong meridional on the second layer line of the α-chitin pattern. From the position of the first maximum on the second (protein) layer line the radius of the equivalent point helix is calculated to be 2·42nm, which should correspond approximately to the mid-point of the protein core. This compares well with calculated mid-point radii of 2·51 and 2·76nm for core-sheath models with core diameters of 2·8 and 3·8nm respectively.

From the above arguments we can propose the core-sheath model shown in fig. 5: the chitin core is surrounded by a helix of six protein subunits repeating in 3·06nm. We have shown the subunits as globular proteins, and if the basic subunit is indeed a protein of molecular weight 27,000, then this would have approximately the dimensions shown. However, a more heterogeneous protein population could also be compatible with the proposed symmetry, and the scattering subunits could contain sections of several protein chains, perhaps with some fibrous character. Further work is necessary on the chemistry of the proteins before we can proceed further in this area.

On closer inspection it can be seen that this treatment of the protein data is an over simplification, in that the layer lines identified as 1, 2, 3... above are in fact split into several components. This indicates that the description of the protein structure as a 6_1 (or 6_5) helix repeating in 3·06nm is only an approximation to a structure with a longer repeat. From our measurements of the splitting of the third layer line, a rough estimate of the actual repeat is 15·3nm, or more probably an integral multiple thereof. This longer repeat could arise from a small additional twist to the protein helix, giving it a non-integral character, or the formation of a coiled-coil structure. We are currently working on a number of other chitin-protein complexes, all of which have the same basic 6_1 helical structure repeating in approximately 3·06nm, but have different layer line splittings and long repeats for the overall protein structure. This long repeat may be significant in view of the numerous electron microscope observations that chitin (and also cellulose) microfibrils tend to show a ribbon like morphology with a slow twist.
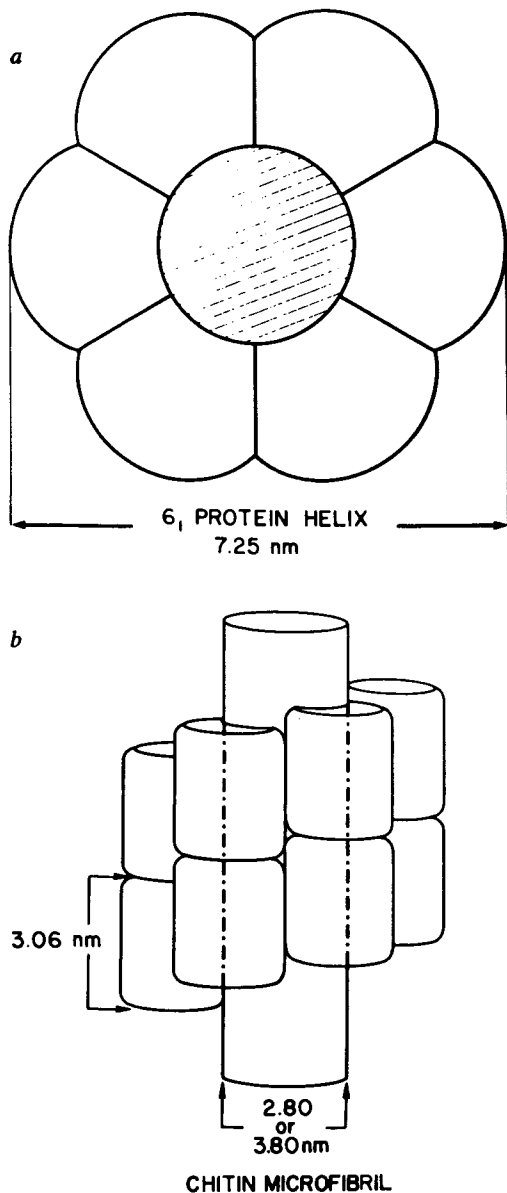
Figure 5. *Proposed arrangement of the protein subunits around the chitin core in* Megarhyssa *chitin–protein complex. Key: a, axial projection and b, projection perpendicular to fiber axis. (Reproduced, with permission, from Ref. 26. Copyright 1980, Academic Press.)*

Reconstruction of Electron Micrographs

Figure 6 shows an electron micrograph of a transverse sec-
tion of *Megarhyssa* ovipositor, in which the protein is stained
with uranyl acetate and lead citrate. Unstained chitin micro-
fibrils are seen in a matrix of stained protein; the fibrils are
arranged hexagonally in the well ordered regions, and have a
center to center distance of 7·5 ± 0·5nm. When we examine the
protein matrix in more detail it is seen to have a granular ap-
pearance, consisting of stained dots which are 2 - 3nm across.
In certain areas there is a suggestion of sixfold symmetry to
these dots, although this is not nearly as perfect as the fibril
arrangement. We have therefore used Fourier reconstruction
methods to see if the micrographs contain general evidence of
protein substructure, and our preliminary results are shown
below.

Figure 7 shows the Fourier reconstruction of a region se-
lected from the micrograph in fig. 6, in which the chitin fibrils
form a well ordered hexagonal array. The procedure used calcu-
lates the Fourier transform which shows the maxima predicted for
the hexagonal lattice, plus additional information or "noise"
between these maxima due to the irregularities in the structure.
If it can be assumed that these irregularities arise due to
artifacts in specimen preparation, then the noise can be ignored
or filtered out, and the image can be resynthesized from the
hexagonal lattice (hk0) maxima. The filtered image is shown
in fig. 7b, based on 46 observed maxima out to d ≃ 20Å, along
with the original unfiltered region of the micrograph. The re-
construction shows an array of black dots around the unstained
chitin fibril, and this array appears to have approximately
sixfold symmetry. Once sixfold symmetry can be assumed it is
possible to identify fibrils for which the immediate protein
environment is particularly well ordered. Figure 8a shows fif-
teen such fibrils selected from different regions of the same
micrograph. These were then reoriented to optimize the match
between them, and then averaged and filtered using the tech-
niques of Frank *et al.* (28) This resultant average is shown in
fig. 8b, and the sixfold arrangement of the protein subunits can
be seen clearly.

Although caution is necessary to avoid over interpreting
such analyses, these results can be taken as conformation of the
sixfold arrangement of the protein subunits. It should be noted
that the models from the two techniques are not exactly the
same; each fibril in figs. 3 and 4 is surrounded by its own
group of subunits, whereas the subunits are shared between the
fibrils in fig. 7. Further work is necessary to resolve the de-
tails of the protein structure, although these two models are
not necessarily inconsistent with each other, given the differ-
ence in the experimental techniques.
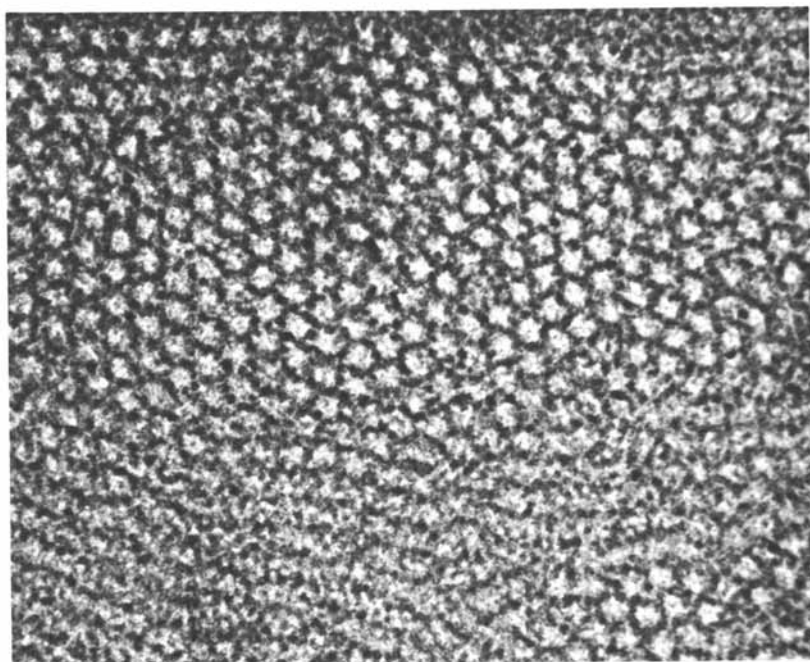
Figure 6.   Electron micrograph of transverse section of Megarhyssa *ovipositor stained with uranyl acetate and lead citrate.*
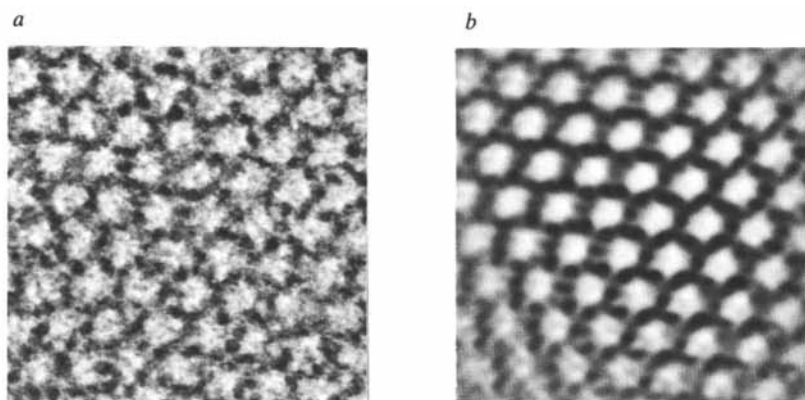
Figure 7. Fourier reconstruction of electron microscope image for Megarhyssa ovipositor. Key: a, selected well-ordered region and b, reconstructed image.
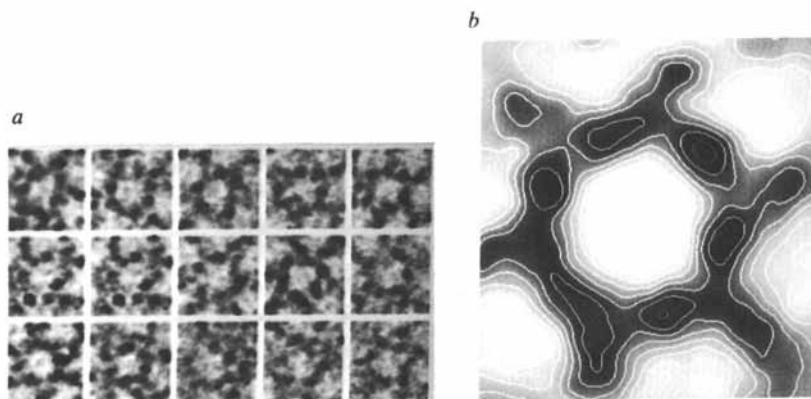
Figure 8. (a) Fifteen selected chitin fibrils with well-ordered surrounding protein and (b) filtered average synthesized from (a).

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Interaction of Synthetic Polymers with Cell Membranes and Model Membrane Systems: Pyran Copolymer[1]

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We have investigated the interaction of pyran copolymer with erythrocyte ghosts, with intact erythrocytes and with liposomes of pure dipalmitoyl-phosphatidylcholine (DPPC). Particular attention was given to the importance of divalent metals (especially Ca^{2+}) in promoting the association of pyran copolymer with cell and lipid surfaces. High sensitivity differential scanning calorimetry shows: i) The perturbation of erythrocyte ghost structure by pyran copolymer is not readily "washed out" by treatment with fresh buffer, ii) This perturbation is not due simply to sequestration of metal ions, iii) Pyran copolymer reverses at least partially the effects of Ca^{2+} added to erythrocyte ghosts, and iv) Pyran copolymer raises the transition temperature of DPPC liposomes in the presence of 5 mM Ca^{2+}, but not in the absence of added Ca^{2+}.

The 1:2 copolymer of divinyl ether and maleic anhydride ("pyran copolymer," I) exhibits a broad range of biological activity, and has been discussed elsewhere in this volume and in earlier reviews (1, 2, 3). Of particular interest is the polymer's antitumor activity; after initial clinical testing and then withdrawal from clinical use due to severe toxicity, the drug is again under evaluation as an agent for the treatment of human cancer.
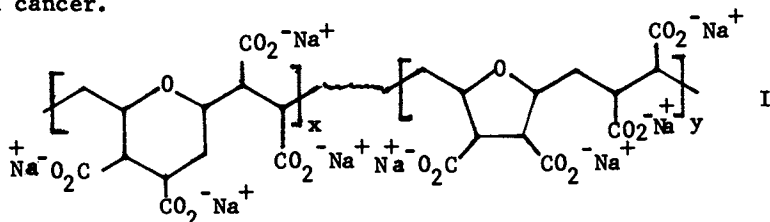
[Structure I is a nominal structure for the 1:2 copolymer of divinyl ether and maleic anhydride, after hydrolysis; in fact, the size of the ether ring is a subject of current investigation (4). Potentiometric titration shows that approximately 3 of the 4 carboxyl groups are ionized at pH 7 (1)].

The mechanism of the antitumor action of pyran copolymer has been studied by a number of groups, and there is now ample experimental evidence supporting the hypothesis that this activity is mediated by macrophages (for a review, see ref. (5)). In addition, Fiel and coworkers (6-9) have studied the interaction of the drug with divalent cations, and have presented evidence for the formation of soluble and insoluble complexes. Recently, these workers have speculated that pyran may function as a macromolecular ionophore, activating macrophages via changes in intracellular ion concentrations (6). Schultz, Papamatheakis and Chirigos (10) have observed that tumor cells aggregate around and adhere firmly to pyran-activated macrophages, in vitro, suggesting structural and functional modification of the macrophage membrane, but the nature of the modification is unknown. Finally, the acute toxicity of the partial Ca^{2+} salt of pyran is significantly less than that of the Na^+ salt, and it has been suggested that divalent metal ions may be important in mediating the interactions of pyran with proteins and with cell surfaces (11).

In this paper, we examine the interactions of pyran copolymer with model biomembranes of two kinds: i) the human red blood cell membrane (or red cell "ghost") and ii) multilamellar suspensions (liposomes) of dipalmitoylphosphatidylcholine (DPPC), a pure synthetic phospholipid. Each of these systems offers advantages in studies of polymer-cell surface interaction: The red cell membrane, while complex, is still the most readily isolated and best understood of the membranes of normal human cells, and its molecular architecture is, in a general way at least, typical of such membranes. The pure phospholipids provide a much simpler biomembrane model, with the prospect of yielding more complete interpretation of experimental observations.

The results described in this paper support the suggestion that divalent metal ions promote the interaction of pyran copolymer with membrane and lipid surfaces.

Experimental

Pyran Copolymer. Pyran copolymer used in calorimetric studies of red cell ghosts was NSC-46015, a gift of Dr. David S. Breslow of Hercules, Inc. For studies of pyran-lipid mixtures, the polymer was prepared by radical copolymerization of maleic anhydride and divinyl ether according to the technique of Breslow (1), using 9/1 acetone/tetrahydrofuran mixture as solvent, and azobisisobutyronitrile as initiator. The inherent viscosity of the sample was 0.189 dl/g (0.5 g/100 ml in 0.05M NaCl, 30°C).

[14]C-labeled copolymer was prepared in similar fashion, using
[2,3-^{14}C]-maleic anhydride (Amersham). Two samples were prepared
- one of specific activity 0.66 mCi/mmol and the other of specific
activity 0.54 mCi/mmol. The specific activity was varied by
changing the ratio of labeled and unlabeled maleic anhydride
used in the copolymerization.

Calorimetry. Human blood (Type A+) was obtained from the
Central Blood Bank of Pittsburgh, PA., and used before becoming
clinically outdated. Red cell (erythrocyte) ghosts were prepared
by the method of Dodge (12), and suspended in 310 ideal millios-
molar (imosm) buffer, either phosphate (a mixture of 0.155 M
NaH_2PO_4 plus 0.103 M Na_2HPO_4) or cacodylate [0.020 M $Na(CH_3)_2AsO_2$,
0.135 M (HCl + NaCl)], at pH 7.4. Pyran copolymer, ethylene-
diaminetetraacetic acid (EDTA) and $CaCl_2$ were introduced into the
ghost suspensions as required by two washings with a solution of
the desired composition, also at pH 7.4. In those experiments
where the concentration of pyran copolymer was sufficient to
depress the buffer pH, readjustment was made by adding aqueous
NaOH. Heat capacity profiles of treated and control ghost
suspensions were recorded on a Microcal, Inc., MC-1 micro-
calorimeter, at a heating rate of 60°C/hr. Control and treated
suspensions were run on the same day, within 24 hr of ghost isola-
tion.

Polymer-lipid mixtures were prepared by hydrating the dry
lipid (L-α-phosphatidylcholine, dipalmitoyl; Sigma Chemical Co.)
in 50 mM Tris (tris(hydroxymethyl)aminomethane) buffer, pH 7.4
containing pyran copolymer at a concentration of 1 mg/ml. The
final lipid concentration was also 1 mg/ml. Ca^{2+} was added in
the form of $CaCl_2$. Thermal transitions of lipid samples were
recorded at a heating rate of 12°C/hr.

Binding of Pyran Copolymer to Intact Erythrocytes. Human
erythrocytes were isolated from whole blood (Type A+) by centri-
fugation, washed 4 times with isotonic saline and then 3 times
with isotonic buffer, pH 7.4. Phosphate buffer was used in
experiments without added Ca^{2+}; cacodylate buffer in experiments
with added Ca^{2+}. The cells were then suspended in buffer at
the desired concentration (5×10^8 cells/cm^3, as determined using
a hemocytometer). [^{14}C]-Pyran copolymer in buffer was added
and the mixture incubated for 10 min at room temperature. The
cells were then sedimented by centrifugation and the supernatant
removed. After 2 washings with buffer (5 ml each), the cell
pellet was dissolved in 1 ml of 10% aqueous Triton X-100 and 10 ml
of Budget Solve (Research Products International) scintillation
cocktail, and the bound radioactivity determined by liquid
scintillation counting, using a Beckman LS7000 Counter. Each
sample was run in quadruplicate, and duplicate control experi-
ments without cells were run with each.

Results and Discussion

Figure 1 shows heat capacity profiles of human erythrocyte ghosts incubated with increasing concentrations of pyran copolymer in 310 imosm phosphate buffer. The five endothermic transitions of the control sample (bottom curve) are labeled A, B_1, B_2, C and D, in accord with earlier work by Brandts and coworkers (13-16) and from this laboratory (17-20). The A transition is assigned quite securely as a partial denaturation of spectrin, the major cytoskeletal protein on the erythrocyte membrane (13). B_1 and B_2 appear to involve proteins and lipids (protein Bands IV.1 and IV.2 in B_1 and Band III in B_2) (14). The C transition exhibits many of the characteristics of a phospholipid phase transition, and D has been assigned to a protein denaturation process (13). Thus the membrane is resolved into a number of "domains," and one can probe the interactions of a polymeric solute with each of these structures.

Pyran copolymer at a concentration of 3.7 mg/ml causes a marked change in the appearance of the heat capacity profile. In fact, each transition is affected, with the possible exception of D. Although there are small changes in the temperatures of the A, B_2 and C transitions, we feel at this point that the most significant perturbation is that of the B_1 transition, which appears depressed in temperature and in amplitude. Neither poly(acrylic acid) nor poly(methacrylic acid) affects the B_1 transition in this way. Recently, we have observed a similar "flattening" of the B_1 transition by poly(α-ethylacrylic acid) at a repeating unit concentration of 10 mM (21), and we are now pursuing possible further parallels in the biological properties of poly(α-ethylacrylic acid) and pyran copolymer. As the concentration of pyran copolymer is reduced (middle curves in Figure 1), the shift in each transition decreases, and no perturbation is apparent at 20 μg/ml.

In trying to characterize this interaction more fully, we determined the reversibility of the pyran-induced perturbation of the heat capacity profile. Figure 2 shows profiles for control ghosts (the lower curve) and for ghosts incubated with 3.7 mg/ml of pyran and then washed with fresh buffer 3 times - a treatment which reverses completely any effects of pH, ionic strength or divalent ions (14). The upper curve shows that the effect of pyran copolymer is partially, but not completely, reversible under these conditions, in that the B_2 and C transitions remain shifted. A small loss in amplitude of B_1 may also be significant. This is consistent with our quantitative binding experiments (discussed below) in which we have observed that a small but measurable amount of pyran remains associated with whole cells after a comparable treatment with fresh buffer.

Effects of Divalent Ions. Because of the known affinity of pyran copolymer for divalent ions (6-9), it seemed plausible that
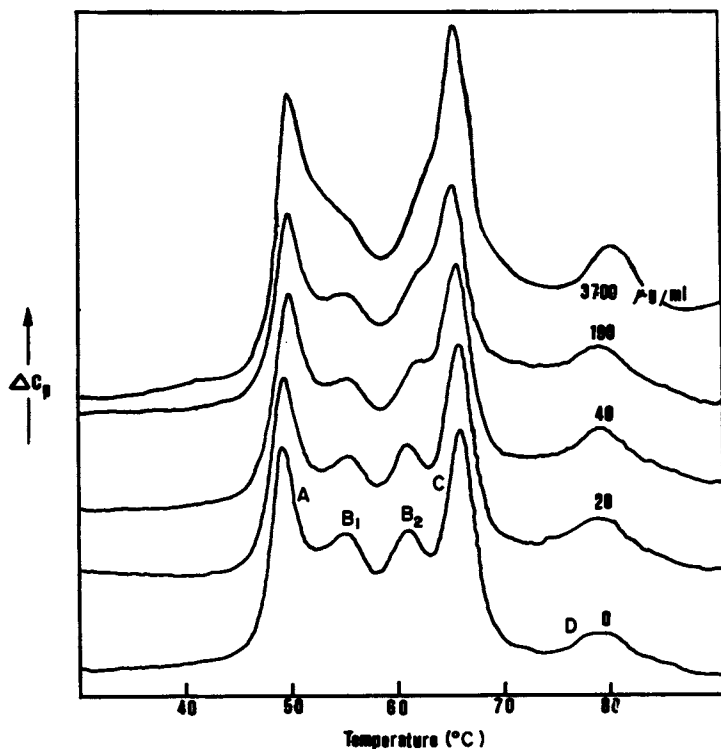
Figure 1.   Heat capacity profiles of erythrocyte ghosts suspended in 310 imosm sodium phosphate at pH 7.4.   Concentration of pyran copolymer in each suspension is indicated.
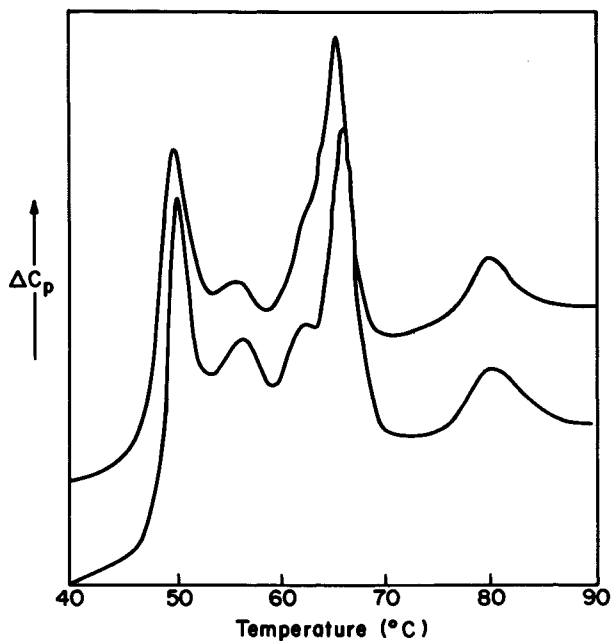
Figure 2.   Reversibility of pyran-induced perturbation of heat capacity profile. Key: lower curve, control profile with ghosts suspended in 310 imosm sodium phosphate; and upper curve, ghosts incubated with 3.7 mg/mL pyran copolymer, then washed three times with fresh 310 imosm phosphate buffer.

the observed perturbation of membrane structure might result
from a depletion of Ca^{2+} or Mg^{2+} bound to the membrane or intro-
duced in the ghost preparation. A typical preparation contains
approximately 0.2 mM Ca^{2+} plus Mg^{2+} by atomic absorption spec-
troscopy. Figure 3 compares the effects of pyran and EDTA,
which should also compete for metal ions. The effects of these
two compounds are clearly different, especially in the region of
the C transition, which is raised in temperature (from 66° to 68°
C) by EDTA, and depressed in temperature by a comparable amount
of pyran copolymer. The effects on the B_1 transition also differ
markedly. On this basis, it seems clear that the interaction of
pyran with the erythrocyte ghost is not simply a sequestration
of membrane-bound metal ions, with the formation of soluble
polymeric metal salts. On the other hand, one cannot rule out
complexation plus association of the complex with the membrane
surface.

The importance of divalent ions shows up most clearly in
quantitative binding experiments using ^{14}C-labeled pyran co-
polymer. In these experiments, whole erythrocytes - not
isolated membranes - were incubated at room temperature for 10
minutes with the desired concentration of pyran copolymer in
buffer. The cells were then sedimented by centrifugation,
washed twice with fresh buffer, disrupted with detergent, and
the radioactivity determined by liquid scintillation counting.
The same procedure was performed as a control in the absence of
cells, in order to check for sedimentation of polymer. The
results are shown in Figure 4.

Figure 4 is a plot of sedimented radioactivity vs the con-
centration of pyran in the incubation mixture, in the presence
of 5 mM Ca^{2+} and in the absence of added Ca^{2+}. These measure-
ments were done in quadruplicate, with 5×10^8 cells/cm^3, and the
error bars are the 95% confidence limits on each value. The
effect of Ca^{2+} is striking, in that approximately 100 times as
much radioactivity is sedimented in the presence of 5 mM Ca^{2+}
as in the absence of added divalent ions. 5 mM represents a
reasonable concentration of divalent ions in human plasma, and
60 μg/ml of pyran (the highest concentration shown) would be
reasonably obtained in clinical use. It is significant that the
increased binding in the presence of Ca^{2+} is observed only when
Ca^{2+} is present also in the wash buffer - washing with Ca^{2+}-free
buffer reduces the sedimented radioactivity to levels obtained
in incubations without added Ca^{2+}.

