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Absolute Configuration of C(1)-C(5) Fragment of AAL-toxin: Conformationally Rigid Acyclic Aminotriol Moiety.

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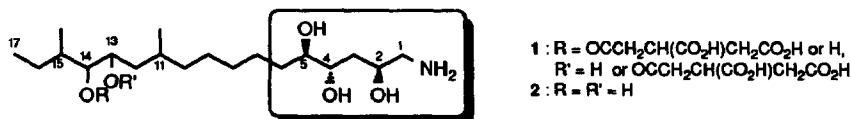
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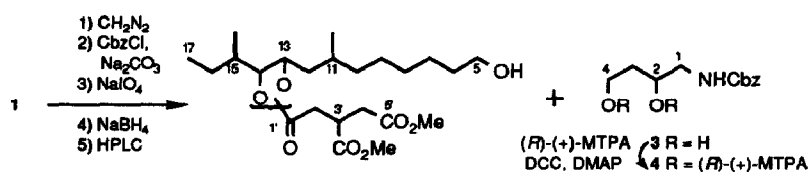
Key Words: AAL-toxin; host-specific toxin; *Alternaria alternata*; absolute configuration

Abstract: Degradation of AAL-toxin **1** a host-specific phytotoxin and synthesis of model aminotriol **7a** - **7d** allowed us to determine the absolute configuration of C(1)-C(5) fragment as *2S*, *4S* and *5R*. Unusually rigid conformation of this acyclic fragment was also discussed.

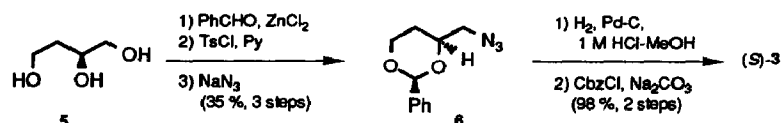
Host-specific toxins (HST) in plant diseases are interesting topics for studying host-parasite interaction.¹ AAL-toxin **1**,^{2a,3} one of HST produced by *Alternaria alternata* f. sp. *lycopersici*, a causal fungus of tomato stem canker disease reproduces similar symptoms to that of the disease for susceptible genotype of tomato leaf in concentrations less than 10 ng/ml. From genetic analysis, Gilchrist *et al.* proposed that a single gene controls sensitivity to the toxin and susceptibility to the fungus.^{4a} The same group also suggested that reduction of toxin activity by L-aspartate and orotate can be explained by inhibition of aspartate carbamoyl transferase,^{4b} and that induction of necrosis in tomato caused by **1** requires *de novo* synthesis and external binding of ethylene.^{4b} Although Bottini *et al.* determined the 2D-structure of **1** by the extensive analysis of MS, ¹H- and ¹³C-NMR spectra,^{2b,2c} relative and absolute stereochemistry of **1** remained to solve. For understanding the mechanism of the host-specificity of **1** in molecular level, the elucidation of 3D-structure and synthetic studies are essential. Here, we describe the absolute configuration and the local conformation of C(1)-C(5) fragment of **1**.



For degradative study, AAL-toxin **1** was isolated from the cultures of *Alternaria alternata* tomato pathotype (*A. alternata* f. sp. *lycopersici*) O-227.³ Since large amount of **1** was not available in our hand, we at first concentrated our attention to the absolute stereochemistry at C-2. Degradation of **1** was carried out as shown in Scheme 1. After methylation, the amino group was protected by carbobenzyloxy (Cbz) group which also served as a chromophore in HPLC separation. Oxidative cleavage between C-4 and C-5, following reduction and HPLC separation gave diol **3**, which was further converted to bis-(*R*)-MTPA ester **4**. Starting from triol (*2S*)-**5**⁵ derived from L-malic acid, authentic (*2S*)-**3** was prepared in five steps as shown in Scheme 2. Independently prepared (*2RS*)-**3**⁶ and the (*2S*)-**3** were further converted to bis-(*R*)-MTPA esters (*2RS*)-**4** and (*2S*)-**4**, respectively. In their ¹H-NMR spectra, the signals of acyloxy methylene and methine protons were well resolved and were observed at 4.26 (CH) and 3.82 ppm (CH₂) in *2R*-isomer, and at 4.13 (CH) and 3.99 ppm (CH₂) in *2S*-isomer. Since the ¹H-NMR spectrum of the bis-(*R*)-MTPA ester **4** obtained from **1** was identical with that of synthetic (*2S*)-**4**, the absolute stereochemistry at C-2 of **1** was determined as *S*.

