

Figure 1. Chy- fragments isolated after chymotrypsin cleavage; T- fragments established by Edman degradation; —, sequences established by Edman degradation; - - -, sequences established by tlc, electrophoresis, and amino acid analysis. †, CNBr cleavage. The degradation fragment marked by an asterisk was isolated and synthesized as the carboxy-terminal homoserine analog.

Peptide fragments isolated after cleavage with chymotrypsin are also shown in Figure 1. The important tripeptide 20–22 (Chy-4), which overlaps two tryptic fragments, was identified by Edman degradation and by synthesis. The tyrosyl-tryptophan bond in the chymotryptic fragment Chy-1 was stable to chymotrypsin both in the natural and in the synthetic tetrapeptide.

Treatment of the tryptic fragment 15–21 with thermolysin⁹ afforded phenylalanylhistidylarginine as well as fragment 16–18. Degradation of fragment 22–32, obtained from trypsin cleavage, with CNBr gave phenylalanylserylglycylhomoserine as well as fragment 26–32. The presence of an amino-terminal phenylalanine in the tetrapeptide was shown by the dansylation procedure for end-group analysis.¹⁰ It was confirmed by degradation with leucine aminopeptidase which also served to demonstrate the presence of serine in position 23. Moreover, the tetrapeptide was found to be indistinguishable from a synthetic specimen. Sequence 26–31 was established by Edman degradation of fragment 26–32. Leucine aminopeptidase confirmed the presence of the terminal sequence Gly-Phe (26–27).

Hydrolysis of thyrocalcitonin with 0.03 *N* HCl at 100° for 9 hr liberated leucine and aspartic acid as the only detectable amino acids. This result is consistent¹¹ with the presence of an Asn-Leu-Asn fragment in the hormone. The tryptic fragment 15–21 (T-2) was also found to be identical with a synthetic⁷ specimen.

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The Reaction of Derivatives of Tryptophan, Tryptamine, and Other Indoles with 2-Hydroxy-5-nitrobenzyl Bromide (Koshland's Reagent)

Sir:

Although interest in the highly selective reagent 2-hydroxy-5-nitrobenzyl bromide (1) for the rapid modification and assay of tryptophan in proteins continues,^{1–7} there is no information on the structure of the reaction products.^{7a} A recent report⁸ on the complex reactions of the reagent with tryptophan, both free and bound in proteins, prompts us to communicate our findings on the reaction of 1 with skatole (2), 2,3-dimethylindole (3), N-acetyltryptamine (4), and N-acetyl-L-tryptophan methyl ester (5).