Complexation with Ca^{2+} seems to cause pyran copolymer to
become "sticky," and to adhere to the red cell surface, but the
meaning of this result is obscured somewhat by the fact that,
at higher concentrations - higher than 60 μg/ml - significant
radioactivity is sedimented in the absence of cells, suggesting
flocculation of pyran/Ca^{2+} microspheres as observed by Fiel and
coworkers. It is possible that we are seeing only a hetero-
flocculation of microspheres plus red cells, and that the
association of pyran with the membrane is nonspecific.
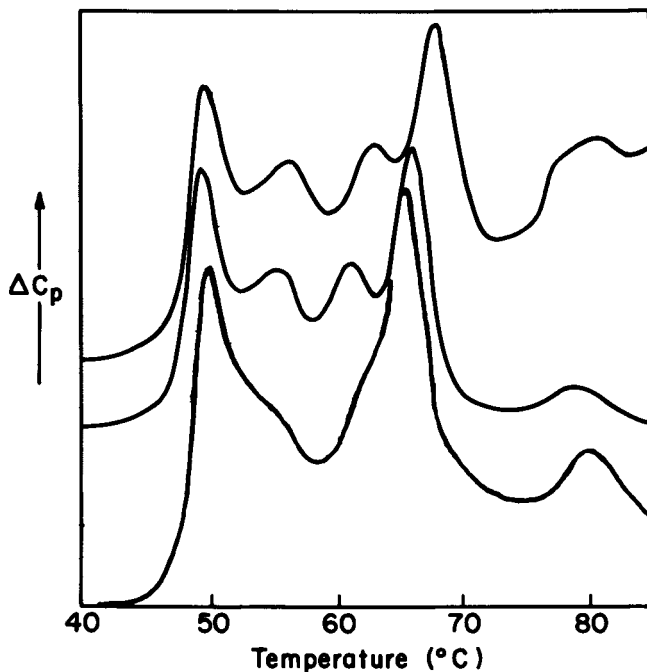
Figure 3.   Comparison of effects of pyran copolymer and EDTA in 310 imosm sodium phosphate.   Key: center curve, control profile; lower curve, ghosts incubated with 3.7 mg/mL pyran copolymer; and upper curve, ghosts incubated with 3.7 mg/mL EDTA.
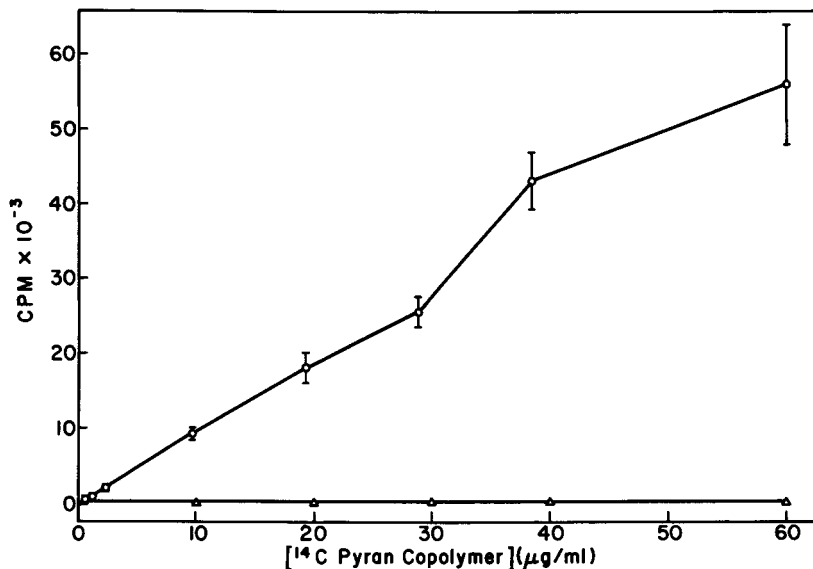
Figure 4.   Effect of Ca^{2+} on "binding" of pyran copolymer to intact erythrocytes. Key: $\triangle$, no Ca^{2+}; and $\bigcirc$, 5mM Ca^{2+}.

   Still, the question of structural perturbation of the mem-
brane by pyran copolymer in the presence of physiological concen-
trations of divalent ions is an important one, so we applied
calorimetry to this problem. Figure 5 shows as the top curve the
heat capacity profile for ghosts in the presence of 5 mM Ca^{2+}.
The A, C and D transitons are apparent, with A centered at about
49.5°C, C at 66°C and D at about 81°C. From Lysko's experiments
at lower Ca^{2+} concentrations, it seems that B_1 and B_2 have merged
with A and C, respectively (14). As pyran is added, the effects
on B_1 and B_2 are reversed, although B_1 is flattened as it is by
pyran in the absence of Ca^{2+}. In addition, the C transition is
raised in temperature by about 3°. This again is not simply a
depletion of membrane Ca^{2+}, since even at 3.7 mg/ml of pyran
(more than a 10-fold excess of pyran carboxylate as compared to
Ca^{2+}) the profile has not returned to its form in the absence of
Ca^{2+}, shown as the bottom curve. Some association is indicated,
and this causes significant modification of the thermal transi-
tion behavior of the membrane.
   The importance of Ca^{2+} is also shown in experiments with pure
phospholipids. In this work, we use "liposomes," or multilayer
suspensions of a single lipid - dipalmitoylphosphatidylcholine
(DPPC) - to approximate the lipid bilayer of the natural membrane.
DPPC was chosen because the bulk of the phospholipids in the
outer monolayer of the red cell membrane are choline lipids.
This then is a much simpler system than is the intact erythrocyte
ghost, and it allows us to examine the interaction of the polymer
or polymer/metal complex with a single membrane component.
   Figure 6 shows our results for DPPC suspended in buffer at a
concentration of 1 mg/ml. The control suspension (lower curve)
shows the normal pre-transition at 35°C and the main melting
transition at 41°C. The main transition, with an enthalpy of
about 8 kcal/mole, represents the introduction of a number of
gauche bond rotations into the saturated acyl chains of the lipid,
which are all trans in the low temperature state. The smaller
pre-transition may be associated with reorientation of the phos-
phocholine headgroup, although there is not complete agreement
on this assignment.
   Addition of pyran copolymer (at a concentration of 1 mg/ml)
to the hydration buffer causes little, if any, perturbation of
the transition behavior - perhaps a slight increase in transition
half-width - whereas the addition of the same amount of pyran in
the presence of 5 mM Ca^{2+} causes each transition to shift up in
temperature - the main transition by 1.2°C, and the pre-transition
by 4°C. The main transition is also broadened, with the half-
width increased from 0.37°C to 0.62°C. Ca^{2+} alone cannot re-
produce this effect; in the presence of Ca^{2+} alone each transition
is shifted by not more than 0.5°C. We feel that this result
suggests a Ca^{2+}-promoted association of pyran copolymer with the
lipid surface. The greater perturbation of the "head-group"
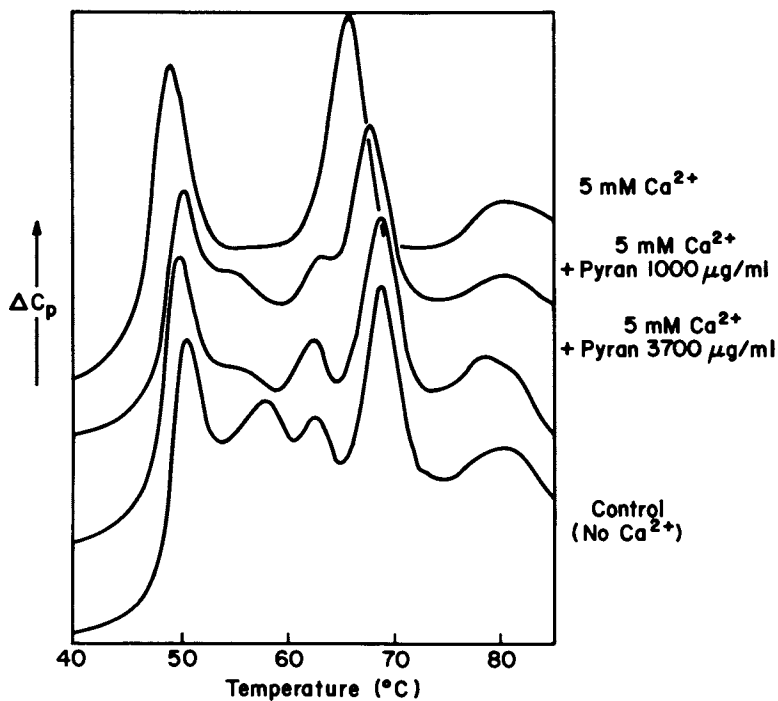
Figure 5.   Heat capacity profiles of erythrocyte ghosts suspended in isotonic caco-dylate buffer at pH 7.4.   The Ca^{2+} and pyran copolymer were added as shown.
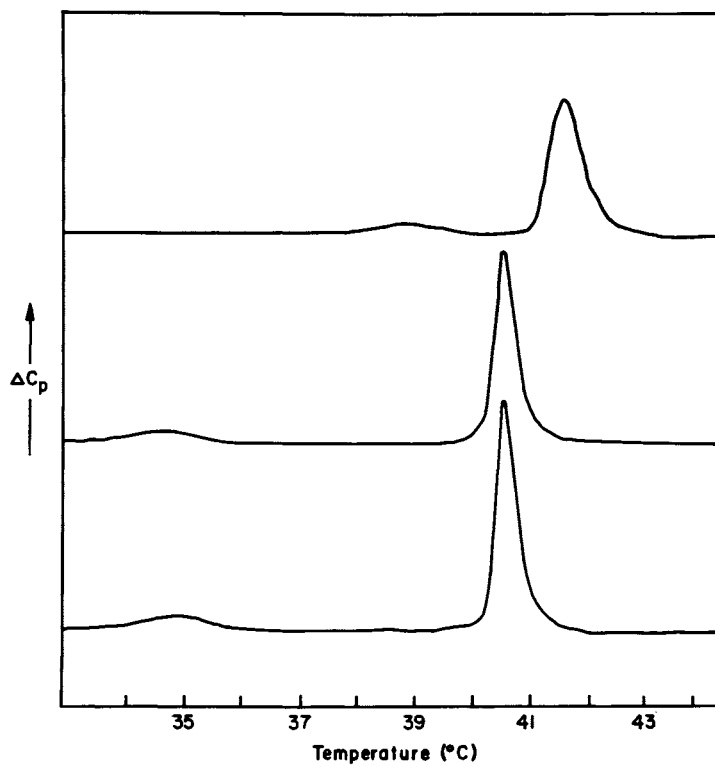
Figure 6. Thermal transition behavior of DPPC in multilamellar suspension in 50mM Tris at pH 7.4. Key: lower curve, DPPC 1 mg/mL; middle curve, DPPC 1 mg/mL + pyran copolymer 1 mg/mL; and upper curve, DPPC 1 mg/mL + pyran copolymer 1 mg/mL + CaCl$_2$ 5mM.

transition vs the hydrocarbon core transition suggests localiza-
tion of the polymer complex in the head-group region at the
surface, and we are pursuing complementary experiments using
lipid monolayers in order to test this hypothesis.

Conclusion

Ca^{2+} at concentrations typical for divalent ions in human
plasma, promotes the association of pyran copolymer with a pure
phospholipid (dipalmitoylphosphatidylcholine) and with intact
erythrocytes. This may suggest a more general mediation by
divalent ions of the cell surface interactions and pharmacological
properties of pyran copolymer.

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Thermodynamic Characterization of Proflavine Binding to DNA

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The interaction between DNA and proflavine was studied by
means of the heating of mixing. From the results, the thermo-
dynamic quantities were estimated for the DNA-proflavine system by
an intercalation process. The thermodynamic results show that the
intercalation process of DNA-proflavine system would be expected
to be dominated by the enthalpy rather than the entropy. In
addition, the heat of interaction depends on GC base composition
of DNA; the heat of interaction between GC base pairs and pro-
flavine is greater than that between AT base pairs and proflavine.

It is well known that aminoacridine dyes are bound to DNA in
two different ways depending on the concentration of dye: the
first type corresponds to strong binding[1] and/or intercalation
process[2] in which dye molecules are bound to the DNA base at
dilute concentration of dye. The other type corresponds to the
weak binding[1] and/or stacking process[2] in which dye molecules
are bound to the outside of DNA molecules without base specificity
as the concentration of dye increases.

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The interaction for the intercalation process seems to be
based on an interaction of flat aromatic dye molecules between
the base pairs of DNA as reported by many investigators[1-7].
However, the exact steric location of the intercalated dye and the
heat of interaction between DNA and dye have not been established
yet.

In this paper, we attempt to elucidate the thermodynamic
quantities of the interaction in the dye binding by intercalation
process and also explore the effect of GC base composition of DNA
on dye binding by measuring the heat of mixing of proflavine (PF)
and Cl. perfringens DNA (DNA I), E. coli DNA (DNA II) and
M. lysodeikticus DNA (DNA III), respectively because the inter-
action for intercalation process is based on the interaction
between dye and base pairs of DNA.

EXPERIMENTAL

Materials: The DNA's employed were Cl. perfringens DNA (31 % GC,
Sigma type XII), E. coli DNA (50 % GC, Sigma type VIII) and
M. lysodeikticus DNA (72 % GC, Sigma type XI) and used without
further purification. The dye used was proflavine (PF) which
was purchased from Aldrich Chemical Co., Inc. All measurements
were in 0.1 mol/l tris-HCl buffer solution (pH 7.60).

Apparatus and Procedure: The calorimeter which was used for the
measurement of the heats of mixing of DNA and PF was a LKB batch
type microcalorimeter (LKB-10700). For calorimetric measurement,
the DNA sample was dissolved into the buffer solution, and equal
volumes (about 1.2 cm^3) of DNA and dye solutions were mixed.
In this case, the concentration of DNA was kept at a definite
value of 4×10^{-4} mol/l of phosphate units of DNA while for the dye
solution various amounts of dye were used. The DNA concentration
was determined from molar extinction coefficient at 260 nm[8]
E_p=7400, 6500 and 7000 for Cl. perfringens DNA (DNA I), E. coli
DNA (DNA II) and M. lysodeikticus DNA (DNA III), respectively.
In order to obtain information about the binding parameter
for the intercalation process, the absorption spectra of
solutions containing a constant concentration of PF and varing
amounts of DNA were measured by using a spectrophotometer
(Perkin-Elmer, Model 552).

RESULTS AND DISCUSSION

Spectral measurement

Typical absorption spectra of solutions containing a definite
concentration of PF and various amounts of DNA are shown in
Figures 1(a), (b) and (c). As seen in Figures 1, the wavelength
of maximum absorbance for each system shifts to red as the
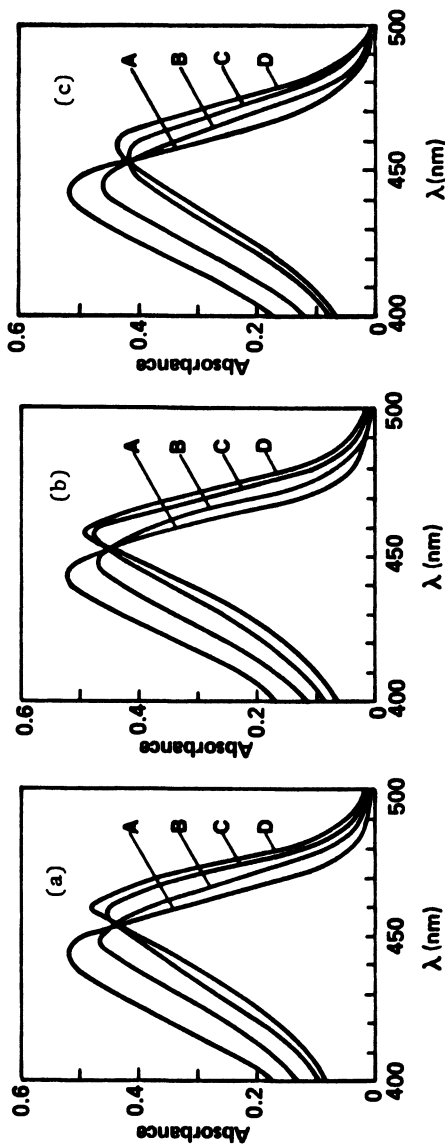
Figure 1. Typical absorption spectra of PF solution (1.6 × 10⁻⁵ mol/L) containing various amounts of DNA.

(a) DNA II–PF system shows DNA concentrations: A, 0; B, 5.36 × 10⁻⁵; C, 1.36 × 10⁻⁴; and D, 5.30 × 10⁻⁴ mol/L. (b) DNA I–PF system shows DNA concentrations: A, 0; B, 3.92 × 10⁻⁵; C, 9.79 × 10⁻⁵; and D, 3.92 × 10⁻⁴ mol/L. (c) DNA III–PF system shows DNA concentrations: A, 0; B, 4.01 × 10⁻⁵; C, 1.15 × 10⁻⁴; and D, 4.95 × 10⁻⁴ mol/L.

concentration of DNA increases and well-defined isosbestic points
for each system are located at about 454 nm which is compatible[16]
with the previous result[9] and results reported by Lee et al.[10].
The percentage of dye bound to DNA was calculated as described by
Peacocke and Skerrett[11]. The plots of r/C_f versus r for each
DNA-PF system are shown in Figures 2(a), (b) and (c). In this
case, r is the amount of PF bound per nucleic acid phosphate, and
C_f is the concentration of free dye. From those plots the
binding parameters were estimated and are given in Table I. It
is apparent that the binding parameters for each system are in
fair agreement, showing that there is no base composition of DNA
selectively in binding between DNA and PF.

Heat of Mixing

     The heats of mixing of DNA and PF measured at 298.15 K are
found to be exothermic, indicating the attractive interaction
between DNA and PF molecules. The results are shown in Figure 3,
where the heat of mixing per mole of dye, ΔH^M is plotted against
the mole ratio of PF to DNA, [dye]/[DNA]. ΔH^M, for each system
shows the sigmoidal curve such as ΔH^M increases (takes a smaller
negative value) as [dye]/[DNA], that is, PF concentration
increases. Those results are compared with the successive stages
of the intercalation process as described by Fredricq and
Houssier[12]. When [dye]/[DNA] is less than about 0.05, the
intercalated dye molecules do not interact with one another, in
the range from 0.05 to 0.13 of [dye]/[DNA] the intercalated dye
molecules do interact, and then dye molecules begin to bind to
the side of DNA molecules when [dye]/[DNA] is more than 0.13.
     To obtain the heat of interaction between DNA and PF from
calorimetric results, the following analysis is used.

Thermodynamic Quantities

     Assuming that the DNA-dye complex is formed by following
reaction process between DNA and dye,

$$\text{DNA} + \text{dye} \rightleftharpoons (\text{DNA} - \text{dye}) \tag{1}$$

the binding constant, K, is expressed as

$$K = \frac{r}{C_f (n - r)} \tag{2}$$
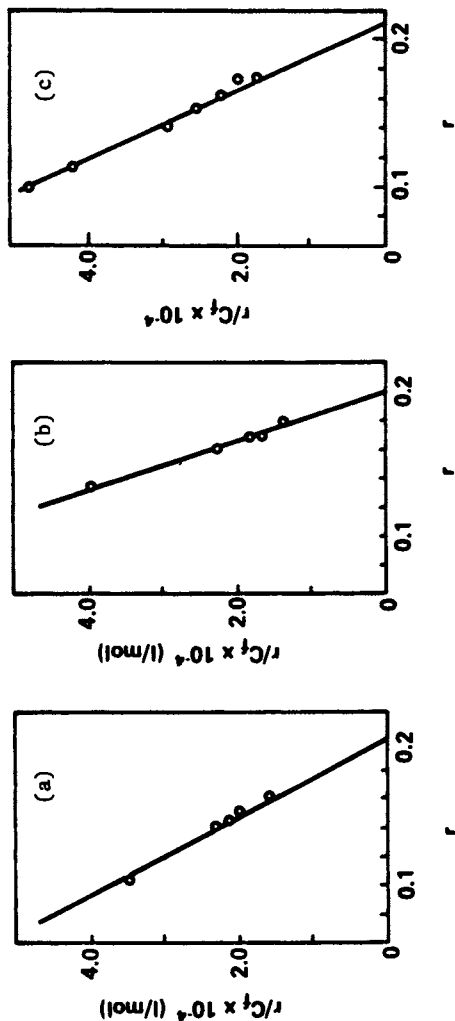
Figure 2. Plots of r/C_f against r for (a) DNA II–PF system, (b) DNA I–PF system, and (c) DNA III–PF system.

Table I   The binding parameters for the intercalation
process of the DNA-PF complexes estimated from spectro-
photometry

DNA	$K \times 10^{-5}$ [a]	n [b]
DNA I (GC 31 %)	5.6	0.20
DNA II (GC 50 %)	4.1	0.20
DNA III (GC 72 %)	6.0	0.21

[a] Binding constant
[b] Number of binding sites per DNA phosphate
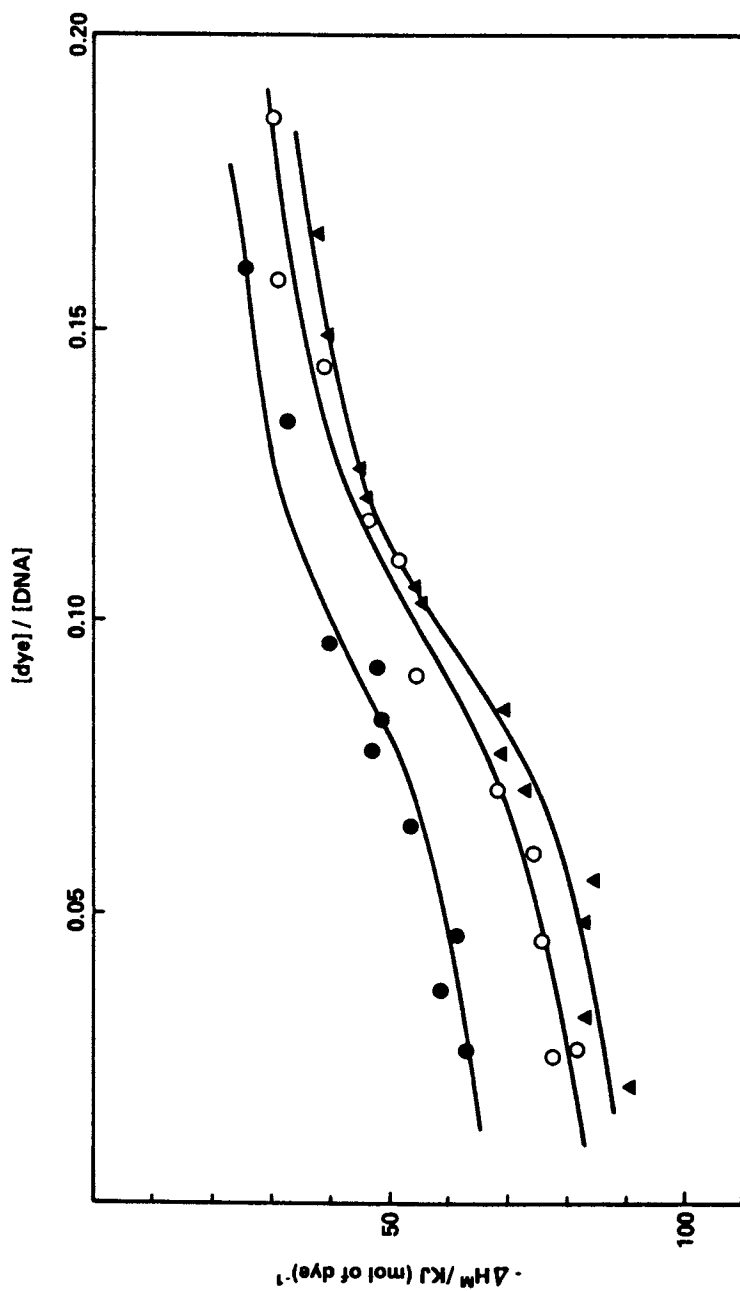
Figure 3. Plots of the heats of mixing, ΔH^M, against the mole ratio of PF to DNA, [dye]/[DNA], for DNA I–PF system (●), DNA II–PF system (○), and DNA III–PF system (▲).

where n is the number of binding sites per DNA phosphate, r the amount of bound dye per DNA phosphate and C_f the concentration of free dye.

From the experimental results of the heat of mixing, the heat of interaction of DNA-dye complex per mole of dye, ΔH, can be expressed as

$$\Delta H = \frac{\Delta Q}{C_b \, V} \qquad\qquad\qquad (3)$$

Where ΔQ is the observed heat, C_b the concentration of bound dye which is correlated with r and the concentration of DNA phosphate P, such as $C_b = rP$, and V the total volume of the mixture of DNA and dye solutions. The total concentration of dye, C, is the sum of C_b and C_f;

$$C = C_b + C_f \qquad\qquad\qquad (4)$$

From Eqs. (2) to (4), ΔQ will be represented by following complicated function of C, in which the negative sign of the right side must be adopted because correct zero value of $\Delta Q/V$ is given at zero of the concentration of dye, C.

$$\frac{\Delta Q}{V} = \frac{\Delta H}{2} \left[Pn + \frac{1}{K} + C - \sqrt{\left(Pn + \frac{1}{K} + C\right)^2 - 4PnC} \; \right] \qquad (5)$$

According to Eq. (5) we are able to calculate the value of $\Delta Q/V$ for a given value of C by using both the values of K and n estimated from spectrophotometry. The ΔH value which gives the best fit between the calculated value and the experimental one of $\Delta Q/V$ is adopted. The ΔH values adopted are about -70, -85 and -90 kJ/mol for DNA I, DNA II and DNA III, respectively. But the calculated value of $\Delta Q/V$ deviates from the experimental value in the concentration region of dye more than about 3×10^{-5} mol/l as shown in the broken lines in Figure 4. Those deviations may be a result of K and n values estimated from spectrophotometry in which the concentration of DNA is different from that in calorimetry.

To improve those discrepancies, K and n can be treated as variable parameters and adjusted in such a way that the best fit between experimental and calculated values of $\Delta Q/V$ is obtained. These quantities are plotted against C as a solid line in Figure 4 and the values of ΔH, K and n estimated are listed in Table II. The K values for each system estimated from calorimetry are in agreement with those from spectrophotometry, but the n values estimated from calorimetry are a little lower than those from spectrophotometry. The decrease in n values may be ascribed to the intermolecular interaction of DNA which increases with the concentration of DNA and brings about the decrease of effective binding sites of DNA.
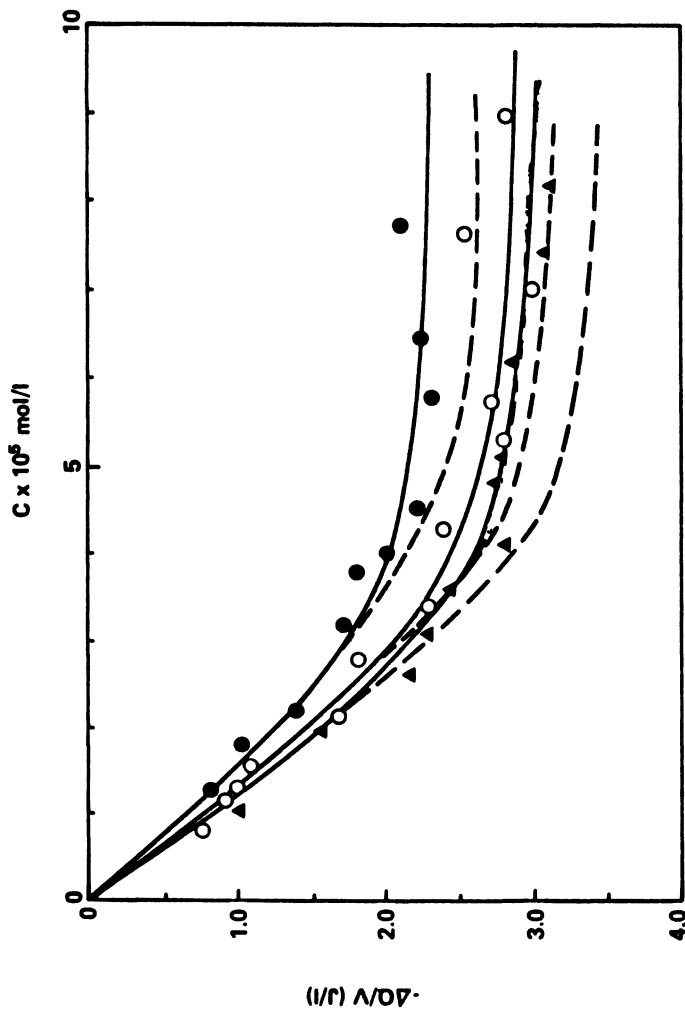
Figure 4. Dependence of the observed heat of mixing, ΔQ/V on the dye concentration, C, for DNA–PF system. Key: (●) DNA I–PF system; (○) DNA II–PF system; and (▲) DNA III–PF system; theoretical ΔQ/V curve calculated with Equation 5 using K and n values in Table I (– – –) and in Table II (———).

