Phosphinyl Acid-Based Bisubstrate Analog Inhibitors of Ras Farnesyl Protein Transferase

Dinesh V. Patel,^{*,†} Eric M. Gordon,[†] Robert J. Schmidt, Harold N. Weller, Marian G. Young, Robert Zahler, Mariano Barbacid, Joan M. Carboni, Johnni L. Gullo-Brown, Lisa Hunihan, Carol Ricca, Simon Robinson,[‡] Bernd R. Seizinger, Anne V. Tuomari, and Veeraswamy Manne

The Bristol-Myers Squibb Pharmaceutical Research Institute, P.O. Box 4000, Princeton, New Jersey 08543-4000

Received September 19, 1994[®]

The rational design, synthesis, and biological activity of phosphonyl- and phosphinyl-linked bisubstrate analog inhibitors of the enzyme Ras farnesyl protein transferase (FPT) are described. The design strategy for these bisubstrate inhibitors involved connection of the critical binding components of the two substrates of FPT (ras protein and farnesyl pyrophosphate, FPP) through a phosphonyl- or phosphinyl-bearing linker. Compound 14, the first example in this series, was found to be a potent FPT inhibitor ($I_{50} = 60$ nM). A further 15-fold enhancement in activity was observed upon replacement of the VLS tripeptide sequence in 14 with VVM (15, $I_{50} = 6$ nM). The phosphinic acid analog 16 ($I_{50} = 6$ nM) was equiactive to phosphonic acid 15. Compounds 14-16 afforded 1000-fold selectivity for FPT against the closely related enzyme geranylgeranyl protein transferase type I, GGT-I [14, I_{50} (GGT-I) = 59 μ M; 15 I_{50} (GGT-I) = 10 μ M; 16 I₅₀(GGT-I) = 21 μ M]. Methyl and POM ester prodrugs 17-19 were prepared and evaluated in whole cell assays and appear to block ras-induced cell transformation, as well as colony formation in soft agar. A distinctive feature of this novel class of potent and selective bisubstrate FPT inhibitors is that they are non-sulfhydryl in nature.

Introduction

Ras genes play a fundamental role in cell proliferation and differentiation, and ras oncogenes have been implicated in the development of a variety of tumors.¹ Ras proteins must become membrane bound in order to exert their normal as well as oncogenic effects. This mandatory membrane binding occurs following a defined sequence of post-translational modifications, namely farnesylation, proteolysis, and carboxymethylation.² Inhibition of S-farnesvlation, which is the first and essential step in this cascade of events, may thus be expected to block the oncogenic activity of ras proteins. This hypothesis has recently led to the search for inhibitors of farnesyl protein transferase $(FPT)^3$ as novel, rationally designed, antitumor agents.⁴

FPT catalyzes the transfer of a farnesyl group from farnesyl pyrophosphate (FPP), an intermediate in the cholesterol biosynthetic pathway, to ras proteins. S-Farnesylation occurs on the cysteine located at the 4th position from the C-terminus of the protein. The "CAAX" motif is a consensus sequence for farnesylation, where C is cysteine, A is an aliphatic amino acid, and X is preferably serine or methionine.³ Since FPT catalyzes a bisubstrate reaction, it provides various opportunities for inhibitor design. Thus, inhibitors based on either of the two substrates can be prepared, and examples of both CAAX-4, 5 and FPP-based⁶ inhibitors have recently appeared in the literature. An alternative and unexplored approach is the design of a collective bisubstrate inhibitor involving the incorporation of critical binding components of both the reacting partners (FPP and ras) in a chemically and biologically stable form. We herein report the design, synthesis, and biological activity of a new class of potent, rationally designed bisubstrate analog inhibitors of FPT.

Bisubstrate Inhibitor Design

A hypothetical "active site model" of FPT was constructed from knowledge of the substrate reaction and structural information about the enzyme (Figure 1).^{6a,7} The model comprises a peptide binding domain that accommodates the critical "CAAX" motif of Ras, and a lipid binding domain in which the hydrophobic farnesyl group of FPP resides. The reaction formally proceeds by a nucleophilic attack by the thiol group of cysteine on the allylic carbon atom of FPP bearing the pyrophosphate group. An enzymic base may facilitate the reaction by abstracting the thiol proton and enhancing the nucleophilicity of the thiol group. A suitable metal ion (e.g., Mg^{2+}) may serve to complex with the PP group in order to neutralize charge and set the stage for displacement reaction. The role of Zn^{2+} ion is unclear; it may be coordinated with the thiol and/or the PP group, and/or with the free CO₂H group of Ras, or it may simply serve to stabilize the heterodimeric form of the enzyme.

The inhibitor design strategy involved hybridization of the two reacting partners into a single chemically and biologically stable entity. In our inhibitor design, the farnesyl group of FPP was retained to preserve putative hydrophobic interactions, and the ras C-terminal tripeptide was chosen as the peptide substrate component.⁸ These two entities were connected via a phosphonic or phosphinic acid linker.⁹ The phosphonic or phosphinic acid group was designed to interact with the zinc and/or magnesium metals associated with the cysteine sulfhydryl at the active site.^{7b} The functional equivalence of mercaptan and phosphonyl or phosphinyl acid moieties has precedent in the area of metalloprotease inhibitors, including inhibitors of the zinc metalloproteases angiotensin-converting enzyme (ACE) and

⁺Current address: Affymax Research Institute, 3410 Central Expressway, Santa Clara, CA 95051. [†]Current address: BASF Bioresearch Corporation, 100 Research

Drive, Worcester, MA 01605

Abstract published in Advance ACS Abstracts, December 15, 1994.



Figure 1. Hypothetical active site model of FPT.

neutral endopeptidase (NEP).⁹ The length of the linker was selected so as to allow conformational flexibility between the two substrate fragments, enabling them to adopt a transition state like geometry. In addition to structural novelty, incorporation of a phosphonic or phosphinic acid in place of the substrate sulfhydryl group avoids biological issues associated with a mercaptan moiety.¹⁰ Additionally, the chemical equivalence of phosphinyl or phosphonyl oxygens avoids the introduction of an additional chiral center. This led to the proposal of evaluating phosphonic and phosphinic acid containing bisubstrate analogs 14-16 as FPT inhibitors.

Chemistry and Biology

Preparation of the phosphonic acid bisubstrate inhibitor 14 began with an Arbuzov reaction of trimethyl phosphite with farnesyl bromide affording the dimethyl phosphonate 2 in good yield (78%). Monohydrolysis of ${f 2}$ under basic conditions followed by O-alkylation of the resulting acid 3 with ethyl bromoacetate gave 4(100%). Conversion of diester 4 to carboxylic acid 5 was cleanly accomplished by careful hydrolysis with 1 equiv of base (91%). EDC-mediated¹¹ coupling of carboxylic acid 5 with the tripeptide H-VLS-OMe (9) gave the diester precursor 11 (50%). Selective monodemethylation at the phosphorus center by heating with sodium iodide in acetone (45%) gave intermediate 17, which upon treatment with 1 N NaOH in dioxane afforded the desired diacid 14 (80%).¹² Coupling of 5 with the tripeptide amine H-VVM-OMe (10) utilizing the BOP¹¹ reagent, followed by similar sequence of reactions as outlined for 14, provided the VVM analog 15. To prepare the phosphinic acid analog of 15, alkylation of phosphinate 6^{13} with farnesyl bromide (85%) was followed by selective monohydrolysis to carboxylic acid 8 (90%). Coupling of 8 with 10 using BOP provided the diester intermediate 13, which was converted by one-step base hydrolysis to the desired analog 16 (Scheme 1).

Compound 14, the first example in this series, was found to be a potent FPT inhibitor $(I_{50} = 60 \text{ nM})$.¹⁴ A further 15-fold enhancement in activity was observed upon replacement of the VLS tripeptide sequence in 14 Scheme 1^a



^a Reagents: (a) $P(OMe)_3$, reflux, 78%; (b) 1 N NaOH, MeOH, quant; (c) K_2CO_3 , DMF, BrCH₂CO₂Et, quant; (d) 1 N NaOH (1.0), MeOH, 91% for **5**, 90% for **8**; (e) NaH, HP(O)(OMe)CH₂CO₂Me **6**, 85%; (f) EDC, HOBt, iPr₂NEt, HCl·H-VLS-OMe **9**, 50%; (g) BOP, iPr₂NEt, HCl·H-VVM-OMe **10**, 63% for **12**, 97% for **13**; (h) NaI, acetone, reflux, 45% for **14**, 52% for **17**; (i) 1 N NaOH, dioxane, 80% for **14**, 83% for **15**, 60% for **16**; (j) TMSBr, BSTFA, 63% or Me₃N, acetone, 47%; (k) 1 N NaOH (1.0), MeOH, 73%; (l) tBuC(O)OCH₂I (10.0), Ag₂CO₃ (2.0), MeP(O)(OMe)₂ (10.0), DMF, 79%; (m) BSTFA, TMSBr, 62%. a: Compounds **17–19** were prepared and evaluated as prodrugs.

with VVM (15, $I_{50} = 4$ nM), a trend that parallels the structure-activity relationship observed for simple tetrapeptide-based inhibitors.⁴ The phosphinic acid analog 16 ($I_{50} = 6$ nM) was essentially equiactive to phosphonic acid 15. This result suggests that the phosphonic ether oxygen of 14 and 15 contributes minimally to binding, and that the binding affinity of the phosphonate moiety is primarily derived from the charged oxygen.

