

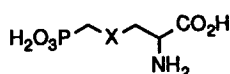


SYNTHESES OF HOMOSERINE PHOSPHATE ANALOGS AS POTENTIAL INHIBITORS OF BACTERIAL THREONINE SYNTHASE

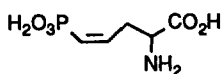
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Abstract: The syntheses of five homoserine phosphate analogs (1-4, Z-5) are described. A brief summary of their potential to inhibit threonine synthase from *Escherichia coli* is given.

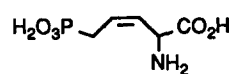
Bacterial threonine synthase (TS) [EC 4.2.99.2], a pyridoxal phosphate (PLP) dependent enzyme, catalyzes the final step in the biosynthesis of L-threonine from L-aspartic acid, i.e. the conversion of L-homoserine phosphate into L-threonine. The reaction involves the non-hydrolytic elimination of phosphate from the PLP bound substrate and subsequent addition of water to a PLP bound vinylglycine intermediate¹⁾. We were interested in homoserine phosphate analogs both as potential inhibitors of TS and also in order to study intermediates of this enzymic reaction by UV spectroscopy.



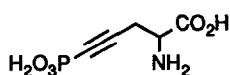
- 1: X = S
 2: X = NH
 3: X = O



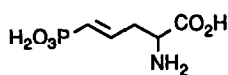
Z-5



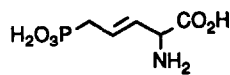
Z-6



4



E-5



E-6

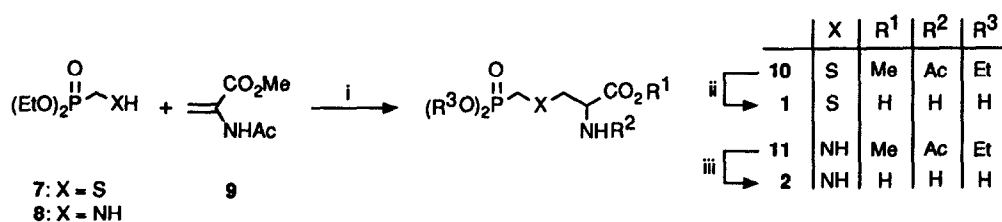
The sulfur containing phosphonic acid 1 has been reported to be a slow binding inhibitor of TS²⁾, and the nitrogen analog L-2 to possess N-methyl-D-aspartic acid (NMDA) antagonistic activity³⁾. Since no experimental details for the synthesis of these two compounds have been published yet, we developed syntheses for racemic and enantiomerically pure phosphonic acid 1, as well as for the racemic analogs 2 and 3.

The pentynoic acid 4 not only resembles homoserine phosphate, it is also a simple derivative of propargylglycine, which is a well known inhibitor of various PLP dependent enzymes⁴⁾.

Amino acid Z-6 represents the C-terminus of two classes of naturally occurring di- and tripeptides, the rhizocticins and the plumbemycins. Z-6^{5,6)} has the L-configuration in the fungicidal rhizocticins⁷⁾, whereas the D-enantiomer is in the plumbemycins^{8,9)}, which are active against bacteria. Moreover, Natchev has also reported that the L-enantiomer of Z-6 displays herbicidal and fungicidal activity⁶⁾. Inhibition studies with

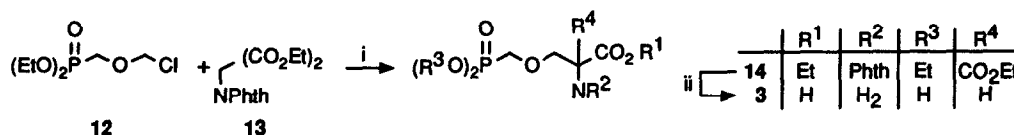
racemic Z-6, performed in our laboratory¹⁰, clearly showed that this compound is an inhibitor of bacterial TS (Table 1). The amino acids E-5¹¹ and E-6¹² have already been described in the literature as NMDA antagonists. Therefore, we undertook the synthesis of the unsaturated amino acid Z-5 to complete the series of pentenoic acids 5 and 6. In this letter we describe the facile syntheses of phosphonic acids 1-4, and Z-5, and give a brief summary of their action on bacterial TS.