Table II   The thermodynamic quantities for intercalation process of DNA - PF system estimated from calorimetry at 298 K

DNA	$K^{a)}$ $l\ mol^{-1}$	ΔG $kJ\ mol^{-1}$	ΔH $kJ\ mol^{-1}$	ΔS $J\ K^{-1}\ mol^{-1}$	$n^{b)}$
DNA I(GC 31 %)	$5.6\ X10^{5}$	-33	-70	-120	0.17
DNA II(GC 50 %)	$4.1\ X10^{5}$	-32	-85	-180	0.18
DNA III(GC 72 %)	$6.0\ X10^{5}$	-33	-90	-190	0.18

mol refers to the number of moles of PF
a) Binding constant
b) Number of binding sites per DNA phosphate

The free energy change, ΔG, and the entropy change, ΔS, for each system are easily calculated from the following equations,

$$\Delta G = -RT \ln K, \qquad\qquad \Delta S = (\Delta H - \Delta G)/T \qquad\qquad (6)$$

and are listed in the third and fifth columns in Table II, respectively.

As seen in Table II, the values of ΔH for each system are different and decrease with decreasing GC base composition of DNA although the values of ΔG are nearly equal. The ΔH dependence on GC base composition of DNA seems to be based on difference of the heat of interaction between GC base pairs and PF and between AT base pairs and PF.

Base composition effect of heat of interaction

Assuming that the three types of DNA/dye interaction (i.e. between GC pairs, AT pairs and GC-AT base pairs) occur in numbers given by the probability of existence of such pairs when DNA molecule corresponds to a random chain, then the heat of interaction, ΔH, is represented as

$$\Delta H = n_A^2 \Delta H_1 + n_B^2 \Delta H_2 + n_A n_B \Delta H_3 \qquad\qquad (7)$$

where ΔH_1, ΔH_2 and ΔH_3 are the heats of interaction between GC base pairs and dye, AT base pairs and dye and GC-AT base pairs and dye, respectively, and n_A and n_B are fractions of GC and AT base pairs in DNA molecule. Since n_B is equal to one minus n_A, ΔH is also written as the function of n_A,

$$\Delta H = (\Delta H_1 + \Delta H_2 - 2\Delta H_3)n_A^2 + 2(\Delta H_3 - \Delta H_2)n_A + \Delta H_3 \qquad (8)$$

The plots of ΔH estimated from calorimetry versus GC fraction of DNA are shown in Figure 5, and the solid line in Figure 5 shows the best fit curve of Eq. (8). From this curve, we can obtain $\Delta H_1 = -96$, and $\Delta H_2 = -50$ kJ/mol.

Conclusion

From the thermodynamic quantities estimated from calorimetry, the intercalation process of DNA – PF system would be expected to be dominated by the enthalpy rather than entropy because of negative value of the heat of interaction and positive value of entropy change for the interaction between DNA and PF. The heat of interaction between DNA and PF is greater than that between DNA and acridine orange[13]. This greater heat of interaction may be based not only on interaction between base pairs of DNA and PF but on hydrogen bonding between phosphate group with negative charge of DNA and amino group on 3 and/or 6 position of PF as reported by Peacocke et al.[14].
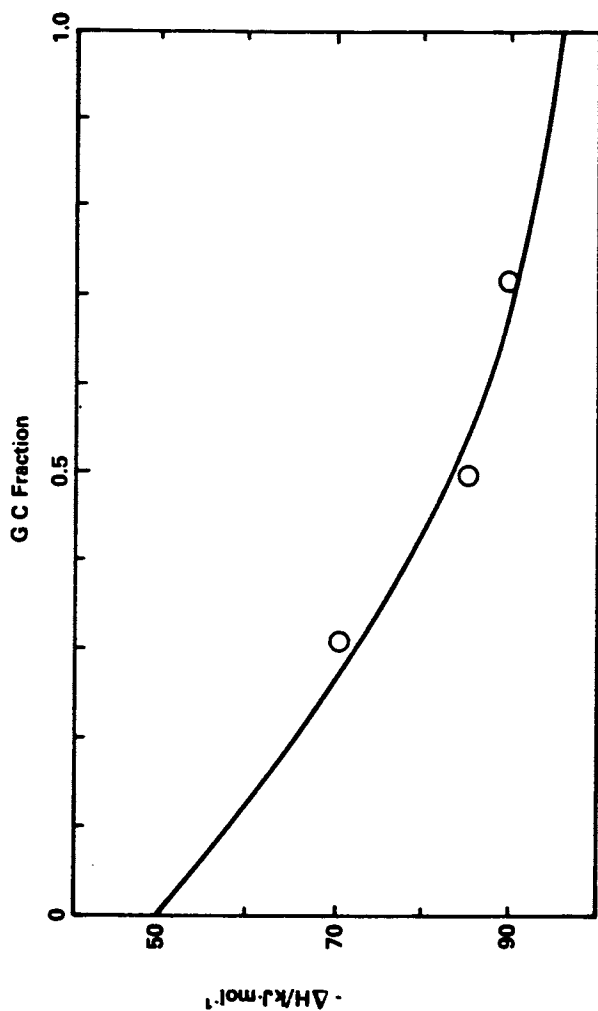
Figure 5.   Dependence of the heat of interaction, ΔH, on GC base composition of DNA.

In addition the dependence of the heat of interaction of GC base concentration of DNA may be based on the difference of the heat of interaction between GC base pairs and PF and AT base pairs and PF because the heat of interaction of GC base pairs and PF is greater than that between AT base pairs and PF.

Acknowledgment

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Polymers for Potential Cancer Therapy: A Brief Review

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A polymeric drug is a polymer that contains a
drug unit either as part of the polymer backbone,
as a terminal group, or as a pendant unit off the
polymer backbone. In some cases, a polymer can
function as a drug even though there are no low
molecular weight analogs of that polymer. Polymeric
drugs can be subdivided into (a) insoluble polymers,
(b) soluble polymers and (c) directed polymers.
Many examples have been reported of polymers which
exhibit antineoplastic activity. In this review,
antineoplastic polymers are subdivided into the
classes: (a) polyanions, (b) nucleic acid analogs
and (c) all other systems. Examples are cited for
each of these different catagories of antineoplastic
polymers.

The potential use of polymeric materials in treating disease
is now well documented by many research papers and several re-
views (1-10). Basically, there are three ways that polymers can
be used in disease therapy: (a) controlled release systems, (b)
mechanical or diffusion controlled pumps and related devices, and
(c) polymeric drugs. A number of other papers in this volume
will consider the controlled release of chemotherapeutic agents
and some recent reviews (11-14) have covered the topics (a) and
(b). This paper will concentrate only on the polymeric drug
approach to the treatment of neoplastic diseases.

Polymeric Drug Therapy

The basic concept of polymeric drug therapy has been review-
ed recently (1,3,5,9). A polymeric drug is a polymer which
contains a drug unit either as part of the polymer backbone, as a
terminal unit, or as a pendant unit off the polymer backbone. In
some cases, the polymer itself functions as the therapeutic agent,

whereas the low molecular weight analogs are inactive. There are
three distinctly different types of polymeric drugs: (a) insoluble
polymers, (b) soluble polymer systems, and (c) directed polymer
systems. In all three types it is assumed that the polymers would
be biologically active rather than being a source of slow release
of the drug. Generally the insoluble polymeric drugs would be
implanted at or near the disease site and would interact with
this disease. Since the material would be fairly insoluble in the
body fluids, there would be very little interaction with other
body tissues or organs. Although this technique would aid in
reducing the toxic side effects of the drug such implantation
would generally require surgery. Drug activity would be prolonged
largely because of the polymer's lack of solubility which would
retard migration and excretion. The insoluble polymeric drugs,
which are usually the easiest of the three types to prepare,
would thus have some advantage over a simple drug although this
advantage would probably be of limited value since surgical
implantation would usually not be desirable.

The soluble polymeric drugs, on the other hand, would be
capable of solution in the body fluids and could be introduced
orally or by injection. In simple, non-polymeric drugs, the only
way to change the solubility characteristics would be to modify
the chemical structure of the drug, itself although this might
also change the therapeutic characteristics as well. Generally
in a soluble polymeric drug system, the therapeutic and solubiliz-
ing parts of the molecule are located in different parts of a
copolymer chain and modification of one portion would have little
effect on the other. Such copolymers could be prepared by
copolymerization of a monomer containing the drug unit with anoth-
er monomer containing groups which would make the copolymer
soluble in the body fluids. Unfortunately, since the resulting
polymeric drug is soluble in the body fluids, it could migrate
throughout the body about as well as the original drug and could
still show toxic side effects. These systems could, however,
exhibit prolonged activity and could be active at a lower total
concentration since the fact that the drug is polymeric would
increase the concentration greatly on a localized scale in the
solution. From the standpoint of ease of usage, these would be
better than the insoluble polymeric drugs.

The directed polymeric drugs would consist of at least three
functional units although lesser number of units might suffice.
Basically directed polymeric drug would contain (a) the drug unit,
(b) a unit to make the copolymer soluble, and (c) a unit to direct
the copolymer to a specific type of diseased tissue or organ. In
the case of an antineoplastic polymer, this directing unit would
be some unit that would tend to migrate selectively into the
tumorous tissue and would thereby carry the copolymer along with
it. These systems could be administered orally or by injection
of a solution and, since the polymer would go selectively to the
diseased site, such directed polymeric drugs would have minimal

interactions with other organs which would result in the reduction of the toxic side effects. Directed polymeric drugs would be the most effective type of polymeric drugs but would be the most difficult type to design.

Numerous antineoplastic polymers have been prepared with varying degrees of activity. For convenience, these polymers are divided here into (a) polyanions, (b) nucleic acid analogs, and (c) other systems. Most of these polymers are of the insoluble or soluble class but much recent advances have been made toward developing directed polymeric drugs. Several workers have utilized controlled release systems for the release of anti-tumor drugs although this is beyond the scope of this present review (11,15).

Polyanionic Antineoplastic Polymers

Numerous polyanionic materials have been examined for potential biological activity (6,16). The rationale for this approach probably arises from the fact that many naturally-occurring, biologically-active polymers, such as heparin, DNA, various proteins, alginic acid and pectin, are polyanionic polyelectrolytes. Statolon, which is a polyanionic, polysaccharide, has been shown to supress leukemia in mice (17). The first synthetic polyanionic polymer to be examined extensively for antitumor activity appears to be sodium poly(vinyl-sulfonate) (I) which inhibited the following types of tumors in mice: adenocarcinoma 775, Ehrlich (ascites), Krebs 2 carcinoma (ascites), L1210 lymphoid leukemia, L5178 lymphatic leukemia and sarcoma 180 (6,18). Subsequently it was shown that poly(acrylic acid) (II), poly(methacrylic acid) (III), and the copolymer of ethylene-maleic anhydride (IV) had antineoplastic activity (19). Unfortunately, most of these polymers were extremely toxic and could not be tested clinically in humans.

$$\{CH_2-CH\}_n$$
$$| $$
$$SO_3^{\ominus} Na^{\oplus}$$

(I)

$$\{CH_2-CH\}_n$$
$$| $$
$$COOH$$

(II)

$$\begin{array}{c} CH_3 \\ | \\ \{CH_2-C\}_n \\ | \\ COOH \end{array}$$

(III)

$$\{CH_2CH_2-CH-CH\}_n$$
$$\underset{O}{\overset{C}{\diagup}}\underset{O}{\overset{}{\diagdown}}\underset{O}{\overset{C}{\diagup}}$$
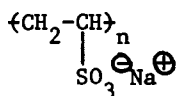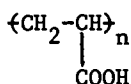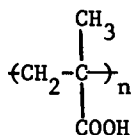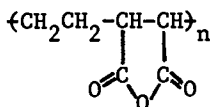
(IV)

One of the most interesting polymers of this type is the 1:2
divinyl ether: maleic anhydride copolymer (DIVEMA) (V) and the
hydrolysis product (VI). Polymer (V) was first reported by
Butler in 1960 (20) and has since been shown to exhibit an incred-
ibly wide spectrum of biological activity including antitumor,
antiviral, antibacterial, antifungal, anticoagulant and anti-
inflammatory activity (6). While this polymer is shown below as
the six-membered pyran ring, this structure has not been fully
proven and some other cyclopolymerization products do give five-
membered rings instead (16,21).

(V)   (VI)
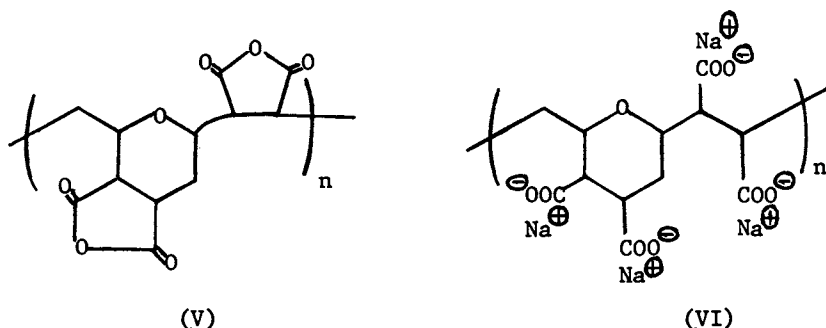
Polymer (VI) has been shown to have antitumor activity
against adenocarcinoma 755, Dunning ascites leukemia, Friend
leukemia virus, and Lewis lung carcinoma. In the latter case,
polymer (VI) showed activity about equal to that of cyclophos-
phamide (an alkylating agent) and was more effective than
6-mercaptopurine (an antimetabolite) (6). The DIVEMA polymer (V)
is also active against some cancer causing viruses such as Friend
leukemia, Moloney sarcoma and Rauscher leukemia (6).
    There has been much speculation concerning the mechanism of
tumor inhibition by these polyanionic systems. Many polyanions
induce interferon production (22) and this fact has also been
suggested as the source of the activity. More recently, it has
been observed that DIVEMA is not cytotoxic to tumor cells (23)
and that only those polymers that activated macrophages showed
antineoplastic activity (24). Inhibition via macrophages is
believed to be the mode of action for DIVEMA (V) if not for all
polyanionic polymers (16).

Nucleic Acid Analogs as Antineoplastic Polymers

A wide variety of polymers have been prepared containing a
nucleic base unit on the polymer chain. In most cases this was
achieved by polymerizing a monomeric derivative of the nucleic
base. The resulting polymers are similar in some respects to
DNA or RNA and could have utility in cancer treatment by interfer-
ing with tumor cell replication.

The most extensive work in this area has been on the vinyl-
type derivatives of uracil and adenine and this research has been
reviewed recently (8,25,26). These monomers include 1-vinyl-
uracil (VII) and 9-vinyladenine (VIII) (27,28); methacryloylethyl
derivatives of uracil (IX) and adenine (X) (29,30); 1-(N-vinyl-
carbamoyl)uracil and its 5-fluoro- derivative (XI) (31-34);
9-(N-vinylcarbamoyl)-methylthiopurine (XII) (34-36); 9-vinyl-6-
alkylthiopurine (XIII) (37); and styrene derivatives of uracil
(XIV) and adenine (XI) (38,39).
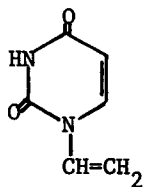
(VII)
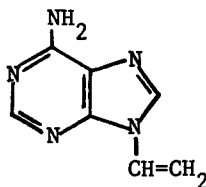
(VIII)
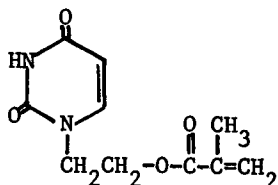
(IX)
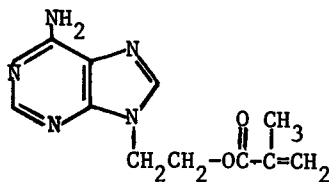
(X)
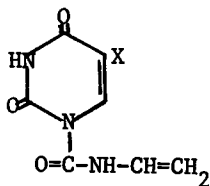
(XI)

(a: X = H)
(b: X = F)
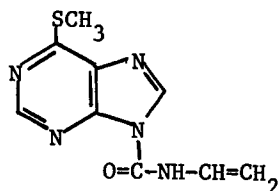
(XII)
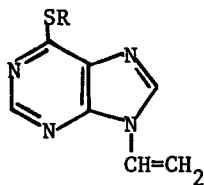
(XIII)
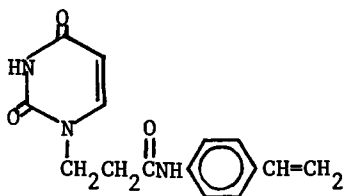
(XIV)
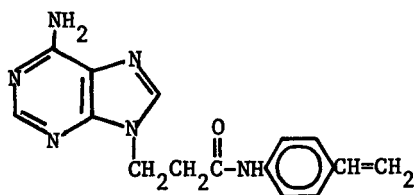
(XV)

Monomers (VII) and (VIII) have been studied the most exten-
sively. The original work on polymerizing (VII) lead to an
intractable, insoluble, cyclopolymerized polymer (40) but sub-
sequent work lead to linear, water soluble polymers (41,42).
Monomer (VIII) readily polymerizes under free radical conditions
to form linear, water soluble polymers (27,43,44). The polymers
of (VII) and (VIII) form complexes with each other in a manner
similar to the natural nucleic acids (25,26) and they do show
antitumor activity against murine leukemia virus (45,46).
    Monomers (XI) and (XII) polymerize readily under free radical
conditions to yield uncrosslinked polymers (31-36). The polymer
from (XIb) was found to inhibit P388 leukemia in mice but it was
uncertain whether this was due to activity of the polymer or to a
slow release of 5-fluorouracil since this polymer (and monomer) do
slowly hydrolyze in an aqueous medium (34). Monomers (IX), (X),
(XIII), (XIV) and (XV) also polymerize but no biological data
have been reported.
    The first 5-fluorouracil containing polymer had the 5-FU
units in the polymer backbone and this structure is shown below
as (XVI). This was claimed to be biologically active (47). More
recently, monomer (XVII) was prepared by the reaction of 5-FU
with methyl fumaroyl chloride. The polymers and copolymers of
(XVII) do show antitumor activity but this may be due to the
hydrolysis of this unit to release 5-FU (48).
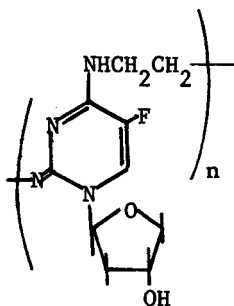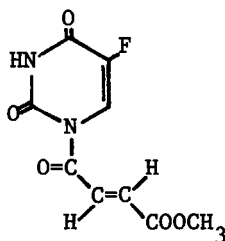
(XVI)   (XVII)

    The first reported example of a polymer containing a 6-mer-
captopurine derivative involved attaching the methylthio deriva-

tive to poly(vinyl alcohol) as shown in (XVIII). This water
soluble polymer does form complexes with uracil containing polymers
but no biological data was reported (49). Many other polymeriz-
able derivatives of nucleic bases have been prepared including
amino acid (50) and NCA derivatives (51) and derivatives of
thymine, hypoxanthine and theophylline (8,25) but no information
regarding possible antineoplastic activity is available.
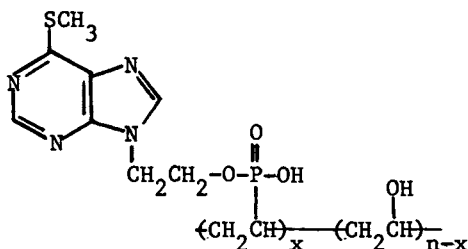
(XVIII)

It is obvious that this approach using nucleic bases can lead
to antineoplastic polymers since several such systems have shown
this activity (34,45,46,48). This is not totally unexpected since
the nucleic base derivatives 5-fluorouracil and 6-mercaptopurine
are antineoplastic agents. The polymers of this type made to date
have been either the insoluble type or the soluble type polymeric
drug systems. This approach should be especially effective if
some directing group is also put into these polymers to carry them
selectively into tumorous tissue. Research along this line is in
progress in our laboratories and some other places as well.

Other Antineoplastic Polymers

Many other polymers with antineoplastic properties have been
prepared. For the most part, these polymers contain a known
antineoplastic compound attached to the polymer chain as a pendant
group. The earliest examples of this type seem to be the aziridine
alkylating group containing polymers (XIX) (52), and (XX) (53),
and (XXI) (54) which contains the cytostatic agent sarcolysin.

(XIX)

(XX)

$$-(CH_2CH)_n-$$
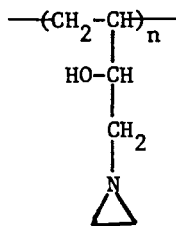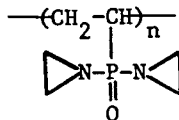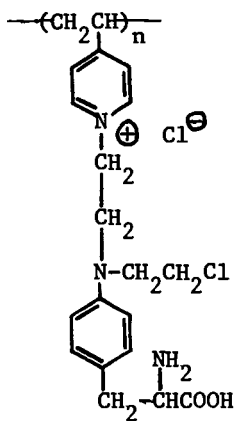
(XXI)

Other potential anticancer polymers also include the tropene derivative (XXII) (55,56) and the derivative of cyclophosphoramide (XXIII) (9,57,58). Monomer (XXIII) has also been copolymerized with solubilizing groups, such as N-vinylpyrrolidone or vinyl-pyridine-N-oxide, and some potential directing groups, which were polymerizable derivatives of sulfa drugs, to attempt to make a directed polymeric antineoplastic agent. This directed polymeric drug was initially claimed to migrate selectively into tumorous tissue (9).
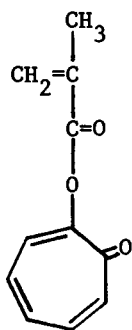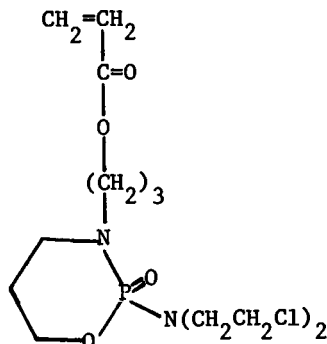
(XXII)   (XXIII)

The antineoplastic agents methotrexate and adriamycin have been bound to various protein substrates (59) and to DNA (60) and longer duration of the biological activity was noted compared to the monomeric drugs. The concept of affinity polymeric drugs, in which the drug unit is attached to some specific polymeric entity with affinity for some particular part of the body, has been reviewed recently (61). Polymeric derivatives of the anticancer

agent <u>cis</u>-dichlorodiammineplatinium II have been prepared having
the structure (XXIV) (<u>62,63</u>) and (XXV) (<u>64</u>). Polymer (XXIV) was
found to be active against mouse P388 leukemia and Ehrlich
ascites tumors (<u>65</u>) while polymer (XXV) was active against mouse
929 tumor cells and HeLa tumor cells (<u>64</u>).
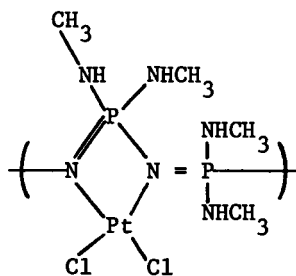
(XXIV)
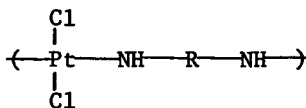
(XXV)

Conclusions

The present, brief review clearly shows that several different
types of polymers can exhibit antineoplastic activity. It is not
possible, at the present time, to state with certainty which
polymeric system(s) will eventually prove to be the most useful
in cancer therapy, but a directed polymeric antineoplastic agent
would probably be the most desirable, if such a system can
actually be synthesized. Some of the areas highlighted in this
review will be considered further in some of the subsequent
chapters of this book.

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The Antitumor and Antiviral Effects of Polycarboxylic Acid Polymers

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Anionic and cationic polyelectrolytes of both natural and synthetic origin have been found to exhibit an inhibitory effect on viruses, bacteria, tumors, and enzymes (1). Polyanions, in particular, have a broad range of biological activity and have received considerable interest in the areas of oncology and virology. The prolonged protective action of synthetic polyanions when given prior to virus inoculation has tremendous clinical potential; consequently, establishing an impetus for assaying the fundamental role of polyanions in controlling host resistance to a variety of pathophysiology.

Synthetic polyanions, in particular, are known to produce a wide spectrum of effects on immune reactivity (1). They have been shown to induce production of interferon (2) modify RES function (3) and to have immunoadjuvant (4) antiviral (5) and antitumor activity (6). The antineoplastic activity of pyran copolymer has been largely attributed to its ability to activate macrophages (7).

One of the most interesting anionic polymers is pyran copolymer which is prepared from divinyl ether and maleic anhydride through copolymerization. It has been under investigation in cancer chemotherapy for several years and is designated as NSC 46015 by the National Cancer Institute. It shows a variety of biological activities and has elicited a considerable amount of interest by several researchers in different areas. Pyran copolymer is an inducer of interferon (8-11); it has activity against a number of viruses (6-14), including Friend leukemia, Rauscher leukemia, Moloney sarcoma, polyoma, vesicular stomatitis, Mengo, encephalomyocarditis, MM, and foot-and-mouth disease; it has antibacterial (21-23) and antifungal activity (21); it stimulates immune response (18-24); it inhibits adjuvant disease (31); it is an interesting anticoagulant (32); and it shows promise in removing plutonium from the liver (33). However, the toxicity of the copolymer, although much lower than that of other anionic polymers which have been investigated, was apparently still too high for it to be used extensively in clinical investigations

(34,35). Recent studies have shown that low molecular weight
fractions and the calcium salt of the polymer are much less tox-
ic. This has stimulated further clinical investigation of this
drug.

Antitumor and Antiviral Activity of Polycarboxylic Acid Polymers

We have found that the structure and the molecular weight of
the polycarboxylic acid polymers play an important role in the
biological activity elicited by these materials (1). These find-
ings have been coorborated by Breslow (36) who has reported that
fractionated pyran shows lower toxicity with lower molecular
weight fractions with very little variation in antitumor activity.
Hodnett (37) has also prepared a number of polycarboxylic acid
copolymers which gave varying activity with structure against
ascites sarcoma 180 tumor. The average increase in survival time
was 25%. He found that the most effective polymers were those
with low molecular weight, high carboxyl group density and low
pKa at physiological pH.
    We have prepared several different molecular fractions of
pyran, poly(acrylic acid-co-maleic acid) PAAMA, poly(maleic acid)
PMA, and poly(acrylic acid-co-3,6-endoxo-1,2,3,6-tetrahydrophtha-
lic acid) BCEP and evaluated their activity against Lewis lung
carcinoma (Table I) and encephalomyocarditis (EMC) virus (Table II).

Table I. Inhibition of Tumor Size with Polymer Molecular Weight[a]

POLYMER[b]	PYRAN (%)	PAAMA (%)	PMA (%)	BCEP (%)
Whole Polymer	78	74	75	30
(Increased life span)	(40)	(33)	(15)	(<10)
1,000-10,000 (A)[c]	89	-	74	22
10,000-30,000 (B)	84	80	72	-
30,000-50,000 (C)	74	-	76	-
50,000-100,000 (D)	70	78	74	30

[a]BDF, mice were inoculated with 10^6 Lewis' lung cells into right-
hind gluteus muscle. Polymers were administered daily by the
intraperitoneal route for 10 consecutive days following tumor
implantation. Primary tumor size was determined on day 14 (% in-
hibition) and increased life span (% ILS) calculated from mean
time to death.

[b]Poly(divinylether-co-maleic anhydride)(Pyran); Poly(acrylic acid-
co-maleic anhydride) (PAAMA); Poly(maleic anhydride) (MA); Poly(

acrylic acid-co-3,6-endoxo-1,2,3,6-tetrahydrophthalic anhydride)
(BCEP).