These inhibitors were evaluated for selectivity using the closely related enzyme geranylgeranyl protein transferase type I (GGT-I),¹⁴ whose preferred CAAX sequences contain hydrophobic amino acids at the X position.¹⁴ Compounds 14–16 were only moderately active against GGT-I [14, I_{50} (GGT-I) = 59 μ M; 15 I_{50} -(GGT-I) = 10 μ M; 16 I_{50} (GGT-I) = 21 μ M], thereby

Phosphinyl Acid-Based Bisubstrate Analog Inhibitors

affording greater than 1000-fold selectivity in favor of the targeted enzyme, FPT.

Having established the *in vitro* potency and selectivity of these inhibitors, we next tested them in cell-based models for inhibition of ras-mediated cell transformation.⁵ Since the parent diacids have very limited cell permeability, prodrugs of the lead compounds 15 and 16 were prepared for their evaluation in whole cell assays. Carboxylic methyl ester 17, the prodrug of phosphonic bisubstrate inhibitor 15, was already encountered as the penultimate precursor in the synthesis of the parent diacid 15. Carboxylic methyl ester 18 in the phosphinic bisubstrate series was prepared from diester 13 via selective demethylation at the phosphorus center by heating with trimethylamine in acetone (47%), or more efficiently by selective hydrolysis (63%) upon treatment with TMSBr in the presence of excess bis-(trimethylsilyl)trifluoroacetamide (BSTFA). Alternatively, careful treatment of diester 13 with 1.0 equiv of 1 N NaOH enabled selective hydrolysis of the carboxylic methyl ester group. The resulting acid intermediate was alkylated with pivaloyloxy methyl iodide under forcing conditions (79%) followed by treatment with BSTFA/TMSBr to provide the POM ester 19 (62%).

The effect of these inhibitors and their prodrugs in whole cells was examined in a ras transformation inhibition (RTI) assay. NIH-3T3 cells transfected with oncogenic H-ras DNA were treated with the appropriate compounds 24 h after transfection, and the effect on transformation was evaluated after 14 days. While the parent diacids 15 and 16 are essentially ineffective, the methyl ester prodrugs 17 and 18 show 75 to 80% decrease in transformed foci at 100 µM concentration (Figure 2). No gross cytotoxicity was observed for any of the compounds at the tested concentrations. The ability of prodrugs 18 and 1915 to inhibit anchorage independent colony growth of ras transformed cells was tested by soft agar growth assay. At 100 µM concentrations, both 18 and 1915 almost completely suppress colony formation of H-ras transformed NIH 3T3 cells in soft agar (Figure 3).

Summary

In summary, the rational design and synthesis of phosphonyl and phosphinyl acids 14-16 constitute the first examples of potent nonsulfhydryl bisubstrate analog inhibitors of FPT.20 Unlike tetrapeptide-based inhibitors, a free sulfhydryl group is not a requirement for the activity of these phosphonic / phosphinic acid bisubstrate *inhibitors.* The presence of a farnesyl group in these molecules may diminish their dependence on the peptide fragment for overall activity and thus enable the discovery of novel FPT inhibitors with reduced peptide character. As anticipated, these inhibitors are selective for FPT when compared to geranylgeranyl transferase GGT-I.¹⁴ The methyl and POM ester prodrugs 17-19 appear to block ras-induced cell transformation in NIH3T3 cells,^{15,21} as well as colony formation in soft agar. The preliminary results reported here support FPT as a target for rational discovery of antitumor agents, a conclusion that has also been derived from the independent work of other research groups.5 Studies describing a thorough biological evaluation of these new compounds, with appropriate controls, will be reported elsewhere.15



No Adds Salts Carrier H-ras Alone Alone DNA





Figure 2. Ras transformation inhibition by compounds 14–18.

Experimental Section

General. All reactions were carried out under a positive pressure of dry argon, unless otherwise specified. Tetrahydrofuran (THF) and diethyl ether were distilled from sodium or potassium/benzophenone ketyl prior to use. TLC was performed using EM Science 5×10 cm plates precoated with silica gel 60 F₂₅₄ (0.25 mm thickness), and the spots were visualized by any of the following: UV, iodine, phosphomolybdic acid (PMA), ceric ammonium sulfate, anisaldehyde, vanillin, or Rydons stain. EM Science silica gel 60 (230–400 mesh ASTM) was used for flash chromatography. A ratio of 25-100:1 silica gel/crude product by weight and a nitrogen pressure of 5-25 psi was normally employed for flash columns. Reverse phase chromatographic purification of final compounds was carried out using CHP20P gel, a 75–150 μ m polystyrene-divinylbenzene copolymer purchased from Mit-

Colony Formation of RAS Transformed Cells in Soft Agar



Untreated Control



Treated with 100 µM Compound 18

Figure 3. Inhibition of colony formation in soft agar by compound 18.

subishi Chemical Industries. Analytical HPLC was performed using two Shimadzu LC-6A pumps with an SCL-6B system controller, a C-R4AX Chromatopac integrator, and an SPD-6AV UV-vis spectrophotometric detector. HPLC columns were from YMC Corp. (YMC S3 120A ODS, 6.0 × 150 mm) and were eluted with gradients of methanol/water containing 0.2% phosphoric acid. Melting points were determined on an electrothermal Thomas-Hoover capillary melting point apparatus and are uncorrected. NMR spectra were recorded on one of the following instruments: JEOL GX-400 operating at 400 (¹H) or 100 MHz (¹³C), JEOL FX-270 operating at 270 (¹H) or 67.8 (13C) MHz and JEOL FX-60Q operating at 15 MHz (¹³C). Chemical shifts are reported in parts per million (ppm) downfield from tetramethylsilane (TMS), and coupling constants (J) are in hertz (Hz). IR spectra were recorded on a Mattson Sirius 100 FT-IR spectrophotometer, and the absorption maxima are reported in cm⁻¹. Mass spectra were recorded on a Finnigan MAT TSQ-4600 mass spectrometer (chemical ionization, CI) or a VG-ZAB-2F mass spectrometer (fast atom bombardment, FAB). High-resolution mass spectra (HRMS) were determined using peak matching techniques versus PEG standards on a VG-ZAB-2F spectrometer. Optical rotations were measured using a Perkin-Elmer model 241 polarimeter and a 10 cm path length optical cell. All compounds were homogeneous by TLC and HPLC. Microanalysis results were adjusted to obtain the best fit assuming nonstoichiometric hydration.