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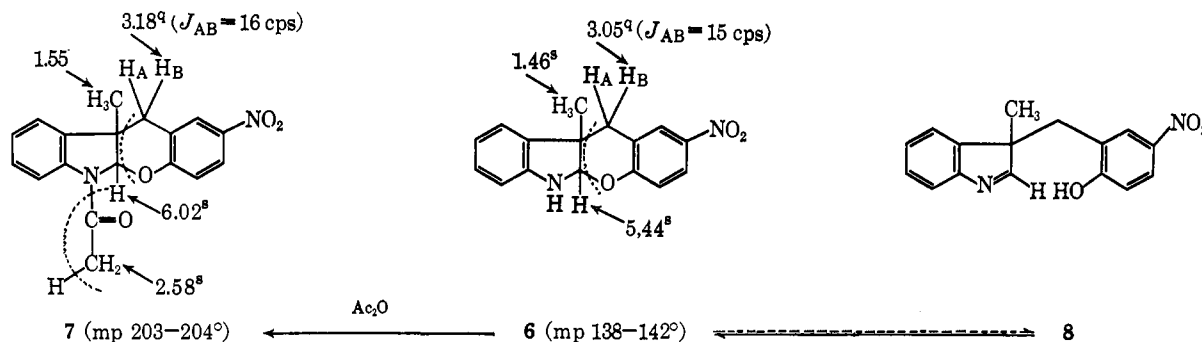
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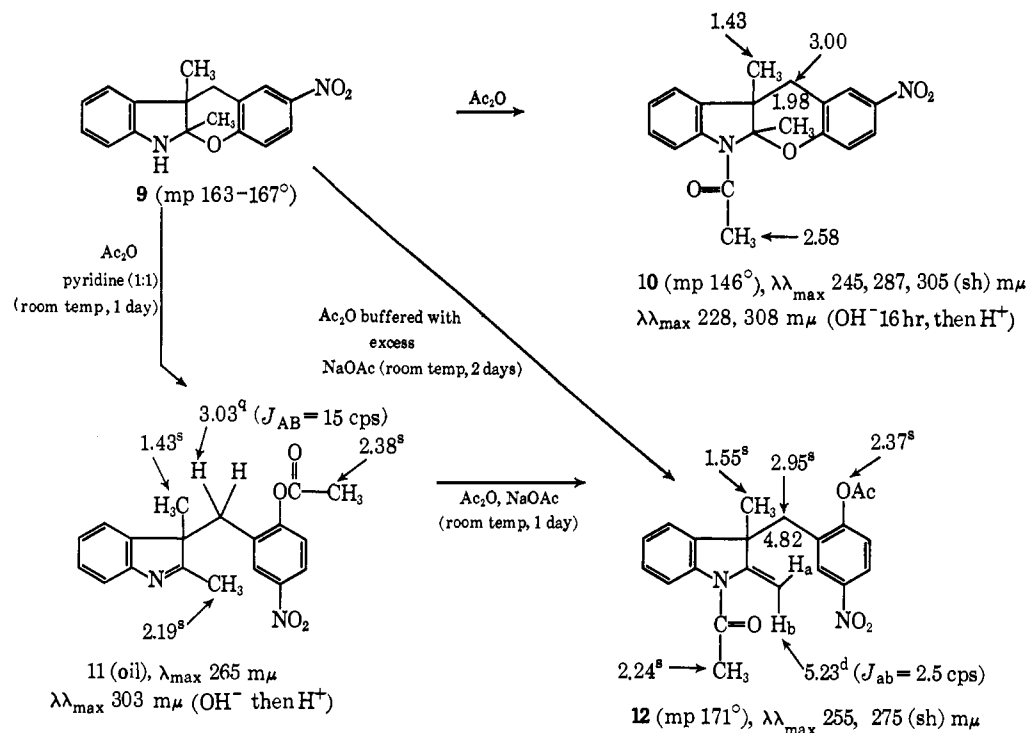
(7a) NOTE ADDED IN PROOF. Under somewhat different conditions 2-alkylation of skatole has been reported: M. Wakselman, G. Decodts, and M. Vilkas, *Compt. Rend.*, **266**, 1090 (1968).

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$\lambda\lambda_{\max}$ 285, 310 m μ (95% EtOH)

λ_{\max} 310 m μ (OH⁻, 12 hr, then H⁺ in 95% EtOH)



When a solution of **1** in dry acetone was added to an equimolar quantity of **2** in 50% acetone–water (final composition 72% acetone), a homogeneous, yellow, crystalline material (mp 138–142° after recrystallization from 80% acetone) separated in 70% yield. Combustion analysis and mass spectrometry (m/e 282, parent molecular ion) established formula C₁₆H₁₄N₂O₃, *i.e.*, a 1:1 reaction product. Principal peaks were observed at m/e 132 and 131 (intensity ratio \cong 3:1), equivalent to the loss from the parent ion of C₇H₅NO₃ and C₇H₆NO₃, respectively. The nmr spectrum (CDCl₃; TMS internal standard) supports structure **6**: a three-proton singlet at 1.46 ppm, a two-proton quartet ($J_{AB} = 15$ cps) at 3.05 ppm, a one-proton singlet at 5.44 ppm, a five-proton multiplet at 6.4–7.0 ppm, and a two-proton multiplet at 7.6–7.9 ppm. D₂O was without effect on this spectrum. The ir spectrum (CHCl₃) exhibited three principal peaks above 1525 cm⁻¹ at 3450 (sharp), 1615, and 1585 cm⁻¹, the latter two being the most intense. On acetylation (Ac₂O, room temperature), a colorless microcrystalline N-acetate, **7** (mp 203–204°), was formed in quantitative yield. The analysis, parent molecular ion (m/e 324), and the ir (1670 cm⁻¹; 3450 cm⁻¹ had disappeared) and nmr

spectra [1.55 (3 H), singlet; 2.58 (3 H), singlet; 3.18 (2 H), quartet ($J_{AB} = 16$ cps); 6.02 (1 H), singlet; 6.8–7.3 (4 H), multiplet; 7.8–8.1 ppm (3 H), multiplet] support structure **7**. The dotted lines show the principal mode of cleavage in the mass spectrometer.