[c]FRACTION A = PM-10 Filtrate passed through Amicon PM-10 filter
and not UM-2
  FRACTION B = PM-30 Filtrate passed through Amicon PM-30 filter
and not PM-10
  FRACTION C = XM-50 Filtrate passed through Amicon XM-50 filter
and not UM-2
  FRACTION D = Whole Polymer passed through Amicon XM-100 filter
and not XM-50

Table II.   Change in Antiviral Encephalomyocarditis Protection with
Polymer Molecular Weight[d]

POLYMER	PYRAN (%)	PAAMA (%)	PMA (%)	BCEP (%)
Whole Polymer	89	90	30	25
A 1,000-10,000	0	0	0	0
B 10,000-30,000	30	29	0	-
C 30,000-50,000	58	54	<10	-
D 50,000-100,000	86	80	26	38

[d]Mice were inoculated intravenously with 10 LD_{50} of encephalomyo-
carditis virus 24 hours after administration (i.v.) of 25 mg/kg
of polymer. Percent protection based on 20 mice/group calculat-
ed from mean time to death.

Polymer structure changes showed different effects in tumor size.
All the polymers reduced the tumor growth by approximately 75%
except for BCEP which only inhibited the growth by 30%. More im-
portantly each polymer had a different effect on the increased
life span (ILS) of the animal with pyran being the most effec-
tive (40% ILS) and BCEP the least (10% ILS). The molecular weight
did not have significant effect on the antitumor activity but did
have a great effect on toxicity of the material which will be dis-
cussed later.
      The polymer structure had a more significant effect on anti-
viral activity (Table II). Pyran and PAAMA produced approximate-
ly the same activity while PMA and BCEP were very poor. Further,
molecular weight had a dramatic effect. The lower molecular
weight fractions (<30,000) were ineffective and only the higher
molecular weight materials (>30,000) were active.
      In a more recent study by Munson et. al. (38), it was shown
that pyran (Table III) elicited better cytotoxicity against Lewis
lung with the lower molecular weight fractions. It was also

Table III.   Relationship Between Molecular Weight, Macrophage
             Activation, Antitumor Activity and Antiviral Activity

MW Pyran Polymer	Percent cytotoxicity[e]		Antitumor[f] Activity P815 mastocytoma % ILS	mg/kg Pyran[g] Required for 100% Protec- tion Against EMC
	Lewis lung	mouse embryo fibroblast		
12,500	100	0	126	<75
15,500	94	0	126	25
21,300	94	0	179	12.5
32,000	81	0	179	6.3
52,600	88	0	162	3.1
NSC46015 (pyran)	88	0	not done	3.1

[e] Peritoneal exudate cells were removed and tested 7 days after i.p. injection of pyran polymers at 25 mg/kg for cytotoxicity against Lewis lung and secondary mouse embryo cells.

[f] Mice were inoculated in the footpad with 5×10^5 P815 mastocytoma cells.

[g] Mice were injected i.v. with pyran polymers and challenged i.v. 24 hours later with 10 LD_{50} of EMC virus.

shown in this study that there was no cytotoxicity of the natural mouse embryo fibroblast with any pyran; thus the activity elicited by pyran was tumor cell specific. Interestingly, increased life span was observed for the high molecular weight pyran in treatment against P815 mastocytoma. Furthermore, much less higher molecular weight (52,600) pyran (3.1 mg/kg) was required to produce 100% protection against EMC whereas it required 75 mg/kg of the 12,500 molecular fraction to produce the same result.

     More recently, we have evaluated a number of maleic anhydride copolymers (Table IV) and 2,3-dicarboxynorborn-5-ene copolymers (Table V) against Ehrlich ascites tumor (33). In both of these systems, very little or no activity was observed for the 10,000-30,000 molecular weight fraction compared to pyran. However, the lower molecular weight fractions (1,000-10,000) all produced good activity and in many cases were better than pyran.

Table IV. Effect of Maleic Anhydride Copolymers Against Ehrlich Ascites Tumor Cells[h]

POLYMER	10,000 - 30,000 Molecular Weight		1,000 - 10,000 Molecular Weight	
	Cells/pc x 10^8	% Control[i]	Cells/pc x 10^8	% Control
Saline	1.52	100	2.17	100
Pyran	0.74	49	.63	30
MA-co-Acrylic Acid	1.8	118	1.25	58
MA-co-STYRENE	0.76	45	0.15	7
MA-co-Methracylic Acid	1.42	93	1.31	60
MA-co-Allyl Phenol	1.31	86	0.35	16
MA-co-Allyl Succinic Anhydride	1.32	87	0.91	42
MA-co-1,3-Dioxepin	1.43	94	0.63	30
MA-co-Isobutenyl Succinic Anhydride	1.21	80	1.33	61

h) 10^6 Ehrlich Ascites cell were inoculated i.p. into mice.

i) Five days after inoculation, animals were sacrificed and total peritoneal exudate cells were counted with a hemocytometer.

Table V. Effect of 2,3-Dicarboxynorborn-5-ene
          Copolymers Against Ehrlich Ascities
          Tumor[j]
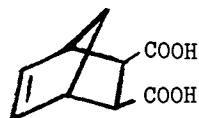
COPOLYMER	10,000 - 30,000 Molecular Weight		1,000 - 10,000 Molecular Weitht	
	Cells/pc x 10^8	% Control[k]	Cells/pc x 10^8	% Control
Control	1.52	100	2.17	100
Pyran	0.74	49	.64	30
Maleic Anhydride	1.64	1.08	0.16	7
Acylic Acid	1.61	1.06	0.48	22
Vinyl Acetate	0.89	41	0.76	34
Vinyl Alcohol	0.85	56	0.71	32

j) 10^6 Ehrlich Ascites cells were inoculated i.p. into mice.

k) Five days after inoculation, animals were sacrificed and total
    peritoneal exudate cells were counted with a hemocytometer.

    Consequently, we are observing two significant effects with
regard to antitumor and antiviral activity. These effects appear
to be related in most cases, to the polymer structure and mole-
cular weight.

Toxicological Effects of Polycarboxylic Acid Polymers

    The toxicity of polycarboxylic acid polymers is more sensi-
tive to polymer structure and molecular weight than antitumor
activity. The acute toxicity is markedly decreased with decreas-
ing molecular weight (Table VI and XI). Further, polyanion poly-
mers have been shown to highly sensitize mice to bacterial endo-
toxin (39); this activity is strikingly molecular weight depen-
dent as shown in Tables VII and XI as well as dose dependent (38).
The enhanced sensitivity to endotoxin by synthetic anionic poly-
mers has been extensively studied, but the mechanism of action is
still not understood (39).

Table VI. Polymer Molecular Weight Effect on Acute Toxicity (LD_{50})[1]

POLYMER	PYRAN	PAAMA	PMA	BCEP
Whole Polymer	74	110	120	150
A 1,000–10,000	120		160	>200
B 10,000–30,000	115	>200	150	-
C 30,000–50,000	95		140	-
D 50,000–100,000	84	180	135	180

1) LD_{50} was determined by administering the polymer in mg/kg intraveneously and the mortality recorded 24 hours later.

Table VII.   Effect of Endotoxin Sensitization With Polymer Molecular Weight[m]

POLYMER	PYRAN	PAAMA	PMA	BCEP
Whole Polymer	0.12	1.0	15	3.0
A 1,000–10,000	15	-	>20	>10
B 10,000–30,000	15	-	>20	>10
C 30,000–50,000	-	>20	>20	>3
D 50,000–100,000	-	>20	>20	>3.0

m) LD_{50} of S. typhosa 0904 lipopolysaccharide 24 hours after a single administration of 24 mg/kg of polymer. The dosage of endotoxin was administered in mg/kg.

Molecular weight toxicities are also observed in the liver (Tables VIII and IX) as well as in the spleen as shown in Table X (1). These effects are also polymer structure and molecular weight dependent. The larger molecular weight polymers seem to be localized in the liver and in the spleen to a greater extent than the lower molecular weight polymers; which appear to affect the reticular and other cells of the spleen as well as the Kupffer cells, mixed microsomal enzymes and other factors in the liver (1).
     In a recent study by Munson, White and Klykken (38), it was reported that the acute toxicity of pyran was a function of salt

form. It was demonstrated that the calcium salts were less toxic
than the sodium salts of pyran and that the toxicity in both
cases was molecular weight dependent (Table XI) as well as dose
dependent (38). In contrast to the significant differences in a-
cute toxicity observed for the two salts, no significant effects
were seen on the sensitization to bacterial endotoxin (Table XI)
clinical chemistry, or antiviral activity (38).

Table VIII. Polymer Molecular Weight Effect on Liver Weight[n]

POLYMER	PYRAN	PAAMA	PMA	BCEP
Whole Polymer	7.8	6.9	7.4	6.4
A 1,000-10,000	5.1	-	7.1	5.8
B 10,000-30,000	5.9	-	6.8	5.9
C 30,000-50,000	-	4.9	6.8	6.2
D 50,000-100,000	-	5.4	7.3	6.3

n) Polymers administered in a dose of 25 mg/kg intravenously.
   Organ weights determined seven days after drug injection and
   expressed as percent of total weight. Liver size of control
   group was 5.4.

Table IX. Hexobarbitol Sleeping Time Effected by Polymer Mole-
           cular Weight[o]

POLYMER	PYRAN	PAAMA	PMA	BCEP
Whole Polymer	98	64	58	62
A 1,000-10,000	43	-	44	38
B 10,000-30,000	47	-	46	38
C 30,000-50,000	-	41	44	40
D 50,000-100,000	-	48	49	44

o) Mice were inoculated with 25 mg/kg i.v. of the polymer.
   Twenty four hours later an anesthetic dose (80 mg/kg) of
   sodium hexobarbital was administered i.v. and duration of
   anesthesia recorded in minutes. Sleeping time of control
   group was 37 minutes.

Table X. Polymer Molecular Weight Effect on Spleen Weight[p]

POLYMER	PYRAN	PAAMA	PMA	BCEP
Whole Polymer	1.08	0.98	.90	0.84
A 1,000–10,000	0.40	–	.60	0.78
B 10,000–30,000	0.44	–	.73	0.78
C 30,000–50,000	–	.41	.84	0.82
D 50,000–100,000	–	.58	1.04	0.83

p) Polymers administered in a dose of 25 mg/kg intraveneously.
    Organ weights determined 7 days after drug injection and ex-
    pressed as percent of total body weight. Spleen size of
    control group was 0.36.

Table XI. Effect of Molecular Weight of Pyran on Acute Toxicity
           and Sensitivity to Bacterial Endotoxin.

POLYMER MW	LD_{50}[q]		LD_{50} of Bacterial[r]	
	Na^+ Salt	Ca^{++} Salt	Endotoxin Na^+ Salt	Ca^{++} Salt
12,500	112	–	24.0	–
15,500	98	190	7.0	10
21,300	94	–	0.8	–
52,600	86	170	0.5	0.5
NSC–46015	78	–	0.3	–
Control	–	–	21.0	21.0

q) Mice were injected i.v. with sodium and calcium salts of
    Pyran. Mortality was observed over a 14 day period.

r) Mice were injected i.v. with 25 mg/kg of pyran followed 24
    hours later by i.s. administration of S. typhosa lysopoly-
    saccharide 0.901. Control vehicles were NaCl and bacterial
    dosage were in mg/kg.

Macrophage Activation by Polycarboxyl Acid Polymer Immunopotentiators

Macrophages are produced by rapidly dividing precursor cells in the bone marrow and the daughter cells are released into the blood-stream. This mononuclear phagocytic system consists of several types of cells widely distributed throughout the body. Once they leave the blood, the macrophage settle in the tissues of the liver, spleen, lymph nodes, lungs, and peritoneum. In these sites, they differentiate into the various types of fixed macrophages (40,41).

Initially, macrophages were considered as the "antigenic garbage disposal unit," concerned only with the phagocytosis and the degradation of antigen. In recent years, however, macrophages have undergone considerable reevaluation with regard to their role in the immune system. The function that macrophage play in the generation of the humoral response has been extensively studied (40) and postulated mechanisms of macrophage activity in antibody production include; antigen processing for "presentation" to the lymphocytes and antigen processing by elaboration of soluble factors. Wagner et al. (42) have also reported that the expression of cell-mediated immunity is primarily dependent on the interaction between T-lymphocytes and macrophages.

It has been observed that under suitable conditions, macrophages can transform into "activated" macrophages, characterized by enlarged cells with undulating (ruffled) plasma membranes. Several lines of evidence indicate that activated macrophages are both qualitatively and quantitatively different from normal macrophages.

Although several biologic reticuloendothelial stimulants such as Mycorbacilerium bovis, bacille Calumette-Guerin, C. parvum and Toxoplasma gondii, are known to enhance macrophage function as well as to induce resistance to tumor growth (43), synthetic polyanionic immunopotentiators such as pyran and polyacrylic acid have been shown to be potent activators as well. Macrophage treated with these polyanionic polymer stimulants have shown cytotoxicity for normal cells (44). It has been demonstrated in our laboratories at MCV/VCU that the immunopotentiator pyran causes an increase in the number of histocytes present in the connective tissue surrounding the Lewis lung tumor of mice (45). In addition, increased numbers of tumor-associated macrophages are correlated with a decrease in metastasis in pyran-treated tumor-bearing mice (46).

It is not clear whether cytotoxicity and cytostasis involve the same function of the macrophages. Cytostasis by pyran-activated macrophages, similar to other stimulants, occurs as early as 4-6 hours after incubation of target cells with the macrophages (47). Although the growth of tumor cells is inhibited, ^{125}IUDR is not released from target cells until 48 hours after initation (48). Similar behavior has also been reported for

macrophages activated by other agents as well (49). The possible involvement of "factors" produced by activated macrophages in the tumorcidal event is receiving considerable attention (50-52).

Recent data which has been reported indicates that macrophages activated by pyran increase biochemical activity, are cytotoxic for tumor cells, and have a new macrophage cell surface antigen (AMØSCA). In contrast, this new macrophage cell surface antigen was not detected on normal macrophages or macrophages elicited by glycogen or thioglycollate, which are known to have increased biochemical activity but are not cytotoxic for tumor cells (53). AMØCSA can be detected within six hours after i.p. inoculation with pyran and its appearance correlated with the progression of macrophages to cytotoxic effector cells (53). The excellent correlation of AMØCSA with cytotoxicity and in vivo antitumor activity provides us with a simple, rapid technique for analyzing macrophage activation. Moreover, since polyanions such as pyran and dextran sulfate have been shown to activate macrophages in vitro (54) this provides a model system for analyzing the chemical moiety associated with these agents which is responsible for anti-tumor activity. Since their ability to activate macrophages has been associated with the anti-tumor activity of these agents this approach could provide a powerful tool for quickly screening potentially active compounds.

The cytotoxicity of activated macrophage has been demonstrated to be confined to only tumor cells (55-58) and has an insignificant or no effect on normal or natural cells such as mouse embryo cells (59) or murine kidney cells (56,58). The mechanism of activation of macrophage or the tumoricidal activity of activated macrophage is not understood. It has been observed that a population of aneuploid Lewis lung carcinoma cells, in the presence of activated macrophage, show reductive antitotic cell division resulting from tumor cells with 50% less DNA per cell (59). This depletion of DNA leads to cell destruction since these cells no longer have the appropriate DNA to synthesize necessary molecules for cell division.

In Table XII, cell cycle analysis shows (60) that in vitro Lewis lung cells are distributed such that 15% are in the resting phase (G_1) 52% in the DNA synthesis phase (S) and 33% are in cell division phases (G_2M). Following 24 hours inculation with pyran activated macrophage, the percent tumor cells in the resting phase (G_1) rose to 74% while synthesis (S) and cell division phases fell to 22% and 4% respectively. The shift to G_1 phase was not seen with Lewis lung cells cultivated with normal macrophage or mouse embryo fibroblasts incubated with activated macrophage.

Table XII. DNA Distribution of Lewis Lung Cell Cultivated With
            Pyran Activated Macrophages

Lewis Lung Cultured[s] With AMØ for	Percent in Phase[t]		
	G_1	S	G_2M
0 hr	15	52	33
4 hr	16	52	32
8 hr	32	47	20
16 hr	62	34	4
24 hr	74	22	4

s) Activated macrophage (AMØ) were obtained from mice injected
    i.p. with 25 mg/kg of pyran 7 days before peritoneal lavage.

t) A pattern recognition program was used to determine G_1, S and
    G_2M phase.

    Animals treated i.p. with pyran produced a 30 fold inhibi-
tion of Ehrlich ascites tumor growth with maximum inhibition of
tumor growth occuring with pyran treatment two days prior to
tumor cell inoculation (Table XIII) (60).
    Similar to the in vitro study using Lewis lung cells, it was
observed that animals pretreated with pyran had no detectable
Ehrlich ascites cells in phase G_2M and only a few in the S phase
after 2 or 6 days (Table XIV) (60) compared to pyran untreated
mice which had 58% in the S phase and 22 in the G_2M phase. Con-
sequently, the use of pyran activated macrophage resulted in a) a
high overall level of tumoricidal activity both in vivo and in
vitro, b) the appearance of a tumor cell population with 50%
of their normal DNA content and c) a shift of tumor cells from
phase G_2M to G_1. These results indicate that a possible mech-
anism of activated macrophage tumoricidal activity involves the
induction of tumor cells with a reduced DNA content.

Table XIII. Inhibition of Ehrlich Ascites Tumor Cell Proliferation in vivo by Pyran Inoculation

Macrophage Inducer	Number of Days Between Inoculation of Macrophage Pyran[u] and EA Cells	Total PEC Counts Per Animal ($\times 10^{-7}$)[v]
Pyran	2	1.6
Pyran	5	18
Pyran	7	28
Tumor Control	-	35

u) At various days, prior to tumor cell inoculation, animals received i.p. inoculation of pyran (25 mg/kg). 1×10^6 EA cells were inoculated i.p. into mice on day zero.

v) Five days after tumor inoculation, animals were killed and the total peritoneal exudate cells (PEC) counted.

Table XIV. Cell Cycle Analysis of Ehrlich Ascites Tumor Cells Before and After in vivo Exposure to Pyran

Days Following Tumor Inoculation[w]	Pyran Treatment[x]	Percent in Phase[y]		
		G_1	S	G_2M
2	-	20.	58	22
2	+	98	2	0
6	-	36	47	17
6	+	98.	2.	0

w) Mice were inoculated i.p. with 1×10^6 EA cells and killed on various days thereafter.

x) Pyran was injected i.p. (23 mg/kg) 2 days prior to tumor inoculation.

y) Cell cycle analysis of peritoneal EA cells was determined with a cell sorter utilizing a pattern recognition program.

Clinical Effects of Polycarboxylic Acid Polymers

Most polyanions are water soluble, which is essential for biological transport and more importantly for systemic administration, since injection of suspensions into blood system can cause "colloido-clasmic shock" or "macromolecular syndrome" with major hypersensitive clinical toxicity. Water-soluble polyanionic polymers can be distributed in a living system by means of blood or lymphatic circulation, by cellular transport through the involvement of mobile phagocytic cells, and absorption on cell surfaces.

In the initial study, pyran copolymer was given to advanced cancer patients who were no longer responsive to other treatment modalities. Survival had to be estimated at one month and patients had to be off other forms of chemotherapy or radiotherapy at least two weeks without signs of narrow depression or active sepsis and liver disease. Pyran copolymer was then given to 62 patients and the consistent limiting toxicity of pyran was transient thrombocytopenia. Cytoplasmic inclusions were seen in circulating leukocytes, nucleated marrow cells, and phagocytic cells of liver and spleen after administration. Although not dose related, fever was seen in 50% of patients.

The recent development of less toxic, lower MV fractions; less sensitive interpreitoneal or intervenous administration; as well as the formulation of the polyanion hydrolyrate as the more compatable calcium salt has stimulated renewed interest in at least two clinical studies now in progress. Future clinical studies should involve investigations of polyanions as immuno-adjuvants for antitumor and antiviral vaccines.

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Polymeric Derivatives Based on *cis*-Diamminedichloroplatinum(II) as Antineoplastic Agents

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Platinum polyamines can be readily synthesized with good control on the chain length through reaction of salts of PtX_4^{-2} with diamines. The

$$PtX_4^{-2} + H_2NRNH_2 \longrightarrow \underset{X}{\overset{X}{\text{Pt}}}\!\!\diagdown_{NH_2-R-NH_2}^{X}$$
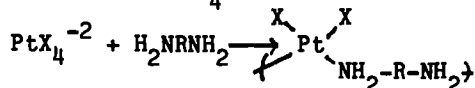
polyamines show good antineoplastic activity against a wide range of tumors including mouse connective tissues, human cervical carcinoma and human amnion cancer cells. Further the vast majority of the polyamines successfully altered the normal replication cycle of the polio virus Type 1 and Encephalomyocarditic virus, strain MM when the former cells were treated with the virus, without destruction of the cells themselves. Mice are able to tolerate large dosages of the polyamines.

Malignant neoplasms are the second leading cause of death in the United States. In 1964 Rosenberg and coworkers discovered

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that bacteria failed to divide, but continued to grow giving
filamentous cells in the presence of platinum electrodes (1). A
major cause of this inhibition to cell division was cis-dichloro-
diamineplatinum II,c-DDP. Much work centered about the clinical
use of c-DDP leading to the licensing of it as an antineoplast
drug. It is currently widely used, in conjunction with other
drugs, in the treatment of a wide variety of tumors (such as 2-5).
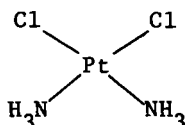
I, C - DDP

Most of the research involving use of platinum compounds as
antitumoral agents has involved either c-DDP itself or struc-
turally close derivatives. Through studies (such as 6-13), it
has been observed that there generally exists a structural window
for activity such that the most active antitumoral compounds are
a. neutral, b. contain two inert (NH_3) and two labile (Cl) ligands
with c. the ligands cis to one another.

The use of c-DDP has been complicated because of negative
side effects including gastrointestinal, hematopoietic, immuno-
suppressive, auditory and renal disfunction (such as 14-17). One
method of overcoming some of these negative side effects is to
limit the movement of c-DDP through inclusion of it into a polymer
thus prohibiting filtration of the polymer by the kidneys, etc.

Two general approaches have been taken. First, attachment
of the c-DDP derivative onto a preformed polymer. This was done
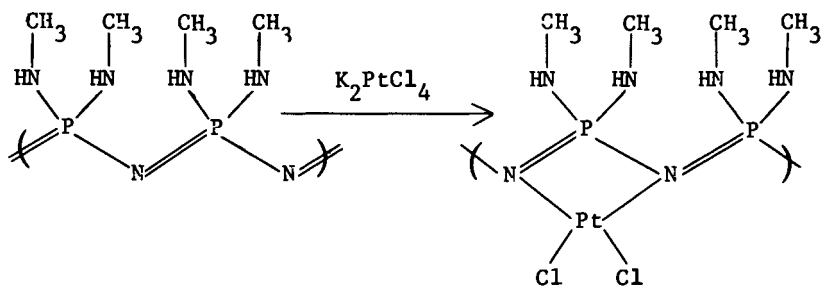
II.   III.

by Allcock and coworkers (such as 18-20). Poly[bis(methylamino)-phosphazene],II, is a water soluble polymer with coordination sites on both the side group and chain nitrogen atoms (13,21). Compound II reacts with K_2PtCl_4 and 18-crown-6-ether in organic media giving a coordination complex of structure III (13,19). Compound III shows tumor inhibitory activity towards mouse P38: lymphocytic leukemia and in the Ehrlich Ascites tumor regression test (19). Work is continuing in this area.

A second approach has the c-DDP derivative included as part of the polymer. In 1977 we initially synthesized the first poly-(cis-dichlorodiamineplatinum II) compound (22). Here we will briefly describe the synthesis of such compounds and describe results of preliminary biological assays concentrating on tests related to their potential use as antitumoral agents.

$$K_2PtCl_4 + H_2NRNH_2 \rightarrow$$
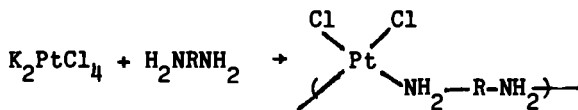
IV.

This paper presents preliminary biological assay results related to potential antineoplastic activity for select platinum polyamines of Form IV.

Synthesis and Physical Characterization

Following are typical synthetic and structural characterization procedures more fully described in references 22 and 23. Potassium tetrachloroplatinate (6.22×10^{-4} moles in 10 ml H_2O) and potassium iodide (4.97×10^{-3} moles in 10 ml H_2O) were separately dissolved, treated for ten minutes on a boiling water bath, mixed, heated 20 minutes longer and filtered to remove KCl. The resulting potassium tetraiodiplatinate solution was then mixed with an aqueous solution of 1,6-hexanediamine (7.83×10^{-4} moles in 10 ml of water) to immediately give a curdy, yellow solid, poly(cis-diiodo-1,6-hexamethylenediamine platinum (II)), DIHP, in 74% yield based on initial potassium tetrachloroplatinate. Elemental analysis (performed by Galbraith Labs., Knoxville, Tenn.) was in agreement with a structure of form IV; %C-found=12.0, theory=12.8; %N-found=4.9, theory=4.9; %H-found=2.9, theory= 2.9; %Pt-found=31.3, theory=34.5; %Cl-found 0.1, theory= 0.0; %I-found=49.9, theory=44.9. The following bands, with assignments, were present in infrared spectra of the product (obtained utilizing Perkin-Elmer 457 and 735B spectrophotometers; all values given in cm^{-1}) N-H stretch -2920, 2820; NH_2 bend -1560; CH_2 bend - 1455; CH_2 rock - 720; and Pt-N stretch - 545 and 460 cm^{-1} (24-28). The infrared spectra for the products derived from

K_2PtBr_4 and K_2PtCl_4 are identical from 4000 to 350 cm^{-1} (Figure 1). The Pt-Halide stretch for Pt-I and Pt-Br fall below 250 cm^{-1}, below the capability of the utilized spectrophotometer. The Pt-Cl band is present at 315 cm^{-1}. The UV spectra of the compounds are consistent with a cis-dihalodiamino-platinum (II) compound with bands at (obtained using a Cary 14 UV-visible spectrophotometer; wavelengths given in nm for chloro product) 250, 310, 340 and 390 with identification based on references 12 and 13. NMR spectra (obtained on a Varian EM360A spectrometer) in d_6-DMSO (using tetramethylsilane as a reference) showed three peaks; a broad band at 1.3 ppm (from the eight inner protons of the hexamethylene chain), a broad multiplet at 2.6 ppm (from the methylene protons) and a multiplet at 3.7 ppm. Interference by DMSO contamination of the d_6-DMSO and the low solubility of DIHP in DMSO - about 2% -prevented adequate integration of the NMR bands. A weight average molecular weight (obtained using a Brice-Phoenix 2000 light scattering photometer) of 4200 was found. Analysis employing coupled thermogravimetric-mass spectroscopy of the analogous chloropolyamine showed m/e ions at 35,37 and 70-74 amu indicative of the presence of the chloride moiety and a fragmentation pattern consistent with the presence of hexamethylenediamine.

The above was repeated except utilizing a more nearly equal molar amount of the $PtCl_4^{-2}$ and diamine (2.84 mmole of each reactant) and led to the synthesis (73% yield) of high molecular weight product with a weight average molecular weight of 1.5×10^6 via light scattering photometry. Thus product chain length can be easily and effectively controlled through control of the ratio of reactants. All of the tested polyamines are polymeric with degrees of polymerization in excess of 100 (23).

Biological Testing

Figure 2 contains structures of platinum polyamines tested along with a sample designation number. Table 1 contains a brief description of cell lines utilized in the present study.