Farnesyl Protein Transferase (FPT) Inhibition Assay. Farnesyl protein transferase was isolated from pig brain as described by Manne et al.^{3a} and further purified approximately 2000-fold relative to the initial crude pig brain cytosol.¹⁶ The PXCR expression vector containing H-ras was kindly provided by Dr. Larry Feig (Tufts University, Medford, MA). Recombinant p21 H-ras was expressed in the E. coli strain PR13Q and processed as described by Farnsworth et al.¹⁷ Following processing and ammonium sulfate precipitation the pellet was resuspended in 50 mM Tris-HCl, pH 7.5 20 mM NaCl, 1 mM DTT, 1 mM PMSF, 1 mM benzamidine, 10 µg/mL soybean trypsin inhibitor, $10 \,\mu\text{M}$ E64, and $1 \,\mu\text{M}$ pepstatin and dialyzed overnight. The recombinant p21 H-ras was then partially purified by FPLC using a DE52 column and a linear NaCl gradient from 20 to 320 mM. Fractions containing the p21 H-ras were visualized by Coomassie blue-stained SDS-polyacrylamide gels and assayed for substrate capacity using the farnesylation assay described below. p21 H-Ras with a purity of >60% was obtained with this single column purification. Additional processing often led to a more pure protein with a reduced capacity to be farnesylated (assumed to be a consequence of the carboxyl terminus cleavage described in Farnsworth et al.¹⁷). Farnesyl protein transferase assays were run in 96-well dishes in a reaction volume of 20 μ L. The final reaction mixture contained 1 μ M [³H] FPP (NEN Dupont), 7 µM p21 H-Ras, 25 mM MgCl₂, 10 mM DTT, 100 mM HEPES 7.4, and serial dilutions of inhibitor usually ranging from 360 to 0.02 μ M. Reactions were started by adding sufficient enzyme to produce approximately 2 pmol of [3H]FPP incorporation in 1 h in the control wells. Following incubation at 37 °C for 1 h the reactions were stopped by adding 90 μ l of 4% sodium dodecyl sulfate (SDS) followed by 90 µL of 30% TCA. Plates were incubated overnight at 4 °C, and then the precipitates were transferred to Millipore multiscreen filtration 96-well plates with 0.65 PVDF membranes (Millipore Corp., Bedford, MA). Following filtration using the multiscreen vacuum manifold, the wells were washed once with 200 μ L of 4% SDS/6%TCA and five times with 200 µL of 6% TCA. Following removal of the bottom seal, excess washing fluid was blotted and the plates were allowed to dry before the filters were punched into 4 mL vials using the multiscreen punch. After incubation at 60-70 °C with 300 µL of Solvable (NEN Dupont), 3 mL of Formula 989 (Dupont) scintillation fluid was added and radioactivity determined by scintillation counting. Dose-response curves for inhibitors used triplicate estimates at each drug concentration, and the IC₅₀ estimations were made from percent control versus log drug concentration plots. Each compound was tested at least twice.

Geranylgeranyl Transferase (GGT-I) Inhibition Assay. GGT-I was purified from porcine brain tissue. Ras-CVLL proteins were expressed in bacteria and purified as described.¹⁸ In order to determine the IC₅₀ values for inhibition, the GGT-I assay was adapted to a 96-well microtiter plate format as described.¹⁸ Briefly, 12 μ M p21 rasCVLL, 0.5 μ M [³H]GGPP (19.3 Ci/mmol; NEN), 10 mM dithiothreitol (DTT), 5 mM MgCl₂, 5 μ M ZnCl₂, 100 mM Hepes, pH 7.4, and partially purified GGTI were incubated in a total volume of 20 μ L at 37 °C for 1 h, and samples were processed as described for the farnesyltransferase assay. All assays were carried out in triplicate and SEMs determined.

Ras Transformation Inhibition (RTI) Assay. The whole cell activity of FPT inhibitors was evaluated by a ras transformation inhibiton (RTI) assay. The RTI assay is based on transformation of mouse NIH 3T3 cells by oncogenic ras DNA transfection. NIH 3T3 cells $(2.5 \times 10^4 \text{ per } 35 \text{ mm well})$ were seeded and allowed to attach overnight. The cells were transfected with linearized plasmid carrying oncogenic H-ras¹⁹ using the calcium phosphate precipitation technique. One day following the transfection, cells were washed and inhibitors were added at the indicated concentrations into the medium. Inhibitors were replenished every 48 h for 8 days along with the change of medium, for a total of four treatments. The degree of transformation was scored after 14 days. In the absence of an inhibitor, transfected cells grow aggressively and initially give the appearance of abnormal "foci" surrounded by normal cells. In the presence of a cell permeable inhibitor, the number of foci are either reduced or transformation is completely inhibited. The percent inhibition and the gross cytotoxicity of inhibitors are evaluated visually by light microscopy.

Soft Agar Growth (SAG) Assay. The SAG assay is based on the ability of ras-transformed cells to grow in soft agar and form discrete colonies in 14-21 days. A bottom agar layer of 0.6% Noble agar (Difco) in culture medium (2 mL, Dulbecco's Modified Eagles Medium (DMEM) from Gibco supplemented with 10% calf serum from Colorado Serum Co.) was formed in 6-well culture plates and allowed to solidify at room temperature for 1-2 h. Cultures were trypsinized, and single cell suspensions were obtained. A top soft agar mixture (2 mL) containing 0.3% Noble agar, 10 000 cells, and culture medium was laid on the solidified bottom agar layer. Initially, inhibitors at the indicated concentrations were incorporated into the soft agar mixture. The cells were cultured for 2 weeks at 37 °C. FT inhibitors were replenished every 2 days for 8 days by overlaying 100 μ L of 20-fold concentrated stock in PBS on the surface of the top agar layer. All the assays are carried out in parallel duplicate wells. At the end of the test period, the number of colonies larger than 0.1 mm in diameter was determined for both treated and untreated samples.

Preparation of (*E,E*)-(3,7,11-Trimethyl-2,6,10-dodecatrienyl)phosphonic Acid, Dimethyl Ester (2). A mixture of farnesyl bromide 1 (14.26 mL, 52.6 mmol) and trimethyl phosphite (6.84 mL, 57.9 mmol) was heated at reflux for 16 h. The excess trimethyl phosphite was removed *in vacuo*, and the residue was purified by flash chromatography, eluting with hexane:ethyl acetate (10:1 to 6:1) to afford 2 (8.03 g, 78%) as an oil which was homogeneous by TLC ($R_f = 0.8$ in hexane: ethyl acetate, 1:1, visualized by ceric ammonium sulfate) and HPLC. Mass spectrum (M + H)⁺: calculated for C₁₇H₃₂O₃P 315.2089, found 315.2099. ¹H NMR (270 MHz, CDCl₃): 1.58 (s, 6H), 1.66 (s, 6H), 2.06 (br m, 8H), 2.56 (dd, 2H, J = 7.6, 21.7), 3.73 (d, 6H, J = 10.5), 5.08–5.18 (m, 3H). ¹³C NMR (68 MHz, CDCl₃): 15.7, 15.9, 17.4, 24.2, 25.4, 26.1, 26.2, 26.4, 39.4, 52.2, 52.3, 111.7, 111.9, 123.5, 124.0, 130.9, 135.0, 140.1, 140.3.

Preparation of (E,E)-(3,7,11-Trimethyl-2,6,10-dodecatrienyl)phosphonic Acid, Monomethyl Ester (3). To a solution of 2~(8.03~g,~25.6~mmol) in methanol (70 mL) was added 1 N NaOH (77 mL, 76.7 mmol), and the reaction was refluxed for 20 h. Methanol was removed in vacuo, and the aqueous solution was diluted with water (40 mL) and extracted with ethyl acetate $(2 \times 40 \text{ mL})$. The organic extracts were discarded, and the aqueous solution was acidified by addition of 1 N aqueous hydrochloric acid (pH = 2.0) in the presence of ethyl acetate (40 mL). After the layers were separated, the aqueous solution was reextracted with ethyl acetate (3×40) mL), and the combined organic extracts were dried over sodium sulfate, filtered, and concentrated in vacuo to give 3 (7.71 g, 100%). ¹H NMR (270 MHz, CDCl₃): 1.59 (s, 6H), 1.67 (s, 6H), 2.06 (br m, 8H), 2.57 (dd, 2H, J = 7.6, 22.2), 3.70 (d, 3H, J = 11.1), 5.08–5.22 (m, 3H). ¹³C NMR (68 MHz, CDCl₃): 15.9, 16.2, 17.6, 24.9, 25.6, 26.3, 26.4, 26.6, 27.0, 39.6, 51.7, 51.8, 111.8, 112.0, 123.7, 124.3, 131.1, 135.1, 140.5, 140.7.Anal. $(C_{16}H_{29}O_3P) C, H.$

Preparation of (E,E)-[[Methoxy(3,7,11-trimethyl-2,6, 10-dodecatrienyl)phosphinyl]oxy]acetic Acid, Ethyl Ester (4). To a solution of 3 (1.5 g, 5 mmol) in DMF (25 mL) was added potassium carbonate (4.15 g, 30 mmol) and ethyl bromoacetate (1.66 mL, 15 mmol). The reaction was stirred at 25 °C for 4 h and filtered through Celite. The precipitate was washed with ethyl acetate, and the filtrate and the washes were combined and washed sequentially with 10% aqueous hydrochloric acid (50 mL) and saturated aqueous lithium chloride $(3 \times 60 \text{ mL})$, dried over sodium sulfate, filtered, and concentrated in vacuo. The residue was purified by flash chromatography, eluting with hexane:ethyl acetate (3:1 to 1:1) to give 3 (1.95 g, 100%), which was homogeneous by TLC (R_f = 0.37 in hexane:ethyl acetate, 1:2, visualized by ceric ammonium sulfate stain). Mass spectrum $(M + H)^+$: calculated for C₂₀H₃₆O₅P 387.2247, found 387.2307. ¹H NMR (270 MHz, CDCl₃): 1.29 (t, 3H, J = 7.0), 1.60 (s, 6H), 1.68 (s, 6H), 2.08 (br m, 8H), 2.68 (dd, 2H, J = 7.6, 22.2), 3.77 (d, 3H, J = 11.1),4.24 (q, 2H, J = 7.0), 4.57 (d,d, 2H, J = 2.9, 11.1), 5.09–5.22 (m, 3H). ¹³C NMR (68 MHz, CDCl₃): 14.0, 15.9, 16.2, 17.6, 25.2, 25.6, 26.3, 26.4, 26.6, 27.3, 52.2, 52.3, 61.3, 62.2, 62.3, 111.6, 111.7, 123.7, 124.2, 131.2, 135.2, 140.8, 141.0, 168.4,168.5.