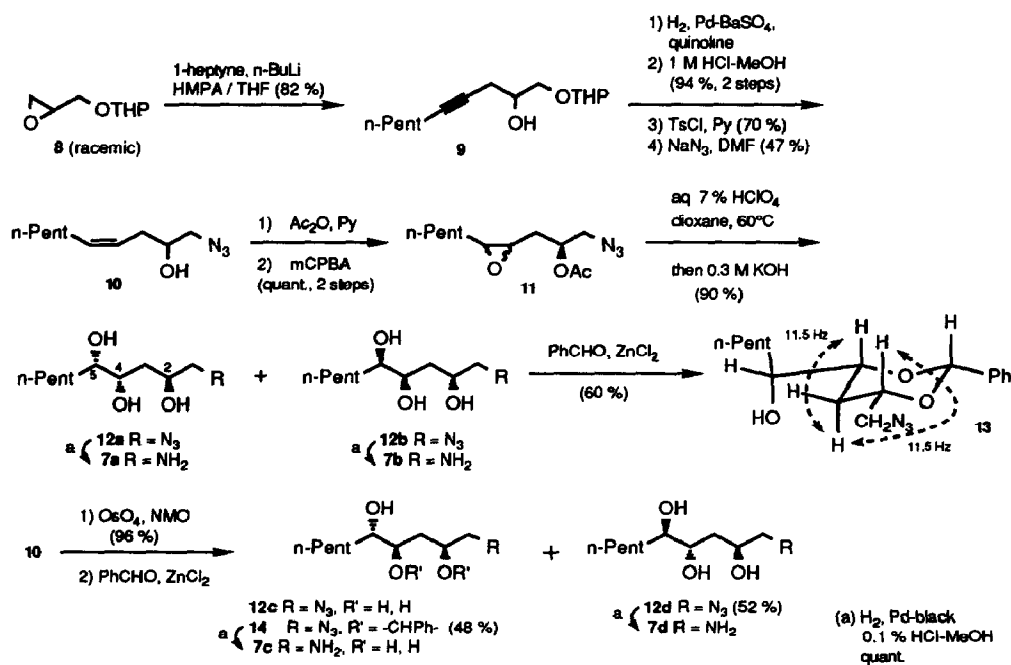


Scheme 1



Scheme 2

Structural difference of two components of AAL-toxin **1**, whose acyl group was substituted at either C-13 or C-14, did not affect the NMR resonances of right half of **1**.^{2c} Therefore, we expected that aminotriols **7a** - **7d** (Scheme 3) are suitable model compounds for determining relative stereochemistries at C-2, C-4 and C-5 of **1**. The synthetic pathway of four possible isomers **7a** - **7d** were shown in Scheme 3. The required carbon skeleton of the model triols were constructed by condensation of epoxide **8** and 1-heptyne in high yield. The coupling product **9** was hydrogenated to corresponding *cis*-olefin which was further converted to azide **10** by deprotection, tosylation of primary alcohol and azidation. From this homoallyl alcohol **10**, 4,5-*syn*- and 4,5-*anti*-diols were prepared by the following transformations. After acetylation, treatment of *O*-acetyl-**10** with *m*CPBA gave diastereomeric epoxides **11**.⁷ These were submitted to sequential hydrolysis with perchloric acid and aq. KOH to afford a 2:1 mixture of two 4, 5-*syn* triols **12a** and **12b**,⁷ which were easily separated by SiO_2 chromatography. The major isomer **12b** was then converted to benzylidene acetal **13**, whose $^1\text{H-NMR}$ spectrum clearly shows 2, 4-*syn* relationship (Scheme 3). Whereas catalytic dihydroxylation with OsO_4 of azide **10** gave hardly separable 4, 5-*anti*-triols **12c** and **12d**. When this mixture was treated with benzaldehyde and ZnCl_2 ,



Scheme 3

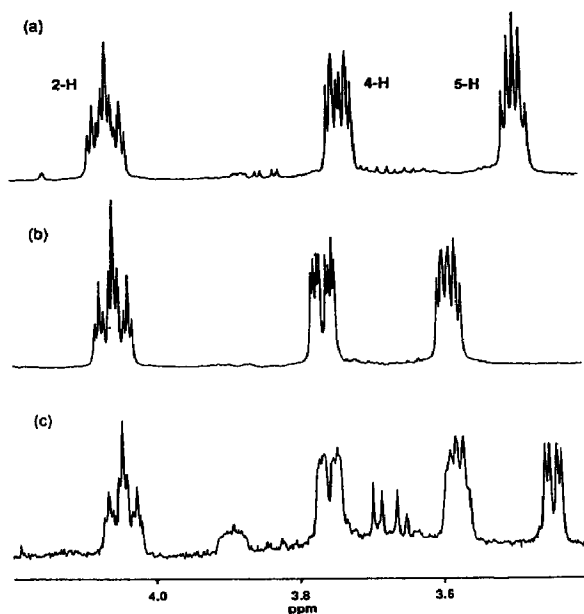


Fig. 1 $^1\text{H-NMR}$ spectra (500 MHz, D_2O) of: aminotriols (a) **12a**; (b) **12d**; (c) AAL-toxin **1**.