Table 1. Cell lines utilized in the study of the inhibitory
nature of platinum polyamines

Abbreviated Designation(s)	Cell Line
L929 and L	Mouse Connective Tissue (Tumor Cells)
WISH	Human Amnion (Transformed Cells)
HeLa	Human Cervical (Cancer Cells)
VERD	African Green Monkey Kidney (Transformed Cells)

In Vitro. Mouse connective tissue (L cells), human cervical carcinoma (HeLa), and human amnion (WISH) cancer cells were used to determine the toxic and nontoxic levels of several of the
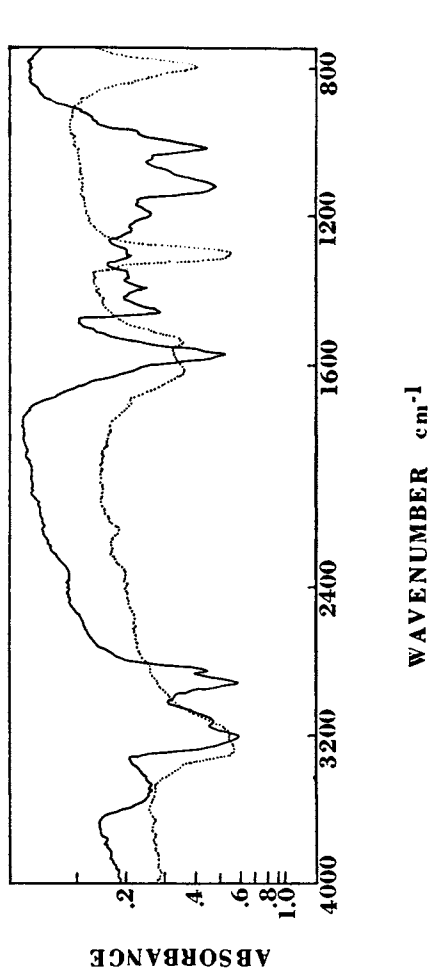
Figure 1. IR spectrum of poly[cis-dichloro(1,6-hexanediamine)platinum(II)] (———) and cis-di-chlorodiaminoplatinum(II) (· · ·).
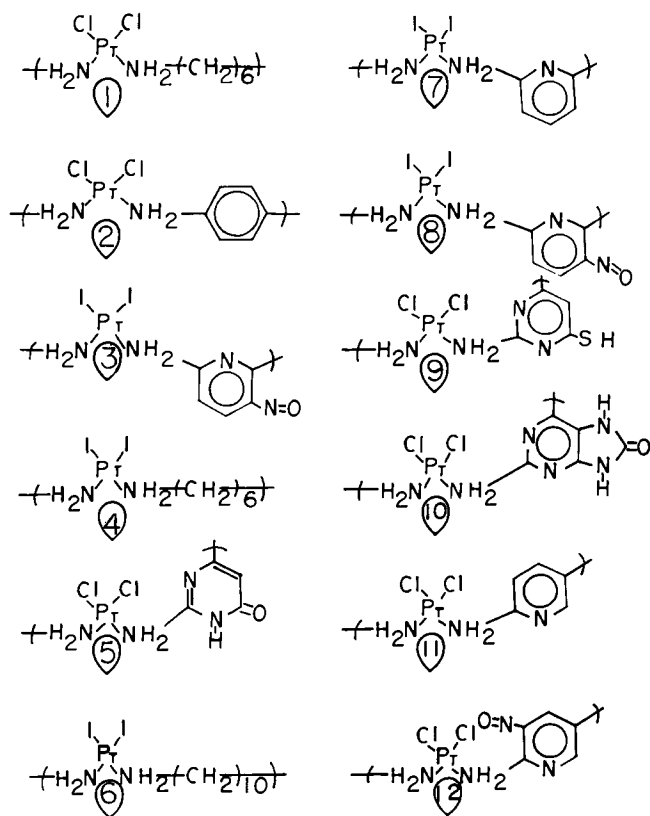
Figure 2. Structures of platinum polyamines utilized in the biological testing.

platinum compounds. Cells were plated into 1 ml wells and, after about 24 hrs, the pure media was removed and 1 ml of media containing the polymer in solution added. The cells were kept under surveillance for a specified time checking for cytopathic effect. The nutrient media contained Dulbeccos modified Eagles medium (DMEM) with 10% calf serum (CS) and 1% penicillin-streptomycin (Pen-Strep). The polymer containing nutrient media contained the afore plus a small amount (less than 1%) of DMSO containing the polyamines. The polyamine remains in solution throughout the testing procedure.

The maximum (threshold) nontoxic concentration level is similar for all compounds tested against tumor and transformed cell lines typically being 1 to 20 μg/ml (Tables 2-4). All of the polyamines exhibit total inhibition (100% cytopathic effect) above 30 μg/ml.

Table 2.   Toxicity of Platinum Compounds on Cells in Monolayer Culture

Pt Compound	Concentration (μg/ml)									
	5		10		20		30		50	
**	HeLa	L929	HeLa	L929	HeLa	L929	HeLa	L929	Hela	L929
1	0	0	0	0	0	0	25	25	100	*100
2	0	0	0	0	0	0	25	25	100	100
3	0	0	0	0	0	0	25	25	100	100
4	0	0	0	0	0	0	25	25	100	100
5	0	0	0	0	0	0	25	25	100	100

*Cytopathic effect after 24 hr of treatment with the indicated concentration of Pt compound.
**Polymer ID number from Figure 2.

Table 3.   Toxicity of Platinum Compounds on Mouse Tumor Cells in Culture.

Polymer Concentration (μg/ml)	Cytopathic Effects After 24hr. Treatment with Polymers						
	6*	7	8	9	10	11	12
60	4	4	4	4	4	4	4
20	4	4	4	4	4	4	4
6.7	3	4	1	+	3	3	3
2.2	1	1	0	0	1	0	0
0.7	0	0	0	0	0	0	0
0	0	0	0	0	0	0	0

Monolayer cultures of L929 (Mouse connective tissue) were treated with the indicated concentrations of each polymer. Toxic effects were recorded at the end of 24 hours.
4 = 100% cell destruction; 3 = 75%, 2 = 50%, 1 = 25%, + = 10%.
*Polymer ID number from Figure 2.

Table 4. Toxicity of Platinum Polymers on Human Amnion (WISH)
          Cells in Culture.

Polymer Concentration (μg/ml)	Cytopathic Effects After 24 Hours Treatment with Polymer		
	1*	2	4
63	4	4	4
21	4	4	4
7.0	2	4	2
2.3	0	0	0
0.80	0	0	0
0	0	0	0

Monolayer cultures of WISH (human amnion) cells were treated with the
indicated concentrations of each polymer. Toxic effects were
recorded at the end of 24 hours. 4 = 100% cell destruction, 3 = 75%,
2 = 50%, 1 = 25%, 0 = 0%.
*Polymer ID Number from Figure 2.

Selected polyamines were tested for their ability to alter the
normal replication cycle of two representative RNA viruses Ence-
phalomyocarditis (EMC strain MM) and Poliovirus Type I. Mono-
layers of L929 and HeLa cells were treated with solutions con-
taining specified polyamines. After incubation the cultures were
washed and then infected with the RNA virus. After a 24 hours
incubation period, the supernatant fluids were collected and
assayed for virus (pfu) content.

The data given in Tables 5 and 6 are offered to illustrate the
variety of results. Thus for Poliovirus, platinum polyamines can
effect an increase or decrease in the growth of the virus or
exhibit no effect at concentrations below that where the HeLa
cells are affected. Similar results are found for the polyamines
in L929 cells infected with EMC strain MM. Thus with
treatment for 24 hours, polymers 9,11 and 12 suppressed viral
replication (Table 6). With the exception of polymer 6, all of
the polymers suppressed EMC viral replication when treatment was
extended to 48 hours. Again viral replication was affected at
polyamine concentrations well below that necessary to inhibit the
L929 cells.

Table 5.   Effects of Platinum Compounds on the Viral Replication
           of Poliovirus in HeLa Cells.

| Concentration | Compound | | | | | |
| | 1[**] | | 2 | | 4 | |
μg/ml	Pfu[*]/ml	%control	Pfu/ml	%control	Pfu/ml	%control
10	3.4×10^6	10	6.5×10^7	200	3.2×10^7	100
20	1.1×10^7	34	3.2×10^7	100	3.2×10^7	100
Control	3.2×10^7	--	3.2×10^7	---	3.2×10^7	---

[*]plaque forming units
[**]Polymer identification number from Figure 2.

Table 6.   Effect of Platinum Compounds on the Replication of
           MM Virus in L Cells

Compound	Length of Treatment 2.2μg/ml for (hr)	24hr Virus Yield (pfu/ml)	% Control
6[*]	24	7.7×10^5	154
	48	4.0×10^5	148
7	24	7.4×10^5	148
	48	1.2×10^5	43
8	24	5.6×10^5	112
	48	2.3×10^5	82
9	24	3.5×10^5	70
	48	1.6×10^5	57
10	24	5.2×10^5	104
	48	1.6×10^5	57
11	24	2.6×10^5	52
	48	1.2×10^5	43
12	24	3.8×10^5	76
	48	2.8×10^5	75
Control	24	5.0×10^5	N/A
	48	2.8×10^5	N/A

[*]Polymer identification number from Figure 2.

Monolayer cultures of L929 cells were treated for 24 and 48 hours with 2.2 µg/ml of each polyamine. The cultures were then washed and infected with MM virus. After a 24 hour incubation period, the supernatant fluids were collected and assayed for virus pfa content.

Thus preliminary studies show that while there was no overt evidence of activity by the platinum polyamines on the cell cultures (CPE) at the employed polymer concentrations, there was biological activity in that the polymers were inhibiting or stimulating viral replication.

The RNA viruses EMC-MM and Poliovirus Type 1 are picoviruses with biological responses similar to other RNA viruses. Biological repsonses to agents by these two viruses is applicable to other RNA viruses. Implications of control of viral activity at low concentrations of drug is applicable to not only (potential) control of tumor growth, but also to control of RNA virus associated diseases including Polio, Encephalipis, the majority of "common colds," and most gastro intestinal viral infections. Thus drug control of RNA viruses is significant with broad implications.

In vivo. Preliminary experiments with mice show that they can tolerate a dosage of 400 µg of compound #4 (highest concentration tested) with no apparent ill effects. This dose is in excess of tenfold greater than that necessary to destroy either HeLa or L929 tumor cells. Further 500 µg of polymer 1 was given intraperitoneally at 3 day intervals to mice for a 15 day period with no apparent ill effects.

Thus mice are able to tolerate relatively high dosages of the platinum polyamines without overt ill effects.

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Controlled Release of Anticancer Agents That Are Complexes of *cis*-Diamminedichloroplatinum(II) and α-Hydroxyquinones

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Complexes between the square planar anticancer
agent cis-diamminedichloroplatinum (II), (cis-Pta$_2$Cl$_2$),
and α-hydroxyquinones are reported. A complex between
cis-Pta$_2$Cl$_2$ and the anthracycline antibiotic, doxoru-
bicin, contains five cis-Pta$_2$Cl$_2$ units per doxorubi-
cin molecule. Ratios of cis-Pta$_2$Cl$_2$ to α-hydroxy-
quinones range from 3/1 for cis-Pta$_2$Cl$_2$/juglone com-
plexes to 6/1 for cis-Pta$_2$Cl$_2$/quinizarin complexes.
Ultraviolet and infrared data are reported for the
complexes as well as thin layer chromatography, con-
ductivity, and molecular weight results. The re-
sults indicate chelation of platinum to oxygen atoms
of the α-hydroxyquinones with concomitant stacking of
cis-Pta$_2$Cl$_2$ units. The reaction leading to dimer-
ization has the potential for further chain exten-
sion although no such compounds have as yet been
isolated. In vivo experiments with cis-Pta$_2$Cl$_2$/
doxorubicin and with cis-Pta$_2$Cl$_2$/quinizarin show the
efficacy of these complexes as a new time-release
system for delivering anticancer agents.

Several papers (1, 2, 3) from this laboratory have dealt with
the development of a system for delivering anticancer agents at a
constant rate over a prolonged period, perhaps as long as several
months. This system consists of incorporating the drug in a poly-
meric matrix, shaping the composite into convenient form (such as
beads or powder) and implanting the composite into the body tissue
of animals. The drugs diffuse continuously from the interior of
the polymer to the outer surface, where they are mechanically
swept away by body fluid. The mechanism of migration of the drug
is that of diffusion and the thermodynamic driving force is the
concentration gradient (3).

Previously reported (2) in vitro and in vivo experiments with composites of poly(lactic acid) (PLA) with cyclophosphamide, with cis-diamminedichloroplatinum (II) (cis-Pta₂Cl₂) (4), and with cis-Pta₂Cl₂-doxorubicin (DOXO) (5) have demonstrated the feasibility of this system, especially their drug release ability and the reduced toxicity of the platinum compound.

During the preparation of the composite containing PLA, cis-Pta₂Cl₂ and DOXO, a distinctive color change to intense purple was observed (2). In view of the enhanced anticancer activity shown by this composite in animals over those containing the separate drugs, it was decided to isolate and characterize the purple material, believed to be a chemical complex of the platinum with DOXO and, in addition, to investigate the efficacy of this complex in in vivo experiments.

To elucidate the binding site, several α-hydroxyquinone analogues of DOXO were complexed with cis-Pta₂Cl₂.

This paper reports the syntheses and characterization of complexes of cis-Pta₂Cl₂ with 5-hydroxy-1,4-naphthoquinone (HNQ), 5,8-dihydroxy-1,4-naphthoquinone (DHNQ), 1,4-dihydroxy-9,10-anthraquinone (DHAQ) and with DOXO. In addition it reports preliminary results of in vivo experiments performed with complexes cis-Pta₂Cl₂/DOXO and cis-Pta₂Cl₂/quinizarin.

Several metal complexes (Be, Co, Cu, Fe, Mg, Ni) with DOXO (6, 7) and with hydroxy-naphthoquinones and -anthraquinones (8, 9, 10) have been recently reported. To our knowledge no study of platinum complexes with α-hydroxyquinones has been published to date, with the exception of an unsuccessful attempt to prepare a Pt/5,8-dihydroxy-1,4-naphthoquinone (8).

Experimental Section

General Information. All reactants were dried at room temperature under vacuum in an Abderhalden apparatus to constant weight. All experiments were carried out in a dry box; continuously under nitrogen. Elemental analyses were performed by Microanalysis, Wilmington, DE, all compounds prepared yielded satisfactory analyses (see Table I). The UV spectra were obtained on samples dissolved in DMF with a NSI Hitachi computerized #100-80 Ultraviolet Spectrophotometer. The IR spectra were recorded on a Perkin Elmer 180 Recording IR Spectrophotometer in CsI pellets (except for DOXO and DOXO complex where KBr discs were used). Conductivity measurements were taken on a Beckman Model RC16B2 conductivity bridge using a Fisher conductivity cell (cell constant = 0.1799, determined on standard KCl solutions). ¹H NMR spectra were obtained in d₇DMF at 90MHz on a Bruker HFX-10 spectrometer. Thin layer chromatographs on silica gel 60 supports from 0.1 to 1.0 mM DMF solutions of starting materials and products were developed in 9/1 acetone/HCl, and visualized with 2% SnCl₂ in 1M HCl spray.

Experiments <u>In Vivo</u>. The standard assay of the Ascites
Sarcoma in the ICR mice was used (<u>2</u>). There were six mice in each
cage, with two cages for the negative controls, two cages for the
positive controls and one cage for each dose level of drug tested.
Tumor cell injections were performed on day 0 and drug injection
on day 1, both given intraperitoneally. Suspensions in carboxy-
methyl cellulose (CMC-7LF, Hercules, Inc., Wilmington, DE) of
complexes, as listed in Table IV, with saline as a carrier were
used.

     Complex I: cis-Pta_2Cl_2/5-hydroxy-1,4-napthoquinone. In a
typical preparation, cis-Pta_2Cl_2 (0.50 g, 1.66 mmol) was dissolved
in 30 ml N,N-dimethylformamide (DMF) and added to HNQ (0.14 g,
0.80 mmol) dissolved in 10 ml DMF. The reactants were combined in
a 120 ml brown glass jar which contained 83.5 gms of 5 mm glass
beads. The jar was sealed under N_2, then placed on a ball mill
(1750 rpm) to rotate for 120 hrs. The ball milled suspension
(brown-purple in color) was centrifuged to remove a small amount
of solids. The main product was precipitated by addition of 80 ml
methylene chloride ($MeCl_2$) to yield a brown-purple solid collected
on a frit-glass filter. Crude product was recrystallized from
10-15 ml DMF with precipitation by excess $MeCl_2$. Yield: 0.59 gm
(90%) of brown-purple product.

Table I Elemental Analyses

	C	H	Cl	N	Pt	
Complex I	11.77	2.27	15.63	8.24	57.37	a
cis-Pta_2Cl_2/HNQ	11.68	2.69	15.71	8.61	55.51	b
$C_{10}H_{22}Cl_{4.5}N_6O_3Pt_3$						
Complex II	9.22	2.16	14.97	8.60	59.94	a
cis-Pta_2Cl_2/DHNQ	8.68	2.40	14.75	7.51	58.68	b
$C_{10}H_{30}Cl_{5.5}N_8O_4Pt_4$						
Complex III	8.38	1.90	21.20	6.98	58.33	a
cis-Pta_2Cl_2/DHAQ	8.68	2.48	20.10	6.52	56.76	b
$C_{14}H_{36}Cl_{12}N_{10}O_4Pt_6$						
Complex IV	16.56	3.32	14.39	7.85	49.50	a
cis-Pta_2Cl_2/DOXO	15.86	3.21	14.15	7.52	47.64	b
$C_{27}H_{56}Cl_8N_{11}O_{11}Pt_5$						

a   Calculated
b   Found

     Complex II: cis-Pta_2Cl_2/5,8-dihydroxy-1,4-naphthoquinone.
Following the procedure described above, 5,8-dihydroxy-1,4-
naphthoquinone (0.039 g, 0.20 mmol) and cis-Pta_2Cl_2 (0.373 g,
1.24 mmol) were condensed to give 0.40 g (yield 97%) of compound
II as navy blue product.

Complex III: cis-Pta₂Cl₂/1,4-dihydroxy-9,10-anthraquinone.
Following the procedure described above, 1,4-dihydroxy-9,10-
anthraquinone (0.062 g, 0.26 mmol) was reacted with cis-Pta₂Cl₂
(0.50 g, 1.66 mmol) to give 0.29 g (yield 52%) III as purple
product.

Doxorubicin Base from Doxorubicin Hydrochloride. Doxorubicin
hydrochloride was converted to the free base by bubbling gaseous
ammonia through a capillary pipet into a stirred suspension of
doxorubicin hydrochloride (0.050 g, 0.08 mmol) in 15 ml of an 8:1
mixture of methylene chloride and methanol at 0°C. The bubbling
of ammonia and the stirring were continued until the mixture was
red-orange. After the ammonium chloride was removed by centrifu-
gation, excess solvent and ammonia were removed on a Rotovap.
DMF was added to give a solution of 0.047 g (0.086 mmole) of
doxorubicin base in 100 ml of solvent.

Complex IV: cis-Pta₂Cl₂/Doxorubicin. Following the proce-
dure described above for complex I, cis-Pta₂Cl₂ (0.104 g, 0.34
mmol) was condensed with doxorubicin (0.047 g, 0.086 mmol) to give
0.103 g (68.7% yield) of compound IV as purple product.

Results and Discussion

Complexes of cis-Pta₂Cl₂ with HNQ, DHNQ, DHAQ and DOXO have
been obtained. All these complexes are similar in physical form:
fine powder (solids) of intense purple to blue color; poorly
soluble in common organic solvents, giving purple or blue solu-
tions; none sublime on heating; all decompose before melting. U.V.
and I.R. spectra of the parent ligands and platinum complexes are
reported in Tables II and III.

α-Hydroxyquinones exhibit a so-called chelated H absorbance
in the UV, caused by H bonding of the α-hydroxyl H to the keto
oxygen of the quinone (11). In our complexes with platinum, this
absorbance disappears, with concurrent appearance of a new absor-
bance at longer wavelengths characteristic of the α-hydroxyquinone
dianion (11).

Retention of the 302 nm band supports the presence of cis-
Pta₂Cl₂ units in the structures of these complexes. It is assumed
that these units are stacked in a way similar to that of the
Magnus' green salt (12). Hydrolysis of a cis-Pta₂Cl₂/DHAQ complex
in dilute HCl, followed by thin layer chromotography has identi-
fied 75% of the Pt containing fraction as pure cis-Pta₂Cl₂, the
other 25% being the fractions which have lost chloride ligand.
Pure quinizarin was also identified as a component of the hydroly-
zate by thin layer chromatography.

The IR spectra of the quinone complexes show a shift of the
ν(C=O) band stretching frequency to lower wavenumbers in comparison

with the corresponding parent hydroxyquinones. This shift indicates coordination or chelation of Pt(II) to oxygen of the hydroxyquinone as shown by Pierpont ($\underline{10}$) for the copper (II) complexes of 1,5-dihydroxy-1,4-naphthoquinone.

Table II   Ultraviolet Spectra of Hydroxyquinone Ligands and Their Corresponding Pt(II) Complexes[1]

Compounds	Absorption (log ε) nm
HNQ	425(3.6), 260(4.2)
HNQ Complex	485, 380, 260(4.2)
DHNQ	515(4.2), 482(4.2), 270(4.4)
DHNQ Complex	620, 570, 302, 271(4.4)
DHAQ	478(3.8), 278(4.0)
DHAQ Complex	580, 540, 302, 278
DOXO	530(2.7), 495(2.9), 475(2.9), 265(3.0)
DOXO Complex	592(2.7), 535(2.8), 506(2.8), 265(3.3)

[1]Absorption maxima cis-Pta_2Cl_2 370,302 nm.

Table III   IR Absorption (cm^{-1}) of hydroxyquinone ligands and their Pt(II) complexes

Cmpds	Chelate OH	Chelate OX	ν(C=O)	C-O-Pt	Pt-O
DHNQ	3010[a]	--	1600	--	--
DHNQ Cmplx	--	2910[b]	1560	1030–1170	445
DHAQ	3000[a]	--	1626,1590	--	--
DHAQ Cmplx	--	2910[b]	1605,1575	1000–1150	445
DOXO	2910	--	1620–1635	--	--
DOXO Cmplx	--	--	1620–1630	1050–1150	450

[a]Hadzi, D.; Sheppard, N., Trans. Far. Soc., $\underline{1954}$, 50, 911
[b]reference 13

Broad, strong absorption bands at 1000–1500 cm^{-1} reported ($\underline{13}$) to be indicative of ether type linkages, are tentatively assumed to be due to C-O-Pt linkages. Absorbance at 445–450 cm^{-1} is analogous to that reported by Nakomoto for Pt-O bonds in Pt(II) acetylacetonates ($\underline{14}$). All complexes also exhibit IR (in cm^{-1}) absorbances reported for cis-Pta_2Cl_2 by Poulet ($\underline{15}$): ν (NH_3): 3280, 3200; δ (NH_3): 1620, 1290; ν (Pt-Cl); 315; ν (Pt-N): 240.
^{1}H NMR spectra of DHAQ (δ = 12.8, chelated OH; δ = 8.3–8.4, aromatic H; δ = 7.5, 2,3H) and cis-Pta_2Cl_2 (δ = 4.2, N-H) were straightforward, however, NMR of the complex showed retention of

the N–H peak of cis-Pta$_2$Cl$_2$ only, plus several unidentified up-field resonances. Experiments are currently underway to show that these unidentified peaks are isotropically shifted DHAQ resonanc s caused by paramagnetism in the complex.

Attempts at obtaining single crystals for X-ray structure determination have failed thus far. Low solubility of the complexes in solvents suitable for vapor pressure osmometry have made molecular weight determination difficult. One result for complex III, cis-Pta$_2$Cl$_2$/DHAQ, in acetone yielded a M.W. of 2270 gm/mole, vs a calculated M.W. from elemental analyses of 2006 gm/mole.

Based on data given, a probable structure of Complex IV, cis-Pta$_2$Cl$_2$/DOXO, is shown in Figure 1. Conductivity of the cis-Pta$_2$Cl$_2$/DHAQ complex increased from zero, immediately after solution in DMF, to values which varied with concentration in the manner of a 2:1 electrolyte (as compared to K$_2$PtCl$_6$ by the method of Feltham and Hayter (16). Probable structure of the electrolyte differs from that in Figure 1 in that the two five-co-ordinate platinum centers become four-co-ordinate through loss of chloride anions, resulting in a dipositive complex ion containing platinum co-ordinated to the hydroxyquinone portion of DOXO. Fractional chlorine content from elemental analyses calculations for HNQ and DHNQ complexes indicates that either one or two chloride ligands are lost from cis-Pta$_2$Cl$_2$ on complexation.

In Vivo Experimental Results

Preliminary in vivo results (see Table IV) show the feasibility of this potential time-release system for delivering anti-cancer agents. cis-Pta$_2$Cl$_2$/DOXO complex was significantly better than the positive control (7 mg cis-Pta$_2$Cl$_2$/kg mouse). The improvement in life span compared with the positive control is substantial and the toxic effect of the platinum compound has certainly been decreased. These tests show that it is possible to use a dose of complex three times the toxic level of cis-Pta$_2$Cl$_2$, because of slow release from cis-Pta$_2$Cl$_2$/DOXO complex.

Preliminary in vivo results (see Table IV) show that the complex formed between cis-Pta$_2$Cl$_2$ and quinizarin was more efficacious than the positive control. It did not have quite the degree of increased life span shown by the cis-Pta$_2$Cl$_2$/DOXO complex. On the other hand, as before with the DOXO complex, fairly high levels of cis-platinum upwards of 20 mg/Kg mouse could be used as compared to a toxic level of 7 mg/Kg mouse for cis-Pta$_2$Cl$_2$ alone.

This paper has reported only products of the reaction which were isolated and characterized. However, the potential for further chelation and therefore doubling or trebling the molecular weight to polymer proportions exists. This suggests the possibility of a polymeric drug moiety which would be inherently slow releasing in vivo. It is planned to continue this line of research in the future.
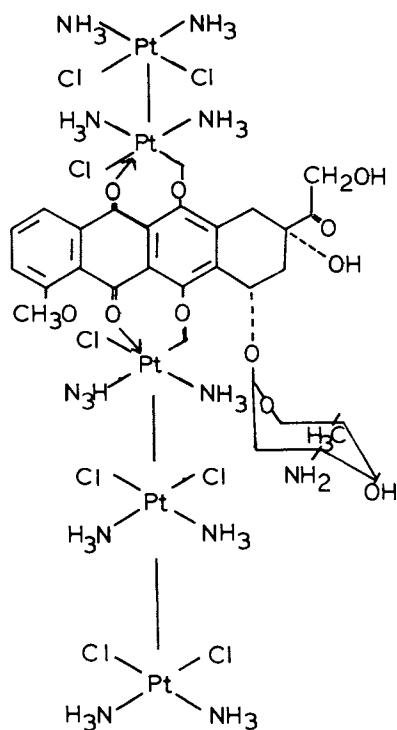
Figure 1. cis-*Diamminedichloroplatinum(II)/DOXO complex.*

Table IV

In Vivo Test Results of Complexes

Compound	Dose mg Compound/ kg Mouse	Avg. Day of Death	% Increased Life Span	Number of Cures[2] (out of 6)
Negative Control[1]	0	17.3	---	-
Positive control cis-Pta$_2$Cl$_2$	7	31.3	80	4
cis-Pta$_2$Cl$_2$ + doxorubicin	5	34.2	97	4
	10	36.0	108	6
	15	34.8	101	1
	20	26.2	51	1
Negative control	0	13.8	---	-
Positive control cis-Pta$_2$Cl$_2$	7	25.0	81	1
cis-Pta$_2$Cl$_2$ + quinizarin	8	28.0	107	1
	12.5	26.0	88	0
	25	16.0	16	0
	50	2.75	-80	
	100	3.8	-72	

[1] Negative control: Mice injected only with cancer cells.

[2] Cure: The National Cancer Institute uses the term "cure" for those animals which live beyond 60 days.