Preparation of (E,E)-[[Methoxy(3,7,11-trimethyl-2,6, 10-dodecatrienyl)phosphinyl]oxy]acetic Acid (5). To a solution of 4 (1 g, 2.59 mmol) in methanol:water (2:1) was added at 0 °C 1 N aqueous sodium hydroxide solution (2.59 mL, 2.59 mmol). After stirring for 3 h at 0 °C, methanol was removed in vacuo, and the aqueous solution was diluted with water (30 mL) and extracted with ethyl acetate (2×20 mL). The organic extracts were discarded, and the aqueous layer was acidified with 1 N hydrochloric acid (pH = 2.0) in the presence of ethyl acetate (40 mL). After the layers were separated, the aqueous solution was reextracted with ethyl acetate $(3 \times 30 \text{ mL})$. The combined organic extracts were dried over sodium sulfate, filtered, and concentrated in vacuo to give 5. Mass spectrum $(M + H)^+$: calculated for $C_{18}H_{32}O_5P$ 359.1988, found 359.2014. ¹H NMR (270 MHz, CDCl₃): 1.59 (s, 6H), 1.67 (s, 6H), 2.07 (m, 8H), 2.73 (dd, 2H, J = 7.6, 22.8),3.79 (d, 3H, J = 11.1), 4.60 (d, 2H, J = 12.3), 5.09-5.18 (m, J = 12.3H). ¹³C NMR (68 MHz, CDCl₃): 16.0, 16.3, 17.7, 25.1, 25.7, 26.4, 26.5, 26.7, 27.2, 38.7, 52.9, 53.0, 62.8, 62.9, 111.1, 111.3,123.7, 123.8, 124.3, 131.3, 135.4, 141.4, 141.6, 170.7.

Preparation of 3-(Methoxyphosphinyl)propanoic Acid, Methyl Ester (6). Compound 6 was prepred by slight modifications of the literature procedure.¹³ Methyl hypophosphite (6 g, 75 mmol) was added to a solution of methyl acrylate (6.8 mL, 75.8 mmol) and diisopropylethylamine (1.34 mL, 7.73 mmol) at 0 °C, after which the mixture was warmed to room temperature and stirred for 24 h. The reaction was quenched with saturated aqueous sodium bicarbonate solution (15 mL) and extracted with chloroform (4 \times 50 mL), and the combined organic extracts were dried (MgSO₄), filtered, and concentrated in vacuo. The residue was purified by vacuum distillation (97-100 °C at 0.7 mmHg) to afford 6 as an oil (5.0 g, 40%). Mass spectrum $(M + H)^+$: calculated for C₅H₁₂O₄P 167.0474, found 167.0472. IR (neat): 3445-2999, 2957, 2373, 1738, 1441, 1244, 1044 cm⁻¹. ¹H NMR (CDCl₃): 2.03-2.15 (m, 2 H), 2.54-2.80 (m, 2 H), 3.72 (s, 3 H), 3.80 (d, 3 H, J = 11.7), 7.19 (d, 1 H, J = 548.2). ¹³C NMR (CDCl₃): 22.9, 24.3, 25.6, 25.7, 51.8, 52.0, 52.7, 52.8, 172.1, 172.3 ppm.

Preparation of (E,E)-3-[Methoxy(3,7,11-trimethyl-2,6, 10-dodecatrienyl)phosphinyl]propanoic Acid, Methyl Ester (7). Sodium hydride (60% suspension in oil, 0.36 g, 9.0 mmol) was added to a solution of 6 (1.5 g, 9.0 mmol, 1.0 equiv) in THF (20 mL) at -5 °C, and the mixture was stirred for 15 min. Farnesyl bromide (2.45 mL, 9.0 mmol) was added dropwise, and the reaction was warmed to room temperature and stirred for 16 h. The reaction was quenched by addition of saturated aqueous ammonium chloride solution (50 mL) and extracted with ethyl acetate (4 \times 50 mL), and the combined organic extracts were dried (MgSO₄), filtered, and concentrated in vacuo. The residue was purified by flash chromatography (eluting with 3:1 hexane/acetone) to afford 7 (2.79 g, 85%). TLC: $R_f = 0.59$ (1:1 hexane/acetone, visualization by PMA). Mass spectrum $(M + H)^+$: 371. ¹H NMR (CDCl₃): 1.60 (s, 6 H), 1.66 (s, 3 H), 1.68 (s, 3 H), 1.97-2.18 (m, 10 H), 2.54-2.66 (m, 4 H), 3.70 (s, 3 H), 3.71 (d, 3 H, J = 10.6), 5.08-5.19 (m, 3 H). ¹³C NMR (CDCl₃): 15.9, 16.2, 17.5, 21.6, 22.9, 25.5, 26.2, 26.3, 26.6, 27.9, 29.2, 39.6, 51.0, 51.1, 51.9, 112.1, 112.3, 123.5,124.1, 131.2, 135.3, 140.6, 140.8, 172.5, 172.7. Anal. (C₂₀H₃₅- $PO_4 \cdot 0.81H_2O) C, H.$

Preparation of (E,E)-3-[Methoxy(3,7,11-trimethyl-2,6, 10-dodecatrienyl)phosphinyl]propanoic Acid (8). Sodium hydroxide (1 N, 7.54 mL, 7.54 mmol) was added to a 0 °C solution of 7 (2.79, 7.54 mmol) in methanol (21 mL) and water (10 mL) and stirred at this temperature for 3 h. The reaction mixture was concentrated in vacuo, and the residue was diluted with water (50 mL) and washed with chloroform (1 \times 50 mL). The organic wash was discarded, and the aqueous solution was acidified by addition of concentrated hydrochloric acid to pH 1.0 and was extracted with methylene chloride $(4 \times 50 \text{ mL})$. The combined organics extracts were dried (MgSO₄), filtered, and concentrated in vacuo to afford $\mathbf{8}$ (2.42 g, 90%), which was homogeneous by TLC ($R_f = 0.63, 6:3:1$ 1-propanol/ammonium hydroxide/water, visualization by PMA). Mass spectrum $(M + H)^+$: 357. ¹H NMR (CDCl₃): 1.60 (s, 6 H), 1.66 (s, 3 H), 1.68 (s, 3 H), 1.96–2.10 (m, 10 H), 2.52-2.70(m, 4 H), 3.72 (d, 3 H, J = 10.6), 5.06-5.30 (m, 3 H). ¹³C NMR Preparation of (E,E)-N-[N-[N-[[[Methoxy(3,7,11-trimethyl-2,6,10-dodecatrienyl)phosphinyl]oxy]acetyl]-Lvaly]-L-leucy]-L-serine, Methyl Ester (11). To a solution of 5 (843 mg, 2.35 mmol) in THF (70 mL) were added sequentially hydroxybenzotriazole (369 mg, 2.35 mmol), diisopropylethylamine (408 μ L, 2.35 mmol), tripeptide 9 (valylleucylserine methyl ester hydrochloride, 864 mg, 2.35 mmol), and N-(3-(dimethylamino)propyl)-N'-ethylcarbodiimide hydrochloride (450 mg, 2.35 mmol) at 0 °C. The reaction mixture was warmed gradually to room temperature, and after stirring for 20 h, the suspension was diluted with ethyl acetate (200 mL) and washed sequentially with saturated aqueous sodium bicarbonate (100 mL), 10% sodium bisulfate (100 mL), and water (3 \times 100 mL). The organic layer was dried over sodium sulfate, filtered, and concentrated in vacuo. The residue was purified by flash chromatography, eluting with hexane:acetone (2:1 to 1:1), to give 11 (775 mg, 50%), which was homogeneous by TLC ($R_f = 0.55$, acetone:hexane, 1:1, visualized by ceric ammonium sulfate). Mass spectrum (M + H)+: calculated for C₃₃H₅₉O₉N₃P 672.3993, found 672.3989. ¹H NMR (270 MHz, CDCl₃): 0.92 (m, 12H), 1.59 (s, 6H), 1.68 (s, 6H), 1.69 (br m, 3H), 2.07 (m, 9H), 2.68 (dd, 2H, J = 7.6, 21.6), 3.76 (s, 3H), 3.78 (d, 3H, J = 10.5), 3.82-4.61 (m, 7H), 5.08-5.16 (m, 3H). ¹³C NMR (68 MHz, CDCl₃): 16.0, 16.4, 17.7, $18.1,\ 18.2,\ 19.2,\ 21.9,\ 22.0,\ 22.9,\ 24.7,\ 25.7,\ 26.4,\ 26.5,\ 26.6,$ 26.7, 30.7, 31.0, 39.7, 40.4, 40.7, 51.9, 52.0, 52.5, 53.2, 53.6, 53.7. 54.9, 58.5, 58.7, 62.6, 63.9, 110.7, 110.8, 110.9, 123.5, 124.3, 131.3, 135.5, 135.5, 141.9, 168.1, 168.2, 170.8, 171.3, 171.3, 172.1.