The upfield shift of the 3-methyl singlet from 2.30 ppm in **2** to 1.46 and 1.55 ppm in **6** and **7** provides compelling evidence for alkylation in the 3 position. Similarly, allyl bromide alkylates 1,2-dimethylindole to a mixture of two bis(alkylated) products, the major component being a 3,3-diallyl derivative.⁹ The presence of a one-proton singlet at 5.44 (**6**) and 6.02 ppm (**7**) supports the carbinolamine structure¹⁰ and rules out the isomeric indolenine **8**, which may well be the initial product of alkylation. Spontaneous cyclization of **8** should lead to the preferred *cis* ring juncture indicated for **6** and **7**.

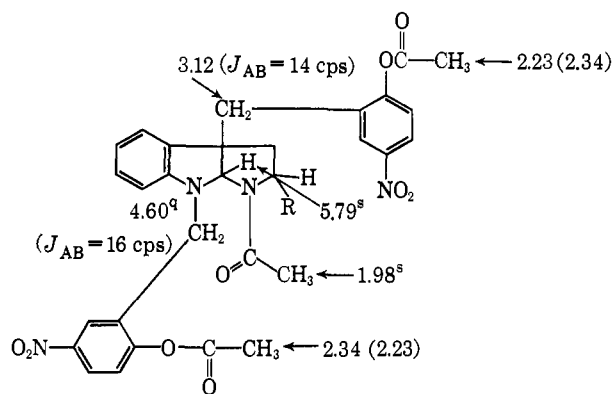
2,3-Dimethylindole (**3**) under the same reaction conditions as **2** formed an initial adduct (**9**) with 1 mole of reagent. Acetylation of this product gave the three products, **10**, **11**, or **12**, depending upon the experimental conditions (see diagram). In each case, the

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product was nearly homogeneous; **10** and **12** were crystallized directly. **11** was an unstable oily material which resisted crystallization but was smoothly converted to **12** by further acetylation with Ac_2O - NaOAc . Pertinent nmr assignments are indicated on the structures.

Preliminary experiments on the alkylation of N-acetyltryptamine (**4**) with 1 equiv of **1** indicated that the major product contained 2 moles of reagent. Accordingly the alkylation was carried out with 2 equiv of **1**. Purification was effected inadvertently during an acetylation attempt with Ac_2O (room temperature) when the major product crystallized directly from the acetylation mixture as a 1:1 complex with Ac_2O (60% yield). This material dissolved readily in CHCl_3 at room temperature and, after 20–30 min, deposited an almost insoluble 1:1 chloroform complex of the composition $\text{C}_{26}\text{H}_{24}\text{N}_4\text{O}_7\text{-CHCl}_3$. An ir spectrum (KBr) indicated associated phenolic stretching vibrations ($3000\text{--}3400\text{ cm}^{-1}$) and a hydrogen-bonded (tertiary?) amide carbonyl (1620 cm^{-1}). This material was acetylated with Ac_2O -pyridine (1:1) to give a quantitative yield of the N,O,O-triacetate **13**, $\text{C}_{30}\text{H}_{28}\text{N}_4\text{O}_9$, obtained solvent free on crystallization from EtOAc -hexane or as a 1:1 CHCl_3 complex from CHCl_3 -hexane. A high-resolution mass measurement of the parent molecular ion indicated m/e 588.178 (calcd 588.183). A low-resolution mass spectrum was consistent with the consecutive loss of two units of m/e 194 each from the parent. The molecular extinction of 35,800 at $400\text{ m}\mu$ in alkaline ethanol is consistent with a molecular weight of 707 for the chloroform complex of **13**, from which base forms 2 moles of the characteristic chromophore of the *p*-nitrophenoxide ion (ϵ_{410} 18,000). Structure **13**, though awaiting confirmation by X-ray crystallography, is preferred to other alternatives such as the open indolenine tautomer with N_6 carrying acyl and benzyl groups.