Acknowledgments

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^{13}C NMR Studies of the Structure of the Divinyl Ether–Maleic Anhydride Cyclic Alternating Copolymer: A Biologically Active Agent

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The copolymer has been shown to contain tetrahydrofuran and tetrahydropyran rings in a ratio of 0.8 to 1. Resonances have been assigned by the use of single frequency off resonance decoupling (SFORD), suppressed nuclear Overhauser effect (NOE) combined with a relaxation reagent, and comparison with model compounds. Published spectra, which have been interpreted as demonstrating the presence of only 5-membered rings, also show the unique 6-membered ring methylene group. Available evidence is insufficient to assign stereochemistry.

Maleic anhydride and divinyl ether form a cyclic alternating copolymer in a 2 to 1 ratio[1]. This polymer has a broad spectrum of biological activities[2], and is at present undergoing preliminary clinical evaluation as an antitumor agent. The structure of the 2 to 1 copolymer has proven to be somewhat elusive. Various techniques have been used in the past to study and characterize this copolymer. Recently, two studies have appeared which focus attention on the ^{13}C and proton NMR properties of the copolymer. Kunitake and Tsukino[3] have presented evidence based on the ^{13}C NMR data that the copolymer may consist primarily of 5-membered or 6-membered bicyclic structures, depending on the method of polymerization; the 6-membered bicyclic structure has been promoted by Butler[4]. The symmetrical or 5-membered ring structure was concluded by Kunitake to be formed when the polymerization was done in chloroform. Moreover, chain methylenes were calculated to be in the cis-configuration on the tetrahydrofuran rings and trans on succinic anhydride units along the chain backbone. When the copolymer was prepared in a mixed-solvent system consisting of acetone and carbon disulfide, the tetrahydropyran bicyclic unit was found to be favored.

Butler and Chu[4] have studied the copolymer prepared in cyclohexanone by means of high (300MHz) and low (60MHz) field

proton NMR spectroscopy. These workers have concluded that the pyran or unsymmetrical structure is exclusively formed. Further, the pyran anhydride ring juncture was concluded to possess a trans-geometry. The proton NMR study was supported by comparison of copolymers prepared in turn with bis(2,2-dideuterovinyl) ether and maleic anhydride, and divinyl ether and 3,4-dideuteromaleic anhydride. These workers further found that the ^{13}C spectra of the copolymers prepared in cyclohexanone were invariant with temperature. This observation was considered by them as additional evidence supporting the energetically favored 6-membered structure. Based on these data, a mechanism using HOMO-LUMO arguments was put forth to rationalize the exclusive formation of the tetrahydropyran structure.

Results

In this paper we present evidence, based primarily on ^{13}C NMR data that, for a copolymer prepared in toluene or benzene, a mixture of the symmetrical and unsymmetrical bicyclic structures is obtained. Our studies rest largely on heretofore unavailable model compounds which are representative of the substructural units.

Although NMR spectra can be taken of the anhydride in acetone, much better spectra can be obtained if the copolymer is dissolved in water (or for the purposes of the NMR experiment in D_2O), as shown in Figure 1. Using the labeling scheme of Kunitake we see, from high field to low, peak A (31.3 ppm), B (35.3 ppm), and C (44.7 ppm); a group of six peaks D-E (50.5-53.1 ppm); a group of four peaks, F (76.6-79.8 ppm); and two types of carbonyl, G (173.9-174.3) and H (177.2 ppm). The two small peaks at 25.3 and 68.5 ppm are due to residual tetrahydrofuran. Note that peak B at 35.3 pm is rather distinct and well resolved from peak A at 31.3 ppm. Dioxane (not shown in Figure 1) was used as an internal chemical shift reference (67.0 ppm).

We have assigned all of the ^{13}C peaks which appear in the spectrum of the copolymer and are in general agreement with the assignments of Kunitake (see Table I). However, based on model compound data (vide infra), we find that peak B is due to a 6-membered tetrahydropyran ring methylene carbon. On the basis of this evidence, as well as additional evidence to be presented, we conclude that the copolymer as prepared in our hands consists of a mixture of 5- and 6-membered rings. We note that the appearance of the same peak, i.e., B in the published spectra of Kunitake, demands that he also found a mixture of structures in the same copolymeric material. From our ^{13}C NMR data, the divinyl ether-maleic anhydride copolymer consists of both 5- and 6-membered cyclic structures in the ratio of about 0.8 to 1.

Assigning the ^{13}C NMR peaks of the copolymer has not been a trivial problem by any means. The assignments rest on a variety of bases. These include ^{13}C shifts for model compounds, SFORD
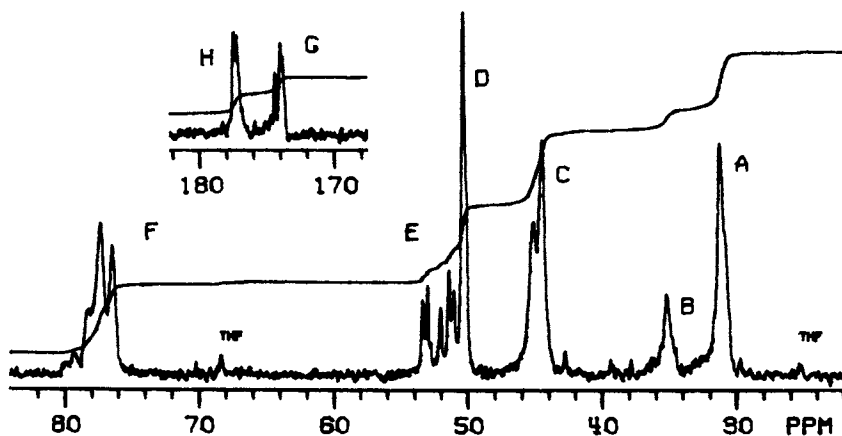
Figure 1. The 90-MHz ^{13}C NMR spectrum of hydrolyzed divinyl ether–maleic anhydride copolymer in D_2O at 58°C taken on a Nicolet NT-360WB.

TABLE I

SUMMARY OF ^{13}C NMR DATA FOR HYDROLYZED DIVINYL ETHER-MALEIC ANHYDRIDE COPOLYMER
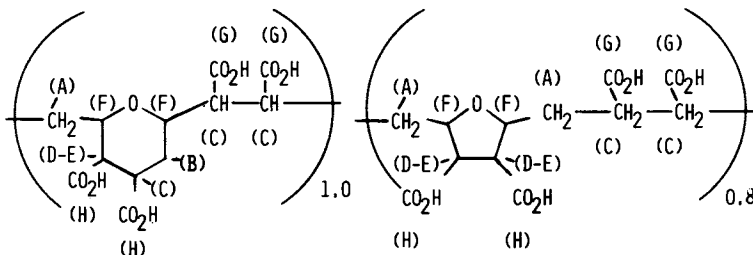
PEAK	A	B	C	D AND E	F	G	H
SHIFT (PPM)	31.3	35.5	44.7	50.5-53.1	76.6–79.8	173.9–174.3	177.2
MULTIPLICITY*	T	T	D	D	D	S	S
NO. OF CARBONS							
CALC.	3	1	5	3	4	4	4
FOUND**	2.6	1	4.6	2.6	3.6	3.6	3.6

(SPECTRA RUN AS D_2O SOLUTIONS; INTERNAL REFERENCE DIOXANE ASSIGNED TO 67.0 PPM.)

* T – TRIPLETT (CH_2) D = DOUBLET (CH) S = SINGLET (C)

** FROM NOE SUPRESSED EXPERIMENTS FOR RATIO OF STRUCTURES SHOWN USING AREA OF PEAK
AT 35 PPM AS ONE CARBON.

spectra, and a quantitative determination of the relative number
of carbons present. The latter was done by means of a
combination of a suppressed NOE and a shift or relaxation agent
experiment.

Table I shows by means of a SFORD experiment that the peaks
at 31.3 and 35.2 ppm appear as triplets and are due to methylene
carbons. The rest of the aliphatic carbons are all doublets or
methine carbons. We find also from suppressed NOE experiments
(Table I) that the peak areas are as follows: the methylene
resonance at 31.3 ppm represents 3 carbons; the methylene
resonance at 35.2 ppm corresponds to one carbon; the methine
resonance at 44.7 ppm represents 5 carbons; the peak appearing at
about 50.5 ppm accounts for 3 carbons; and the methine resonance
at about 77 ppm represents 4 carbons. The number of carbons
present can be assigned either by assuming that the resonance at
77.4 ppm represents 4 carbons or that the resonance at 35.3 ppm
represents one carbon.

Discussion

In order to assign the ^{13}C resonances unequivocally, two
model compounds were prepared, a 2,6-dimethyltetrahydropyran-
3,4-dicarboxylic acid and a 2,5-dimethyltetrahydrofuran-
3,4-dicarboxylic acid; the syntheses and ^{13}C structural
assignments are shown in Figure 2. One distinct stereoisomer of
the tetrahydropyran derivative was produced as a result of the
synthesis. The relative stereochemistry of the four functional
groups of the tetrahydropyran has been established by means of a
proton NMR study. On the basis of the NMR coupling constants and
the Karplus relationship, we have determined that this tetra
substituted tetrahydropyran is in the all trans-configuration.

The second model compound, 2,5-dimethyltetrahydrofuran-
3,4-dicarboxylic acid, was found to consist of three isomers
(plus a partially hydrogenated material). We were unable to
assign the stereochemistry of this model unequivocally. We also
used as a model 2,3-dimethylsuccinic acid which was a mixture of
the racemic and meso forms. On this basis, one can show that the
resonances of the carbons alpha to the anhydride or acid group
along the backbone occur at 44.6 ppm and are approximately equal
to that found in the copolymer.

Upon close inspection of the assignments for the copolymer
(shown in Figure 1), one can see that the carbon resonances of
the 5- and 6-membered rings are overlapped. The only unique
carbon resonance is that of the 6-membered ring, and corresponds
to peak B in the spectrum of the copolymer. Thus, with the aid
of the model compounds and the relative areas found in the
copolymer, it is possible to show that the copolymer must consist
of a mixture of 5- and 6-membered rings in an approximate 0.8 to
1 ratio.

We feel that the fine structure associated with some of the
peaks in the ^{13}C spectrum of the copolymer is undoubtedly due

A
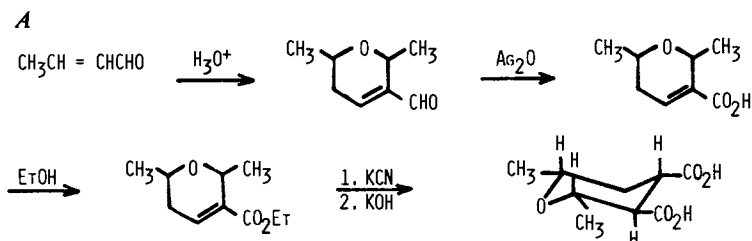
^{13}C CHEM SHIFTS (22.6 MHz): IN D_2O 19.8 (CH_3 ON C-2), 21.2 (CH_3 ON C-6), 34.6 (C-5), 44.3 (C-4), 51.8 (C-3), 74.1 (C-6), 74.9 (C-1), 177.3 (C=O ON C-3), AND 177.8 PPM (C=O ON C-4). ^1H CHEM SHIFTS OF RING PROTON (90 MHz) FOR DIETHYL ESTER IN C_6D_6: 1.42 (H-5A, $J_{4,5A}$ = 12.5 Hz, $J_{5E,5A}$ = -12.5 Hz, $J_{5A,6}$ = 11.5 Hz), 2.08 (H-5E, $J_{4,5E}$ = 3.8 Hz, $J_{5E,6}$ = 2.0 Hz), 2.36 (H-3, $J_{2,3}$ = 9.5 Hz, $J_{3,4}$ = 11.3 Hz), 2.99 (H-4), 3.62 (H-6) AND 3.64 PPM (H-2).

B

MIXTURE OF 3 STEREOISOMERS
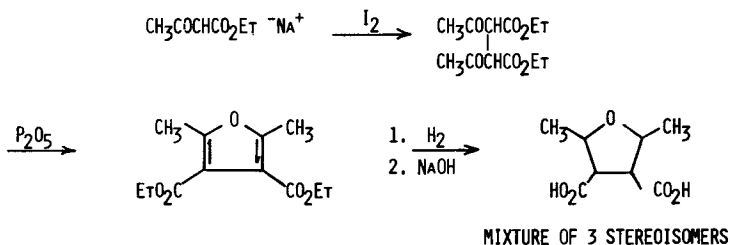
^{13}C CHEM SHIFTS (22.6 MHz) IN D_2O: 17.0-20.9 (CH_3 ON C-2 AND C-5), 50.7-54.7 (C-2 AND C-4) 74.5-78.3 (C-2 AND C-5), AND 176.1-176.9 PPM (C=O ON C-2 AND C-4)

Figure 2. Synthetic schemes and NMR data for model compounds. Key: A, 2,6-dimethyltetrahydropyran-3,4-dicarboxylic acid and B, 2,5-dimethyltetrahydrofuran-3,4-dicarboxylic acid.

to stereochemical differences at ring junctures. Both Butler and
Kunitake make some stereochemical assignments based on what we
feel is rather tentative data. On the basis of our model
compound data, we would agree that stereochemical implications
are contained within the ^{13}C data, but we do not feel that
anyone at this time has a firm enough spectroscopic basis to make
absolute pronouncements about stereochemistry.

Relevance of Models. In our studies, we measured the spectra
of model compounds as opposed to the approach of Kunitake, in
which he calculated chemical shifts utilizing more generalized
methods. We do not feel that a calculation allows one to make a
fine enough distinction, particularly when the models used as a
basis for the calculation are well removed from those needed for
the actual structure study. As an example, Kunitake argues that
for a succinic anhydride substructure, carbons 2 and 3 of the
succinate ring would be expected to resonate at 38 ppm. The
method of calculation used to arrive at this shift is shown in
Figure 3. The 38 ppm value represents an upfield shift of -5.1
ppm to form the cis compound. Using the same reasoning the
anhydride carbonyls and the 2,3-cis-diisopropyl anhydride are
found at 169.1 ppm for an upfield shift of -2.1 ppm for the
carbonyls. The calculation for the cis structure is discounted
by Kunitake, since in the actual polymer no similar shifts are
found. The values calculated by Kunitake for the trans compound
are 171.2 for the carbonyls and 43.1 ppm for carbons 2 and 3 of
the ring. These are observed to be close to that found in the
anhydride form of the copolymer, i.e., peak C at 43.6 ppm and
peak G at 171.3 ppm.

We have obtained a commercial sample of the
2,3-dimethylsuccinic acid, a mixture of the racemic and meso
forms. We have converted the acid to the anhydride and
determined the ^{13}C spectrum without separation of the two
isomers. The chemical shifts of the anhydride may be readily
assigned from the peak intensities and the data of Ernst and
Trowitzsch[5] and are shown in Figure 4. This is further
supported by the fact that in the meso or cis form, the methyl
carbons would be expected to be eclipsed and show a compression
or steric shift. When we performed a calculation analogous to
that of Kunitake as shown in Figure 5 but starting directly from
the cis-2,3-dimethylsuccinic anhydride, we found that the
predicted shifts for the cisoid form represented by the
2,3-diisopropyl structure were 33.5 and 171.0 ppm. Thus the
carbonyl shift is in poor agreement with that calculated by
Kunitake and is certainly in the range of that found for the
actual polymer. On the strength of this evidence, we feel that
cisoid backbone structures can not be ruled out using this type
of argument.

The approach of extending data to account for major structure
differences between a model and an actual compound is frequently
used, and in fact, often valid. However, the above exercise
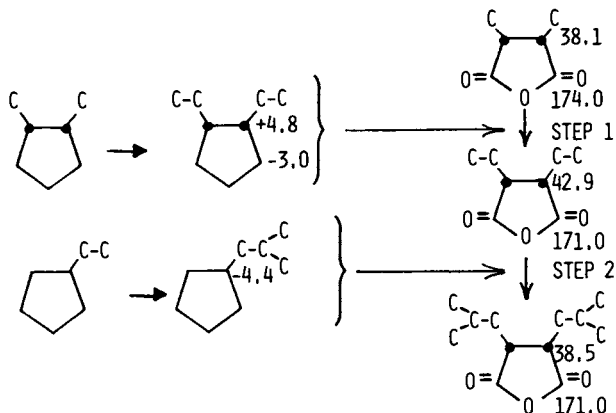
Figure 3. Scheme used by Kunitake (3) to estimate chemical shifts of succinate ring on copolymer backbone.
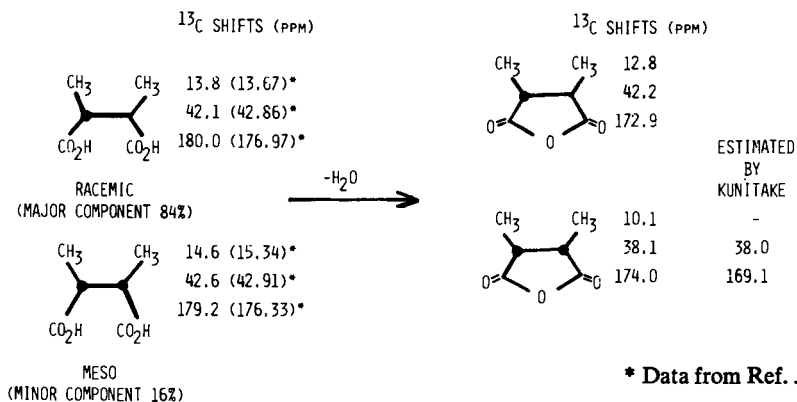
* Data from Ref. 5.

Figure 4. ^{13}C NMR data obtained for dimethyl succinic acid and anhydride. These data represent an improved model for estimating chemical shifts in a five-membered ring anhydride structure.
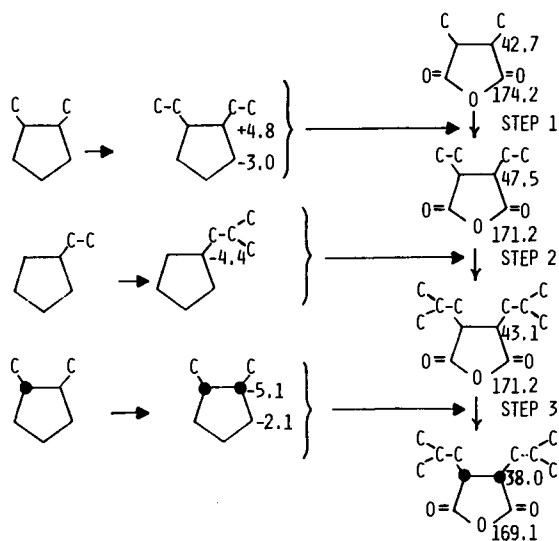
Figure 5. Improved procedure to estimate chemical shifts of succinate ring on copolymer backbone. This scheme is similar to that shown in Figure 3, but uses data from Figure 4. The results presented here show the danger in using such methods to make fine distinctions.

illustrates that such an approach is not without danger. Fine distinctions of the order of 1-2 ppm may be uncertain, and indeed unless one uses a model which is very close in structure, such calculations are likely to be misleading.
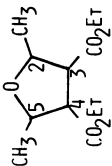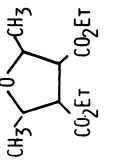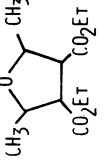
For example, Kunitake used the shifts on going from a <u>trans</u>- to a <u>cis</u>-1,2-dimethylcyclopentane as representative of that which would occur in an analogous 2,3-dimethylsuccinic anhydride. Thus he estimated a shift of -5.1 ppm for C-2 (and C-3) and -2.1 ppm for the carbonyl carbon. We found that the actual shifts on going from <u>trans</u>- to <u>cis</u>-2,3-dimethylsuccinic anhydride are -4.1 ppm (C-2 and C-3) and +1.1 ppm (carbonyls).

<u>Five Membered Ring Model</u>. The 2,5-dimethyl-3,4-tetra-hydrofuran dicarboxylic acid was synthesized by means of the scheme shown in Figure 2b. The synthesis of this compound was complicated by the rather severe conditions necessary to accomplish the final hydrogenation step. These conditions were hydrogenation for 21 hours at 169-190°C at 3000 psi using a mixture of ruthenium on carbon and ruthenium chloride catalyst. This treatment gave rise to a mixture of isomers as well as an incompletely hydrogenated product. Nevertheless, the ^{13}C NMR spectra of the crude reaction product allowed the general assignment of the ring spectra of the 2,5-dimethyl-3,4-tetrahydrofuran to be readily made. These assignments are shown in Figure 2b. The ^{13}C shift of the ring carbons, C-2 and C-5, are of course of primary interest as models for the divinyl ether maleic anhydride copolymer. However, there is also a substantial interest in the sidechain carbons since it is through these shifts that the relative stereochemistry of the isomers is potentially established.

Now we could readily see from the ^{13}C NMR spectrum that there were three isomers present for this model compound. Upon fractionation of the mixture by gas chromatographic separation, we could establish that of the three isomers, two were symmetrical and one was nonsymmetrical by means of the ^{13}C NMR spectrum. These model compounds contain four asymmetrical centers. This in turn leads to a maximum of 2^N or 16 compounds where N=4. This, of course, represents a potential maximum of 8 <u>dl</u> pairs. Elimination of two possible isomers because of the additional symmetry due to having the groups on carbons 2 and 5 and the groups on carbon 3 and 4 identical, then leaves only six possible relative stereoisomers. Formally, this would be 5 <u>dl</u> pairs and one meso compound. This means, then, as a maximum or for the worst possible case, there are 20 possible configurations of 3 out of 6 isomers to consider. The figure 20 comes from the number of ways in which one may take 6 items 3 at a time.

During the course of a gas chromatographic workup of a mixture of the tetrahydrofuran diethylester we noticed that upon washing the crude mixture successively with NaOH, acid, and water that one of the GC peaks, F, decreased while one, A, grew. The remaining GC peaks remained approximately the same. These data

TABLE II

^{13}C AND PARTIAL ^1H NMR SHIFTS FOR STEREOISOMERS OF ETHYL 2,5-DIMETHYL -3,4-TETRAHYDROFURANDICARBOXYLATE

ISOMER	ELEMENT	CARBON NO. (PPM)									
		2	3	4	5	2,Me	5,Me	3,C=O	4,C=O	OCH_2-	-CH_3
GC FRACTION E (SYMMETRICAL)	^{13}C	77.14	50.0	50.0	77.14	17.4	17.4	171.3	171.3	60.5	14.2
	^1H	4.60	3.64[A]	3.64[A]	4.60	–	–	–	–	–	–
GC FRACTION F (SYMMETRICAL)	^{13}C	75.1	51.0	51.0	75.1	17.2	17.2	170.4	170.4	60.5	14.2
	^1H	CA. 4.3	3.32[B]	3.32[B]	4.28	–	–	–	–	–	–
GC FRACTION A (UNSYMMETRICAL)	^{13}C	76.2	52.1[C]	54.0[C]	77.9	17.2	20.0	171.8	172.1	60.6	14.2
	^1H	–	3.13[D]	3.46[E]	–	–	–	–	–	–	–

A) J = 2.07 AND 5.64 Hz
B) J = 1.91 AND 5.28 Hz
C) MAY BE REVERSED
D) J = 6.65 AND 8.39 Hz
E) J = 6.65 AND 7.8 Hz

indicate that one of the trapped isomers was interconverted to another upon treatment with either base or acid. This information suggests that an epimerization is occurring at the C-3 or C-4 carbon to interconvert one of the model isomers. These data, then substantially reduce the problem of characterizing mixtures of stereoisomers. When all the constraints or degrees of freedom are imposed upon the problem, we find that we still must consider 12 separate cases.

Conclusions

Unfortunately, our carbon-13 and proton data (Table II) taken together with other data for the model compounds, do not allow an unambiguous assignment of the 3 stereoisomers which we have obtained. For the present, then, we must be content to leave the relative stereochemistry of the 5-membered ring model compounds in question. For now, we can only say that the fine structure in the copolymer spectra are certainly well within isomers shifts seen for the model compound. However, based on the fact that the relative stereochemistry of the tetrahydrofuran models are unknown and that we have only one pyran isomer, we are unable to make a definite assignment as to the relative stereochemistry of ring junctions in the original copolymer.

Finally, our studies show all divinyl ether–maleic anhydride copolymers examined to date to be a mixture of the 5-membered and 6-membered bicyclic structures. The primary evidence for this, of course, is the presence of peak B at ca. 35 ppm which is assigned to the methylene carbon on a tetrahydropyran ring. Major features of the spectrum can be readily explained based on the available data present by both us and Professor Kunitake. However, still in question is the meaning of the fine structure contained particularly in peaks D, E, and F with respect to the relative stereochemistry of the copolymer as produced.

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Interaction of Methotrexate (Polylysine) with Rat Liver Tumor Cells

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During the past decade interest has developed in the possible therapeutic use of carrier-bound drugs to secure greater inter-action between the drug and its target site of action. Examples of carriers include liposomes (1), polypeptides (2), polysaccharides (3) and antibodies (4), and of drugs daunomycin (5), methotrexate (MTX) (6), actinomycin D and cytosine arabinoside (7). In this laboratory MTX, a potent inhibitor of one-carbon metabolism, coupled to albumin (BSA) has been shown to be equally effective to free MTX in the treatment of murine L1210 leukemia (8) and superior to the free drug in controlling the Lewis lung carcinoma (9). A recent examination of the interaction of this carrier-bound drug with L1210 cells in vitro has suggested that the nature of the carrier is important for interaction with the cell surface and that a carrier-bound drug might be absorbed into the cell by an alternate mechanism to that of the free agent (10). Because of the thera-peutic significance of these observations, it was decided to inves-tigate the interaction of MTX-BSA with a non-systemic cellular source such as the MTX transport-resistant Reuber H35 hepatoma cell line, which had been developed in one of the authors' laboratories (11). Unfortunately, in constrast to L1210 cells MTX-BSA was inef-

fective in supressing growth of the liver tumor cells. However, because of the observations that carrier structure can determine cellular interaction, and that polylysine possesses unique cellular absorptive properties (12), MTX was bound to both L- and D- forms of this carrier and the products were tested with the hepatoma cells in vitro. The comparative toxicities of the polymer-bound and free drugs with both parental and MTX transport-resistant cell lines were measured, and experimental results were obtained, which showed how differential cell rescue with reagents which usually overcome blockade of the one-carbon metabolic pathway could also afford similar results with cells treated with the drug-carrier complex.

Experimental

Materials. The following were obtained from commercial sources: Swims medium S77, horse serum, and fetal calf serum (Grand Island Biologicals); MTX (Lederle Laboratories) was purified by DEAE cellulose prior to use (11); crystalline BSA, poly-L-lysine (M_r ~35,000), hypoxanthine and thymidine (Sigma Chemical Co.); 1-ethyl-3-(3'-dimethylaminopropyl)carbodiimide hydrochloride (EDC) (Story Chemicals); folinic acid (ICN Pharmaceuticals); Blue Dextran (Pharmacia). The concentrations of the folate derivatives and MTX were determined by their respective extinction coefficients (13). MTX-BSA was synthesized according to a previously described procedure (8) in which MTX is coupled via a terminal carboxyl group to - amino groups contained in the albumin molecule.

MTX(poly-L- and poly-D-lysine) were synthesized according to a procedure first outlined by Ryser and Shen (14). 0.1 ml of a neutral aqueous solution of MTX (13 mg/ml) and 0.05 ml of an aqueous solution of EDC (200 mg/ml) were added to the polylysine samples (~ 10 mg) dissolved in 0.5 ml of 0.01 M potassium phosphate, pH 7.2, containing 0.15 M sodium chloride. After stirring in the dark at room temperature for 1.5 hours, the products were applied to columns of Sephadex G-25 (2.2 x 90 cm) which had been pre-equilibrated with the reaction buffer solution. Elution was carried out with the same buffer and the eluent absorbance was monitored at 280 nm. Fractions comprising the initial peak absorbance were combined and dialysed against water (6 liters) then lyophilized. The yellowish flocculent residues (wt. ~7 mg) contained ~5 mole MTX/mole polylysine when estimated by MTX absorbance at 370 nm ($\varepsilon_{0.1 \text{ N NaOH}}$ = 7,100 (13)).