The syntheses of compounds 1 and 2 are described in scheme 1. Michael addition of phosphonates 7¹³ and 8¹⁴ to methylacetamido-acrylate 9 in the presence of triethylamine in refluxing methanol¹⁵ yielded the protected amino acids 10 and 11, respectively. Deprotection of the sulfur compound 10 in refluxing 6N hydrochloric acid¹⁶ and subsequent purification by ion exchange chromatography¹⁶ afforded racemic phosphonic acid 1¹⁷ as an amorphous solid.



Scheme 1. *Reagents and conditions*: i. Et₃N, MeOH, reflux, for 10: 8h :62%; for 11: 2d :73%; ii. 1) 6N HCl, reflux, 8h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O :56%; iii. 1) 3.5N NaOH, reflux, 16h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O → 2N pyridine in H₂O; 3) TMSBr, pyridine, CH₃CN, 20°C, 2d; 4) Dowex 50X-200, H⁺-form, eluent: H₂O → 2N pyridine in H₂O :85%.

The attempted deprotection of the nitrogen analog 11 under these conditions resulted mainly in a fragmentation of the molecule in a retro Michael sense, the amino acid 2 being obtained in only 29% yield. Alternatively, in a two step sequence with refluxing 3N sodium hydroxide followed by trimethylbromosilane/pyridine in acetonitrile¹⁶ all protecting groups were removed. After purification by ion exchange chromatography the amino acid 2¹⁷ was obtained as a white powder in 85% yield. Since the synthesis of the oxygen analog 3 via the Michael additon route described above failed, the reaction sequence shown in scheme 2 was pursued.



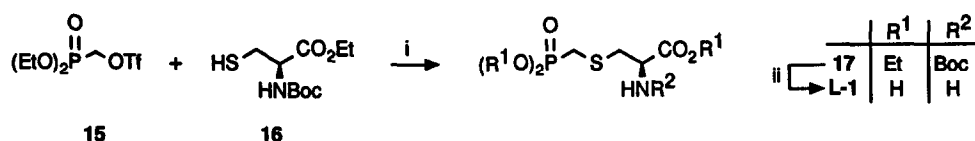
Scheme 2. *Reagents and conditions*: i. 13, NaH, DMF; 2) 12, 80-120°C, 5h :52%; ii. 1) 6N HCl/HOAc (5/1, v/v), reflux, 8h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O :56%.

Reaction of α-chloroether 12¹⁸ with the sodium salt of amino-malonate 13¹⁹ in DMF²⁰ afforded 14. Deprotection and decarboxylation could be achieved by refluxing compound 14 in 6N hydrochloric acid/acetic

acid²¹), and after purification by ion exchange chromatography the target compound **3**¹⁷ was obtained as a white, amorphous solid.

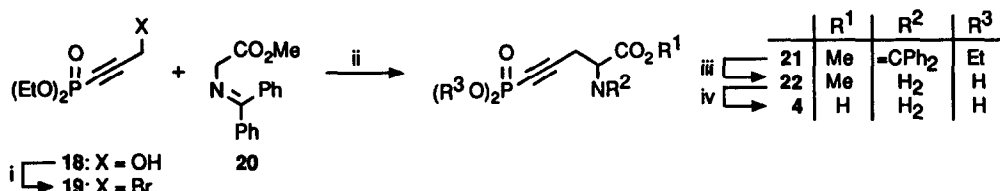
A similar reaction sequence could be used for the preparation of the L-enantiomer of the sulfur containing amino acid **1** (scheme 3). Alkylation of the protected L-cysteine derivative **16**²² with triflate **15**²³ in THF in the presence of triethylamine²³ afforded cleanly the protected phosphonic ester **17**. Deprotection and purification was carried out in the same manner as described for the racemic derivative **10**.

Using this reaction sequence, it should be possible to prepare the phosphonic acids **2** and **3** in enantiomerically pure form as well.