2, 4-*syn* isomer **12c** predominantly yielded acetal **14** while unreacted 2, 4-*anti*-isomer **12d** was recovered. The compounds **12a**, **12b**, **12d** and **14** were hydrogenated under acidic condition to afford the corresponding aminotriols **7a** - **7d**⁸ as hydrochloride salts. The stereospecific transformations from **10** to the **7a** - **7d** and the NMR analysis of benzylidene acetals **13** and **14** enabled us to assign the relative stereocenters of aminotriols **7a** - **7d**.

Among the synthesized aminotriols, 2, 4-*syn* isomers **7b** and **7c** were easily excluded by the comparison of their $^1\text{H-NMR}$ spectral data⁸ to that of aminopentol **2**.^{2b,8} Although NMR spectra of the remaining 2, 4-*anti*-aminotriols^{8,9} were very similar, inspection of the spectra (Fig.1) allowed us to determine that the relative stereochemistry of the right part of **1** is the same as that of **7d** (2, 4-*anti*-4, 5-*anti*). In addition, $^{13}\text{C-NMR}$ data supported this conclusion⁹; the resonances at 74.81 (C-5) and 35.88 ppm (C-3) in **7d** were nearly identical to that of **2**^{2b,8} while those signals in **7a** were observed at 75.54 and 37.31 ppm. On the basis of *J*-values in the $^1\text{H-NMR}$ spectrum of **1**, Bottini *et al.* proposed the relative stereochemistries at C-2, C-4 and C-5 as all-*S* or all-*R* (2, 4-*anti*-4, 5-*syn*).^{2b} Our data clearly indicate that 4, 5-*syn* **7a** and 4, 5-*anti* **7d** cannot be distinguished without comparing spectral data of the synthetic samples. Furthermore, the results of spectral analysis of the model aminotriol **7d** enabled us to conclude rigorously that the conformation of left part of **1** does not affect that of the right part in aqueous solution.

With the absolute configuration of fragment C(1)-C(5) of **1** established, the conformation of this part was examined next. From the calculated dihedral angles using Karplus equation and NOE studies for **7d**, the conformation of the right fragment of **1** was proposed. MM2 calculation of this conformation was undertaken and the energy-minimized conformation was shown in Fig. 2. This 3D-structure indicated the hydrogen bonds to be between $\text{NH} \rightarrow 2\text{-O}$ and $4\text{-OH} \rightarrow 5\text{-O}$ even in aqueous solution. Thus, in order to avoid steric hindrance between two pseudo-cyclic systems, the right part of **1** predominates the conformation as shown in Fig. 2. Previous conformational studies of acyclic polyols¹⁰ in D_2O showed that they exist as several energetically similar conformers. The conformation of aminotriol part of **1** is therefore a relatively unusual case in an acyclic system. The role of this conformer to biological activity is interesting.

In conclusion, we have determined the absolute stereochemistry of C(1)-C(5) fragment of **1**. The synthetic route described above can be applied to prepare a variety of analogues of **1**. Currently, we are undertaking to determine the remaining absolute structure of **1**.

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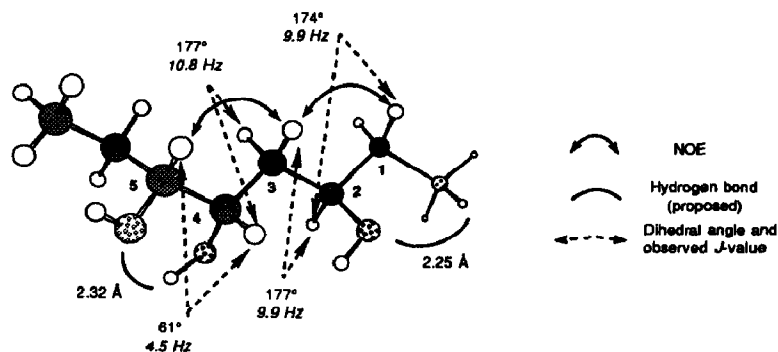


Fig. 2 The proposed conformation of fragment C(1)-C(5) of AAL-toxin 1.