Cell culture. H-11-EC3 (designated H35 cells) derived from the Reuber H35 hepatoma (15,16) were grown as previously described and the development of resistant sublines was outlined in the same publication (11). MTX-resistant lines are indicated by the designation H35R with the subscript following the R indicating the concentration of MTX, in μM, to which the cells are resistant. All the cells employed in this study were resistant solely through defective transport and showed no more than 50% increase in dihy-

drofolate reductase relative to control cultures (21). Hepatocytes isolated from male rats of the Fisher strain by collagenase perfusion and mechanical dispersion were plated at a density of 7.5×10^5 cells/ml in L-15 medium containing 15% FBS and 10 mU/ml insulin on petri dishes according to a procedure outlined earlier by Tarentino and Galivan (17).

Growth Studies. The cells were cultured at a density of 5×10^4 cells/ml of Swims S77 medium with 20% horse serum and 5% fetal calf serum in a 95-5% air-CO_2 mixture at 37°C. The indicated additions were made at this time to duplicate samples and the cultures were allowed to incubate for 72 hours after which the cells were released from the plates with 0.05% trypsin and counted with a ZB1 Coulter counter (11).

Results

Treatment of H35 cells with MTX-BSA. The comparative growth rates of H35 cells plated at 2×10^5 cells/dish in the presence of various levels of MTX-BSA are shown in Table I. Clearly the growth of wild-type cells is retarded by micromolar quantities of MTX-BSA with 40 μM MTX-BSA leading to a complete cessation of growth. The H35R$_1$ (see EXPERIMENTAL for subscript definition) cells, however, are unaffected by this concentration of carrier-bound drug and demonstrate a growth response comparable to that of the untreated H35 cells.

Treatment of H35 cells with MTX(poly-L-lysine). The influence of various concentrations of MTX(poly-L-lysine) on both H35 and H35R$_{0.3}$ hepatoma cells is illustrated in Figure 1. The effect is also compared to that of the free drug. It is apparent that MTX shows little restraint on the growth of the transport-resistant mutants at levels up to 1 μM in contrast to the wild-type cells which submit to nanomolar concentrations of the drug ($I_{50} = 14$ nM). However, when MTX is bound to the poly-L-lysine carrier, both cell lines show similar characteristics with $I_{50} \sim 70$ nM. The concentrations of poly-L-lysine used in the experiments show little influence on the cell growth.

Prevention of MTX and MTX(poly-L-lysine) toxicity to H35 cells. Table II shows the differential effects which are observed when H35 and H35R$_1$ cells treated with MTX (0.1 μM) and MTX(poly-L-lysine) (1.0 μM) are rescued with folinic acid (100 μM) or thymidine/hypoxanthine (500/100 μM). The H35 cells are sensitive to each drug and respond in the case of MTX and MTX(poly-L-lysine) to supplementation with folinic acid although those treated with the free drug give a better response than those treated with the carrier-bound agent (92% as opposed to 69% of normal growth rate). The H35R$_1$ cells, unaffected by MTX, but susceptible to MTX(poly-L-lysine) do not respond to folinic acid. However, both cell types show a moderate response to thymidine/hypoxanthine rescue.

TABLE I: The comparative growth of resistant and
non-resistant H35 cells treated with MTX-BSA.

Cell type	Concentration MTX-BSA	Cells/plate at 72 hr
	(μM)	
H35	0	2.7×10^6
H35	10	9.4×10^4
H35	40	7.3×10^4
H35R	0	2.2×10^6
H35R	10	2.2×10^6
H35R	40	1.8×10^6

TABLE II: The effect of folinic acid and
thymidine/hypoxanthine on H35 and $H35R_1$ cells
treated with MTX(poly-L-lysine) and MTX.

MTX (poly-L-lysine) (1.0 μM)	MTX (0.1 μM)	Folinic Acid (100 μM)	Thymidine/ Hypoxanthine (500 μM/100 μM)	Cells[1] (%)
		H35		
+				9
+		+		60
		+		107
+			+	36
			+	78
	+			6
	+	+		92
		$H35R_1$		
+				15
+		+		20
		+		98
+			+	40
			+	87

[1]Number of cells are expressed relative to the untreated
cultures.
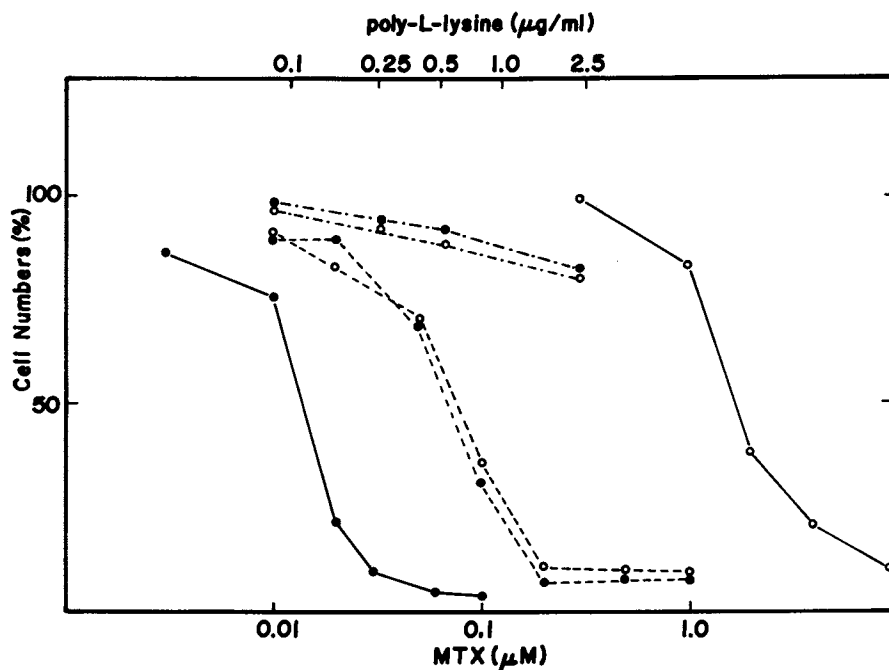
Figure 1. Comparative growth curves of resistant (○) H35R$_{0.3}$ and sensitive (●) H35 cells when treated with MTX (———), MTX(poly-L-lysine) (– – –), and (poly-L-lysine) (–●–) alone at the levels indicated. The growth rates are expressed as a percentage of those observed with untreated cell samples. (Reproduced, with permission, from Ref. 23. Copyright 1981, American Society for Pharmacology and Experimental Therapeutics.)

Measurement of dihydrofolate reductase and thymidylate synthe-
tase activities in MTX and MTX(poly-L-lysine) treated H35 cells.
Extracts of the untreated cells were prepared from two 60 mm plates,
and four 100 mm plates were used for the extracts of inhibited
cells. Dihydrofolate reductase activity was measured by a modifi-
cation (18) of the method of Mathews (19) and thymidylate synthe-
tase in the intact cell by the release of ^{3}H from [5-^{3}H] deoxyuridine
as described previously (20). When either MTX transport-defective
or nondefective cells were treated with MTX and MTX(poly-L-lysine),
it was found that both drugs, administered at the concentrations
shown in Figure 2, led to elimination of both dihydrofolate re-
ductase and thymidylate synthetase activities.

Sephadex chromatography of H35R$_{0.3}$ cell extracts after expos-
ure to [^{3}H]MTX(poly-L- and poly-D-lysine). After 72 hours in cul-
ture thirty 60 mm plates of H35 cells were exposed for a further 24
hours to either [^{3}H]MTX(poly-L- or -D-lysine) (0.75 mM, 3 x 10^5-
dpm/nmol). At the end of the incubation the plates were placed on
ice and washed x 4 with 4 ml ice-cold Hanks balanced salt solution
then scraped into two 1 ml aliquots of isotonic saline, buffered
with 0.01 M potassium phosphate, pH 7.4. The cells were broken by
20 strokes with a Dounce homogenizer and centrifuged at 15,000 x g
for 20 min. 1 mg of Blue Dextran 200 (M$_r$ = 2 x 10^6) was added to the
extracts and each of the mixtures (poly-L- or poly-D-) was chroma-
tographed, as described previously (20), on Sephadex G-75 (Figure
3). When mixtures of [^{3}H](poly-L- or -D-lysine) and Blue Dextran
2000 were mixed in separate experiments and applied to the same
column, radioactivity eluted only in the void volume corresponding
to the first peak. In the MTX(poly-L-lysine) experiment the second
peak was identified by its enzymatic activity as dihydrofolate re-
ductase and the third peak corresponded to a mixture of higher
molecular weight γ-glutamyl conjugates of MTX and free MTX.
When the contents of peak 2 were boiled at pH 8.0 and applied
to a second column of Sephadex G-25 the profile observed in Figure
3A was obtained whose peaks corresponded to known standards for
free MTX and polyglutamate forms of MTX. Similar results were
obtained when an aliquot of combined fractions 55 to 77 were applied
to the Sephadex G-25 column (Figure 3B).
In contrast when the cell extracts from the [^{3}H]MTX(poly-D-
lysine) experiment were applied to the Sephadex G-75 column only
the undegraded complex with Blue Dextran 200 was observed. When the
experiment was repeated with the non-resistant H35 cells, similar
results were obtained.

Uptake of MTX and MTX(poly-L-lysine) by hepatocytes, H35 and
H35R$_1$ cells in culture. Hepatocytes or hepatoma cells maintained
in culture (11) were treated with [^{3}H]MTX(poly-L-lysine) or
[^{3}H]MTX. At the indicated times (Figures 4 and 5) cells were
washed, and the monolayer was measured for radioactivity (11).
Clearly higher concentrations of carrier-bound rather than free
drug remained associated with the cells in every case.
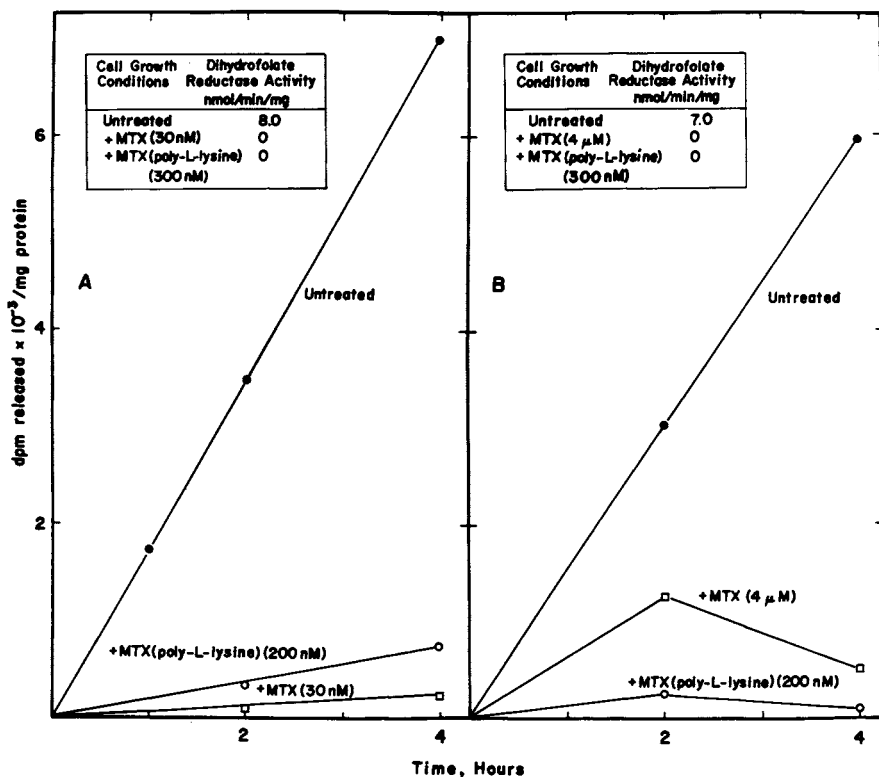
Figure 2. Inhibition of dihydrofolate reductase and de novo *thymidylate biosynthesis in (A) nonresistant H35 and (B) resistant H35R$_{0.3}$ hepatoma cells, after treatment with MTX and MTX(poly-L-lysine). (Reproduced, with permission, from Ref. 23. Copyright 1981, American Society for Pharmacology and Experimental Therapeutics.)*
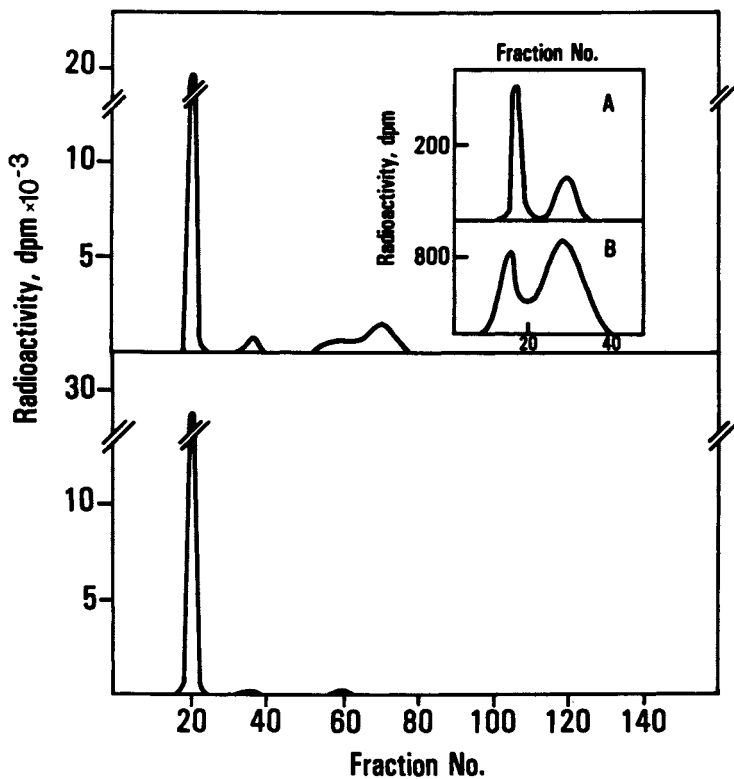
Figure 3.   Sephadex G-75 chromatography of $H35R_{0.3}$ cell extracts after exposure to [^3H]MTX(poly-L-lysine) (upper panel) and [^3H]MTX(poly-D-lysine) (lower panel). Key: A, rechromatography of Sephadex G-25 of the contents of the second peak after boiling at pH 8.0 and B, chromatography on Sephadex G-25 of an aliquot from the combined fractions 55 to 77.
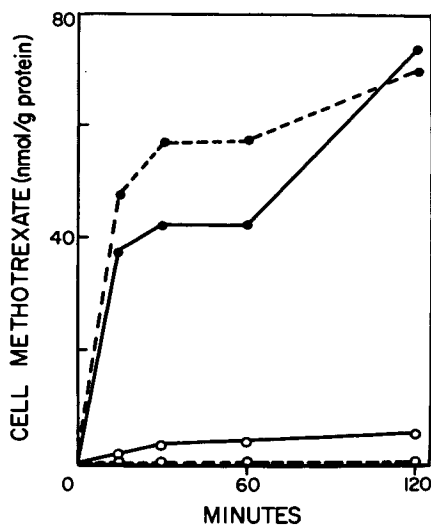
Figure 4.   Uptake of [³H]MTX (○) and [³H]MTX(poly-L-lysine) (●) by H35 (——) and H35R₁ (– – –) cells in culture. The medium concentration of MTX in each case was 0.3 μM.
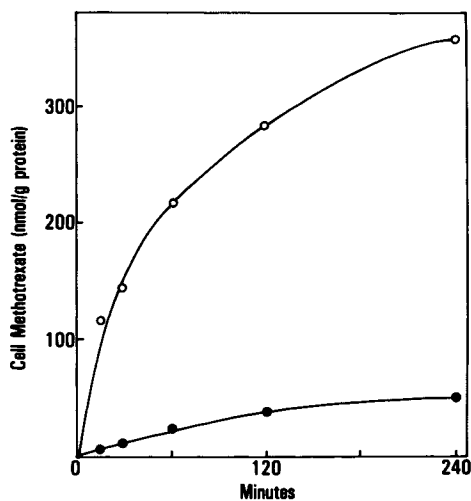
Figure 5.   Uptake of [³H]MTX (●) and [³H]MTX(poly-L-lysine) (○) by hepatocytes in culture. The medium concentration of MTX in each case was 1 μM.

Toxicity of poly-L- and -D-lysine derivatives to rat hepato-
cytes, H35 and H35 R$_{0.3}$ cells in culture. Cultured hepatocytes or
hepatoma cells were treated with differing levels of either poly-L-
or poly-D-lysine. The effect of these agents on cell viability is
shown in Figures 6 and 7. It is apparent that both D- and L-
polymers show limited toxicity towards the hepatoma cell lines, but
that the D-isomer shows considerable toxicity towards the hepato-
cytes. Figure 7 also shows that the MTX derivatives of both D- and
L-carriers are equally toxic to the non-substituted agent when in-
teracting with the hepatocytes. It should also be noted that MTX
linked to poly-D-lysine does not enhance the toxicity of the car-
rier to H35 cells.

Effect of MTX(poly-D- and poly-L-lysine) on rat hepatocyte
dihydrofolate reductase. Hepatocytes were maintained in culture
for 48 hours prior to measurement of viability and dihydrofolate
reductase activity (Table III). Even at concentrations up to 2 μM
MTX(poly-D-lysine) causes little inhibition of cellular dihydrofo-
late reductase, however, it shows considerable cellular toxicity.
In contrast MTX(poly-L-lysine) at these levels maintains good cel-
lular viability with somewhat higher but not complete inhibition of
the reductase. This clearly demonstrates that the toxicity of
MTX(poly-D-lysine) is exerted through the carrier by some, as yet,
unidentified mechanism and that little MTX is released to interact
with the reductase. The comparative figures for treatment with
free MTX indicate complete inhibition of enzyme activity. Since
normal hepatocytes consist of a non-dividing population, extensive
inhibition of dihydrofolate reductase might not be expected to in-
terfere with cell viability.

Effect of MTX(poly-L-lysine) pulse doses on hepatoma cell
growth and dihydrofolate reductase. Cells were maintained in cul-
ture for 72 hours prior to measurement and the pulse dose was added
between 24 and 48 hours. The results are shown in Table IV where it
is apparent that 2.8 μM MTX(poly-L-lysine) is effective in reducing
cell growth of both defective and non-defective cell lines, and
also eliminates dihydrofolate reductase activity.

Discussion

The results described in this report amplify observations that
carrier structure is integral to the concept of polymer-mediated
drug delivery (10). Two general features begin to emerge from our
studies on carrier-bound drug therapy: (1) that in a whole animal
experiment the carrier-bound agent ensures elevated levels of the
drug are retained compartmentalized within a body cavity thus pro-
viding a source of continous high level drug delivery (8,9), and (2)
that polymer-bound drugs can overcome transport blocks which cer-
tain cells use to prevent passage of the free drug into the cell.
Because earlier studies had shown the effectiveness of MTX-BSA in
the treatment of certain mouse tumors (8,9), we considered extend-
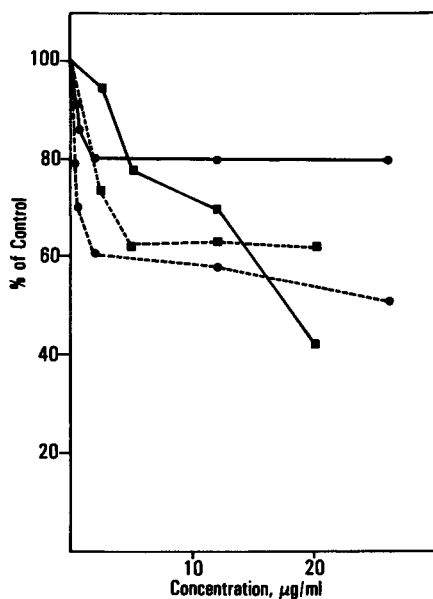
Figure 6. Toxicity of poly-D-lysine (■) and poly-L-lysine (●) against H35 (——)
and H35R$_{0.3}$ (– – –) rat hepatoma cell lines.
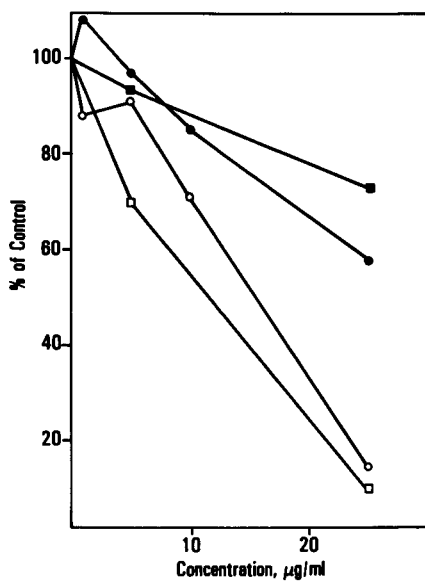
Figure 7. Toxicity of poly-L-lysine (●), MTX(poly-L-lysine) (■), poly-D-lysine
(○), and MTX(poly-D-lysine) (□) to rat hepatocytes in culture.

TABLE III: Effect of MTX(poly-D-lysine) and MTX(poly-L-
lysine) on rat hepatocyte dihydrofolate reductase.

DRUG ADDED	CONCENTRATION MTX, polylysine		DHFR	CELL VIABILITY
	μM	μg/ml	%	%
Control			100	100
MTX-poly-L-lysine	0.2	2.5	58	95
	2.0	25	26	75
MTX-poly-D-lysine	0.2	2.5	80	70
	2.0	25	74	10
MTX	0.2		10	100·
MTX	2.0		0	92

Measurements were made after 48 hours in culture with drug
addition after 24 hours.

ing its application to the growth control of a chemically induced
hepatoma cell line and its transport-resistant counterpart. The
reagent, however, proved ineffective and alternate carriers had to
be considered. The demonstrated toxicity of MTX(poly-L-lysine)
against Chinese hamster ovary cells (14) suggested this amino-acid
polymer as a possible effective delivery system. The results de-
monstrate that MTX(poly-L-lysine) is also effective in this case
and that it bypasses the transport defect, as was also recently
observed by Shen and Ryser in L929 mouse fibroblasts (24). MTX-
(poly-L-lysine), however, is slightly less effective than MTX
against the parent cell line. It has also been shown that the cells
subject to chemotherapy with the MTX(polylysine) complexes can be
rescued with hypoxanthine/thymidine but only the non-resistant
cells can be rescued with folinic acid. The observation that
folinic acid does not reverse toxicity in the resistant cells is
consistent with the transport mutants not being able to take up
either MTX or the reduced folates (21) as has already been shown
with MTX or pyrimethamine toxicity in MTX transport-resistant L-
5178 lymphoblasts (22). Although MTX(poly-D-lysine) is toxic to
H35 cells and hepatocytes, it is no more so than poly-D-lysine
itself. The experiments in which the lysates of hepatoma cells
exposed to MTX(poly-L and -D-lysine) derivatives are examined by
Sephadex chromatography (Figure 3) amplify this observation. Only
with the L-isomeric carrier are both free MTX and enzyme-bound drug
found in the cytoplasm. With MTX(poly-D-lysine) only an unchanged
high-molecular weight species can be identified.

Experiments which examine the comparative toxicities of the poly-L- and -D- carriers for hepatocytes and hepatomas indicate a low level of toxicity with the hepatoma. However, an elevated toxic effect is observed for the hepatocyte with the D-isomeric carrier. When the drug is bound to the carrier little alteration is observed in the toxic response of the hepatocyte suggesting that in its resting state it has little requirement for dihydrofolate reductase. The results shown in Table III illustrate this point more clearly where only 26% of reductase activity remains after treating the hepatocytes with 2 micromolar MTX(poly-L-lysine) yet 75% cellular viability remains. This contrasts with the response of the hepatoma cell line (Table IV), where both dihydrofolate reductase and ~90% cell growth of both transport defective and non-defective cell lines is eliminated by treatment with 2.8 micromolar MTX(poly-L-lysine).

Although *in vitro* experiments do not necessarily predicate successful experiments in the whole animal the differential response of the hepatocytes and hepatoma cells to the MTX(poly-L-lysine) suggests that treatment of rats bearing similar liver tumors could be successful.

TABLE IV: Effect of a pulse dose of MTX(poly-L-lysine) on hepatoma cell growth and dihydrofolate reductase activity.

CELL TYPE	DRUG ADDITION	MTX CONC^N	DHFR	CELL GROWTH
		µM	%	%
H35			100	100
	MTX-poly-L-lysine	2.8[a]	0.3	7.6
H35R$_{0.3}$			100	100
	MTX-poly-L-lysine	2.8[a]	0.1	11

Cells were cultured for 72 hours prior to measurement.

[a]Pulse dose added between 24 and 48 hours.

Legend of Symbols

MTX, Methotrexate; BSA, bovine serum albumin; EDC, 1-ethyl-3(3'-dimethylaminopropyl)carbodiimide hydrochloride.

Acknowledgments

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Evidence of Cooperativity and Allosterism in the Binding of Various Antibiotics and Carcinogens to DNA

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        The interaction of a number of drugs
(actinomycin D, adriamycin, daunorubicin, and DHAQ)
and carcinogens (NQO and HAAF) with DNA is studied
by optical titration and/or phase partition. In each
case pronounced curvature of the Scatchard plots is
observed at low levels of bound drug, indicative of
a cooperative binding process. The overall shapes of
the binding isotherms are also consistant with a
drug induced allosteric change in the solution
structure of the DNA. Salt concentration influences
the magnitude and curvature of the binding
isotherms. The addition of small concentrations of
daunorubicin affects the shape of the actinomycin D
binding isotherm, in a manner which suggests that
both drugs are able to induce similiar allosteric
changes in the DNA. The dissociation kinetics of
actinomycin D from calf thymus DNA is shown to fit a
multi-exponential decay curve, whereas the
dissociation from poly(dGdC)·poly(dGdC) shows a
single lifetime. The dissociation of the ternary
complex actinomycin D:daunorubicin: poly(dAdT)·
poly(dAdT) is best described by two time constants,
while the dissociation of daunorubicin from the same
polymer in the absence of actinomycin D is
characterized by a single time constant.

During recent years a host of solid state and solution
experiments have shown that DNA may exist in a variety of
conformations (e.g.,1-10 and references therein). Factors
affecting the solution conformation of DNA include salt
concentration (7,8,9), nature of the solvent (10) and base content
(i.e. AT/GC ratio)(3). Pohl et al. (7) showed that at high salt
concentrations the solution structure of the heteropolymer
poly(dGdC)·poly(dGdC) is altered to the so-called "Z" conformation
or left-handed helix, as evidenced by an inversion of the circular
dichroism spectrum. The binding of ethidium to the high salt form
of the polymer is a markedly cooperative process (7). Ethidium
binding to the high-salt conformation of this polynucleotide
alters its conformation to resemble the low salt ethidium:
polynucleotide complex.

Crothers and coworkers (1,11) have shown that the binding of
the oligopeptides netropsin and distamycin to calf thymus DNA is a
cooperative process. These results have been interpreted in terms
of a drug induced allosteric change in the conformation of the
DNA. This report shows that several important antitumor drugs
whose chemical structures are shown in Figure 1 (i.e. actinomycin
D, adriamycin, daunorubicin, DHAQ) and two carcinogens (NQO and
HAAF) bind cooperatively to DNAs. Factors affecting the solution
structure of DNA are shown to alter the degree of cooperativity.
The presence of cooperativity and/or allosteric transitions in the
binding of these ligands to DNA may be an important aspect of
their pharmacological activity.