Preparation of (E,E)-N-[N-[N-[[[Methoxy(3,7,11-trimethyl-2,6,10-dodecatrienyl)phosphinyl]oxy]acetyl]-Lvalyl]-L-valyl]-L-methionine, Methyl Ester (12). (Benzotriazol-1-yloxy)tris(dimethylamino)phosphonium hexafluorophosphate (BOP) (0.442 g, 1.0 mmol) was added to a solution of 5 (0.358 g, 1.0 mmol) and tripeptide 10 (valylvalylmethionine methyl ester hydrochloride, 0.398 g, 1.0 mmol) in acetonitrile (5 mL) and DMF (3 mL). Diisopropylethylamine (0.349 mL, 2.0 mmol) was added, and the reaction was stirred for 16 h at room temperature. The reaction was quenched by addition of hydrochloric acid (1 N, 50 mL), and the mixture was extracted with ethyl acetate $(4 \times 50 \text{ mL})$. The combined organic extracts were washed sequentially with aqueous sodium carbonate solution (10%, 50 mL) and aqueous lithium chloride (10%, 2×100 mL), dried (MgSO₄), filtered, and concentrated in vacuo. The residue was purified by flash chromatography (eluting with 3:2 hexane/acetone) to afford 12 (0.44 g, 63%) as an oil, which was homogeneous by TLC ($R_f =$ 0.42, 1:1 acetone/hexane, visualization by PMA). Mass spectrum $(M + H)^+$: calculated for $C_{34}H_{60}N_3O_8PS$ 702.3917, found 702.3928. $[\alpha]_D = -20.5^\circ$ (c = 1.06, CHCl₃). IR (CH₂Cl₂ film): 3279, 2963, 1751, 1640, 1549 cm⁻¹. ¹H NMR (CDCl₃): 0.93-0.97 (m, 12 H), 1.59 (s, 6 H), 1.67 (s, 6 H), 1.97-2.19 (m, 12 H), 2.07 (s, 3 H), 2.47 (t, 2 H, J = 7.0), 2.67 (dd, 2 H, J = 7.6, 22.3), 3.73, 3.75 (s, 3 H), 3.77, 3.78 (d, 3 H, J = 10.6, 11.2, diastereomers), 4.36-4.69 (m, 5 H), 5.17-5.20 (m, 3 H), 7.08-7.42 (m, 3 H). ¹³C NMR (CDCl₃): 15.4, 16.0, 16.4, 17.7, 18.1, 18.3, 19.1, 19.2, 25.7, 26.4, 26.7, 29.3, 29.9, 30.9, 31.1, 39.7 51.5, 52.5, 58.4, 58.5, 58.7, 63.9, 64.0, 111.0, 111.1, 111.2, 123.6,124.3, 131.3, 135.5, 167.7, 167.7, 170.9, 171.1, 172.1 ppm.

Preparation of (E,E)-N-[N-[N-[3-[Hydroxy(3,7,11-trimethyl-1-oxo-2,6,10-dodecatrienyl)phosphinyl]-1-oxo-propyl]-L-valyl]-L-walyl]-L-methionine, Methyl Ester (13). Disopropylethylamine (0.349 mL, 2.0 mmol) was added to a solution of 8 (0.356 g, 1.0 mmol and tripeptide 10 (valylvalyl-methionine methyl ester hydrochloride, 0.398 g, 1.0 mmol) in acetonitrile (12 mL) and DMF (4 mL) at room temperature. To the mixture was added BOP (0.443 g, 1.0 mmol), and the reaction mixture was stirred an additional 16 h. The reaction was quenched by addition of 1 N hydrochloric acid (50 mL) and extracted with ethyl acetate (4 \times 50 mL); the combined organic extracts were washed with 10% aqueous sodium carbonate (50 mL) and 10% aqueous lithium chloride (3 \times 50

mL), dried (MgSO₄), filtered, and concentrated in vacuo. The residue was purified by flash chromatography (eluting with 2:1 acetone/hexane) to afford 13 (0.350 g, 50%) as an oil, which was homogeneous by TLC ($R_f = 0.8\bar{6}$, 2:1 acetone:hexane, visualization by PMA). Mass spectrum $(M + H)^+$: calculated for C₃₅H₆₂N₃O₇PS 700.4124, found 700.4150. IR (CH₂Cl₂ film 1638 cm⁻¹. $[\alpha]_D = -38.4^{\circ} (c = 0.19, CH_3OH)$. ¹H NMR (CD₃-OD): 0.90-1.00 (m, 12 H), 1.60 (s, 6 H), 1.62 (s, 3 H), 1.66 and 1.69 (s, 3 H, diastereomers), 1.93-2.18 (m, 14 H), 2.06 (s, 3 H), 2.48-2.63 (m, 6 H), 3.71 (s, 3 H), 3.72 (d, 3 H, J = 10.6), 4.13 (d, 1 H, J = 8.8), 4.17 (d, 1 H, J = 7.6), 4.56–4.61 (m, 1 H), 5.08-5.11 (m, 3 H). ¹³C NMR (CDCl₃): 15.0, 15.8, 16.2, 17.5, 18.0, 18.2, 18.3, 18.8, 19.0, 25.5, 26.3, 26.5, 27.3, 29.1, 29.8, 30.7, 31.7, 39.5, 51.4, 51.5, 51.6, 52.3, 53.6, 58.7, 58.9, 59.6, 69.5, 111.5, 111.6, 123.4, 124.1, 131.1, 135.3, 141.1, 141.3, 171.9, 172.0, 172.1, 172.7 ppm.

Preparation of (E,E)-N-[N-[N-[[[Hydroxy(3,7,11-trimethyl-2,6,10-dodecatrienyl)phosphinyl]oxy]acetyl]-Lvalyl]-L-leucyl]-L-serine, Disodium Salt (14). To a solution of 11 (161 mg, 0.240 mmol) in acetone (2 mL) was added sodium iodide (181 mg, 1.20 mmol). After refluxing for 8 h, the reaction was diluted with ethyl acetate (50 mL) and was washed sequentially with 10% aqueous sodium thiosulfate (30 mL) and saturated sodium chloride (30 mL). The aqueous washes were combined and extracted with chloroform:methanol (10:1, 6×50 mL). The ethyl acetate and chloroform: methanol extracts were combined, dried over sodium sulfate, filtered, and concentrated *in vacuo*. The crude material was purified by flash chromatography, eluting with chloroform: methanol (100:0 to 90:10), to give the phosphonic acid carboxylic methyl ester intermediate (74 mg, 45%). TLC: $R_f =$ 0.92 (1-butanol:acetic acid:water:ethyl acetate, 1:1:1:1, visualized by ceric ammonium sulfate). ¹H NMR (400 MHz, CD₃-OD): 0.96 (m, 12H), 1.58 (s, 6H), 1.66 (s, 6H), 1.58-1.73 (m, 3H), 1.94-2.17 (m, 9H), 2.47 (dd, 2H, J = 7.7, 21.8), 3.74 (s, 3H), 3.86 (m, 2H), 4.20-4.56 (m, 5H), 5.06-5.32 (m, 3H). ¹³C NMR (100 MHz, CD₃OD): 16.2, 16.6, 17.8, 18.5, 19.9, 22.4, 23.4, 25.9, 26.0, 27.8, 32.6, 40.9, 41.1, 41.5, 48.1, 48.4, 48.7, 49.0, 49.3, 49.6, 49.9, 50.0, 52.9, 53.5, 56.3, 59.4, 62.7, 64.3, 64.4, 117.3, 117.5, 125.3, 125.4, 132.1, 136.1, 138.5, 138.7, 171.9, 172.7, 172.8, 173.6, 174.5 ppm.