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- Identification based on chromatographic behavior: Kohmoto, K.; Verma, V. S.; Nishimura, S.; Takagi, M.; Scheffer, R. P. *J. Fac. Agric., Tottori Univ.* 1982, 17, 1-8; Identification using spectral data will be reported elsewhere.
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- The racemic **3** was prepared from 4-amino-3-hydroxybutanoic acid in three steps (i, CH_2N_2 , ii, $\text{BH}_3\text{-Me}_2\text{S}$, iii, CbzCl , Na_2CO_3 , 23 % overall).
- For non protected epoxide *O*-deacetyl-**11**, the epoxide opening was attempted under acidic and basic conditions. In both cases, unidentified by-products were predominated. Since the epoxidation of *O*-acetyl-**10** gave a 1:1 diastereomer mixture, predominant formation of 2, 4-*syn*-diol **12b** could be explained by acyl group assisted epoxide opening in thermodynamically controlled manner.
- $^1\text{H-NMR}$ data (500 MHz, D_2O): **7a**; δ 4.06 (1H, dddd, $J = 3.3, 3.3, 9.6, 9.6$ Hz, 2-H), 3.74 (1H, ddd, $J = 3.3, 4.1, 9.6$ Hz, 4-H), 3.49 (1H, ddd, $J = 4.1, 4.1, 8.2$ Hz, 5-H), 3.15 (1H, dd, $J = 3.3, 13.1$ Hz, 1-Ha), 2.93 (1H, dd, $J = 9.6, 13.1$ Hz, 1-Hb), 1.64 (1H, ddd, $J = 3.3, 9.6, 14.5$ Hz, 3-Ha), 1.59 (1H, ddd, $J = 3.3, 9.6, 14.5$ Hz, 3-Hb); **7b**; δ 4.07 (1H, dddd, $J = 3.2, 6.4, 6.4, 9.6$ Hz, 2-H), 3.70 (1H, ddd, $J = 4.0, 6.4, 6.4$ Hz, 4-H), 3.53 (1H, ddd, $J = 4.0, 4.0, 8.5$ Hz, 5-H), 3.19 (1H, dd, $J = 3.2, 13.1$ Hz, 1-Ha), 2.96 (1H, dd, $J = 9.6, 13.1$ Hz, 1-Hb), 1.77 (2H, t, $J = 6.4$ Hz, 3-H); **7c**; δ 4.09 (1H, dddd, $J = 3.0, 6.4, 6.4, 9.6$ Hz, 2-H), 3.69 (1H, ddd, $J = 3.3, 4.6, 9.5$ Hz, 4-H), 3.56 (1H, ddd, $J = 3.4, 4.6, 9.1$ Hz, 5-H), 3.19 (1H, dd, $J = 3.0, 13.1$ Hz, 1-Ha), 2.96 (1H, dd, $J = 9.6, 13.1$ Hz, 1-Hb), 1.81 (1H, ddd, $J = 3.3, 5.4, 14.6$ Hz, 3-Ha), 1.70 (1H, ddd, $J = 6.4, 9.5, 14.6$ Hz, 3-Hb); **7d**; δ 4.06 (1H, dddd, $J = 2.9, 2.9, 9.9, 9.9$ Hz, 2-H), 3.76 (1H, ddd, $J = 2.0, 4.3, 10.8$ Hz, 4-H), 3.59 (1H, ddd, $J = 3.7, 4.3, 7.6$ Hz, 5-H), 3.16 (1H, dd, $J = 2.9, 13.1$ Hz, 1-Ha), 2.93 (1H, dd, $J = 9.9, 13.1$ Hz, 1-Hb), 1.67 (1H, ddd, $J = 2.0, 9.9, 14.5$ Hz, 3-Ha), 1.54 (1H, ddd, $J = 2.9, 10.8, 14.5$ Hz, 3-Hb); **2^{2b}**; δ 4.063 (1H, dddd, $J = 3.0, 3.0, 9.9, 9.9$ Hz, 2-H), 3.739 (1H, ddd, $J = 2.0, 4.5, 10.8$ Hz, 4-H), 3.556 (1H, ddd, $J = 4.5, 6-8, 6-8$ Hz, 5-H), 3.164 (1H, dd, $J = 3.0, 13.1$ Hz, 1-Ha), 2.938 (1H, dd, $J = 9.9, 13.1$ Hz, 1-Hb), 1.650 (1H, ddd, $J = 2.0, 9.9, 14.8$ Hz, 3-Ha), 1.533 (1H, ddd, $J = 3.0, 10.8, 14.8$ Hz, 3-Hb).
- $^{13}\text{C-NMR}$ data (67.5 MHz, D_2O) **7a**; δ 75.54, 70.02, 65.02, 45.22, 37.31, 32.01, 31.20, 24.82, 22.11, 13.52; **7b**; δ 73.70, 71.00, 65.96, 44.48, 37.22, 32.10, 31.22, 24.93, 22.15, 13.54; **7c**; δ 74.60, 71.74, 66.24, 44.36, 36.18, 31.40, 31.20, 24.90, 22.11, 13.52; **7d**; δ 74.81, 70.48, 65.00, 45.24, 35.88, 31.51, 31.19, 24.93, 22.09, 13.52; **2^{2c}**; δ 74.72 (C-5), 70.49 (C-4), 64.97 (C-2), 45.28 (C-1), 35.74 (C-3), 31.49 (C-6), 25.70 (C-7).
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