Crothers and coworkers (1,11) have also shown that the
binding of ethidium to calf thymus DNA in the presence of
distamycin (r(distamycin) = 0.037) is cooperative (1), whereas
the binding of ethidium in the absence of distamycin is non-
cooperative. We report that the binding of daunorubicin to calf
thymus DNA (r(daunorubicin) = 0.037) appears to alter the
structure of the DNA as evidenced by the non-cooperative isotherm
of actinomycin D.

Müller and Crothers (12) have shown that the dissociation of
actinomycin D from DNA is a multistep process. The relaxation
curve for the dissociation of actinomycin D from calf thymus DNA
requires a minimum of three exponentials to fit the decay curve
mathematically. They associated the multistep dissociation
kinetics to the presence of the pentapeptide rings of actinomycin
D which are essential for the pharmacological activity of the
drug. Recently, Krugh et al. (13) showed that the dissociation of
actinomycin D from poly(dGdC)·poly(dGdC) was characterized by a
single exponential decay. This and other related experiments have
led Krugh and coworkers (unpublished results) to conclude that the
different dissociation lifetimes on native DNA result from the
dissociation of actinomycin D from different DNA binding sites.
Presented below is a detailed examination of the dissociation of
actinomycin D from both calf thymus DNA and poly(dGdC)·poly(dGdC).
Krugh and Young (14) have shown that daunorubicin facilitates the

ACTINOMYCIN D

R = H DAUNORUBICIN
R = OH ADRIAMYCIN
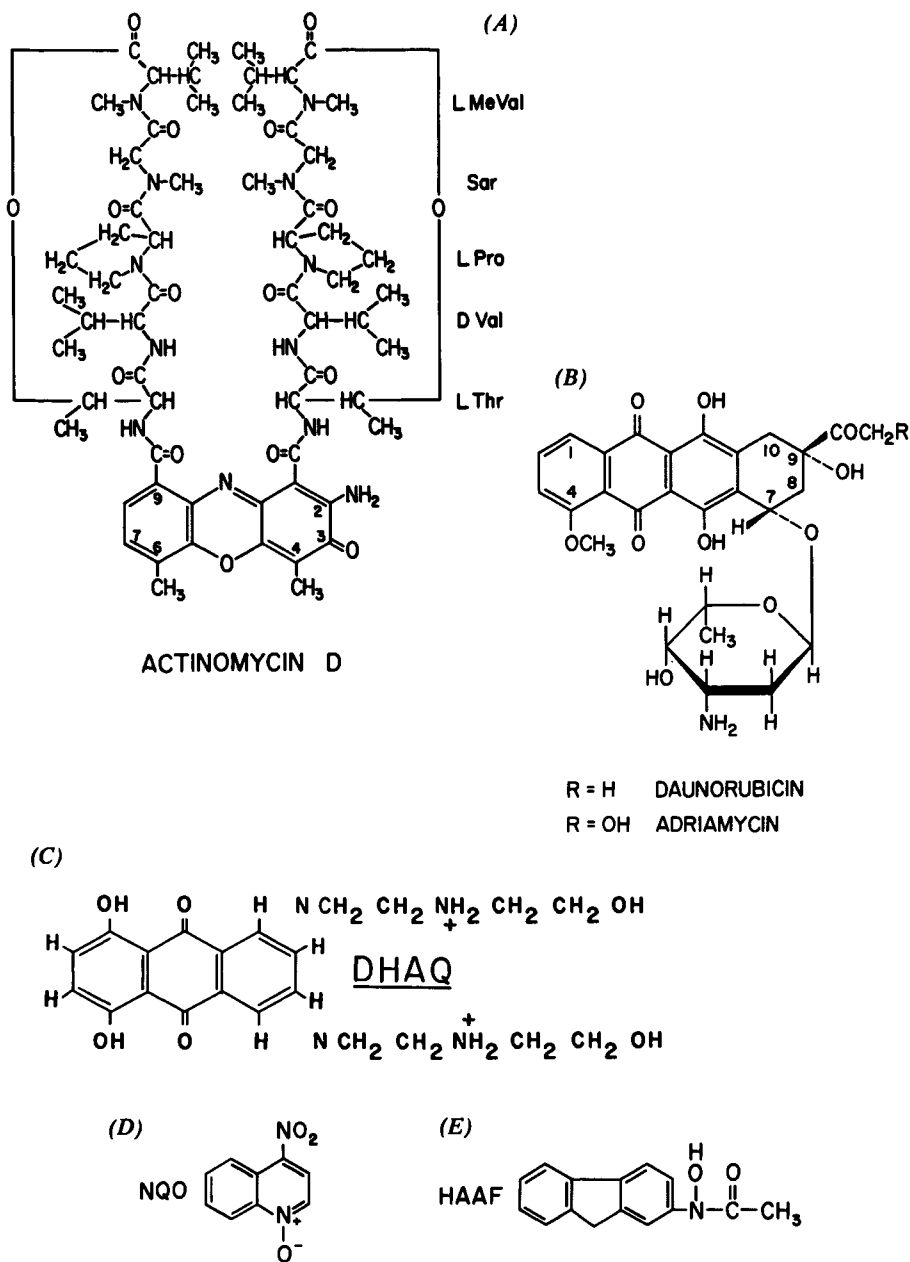
Figure 1. Chemical structure of (A) actinomycin D (key: Thr, threonine; Val, valine; Pro, proline; Sar, sarcosine; and MeVal, methylvaline); (B) adriamycin and daunorubicin; (C) DHAQ; (D) NQO; and (E) HAAF.

binding of actinomycin D to poly(dAdT)·poly(dAdT), a polymer to which actinomycin D binds only weakly. This paper shows that the dissociation kinetics of daunorubicin, in the presence and absence of actinomycin D, reflects the synergistic interaction of these two drugs as mediated by poly(dAdT)·poly(dAdT).

Materials and Methods

1,4-dihydroxy-5,8-bis[[2-[(2-hydroxyethyl)amino]ethyl]amino]-9,10-anthracenedione (DHAQ) a gift from Dr. H. Cox (University of Kansas Medical Center). Actinomycin D was obtained from Merck. Daunorubicin and adriamycin were from the Natural Products Branch, National Cancer Institute. ^{3}H-NQO (4-nitroquinoline-1-oxide) was from the laboratory of Dr. I. Tinoco, Jr. (U.C. Berkeley), ^{14}C-HAAF (N-Hydroxy-N-acetyl-2-amino-fluorene was from the ICN Radiochemicals Division, and ^{3}H-actinomycin D was from Amersham. Calf thymus DNA (Worthington), was prepared according to the procedure of Müller and Crothers (15). Poly(dAdT)·poly(dAdT) and poly(dGdC)·poly(dGdC) were purchased from PL Biochemicals. DNA concentrations were determined absorptiometrically, and expressed in basepairs.

Free and bound concentrations of DHAQ were determined by optical titration using a 10-cm quartz cell. A known volume and concentration of DNA was titrated with a concentrated solution of DHAQ. The increase in the absorbance was read directly from the digital display of a Cary 219 spectrophotometer. Buffer solutions were pH=6.4, 0.052M Na phosphate, 0.1M NaCl, and 0.001M NaEDTA.

The binding of actinomycin D, adriamycin, daunorubicin, HAAF and NQO to nucleic acids was examined utilizing the phase partition technique which allows the bound and free drug concentrations to be determined by direct measurements. Phase partition analysis (16-19, and references therein) consists of the equilibration of an aqueous phase (with or without DNA present) with an immiscible organic phase. The drug may be initially present in either phase. The partition coefficient, P (drug concentration in organic phase/drug concentration in aqueous phase) is determined in the absence of DNA. Upon addition of DNA to the aqueous phase, both phases are equilibrated and seperated. The concentration of the drug in the organic phase is measured and used to calculate the free drug concentration (cf) = [organic drug concentration]/ partition coefficient, P. Drug concentrations in both the organic and aqueous phase can be calculated directly by either scintillation counting or spectrophotometric measurements, as appropriate.

Binding isotherms for both adriamycin and daunorubicin were obtained by phase partition analysis utilyzing both 1-nonanol and chloroform as the organic phase. Aqueous buffer solutions were pH=7, 0.01M Na phosphate, 0.1M NaCl, and 0.001M NaEDTA. Aqueous and organic phases were equilibrated directly in teflon stoppered fluorimeter cells. Organic phase drug concentrations were

determined by direct spectrofluorimetric measurements. Bound drug concentrations were calculated from the total drug concentration in the aqueous phase obtained spectrofluorimetrically after dissociation of the DNA-drug complex by the addition of dimethylsulfoxide (1:1 v/v).

The organic phase for radioactively labeled actinomycin D, NQO, and HAAF was a 1:1 mixture of n-chloroheptane and cyclohexane. For the experiments with HAAF and NQO the aqueous phase was a buffer of 0.1M NaCl, 0.01M Na cacodylate, 0.001M NaEDTA, pH=7. For actinomycin D the buffer was 0.01 M Na phosphate, 0.001M NaEDTA, pH=7, with the addition of NaCl as specified in the Results and Discussion section.

Actinomycin D dissociation kinetics were measured on a Cary 219 spectrophotometer equipped with a magnetic stirrer and thermostated cell holders. Sodium dodecyl sulfate (SDS) was used to sequester dissociating actinomycin D, and the resulting increase in absorbance was monitored at 452 nm as a function of time. Stop-flow studies (daunorubicin and daunorubicin/ actinomycin D) were conducted with a Durrum-Gibson Model 110 stopped-flow spectrophotometer equipped with a dual detector accessory and a Tektronix storage oscilloscope interfaced with a PDP 11/34 computer. Experiments were done in a 0.01M Na phosphate buffer, 0.1M NaCl, 0.001M NaEDTA, pH=7. Dissociation time constants were computed with a multiexponential analysis computer program.

Results and Discussion

Cooperative Binding. The binding isotherm for the interaction of DHAQ with calf thymus DNA (Figure 2) shows an initially increasing slope which is indicative of cooperative binding of DHAQ. At higher r values the binding isotherm changes, and appears to reflect a typical neighbor exclusion binding isotherm.

For DHAQ, increasing either the ionic strength of the buffer (0.2 (0.10M NaCl) to 0.6 (0.50M NaCl)) or the temperature (23 °C to 37 °C) diminishes the initial increasing slope of the DHAQ/DNA binding isotherm. At an ionic strength of 0.6, the slope is only slightly positive in the initial portion of the binding isotherm. Increasing the temperature, at any ionic strength, decreases the initial slope and shows a minimally decreasing Scatchard plot at low concentrations of bound DHAQ. Increasing the pH from 6.4 to 7.1 does not affect the shape of the binding isotherm at low ionic strength. Also, saturating the buffer with 1-nonanol (an organic solvent used in partition analysis experiments) does not affect the shape or magnitude of the binding isotherm.

The binding isotherms for the interaction of the anthracycline antibiotics, adriamycin and daunorubicin, with calf thymus DNA, measured by phase partition techniques, are shown in Figure 3. Both drugs show initially increasing binding isotherms, indicative of a cooperative binding process, and reach a maximum
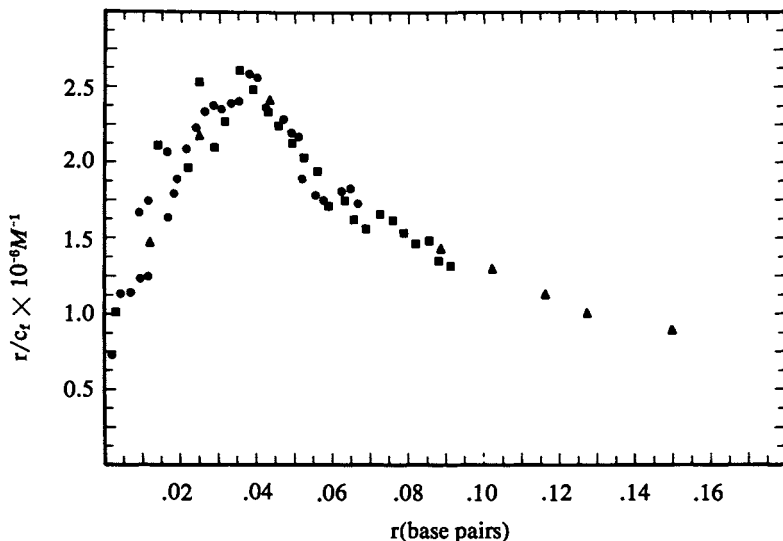
Figure 2.   The binding of DHAQ by calf thymus DNA was evaluated by absorption spectroscopy.

The buffer was sodium phosphate (0.052M), sodium chloride (0.10M), and NaEDTA (0.001M) at pH 6.4 and final ionic strength 0.201. The wavelength of choice was the free drug maxima of 660 nm. Concentrations of the DNA (base pairs) ranged from 7 to 8 × 10⁻⁶M, while the drug concentration ranged from 0.3 to 8.0 × 10⁻⁷M.
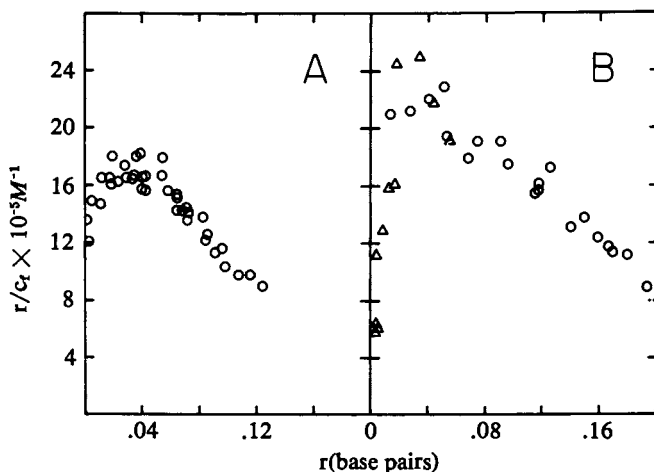
Figure 3.   Scatchard plot of the equilibrium binding of calf DNA with (A) daunorubicin and (B) adriamycin at 22°C in 0.01M sodium phosphate, 0.1M sodium chloride, and 0.001M NaEDTA at pH 7.0.

Key: ○ and △, binding isotherms obtained using 1-nonanol and chloroform, respectively. Bound and free drug concentrations were determined by spectrofluorometric measurements, and excitation and emission wavelengths were 475 and 595 nm, respectively. Drug concentrations ranged from 0.1 to 0.7 μM, while DNA concentrations ranged from 0.15 to 15μM (base pairs).

at r = 0.035-0.040. The cooperativity of both drugs diminish (loss of the initial positive slope) as the sodium chloride concentration (0.10M - 0.01M) is decreased, thus indicating that cooperativity is dependent on the ionic strength.

The binding of actinomycin D to calf thymus DNA (0.01M, 0.1M, and 0.2M NaCl) and to poly(dGdC)·poly(dGdC) (0.1M NaCl) (Figure 4) was also examined by phase partition techniques. For both DNAs, a cooperative process was seen. In the case of calf thymus DNA, the decreasing salt concentration appears to increase the degree of cooperativity.

The effects of salt concentration on the binding of various drugs to DNA have been well documented at higher r values (21-24). Studies by Record et al. (25) and Manning (26) have shown that the counterions of DNAs are "obligate participants" in the interaction of charged ligands with DNA. Although the observed magnitude of the binding isotherms obtained is decreased with increasing [Na^+], the effect of salt on the binding at low r values, where the cooperative binding and allosteric transitions are found, appears to be dependent upon the particular drug being examined. The complex interrelationship between salt concentration and cooperativity is shown by the observation that decreasing the salt concentration may either increase or decrease the curvature in the initial portion of the binding isotherm. For example, DHAQ shows an increasing degree of cooperativity as the salt concentration is decreased, whereas decreasing the salt concentration decreases the degree of cooperativity for both adriamycin and daunorubicin.

The curvature in the HAAF and NQO binding isotherms (Figure 5) indicates a cooperative binding of these carcinogens to ϕX174 DNA (19). A long range allosteric change in the solution structure of the DNA can explain the observed cooperative binding of these carcinogens. An equally plausible interpretation is that of a cluster model in which two carcinogen molecules bind "together" or cluster into a localized site on the DNA (19).

Krugh and Young (14) have shown shown that daunorubicin facilitates the binding of actinomycin D to poly(dAdT)·poly(dAdT). The following experiments illustrate the interaction of two drugs mediated by a polynucleotide helix. The binding of ^{3}H-actinomycin D to calf thymus DNA at 0.02M NaCl in the presence of fixed concentrations of daunorubicin (r(daunorubicin) = 0.015 or r(daunorubicin) = 0.037) was studied by phase partition. The increasing slope in the binding isotherm of actinomycin D (Figure 3) is diminished when the DNA is pre-equilibrated with daunorubicin, (r(daunorubicin) = 0.015) (Figure 6). At higher concentrations of daunorubicin (r(daunorubicin) = 0.037) the initial increasing slope in the actinomycin D binding isotherm is no longer visible. These two r values correspond to the mid-point of the increasing slope and the peak maximum of the daunorubicin: calf thymus DNA binding isotherm. These results suggest a similar mode of binding for these two antitumor drugs. The kinetic studies
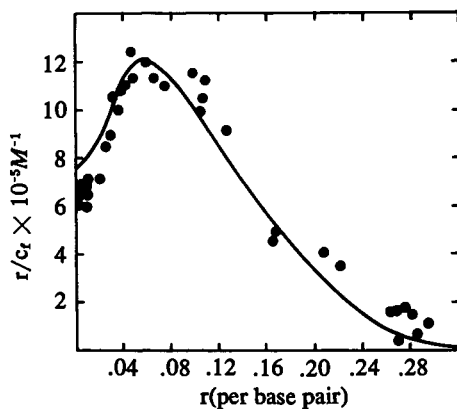
Figure 4. Scatchard plot for the binding of actinomycin D to poly(dGdC) · poly-(dGdC) in 0.01M sodium phosphate, 0.1M sodium chloride, and 0.01M NaEDTA at pH 7.0 and 5°C. Concentrations of the poly(dGdC) · poly(dGdC) (base pairs) ranged from 16μM to 2.3mM, while the actinomycin D concentrations ranged from 0.35μM to 0.64mM.
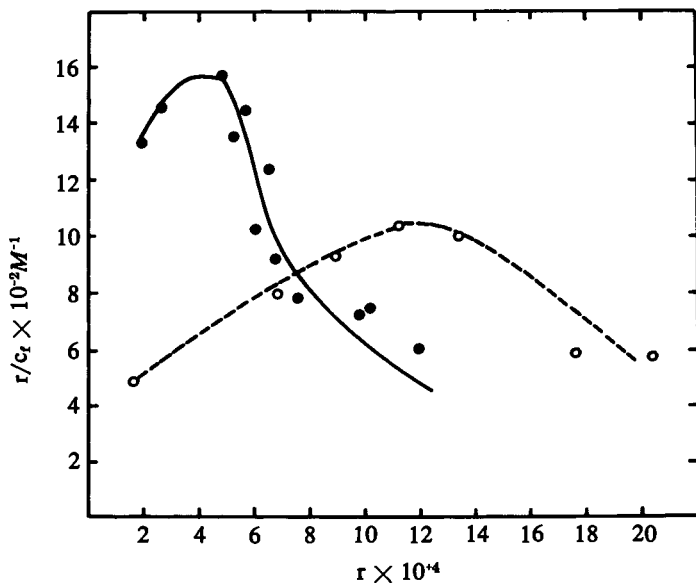
Figure 5. Scatchard plots for carcinogens binding to φX174RF DNA. Key: ●–●, HAAF + φX174RF; and ○–○, NQO + φX174RF; [φX174RF] = 0.1–1.0 × 10⁻⁴M; [HAAF]ₜ = 0.1–7.0 × 10⁻⁵M; and [NQO]ₜ = 0.05–5.0 × 10⁻⁵M. DNA concentrations are in terms of base pairs. The curves through the data are for viewing purposes.
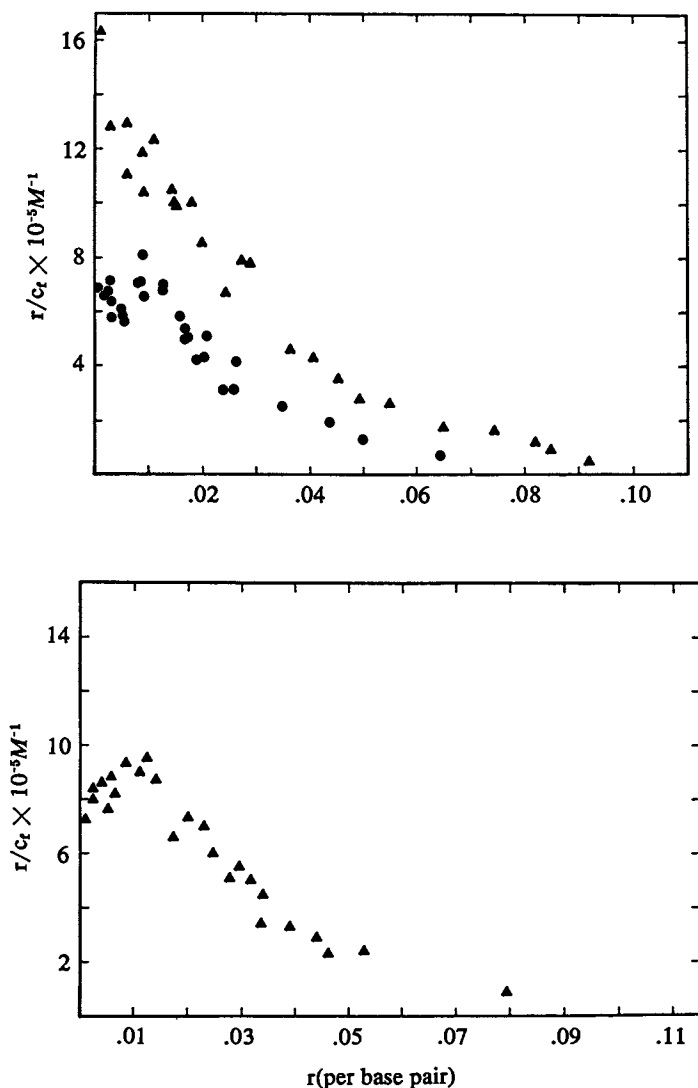
Figure 6.   Scatchard plot for the binding of actinomycin D to calf thymus DNA: daunorubicin complex.

Equilibrium binding was measured by the solvent partition method in 10mM sodium phosphate buffer, pH 7.0, 22°C in 0.02M NaCl using ^3H-actinomycin D, partitioning against 1-chloroheptane:n-hexane (1:1). Key ●, DNA daunorubicin (r = 0.037) and ▲, DNA:daunorubicin (r = 0.015); $[DNA]_T = 0.3–1.5 \times 10^{-4}M$ (base pairs); $[dauno-rubicin]_T = 0.45–5.4 \times 10^{-6}M$; $[actinomycin D]_T = 0.02–1.0 \times 10^{-5}M$.

on daunorubicin:actinomycin D mixed-drug dissociation from
poly(dAdT)·poly(dAdT) also shows that the presence of bound
daunorubicin influences the binding of actinomycin D.

Kinetics. The dissociation of actinomycin D from poly(dGdC)·
poly(dGdC) is well described by a single exponential decay (Table
I). The dissociation of actinomycin D is shown to be dependent on
the phosphate to drug ratio (P/D ratio). At high P/D (low degree
of saturation) the dissociation lifetime is about 900 sec. At
higher degrees of saturation the lifetime increases until a final
value of about 1400 sec is reached. Neither the concentration of

Table I
Dissociation lifetime of actinomycin D from Poly(dGdC)·Poly(dGdC)
as a function of phosphate/drug ratio (T=22°C) and temperature.

P/D	sec	Temp (°K)	sec
3	1400	290.2	2468
10	1414	295.2	1376
20	1047	300.0	907
30	944	302.1	670
40	914	305.2	529
50	861	310.1	296

SDS nor the pH affects the actinomycin D dissociation lifetime;
however, the dissociation time is markedly dependent on both
temperature (Table I) and salt concentration (Table II).
Decreasing either the salt concentration or the temperature
increases the dissociation time, but in neither case does the
dissociation change from a single exponential. The divalent metal
ion, Mg^{++}, is more effective in lowering the dissociation lifetime
than Na^+ (Table II). This suggests that a divalent ion is more
effective in lowering the activation barrier than a monocation.

Table II
Dissociation lifetime of actinomycin D from Poly(dGdC)·Poly(dGdC)
as a function of Na^+ and Mg^{++} concentration.

$[Na^+]$ (M)	dissociation lifetime (sec)	$[Mg^{++}]$ (M)	dissociation lifetime (sec)
0.053	2624	0.00	2624
0.103	1720	0.02	1098
0.150	1428	0.04	794
0.202	1365	0.08	618
0.322	1261		

The dissociation kinetics of actinomycin D from calf thymus DNA is best fit by at least three exponential decay time constants (Table III), indicating that native DNA has a heterogeneous set of binding sites. The slow dissociation lifetime is associated with the pharmacological activity of this drug and the strongest binding sites. Varying the P/D ratio does not significantly alter this slow dissociation lifetime; however, the percentage of absorbance change associated with this lifetime does depend upon the P/D ratio. At low degrees of saturation the slow dissociation component is the dominant one, accounting for almost 70% of the absorbance change. As the degree of saturation is increased the percentage absorbance associated with this component decreases to about 20% of the total absorbance change. This is due to an increased population of faster dissociating binding sites. Thus the kinetic experiments provide direct evidence for heterogeneity of the binding sites and suggest that the cooperative binding of actinomycin D is associated with the high affinity sites.

Table III.
Relative amplitudes of τ_1, τ_2, and τ_3 for actinomycin D dissociation from calf thymus DNA.

P/D ratio	τ_1 (sec)	τ_2 (sec)	τ_3 (sec)	relative amplitude		
				A (%)	A(%)	A(%)
5	10	70	1600	37.7	43.7	18.6
10	25	90	1410	36.5	38.2	25.4
15	25	80	1292	31.8	41.2	26.9
30	25	81	1250	21.3	40.1	38.5
100	42	216	1527	23.7	15.2	61.1
200	48	305	1517	20.5	12.5	66.6
250	60	486	1600	21.9	10.8	67.3

The dissociation kinetics of daunorubicin and daunorubicin: actinomycin D from poly(dAdT)·poly(dAdT) were studied using stopped-flow techniques. The data is summarized in Table IV. The dissociation of daunorubicin from poly(dAdT)·poly(dAdT) is monophasic with a lifetime of 0.2 sec. In the ternary complex (daunorubicin:actinomycin D:poly(dAdT)·poly(dAdT)) the dissociation of the daunorubicin is characterized by two exponential decays, yielding lifetimes of 0.2 sec and 5 sec. A single 5 sec dissociation time was observed for actinomycin D. This data is consistent with a model in which daunorubicin is bound to isolated DNA binding sites and dissociates with its normal lifetime of 0.2 sec. Whereas daunorubicin and actinomycin D molecules, residing at "adjacent sites", dissociate with a time constant of approximately 5 sec. These results are indicative of a cooperative interaction between daunorubicin and actinomycin D when the two are bound to poly(dAdT)·poly(dAdT).

Table IV
Dissociation of Poly(dAdT)· Poly(dAdT):drug complexes at 22 °C.

Poly(dAdT)·Poly(dAdT) :drug complex	Ratio base pairs :drug	τ_1 sec	τ_2 sec
Daunorubicin	20:1	0.2	---
Actinomycin D	20:1	0.7	---
Daunorubicin: Actinomycin D	20:1:1	0.2	5.0

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