13, R = H (mp $164\text{--}167^\circ$; CHCl_3 adduct, mp $96\text{--}100^\circ$)

14, R = COOMe (mp $95\text{--}105^\circ$)

$\lambda\lambda_{\text{max}}$ 252, 270 (sh) $\text{m}\mu$

$\lambda\lambda_{\text{max}}$ (OH^- then H^+) 246, 315 $\text{m}\mu$

Three products resulted when N-acetyl-L-tryptophan methyl ester (**5**) was treated with 2 equiv of the reagent and then acetylated with Ac_2O -pyridine (1:1 room temperature, 3 days). The major component (37% after silica chromatography) had the composition $\text{C}_{32}\text{H}_{30}\text{N}_4\text{O}_{11}$ (mp $95\text{--}105^\circ$ after crystallization from ether-ligroin). The near identity of its uv spectrum with **13**, before and after base treatment, and many close similarities in its nmr spectrum (**13**) suggest the analogous

structure **14**. Such a dihydroindole system would no longer show the reactivity that bound tryptophan shows toward N-bromosuccinimide.⁶

Acknowledgment. We wish to acknowledge our indebtedness to Dr. G. W. A. Milne for determining the exact mass for compound **13** and to Dr. Takashi Tokuyama for obtaining several of the nmr spectra.

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The Selective Photolysis of Dihydrothymidine

Sir:

Of all the building stones of DNA, thymine most easily undergoes photochemical transformations, such as dimerization^{1–3} or possibly reduction,⁴ events which are of genetic and mutagenic interest. This report shows that dihydrothymidine, in contrast to dihydrouridine, undergoes a selective “photochemical hydrolysis” of the Ciamician–Silber type⁵ which, though well known for homocyclic systems,⁶ is here demonstrated for the first time for a heterocyclic representative of biological significance.

In a typical run, a $2 \times 10^{-2}\text{ M}$ aqueous unbuffered solution of dihydropyrimidine (**1a** or **1b**) was irradiated with a 250-W high-pressure mercury lamp (Hanovia S654-36, no filter) which was surrounded by a cylindrical water-cooled quartz jacket. Two semicircular quartz vessels which surrounded the cooler and contained the sample were 3 cm from the light source. Irradiation of 1-dihydrothymidine (**1b**)⁷ for 30 hr led to complete disappearance of the uv absorption at $230\text{ m}\mu$. The photoproduct, a yellow oil, was lyophilized, chromatographed over silica gel, and eluted with chloroform-methanol (9:1). The three major fractions yielded (*S*)-(-)-dihydrothymine, mp 263° (**1a**),⁷ *n*-propylurea (**VIa**, 6%), mp 109° (*p*-nitrobenzoate mp $170\text{--}171^\circ$ dec), and N_1 -deoxyribosyl- N_1 -*n*-propylurea (**VIb**, 64%), which on acid hydrolysis gave *n*-propylurea, mp 109° , and deoxyribose (diphenylamine test). Dihydrothymine on photolysis gave 75% *n*-propylurea (**VIa**) and 5% urea. Under identical conditions the photolysis of dihydrouridine led only to minor cleavage of the ribosyl residue and to the isolation of 5% of dihydrouracil.

There are two likely reaction mechanisms. Pathway A would begin with homolytic cleavage between the carbonyl group (position 4) and the ureido nitrogen

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The chemical evidence for the *S* configuration of the asymmetric center at C-4 of 1-dihydrothymidine has been supplemented and confirmed by a complete X-ray structure analysis (I. L. Karle, Naval Research Laboratory). There is evidence that exposure to radiation by X-ray leads to partial conversion of dihydrothymidine to thymidine, and probably of dihydrothymine [cf. S. Furberg and L. H. Jensen, *ibid.*, **90**, 470 (1968)] to thymine.