Scheme 3. Reagents and conditions: i. Et₃N, THF, 5°C, 2h → 20°C, 2d :75%; ii. 1) 6N HCl, reflux, 8h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O :70%.

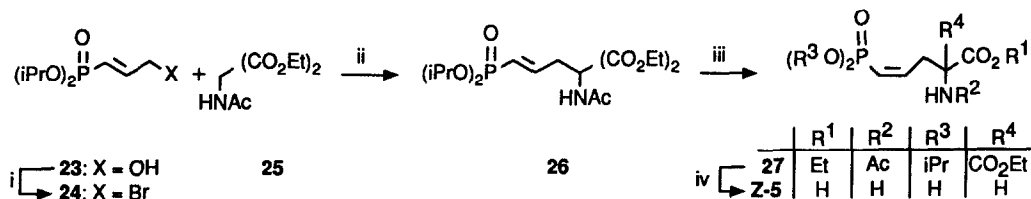
The synthesis of pentynoic acid **4** is outlined in scheme 4. Bromination of propargylalcohol **18**²⁴ with triphenylphosphine/ tetrabromomethane in dichloromethane²⁵ yielded propargylic bromide **19**. Its reaction with the anion²⁶ of glycine derivative **20**²⁷, and subsequent cleavage of the phosphonic esters and the imine with trimethylbromosilane/pyridine afforded cleanly methyl ester **22**. After saponification of the ester with 2N LiOH and purification by ion exchange chromatography, amino acid **4**¹⁷ was obtained as a white amorphous solid.



Scheme 4. Reagents and conditions: i. PPh₃, CBr₄, CH₂Cl₂, 5°C, 18h :50%; ii. 1) **20**, LiHMDS, THF, -78°C; 2) **19**, THF, -78°C → 0°C, 4h; iii. Me₃SiBr, pyridine, CH₂Cl₂, 20°C, 1d :67% (based on **19**); iv. 1) 2N LiOH, 20°C, 18h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O :81%.

The synthesis of amino acid **Z-5** (scheme 5) was accomplished in a similar manner to the synthesis of pentynoic acid **4**. After bromination of the allylic alcohol **23**²⁸, the resulting bromide **24** was alkylated with the anion of aminomalonnate **25**¹⁹ to yield E-olefin **26**. Isomerization of the double bond by irradiation with a Hg medium pressure lamp (Philips HPK 125W) for 2d in a quartz vessel gave a 3:1 mixture of isomers **26** and

27, that could be separated by chromatography on SiO₂. Complete deprotection and decarboxylation of Z-isomer 27 was achieved in refluxing 6N hydrochloric acid. Purification by ion exchange chromatography afforded the desired amino acid Z-5¹⁷⁾ as an amorphous solid.



Scheme 5. Reagents and conditions: i. PPh₃, CBr₄, CH₂Cl₂, 5°C, 18h :47%; ii. 1) 25, NaH, DMF; 2) 24, DMF, 80-100°C, 8h :27%; iii. hv, C₆H₆, 2d, 27:16% and 26: 43%; iv. 1) 6N HCl, reflux, 8h; 2) Dowex 50X-200, H⁺-form, eluent: H₂O :52%.

The amino acids described above, as well as the already known unsaturated amino acids E-5, Z-6, and E-6 were tested for their potential to inhibit TS from *Escherichia coli* [EC 4.2.99.2] (table 1). Compounds 3, 4, and Z-5 did not inhibit the enzyme at all, although the unsaturated amino acids 4 and Z-5 did bind to the enzyme, as was shown by UV-spectroscopy¹⁰⁾. The sulfur compound 1 and olefin E-5 were competitive inhibitors. Amino acids 2, Z-6, and E-6 were found to be irreversible inhibitors of the bacterial enzyme. Detailed experimental, kinetic, and spectroscopic data, as well as a discussion of these results will be published later¹⁰⁾.