To a solution of the above intermediate (74 mg, 0.109 mmol) in dioxane:water (2:1, 1.5 mL) was added 1 N sodium hydroxide (0.12 mL, 0.12 mmol). After stirring for 2 h, the dioxane was removed *in vacuo*. The aqueous solution was purified on CHP 20P (6 mL), eluting with water:acetonitrile (100:0 to 0:100). Fractions containing the major product were combined, concentrated, and lyophilized to give 14 (60 mg, 80%) as a white powder, mp 150–152 °C. TLC: $R_f = 0.81$ (BuOH:HOAc: H₂O:EtOAc, 1:1:1:1, visualized by ceric ammonium sulfate). Mass spectrum (M + Na)⁺: calculated for C₃₁H₅₄N₃O₉PNa 666.3501, found 666.3495. ¹H NMR (400 MHz, DMSO-*d*₆): 0.83 (m, 12H), 1.55 (s, 6H), 1.58 (m, 3H), 1.62 (s, 3H), 1.23–1.66 (m, 3H), 1.93–2.23 (m, 9H), 2.27 (dd, 2H, J = 6.41, 20.94), 3.62 (br m, 2H), 3.99–4.39 (5H), 5.04–5.22 (m, 3H). Anal. (C₃₁H₅₂N₂O₉Na₂'3.36H₂O) C, H, N.

Preparation of (E,E)-N-[N-[N-[[[Hydroxy(3,7,11-trimethyl-2,6,10-dodecatrienyl)phosphinyl]oxy]acetyl]-Lvalyl]-L-valyl]-L-methionine, Disodium Salt. Sodium hydroxide (1 N, 0.567 mL, 0.567 mmol) was added to a solution of 17 (0.15 g, 0.218 mmol, 1.0 equiv) in dioxane (11 mL), and the mixture was stirred at room temperature for 16 h. The reaction was concentrated in vacuo, and the residue was purified on a column of CHP-20 gel (gradient elution 0-50%aqueous acetonitrile). Fractions containing the major product were combined and concentrated in vacuo, and the residue was dissolved in water (10 mL), millipore filtered, and lyophilized to afford 15 (0.13 g, 83%) as a white powder, mp > 220 °C dec. TLC: $R_f = 0.70$ (6:3:1 1-propanol/ammonium hydroxide/ water, visualization by PMA). Mass spectrum $(M + Na)^+$: 696. IR (KBr): 1645 cm⁻¹. ¹H NMR (CD₃OD): 0.94–1.00 (m, 12 H), 1.59 (s, 6 H), 1.66 (s, 6 H), 1.93-2.16 (m, 12 H), 2.05 (s, 3 H), 2.41-2.51 (m, 4 H), 4.14-4.36 (m, 5 H), 5.09-5.11 (m, 2 H), 5.31-5.34 (m, 1 H). $[\alpha]_D = -19.8^{\circ}$ (c = 0.52, CH₃OH). Anal. (C₃₂H₅₄N₃O₈PSNa·1.94H₂O) C, H, N.

Preparation of (E,E)-N-[N-[N-[3-[Hydroxy(3,7,11-trimethyl-1-oxo-2,6,10-dodecatrienyl)phosphinyl]-1-oxopropyl]-L-valyl]-L-valyl]-L-methionine, Disodium Salt (16). Sodium hydroxide (1 N, 0.412 mL, 0.412 mmol) was added to a solution of 13 (0.131 g, 0.187 mmol,) in dioxane (2 mL), and the mixture was stirred at room temperature for 72 h. The reaction mixture was concentrated in vacuo, and the residue was purified by chromatography on CHP-20P gel (eluting sequentially with water (75 mL), 2.5%, 5.0, 7.5%, and 10% aqueous acetonitrile). Fractions containing the major product were combined and concentrated in vacuo, and the residue was dissolved in water (10 mL), millipore filtered, and lyophilized to afford 16 (0.08g, 60%) as a white powder, mp > 195 °C dec. TLC: $R_f = 0.78$ (6:3:1 1-propanol/ammonium hydroxide/water, visualization by PMA). IR (KBr): 1638 cm⁻¹. Mass spectrum $(M + Na)^+$: 694. ¹H NMR (CD₃OD): 0.94 (m, 12 H), 1.59 (s, 6 H), 1.66 (s, 6 H), 1.72-1.75 (m, 2 H), 1.95-2.11 (m, 12 H), 2.05 (s, 3 H), 2.30–2.49 (m, 6 H), 4.15 (d, 1 H, J = 7.6), 4.19 (d, 1 H, J = 7.7), 4.29 (t, 1 H, J = 7.2), 5.08–5.12 (m, 2 H), 5.35 (br m, 1 H). $[\alpha]_D = -27.1^\circ$ (c = 0.17, CH₃OH). Anal. (C₃₃H₅₆N₃O₇PSNa₂·2.71H₂O) C, H, N.

Preparation of (E,E)-N-[N-[N-[[[Hydroxy(3,7,11-trimethyl-2,6,10-dodecatrienyl)phosphinyl]oxy]acetyl]-Lvalyl]-L-valyl]-L-methionine, Methyl Ester (17). Sodium iodide (0.915 g, 6.1 mmol) was added to a solution of 12 (0.856 mmol)g, 1.22 mmol) in acetone (20 mL), and the mixture was heated at reflux for 20 h. The reaction was concentrated in vacuo, the residue was dissolved in aqueous sodium thiosulfate solution (5%, pH 2, 50 mL), and the solution was saturated with sodium chloride. The aqueous solution was extracted with chloroform (10 \times 100 mL), and the combined organic extracts were dried ($MgSO_4$), filtered, and concentrated in vacuo. The residue was purified by successive triturations with diethyl ether (1 \times 50 mL), hexane (1 \times 50 mL), and ethyl acetate $(1 \times 50 \text{ mL})$ to afford 17 (0.436 g, 52%) as a solid, mp 174-178 °C. Mass spectrum $(M + H)^+$: 688. TLC: $R_f = 0.79$ (6:3:1 1-propanol/ammonium hydoxide/water, visualization by PMA). ÎR (KBr): 1640 cm⁻¹. ¹H NMR (CD₃OD): 0.92-1.00 (m, 12 H), 1.60 (s, 6 H), 1.72 (s, 3 H), 1.75 (s, 3 H), 1.94-2.13 (m, 12 H), 2.06 (s, 3 H), 2.52 (m, 4 H), 3.73 (s, 3 H), 4.12-4.75(m, 5 H), 5.14-5.33 (m, 3 H). $[\alpha]_D = -49.2^\circ$ (c = 1.04, CHCl₃). Anal. $(C_{33}H_{58}N_3O_8PS H_2O)C, H, N.$

Preparation of (E,E)-N-[N-[N-[3-[Hydroxy(3,7,11-trimethyl-1-oxo-2,6,10-dodecatrienyl)phosphinyl]-1-oxopropyl]-L-valyl]-L-valyl]-L-methionine, Methyl Ester, Monosodium Salt (18). Trimethylamine was bubbled into a 0 °C solution of 16 (0.50 g, 0.73 mmol) in acetone (50 mL) until the solution was saturated, and then the mixture was stirred for 0.5 h. The reaction mixture was concentrated in vacuo, and the water soluble trimethylammonium salt was passed through a Dowex Na⁺ ion exchange column (eluting with 20% aqueous acetonitrile). The appropriate fractions were combined and concentrated in vacuo, and the residue was dissolved in water (1 mL), millipore filtered, and lyophilized to afford 18 (0.40 g, 78%) as a white powder, mp > 200 °C dec. TLC: $R_f = 0.\overline{78}$ (6:3:1 1-propanol/ammonium hydroxide/ water, visualization by PMA). IR (KBr): 1740 cm⁻¹. ¹H NMR $(CD_3OD): 0.91-0.98$ (m, 12 H), 1.59 (s, 6 H), 1.66 (s, 6 H), 1.71-1.76 (m, 2 H), 1.94-2.09 (m, 12), 2.06 (s, 3 H), 2.31-2.58 (m, 6 H), 3.70 (s, 3 H), 4.12 (d, 1 H, J = 8.1), 4.15 (d, 1 H, J = 8.1)J = 7.7), 4.56 (br m, 1 H), 5.09–5.12 (m, 2 H), 5.32 (br m, 1 H). $[\alpha]_D = -43.3^\circ$ (c = 0.145, CH₃OH). Anal. (C₃₄H₅₉N₃O₇-SPNa-2.49H₂O) C, H, N.

Preparation of (E,E)-N-[N-[N-[N-[Hydroxy(3,7,11-trimethyl-1-oxo-2,6,10-dodecatrienyl)phosphinyl]-1-oxopropyl]-L-valyl]-L-walyl]-L-methionine, (2,2-Dimethyl-1-oxopropoxy)methyl Ester (19). Sodium hydroxide (1 N, 14.3 mL, 14.3 mmol) was added to a -26 °C solution of 16 (10.0 g, 14.3 mmol) in methanol:water (115 mL:5 mL), after which the mixture was warmed to room temperature and stirred for 48 h. Another aliquot of sodium hydroxide (1 N, 3.6 mL, 3.6 mmol) was added, and the reaction mixture was stirred at room temperature for 6 h. The reaction was quenched carefully by addition of concentrated hydrochloric acid and was extracted with methylene chloride (6 \times 700 mL). The combined organic extracts were dried (MgSO₄), filtered,

and concentrated in vacuo to afford the methyl phosphinyl ester carboxylic acid intermediate (7.2 g, 73%, TLC R_f 0.84 (6:3:1 1-propanol:ammonium hydroxide:water). Iodomethyl pivalate (2.4 g, 10.0 mmol) was added to a mixture of the above intermediate (0.685 g, 1.0 mmol), silver carbonate (0.552 g, 2.0 mmol), and dimethyl methylphosphonate (1.1 mL, 10.0 mmol) in DMF (10 mL), and the mixture was stirred at room temperature for 2 h. The reaction was quenched by addition of 10% aqueous lithium chloride (50 mL) and extracted with methylene chloride (4 \times 50 mL). The combined organic extracts were washed sequentially with 10% aqueous lithium chloride $(3 \times 75 \text{ mL})$ and aqueous NaHSO₄ $(2 \times 150 \text{ mL})$, dried $(MgSO_4)$, filtered, and concentrated in vacuo. The residue was purified by flash chromatography (eluting with 1:1 to 3:1 acetone/hexane) to afford the methyl phosphinyl pivaloyl carboxyl diester intermediate (0.632 g, 79%, TLC $R_f = 0.76$ (3:1 acetone/hexane). Mass spectrum $(M + H)^+$: 800. IR (CH₂-Cl₂ film): 1757 cm⁻¹. ¹H NMR (CDCl₃): 0.86-0.94 (m, 12 H), 1.20 (s, 9 H), 1.59 (s, 6 H), 1.68 (s, 3 H), 1.68 (s, 3 H), 2.06-2.19 (m, 14 H), 2.06 (s, 3 H), 2.65 (m, 6 H), 3.68 (dd, 3 H, J =10.6, 11.1), 4.45 (m, 1 H), 4.61 (m, 2 H), 5.08-5.18 (m, 3 H), 5.71 (d, 1H, J = 5.3), 5.85 (d, 1 H, J = 5.6), ¹³C NMR (CDCl₃): 15.15, 15.92, 16.30, 17.59, 17.91, 18.00 18.43, 18.51, 19.03, 19.12, 22.58 (d, J = 90), 25.60, 26.37, 26.64, 26.75, 27.67, 29.20,29.80, 30.72, 31.67, 38.61, 39.62, 39.68, 51.28, 51.37, 53.79, 57.04, 58.83, 69.40, 79.76, 112.20, 123.54, 124.20, 131.20, 135.34, 140.70, 170.50, 171.22, 171.31, 171.51, 210.70. Anal. (C40H70N3O9PS) C, H, N.

Bis(trimethylsilvl)trifluoroacetamide (1.93 mL, 7.26 mmol) was added to a solution of the above mixed diester intermediate (0.29 g, 0.363 mmol) in methylene chloride (6 mL), and the resulting mixture was stirred for 0.5 h. Trimethylsilyl bromide (0.115 ml, 0.871 mmol) was then added, and the mixture was stirred at room temperature for 1 h. The reaction mixture was concentrated in vacuo, the residue dissolved in methanol, (5 mL), and the mixture concentrated again in vacuo. The residue was purified by successive triturations with diethyl ether (3 \times 20 mL) and petroleum ether (4 \times 20 mL), and the resulting solid was vacuum dried to afford 19 (0.17 g, 62%), mp 165–171 °C. TLC: $R_f = 0.73$ (6:3:1 1-propanol/ammonium hydroxide/water, visualization by PMA). Mass spectrum $(M + H)^{+:}$ 786. IR (KBr): 1638, 1757 cm⁻¹. $[\alpha]_{\rm D} = -47.9^{\circ} (c = 0.19, \text{ MeOH})$. ¹H NMR (CD₃OD): 0.83-1.13 (m, 12 H), 1.18-1.28 (s, 9 H), 1.59-1.77 (m, 12 H), 1.97-2.28 (m, 14 H), 2.06 (s, 3H), 2.51-2.75 (m, 6 H), 4.17-4.31 (m, 2 H), 4.63–4.75 (m, 1 H), 5.15–5.23 (m, 3 H), 5.72 (d, 1 H, J = 5.2), 5.83 (d, 1 H, J = 5.6). Anal. (C₃₉H₆₈N₃PS·0.22H₂O) C, H, N.

Acknowledgment. We wish to thank Drs. Jack Chabala and John Hunt at BMS for helpful discussions and Ms. Debra Skeens at Affymax for her help in the preparation of the manuscript.

References

- (1) Barbacid, M. Ras genes. Annu. Rev. Biochem. 1987, 56, 779-827.
- (2) (a) Casey, P. J.; Solski, P. A.; Der, C. J.; Buss, J. E. p21ras is modified by a farnesyl isoprenoid. *Proc. Natl. Acad. Sci. U.S.A.* **1989**, *86*, 8323-8327. (b) Der, C. J.; Cox, A. D. Isoprenoid Modification and Plasma Membrane Association: Critical Factors for Ras Oncogenicity. *Cancer Cells* **1991**, *3*, 331-340. (c) Marshall, C. J. Protein Prenylation: A Mediator of Protein-Protein Interactions. *Science* **1993**, *259*, 1865-1866.
- (3) (a) Manne, V.; Roberts, D.; Tobin, A.; O'Rourke, E.; De Virgilio, M.; Meyers, C.; Ahmed, N.; Kurz, B.; Resh, M.; Kung, H.-F.; Barbacid, M. Identification and preliminary characterization of protein-cysteine farnesyltransferase. Proc. Natl. Acad. Sci. U.S.A. 1990, 87, 7541-7545. (b) Schaber, M. D.; O'Hara, M. B.; Garsky, V. M.; Mosser, S. D.; Bergstrom, J. D.; Moores, S. L.; Marshall, M. S.; Friedman, P. A.; Dixon, R. A. F.; Gibbs, J. B. Polyisoprenylation of Ras in Vitro by a Farnesyl-Protein Transferase. J. Biol. Chem. 1990, 265, 14701-14704. (c) Reiss, Y.; Seabra, M. C.; Goldstein, J. L.; Brown, M. S. Purification of ras Farnesyl Protein Transferase. METHODS: A Companion to Methods in Enzymology 1990, 1, 241-245.
 (4) (a) Reiss, Y.; Goldstein, J. L.; Seabra, M. C.; Casey, P. J.; Brown,
- (4) (a) Reiss, Y.; Goldstein, J. L.; Seabra, M. C.; Casey, P. J.; Brown, M. S. Inhibition of Purified p21^{ras} Farnesyl Protein Transferase by Cys-AAX Tetrapeptides. *Cell* **1990**, *62*, 81–88. (b) Gibbs, J.

B. Ras C-Terminal Processing Enzymes- New Drug Targets? Cell 1991, 65, 1-4. (c) Reiss, Y.; Stradley, S. J.; Gierasch, L. M.; Brown, M. S.; Goldstein, J. L. Sequence requirement for peptide Brown, M. S., Goldstein, J. L. Sequence requirement for peptide recognition by rat brain p21^{ras} protein farnesyltransferase. *Proc. Natl. Acad. Sci. U.S.A.* 1991, 88, 732–736. (d) Goldstein, J. L.; Brown, M. S.; Stradley, S. J.; Reiss, Y.; Gierasch, L. M. Nonfarnesylated Tetrapeptide Inhibitors of Protein Farnesyl-