Table 1: Kinetic constants^{a)} for inhibition^{b)} of threonine synthase from *Escherichia coli*

compound	K _i / [M]	k _{inact} / [min ⁻¹]
1	3.3 · 10 ⁻⁵	d)
L-1	1.1 · 10 ⁻⁵	d)
2	5.9 · 10 ⁻⁵	1.44
3	c)	
4	c)	
Z-5	c)	
E-5	5.4 · 10 ⁻⁴	d)
Z-6	1.0 · 10 ⁻⁴	1.50
E-6	4.0 · 10 ⁻⁴	0.25

a) mean of two determinations; b) for experimental conditions see ref. 10, 29);
c) no inhibition at 10mM inhibitor concentration; d) competitive inhibitor

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17. ¹H NMR spectra were recorded at 300 MHz and ¹³C NMR spectra at 75 MHz, respectively. Dioxan was used as an internal reference for spectra taken from aqueous solutions. The reported compounds possess the following physical data:

- 1: ^1H NMR (D_2O) δ 2.76 (m, 2H), 3.14 (dd, $J = 15, 8$ Hz, 1H), 3.32 (dd, $J = 15, 5$ Hz, 1H), 4.21 (dd, $J = 8, 5$ Hz, 1H); ^{13}C NMR (D_2O) δ 28.06 (d, $J = 138.9$ Hz), 34.15 (d, $J = 5.3$ Hz), 53.26, 171.71; IR (KBr) ν 1730, 1620, 1510, 1120, 1040, 915 cm^{-1} .
- L-1: $[\alpha]_{\text{D}}^{20} -16.0^\circ$ ($c = 0.52, \text{H}_2\text{O}$); spectroscopic data identical with racemic 1.
- 2: mp 240°C (dec.); ^1H NMR ($\text{D}_2\text{O} + \text{NaOD}$) δ 2.55 (dd, $J = 14, 14$ Hz, 1H), 2.64 (dd, $J = 14, 14$ Hz, 1H), 2.73 (dd, $J = 12, 8.5$ Hz, 1H), 2.89 (dd, $J = 12, 5$ Hz, 1H), 3.41 (dd, $J = 8.5, 5$ Hz, 1H); ^{13}C NMR ($\text{D}_2\text{O} + \text{NaOD}$) δ 47.61 (d, $J = 137.4$ Hz), 54.90 (d, $J = 12.8$ Hz), 55.18, 181.59; IR (KBr) ν 1642, 1180, 1118 cm^{-1} .
- 3: ^1H NMR (D_2O) δ 3.74 (m, 2H), 4.02 (dd, $J = 11, 3.5$ Hz, 1H), 4.08 (dd, $J = 11, 5$ Hz, 1H), 4.27 (dd, $J = 5, 3.5$ Hz, 1H); ^{13}C NMR (D_2O) δ 53.28, 67.06 (d, $J = 157.8$ Hz), 69.76 (d, $J = 13.6$ Hz), 170.14; IR (KBr) ν 1745, 1600, 1525, 1270, 1130, 1050, 940 cm^{-1} .
- 4: ^1H NMR (D_2O) δ 3.04 (d, $J = 6$ Hz, 1H), 3.06 (d, $J = 6$ Hz, 1H), 4.23 (dd, $J = 6, 6$ Hz, 1H); ^{13}C NMR (D_2O) δ 20.34 (d, $J = 3.8$ Hz), 50.97, 80.19 (d, $J = 157.8$ Hz), 88.89 (d, $J = 47.6$ Hz), 169.97; IR (KBr) ν 2205, 1730, 1610, 1515, 1150, 1040, 930 cm^{-1} .
- Z-5: ^1H NMR (D_2O) δ 3.00 - 3.22 (m, 2H), 4.09 (dd, $J = 7, 5.5$ Hz, 1H), 6.04 (d, $J = 17.5, 7.5$ Hz, 1H), 6.22 (dddd, $J = 45.5, 16, 7.5, 7$ Hz, 1H); ^{13}C NMR (D_2O) δ 30.43 (d, $J = 9.1$ Hz), 51.96, 127.69 (d, $J = 171.4$ Hz), 138.52, 171.60; IR (KBr) ν 1730, 1630, 1520, 1130, 1045, 940 cm^{-1} .
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