- Nonfarnesylated Tetrapeptide Inhibitors of Protein Farnesyl-transferase. J. Biol. Chem. 1991, 266, 15575-15578.
 (5) (a) Kohl, N. E.; Mosser, S. D.; deSolms, S. J.; Giuliani, E. A.; Pompliano, D. L.; Graham, S. L.; Smith, R. L.; Scolnick, E. M.; Oliff, A.; Gibbs, J. B., Selective Inhibition of ras- Dependent Transformation by a Farnesyltransferase Inhibitor. Science 1993, 260, 1934-1937. (b) James, G. L.; Goldstein, J. L.; Brown, M. S.; Rawson, T. E.; Somers, T. C.; McDowell, R. S.; Crowley, C. W.; Lucas, B. K.; Levinson, A. D.; Marsters, J. C. Jr. Benzodiazepine Peptidomimetics: Potent Inhibitors of Ras Far-nesylation in Animal Cells. Science 1993, 260, 1937-1942. (c) Graham, S. L.; deSolms, S. J.; Guiliani, E. A.; Kihl, N. E.; Mosser, S. D.; Oliff, A. I.; Pompliano, D. L.; Rands, E.; Breslin, M. J.; Deana, A. A.; Garsky, V. M.; Scholz, T. H.; Gibbs, J. B.; Smith, R. L. Pseudopeptide Inhibitors of Ras Farnesyl-Protein Transferase. J. Med. Chem. 1994, 37, 725-732.
 (6) (a) Pompliano, D. L.; Rands, E.; Schaber, M. D.; Mosser, S. D.;
- (a) Pompliano, D. L.; Rands, E.; Schaber, M. D.; Mosser, S. D.; Anthony, N. J.; Gibbs, J. B. Steady-State Kinetic Mechanism of (6)Ras Farnesyl:protein Transferase. Biochemistry 1992, 31, 3800-3807. (b) Singh, S. B.; Zink, D. L.; Liesch, J. M.; Goetz, M. A., Jenkins, R. G.; Nallin-Olmstead, M.; Silverman, K. C.; Bills, G. F.; Mosley, R. T.; Gibbs, J. B.; Albers-Schonberg, G.; Lingham, R. B. Isolation and Structure of Chaetomellic Acids A and B from Chaetomella acutiseta: Farnesyl Pyrophosphate Mimic Inhibitors of Ras Farnesyl-Protein Transferase. Tetrahedron 1993, 49, 5917-5926.
- (a) Reiss, Y.; Seabra, M. C.; Armstrong, S. A.; Slaughter, C. A.; Goldstein, J. L.; Brown, M. S. Nonidentical Subunits of p21^{H-ras} Farnesyltransferase. J. Biol. Chem. 1991, 266, 10672-10677. (b) Reiss, Y.; Brown, M. S.; Goldstein, J. L. Divalent Cation and Prenyl Pyrophosphate Specificities of the Protein Farnesyltransferase from Rat Brain, a Zinc Metalloenzyme. J. Biol. Chem. 1992, 267, 6403-6408.
- Literature (refs 4a,c) as well as in house data (Devergilio, M.; Robinson, S.; Meyers, C. M., unpublished work) suggested that a CAAX tetrapeptide sequence was sufficient for good binding, and that an amino group at the N-terminus was not required for activity.
- Petrillo, E. W., Jr.; Ondetti, M. A. Angiotensin-Converting Enzyme Inhibitors: Medicinal Chemistry and Biological Actions.
- Med. Res. Rev. 1982, 2, 1-41. (10) Gavras, H.; Gavras, I. Angiotensin Converting Enzyme Inhibitors. Properties and Side Effects. Hypertension 1988, 11 (3), Suppl. 11, 37-41.
- (11) Abbreviations: ethyl (3-(dimethylamino)propyl)carbodiimide hy-drochloride (EDC), (benzotrialzol-1-yloxy)tris(dimethylamino)phosphonium hexafluorophosphate (BOP).
- (12) Attempts at direct, one-step base hydrolysis of diester 11 resulted in a mixture of products, arising from competing cleavage of both the methyl group and the linker moiety attached on the phosphorous. The phosphonic monoacid 17 arising from mon-odealkylation of 11 is resistant to further base-catalyzed hydrolysis, thereby solving the problem of the cleavage of the linker group during the base hydrolysis of the carboxylic methyl ester group of 17 in the second step.
- (13) Phosphinate diester 6 was prepared by slight modifications of the literature procedure. Gallagher, M. J.; Sussman, J. Orga-nophosphorous Intermediates V. Some Michael Additions of

Methyl Hypophosphite. A Simple Route to Functionalized Phos-(14) (a) Epstein, W. W.; Lever, D. C.; Rilling, H. C. Prenylated

- proteins: Synthesis of geranylgeranylcysteine and identification of this thioether amino acid as a component of proteins in CHO cells. *Proc. Natl. Acad. Sci. U.S.A.* **1990**, *87*, 7352–7354. (b) Moores, S. L.; Schaber, M. D.; Mosser, S. D.; Rands, E.; O'Hara, M. B.; Garsky, V. M.; Marshall, M. S.; Pompliano, D. L.; Gibbs, J. B. Sequence Dependence of Protein Isoprenylation J. Biol. Chem. 1991, 266, 14603-14610. (c) Seabra, M. C.; Reiss, Y.; Casey, P. J.; Brown, M. S.; Goldstein, J. L., Protein Farnesyltransferase and Geranyltransferase Share a Common a Subunit. Cell 1991, 65, 429-434. (d) Casey, P. J.; Thissen, J. A.; Moomaw, J. F. Enzymatic modification of proteins with a geranylgeranyl isoprenoid. Proc. Natl. Acad. Sci. U.S.A. 1991, 88, 8631–8635. (e) Yokoyoma, K.; Goodwin, G. W.; Ghomashchi, F.; Glomset, J. A.; Gelb, M. H. A protein geranylgeranyltransferase from bovine brain: Implications for protein prenylation specificity. Proc. Natl. Acad. Sci. U.S.A. 1991, 88, 5302-5306. (f) Kinsella, B. T.; Erdman, R. A.; Maltese, W. A. Posttranslational modification of Ha-ras p21 by farnesyl versus geranylgeranyl isoprenoids is determined by the COOH-terminal amino acid. *Proc. Natl. Acad.* Sci. U.S.A. 1991, 88, 8934-8938. (g) Moomaw, J. F.; Casey, P. J. Mammalian Protein Geranylgeranyltransferase: subunit composition and metal requirements. J. Biol. Chem. 1992, 267, 17438-17443.
- Mane, V.; Yan N.; Carboni, J. M.; Tuomari, A. V.; Ricca, C. S.;
 Gullo-Brown, J.; Andahazy, M. L.; Schmidt, R. J.; Patel, D. V.;
 Gordon, E. M.; Zahler, R.; Weinmann, R.; Der, C. J.; Cox, A. D.;
 Hunt, J. T.; Barbacid, M.; Seizinger, B. R. Bisubstrate Inhibitors of Farensyltransferase: A novel class of specific inhibitors of ras transformed cells. Submitted for publication to Oncogene.
- (16) Saini, S. S.; Manne, V., manuscript in preparation.
 (17) Farnsworth, C. L.; Marshall, M. S.; Gibbs, J. B.; Stacey, D. W.; Feig, L. A. Preferential inhibition of the oncogenic form of Ras by mutations in the GAP binding/"effector" domain. Cell 1991, 64, 625–63. (18) Manne, V.; Ricca, C. S.; Brown, J. G.; Tuomari, A. V.; Yan, N.;
- Matel, D.; Schmidt, R.; Lynch, M. J.; Ciosek, C. P.; Carboni, J. M.; Robinson, S.; Gordon, E. M.; Barbacid, M.; Seizinger, B. R.; Biller. S. A. Ras farnesylation as a target for novel antitumor agents: Potent and selective farnesyl diphosphate analog inhibitors of farnesyltransferase. Drug. Dev. Res. In press. (19) Sukumar, S.; Carney, W. P.; Barbacid, M. Independent molecular
- pathways in initiation and loss of hormone responsiveness of breast carcinomas. Science 1988, 240, 524-526.
- (a) A preliminary account of this work was presented at the following: Patel, D. V.; Weller, H. N.; Schmidt, R. J.; Gordon, E. M.; Miller, A. V.; Young, M. G.; Zahler, R.; Barbacid, M.; Gullo-Brown, J. L.; Ricca, C.; Robinson, S.; Tuomari, V. A.; Yan, N.; Manne, V. Design, Synthesis, and biological activity of why burther of the second seco (20)substrate based ras protein farnesyl transferase inhibitors. 207th American Chemical Society National Meeting, San Diego, CA, March 13-17, 1994, paper MEDI 279. (b) After the submission of this manuscript, the following article was published by our colleagues from Bristol-Myers Squibb: Bhide, R. S.; Patel, D. V.; Patel, M. M.; Robinson, S. P.; Hunihan, L. W.; Gordon, E. M. Rational design of potent carboxylic acid based bisubstrate inhibitors of ras farnesyl protein transferase. Bioorg. Med. Chem. Lett. 1994, 4, 2107-2112.
- (21) Preliminary experiments show that 18 and 19 have no effect on NIH3T3 cells transformed with the v-Raf oncogene which acts independently of Ras (unpublished observations).

JM940623L