

52. Further Studies on the Biosynthesis of Tabtoxin (Wildfire Toxin): Incorporation of [2,3-¹³C₂]Pyruvate into the β-Lactam Moiety

by Patricia Roth, Alfons Hädener, and Christoph Tamm*

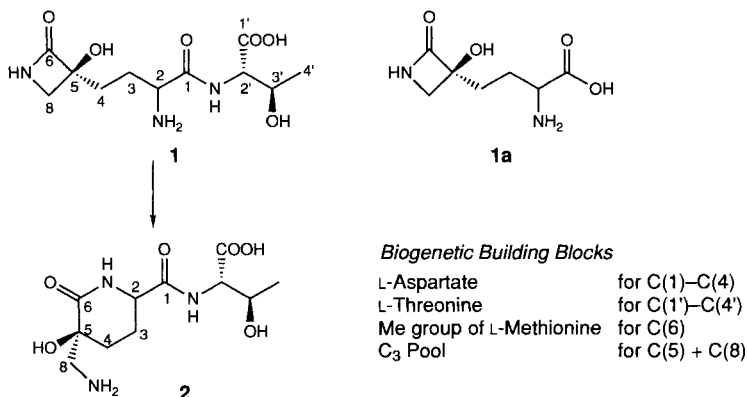
Institut für Organische Chemie der Universität, St. Johannis-Ring 19, CH-4056 Basel

(5.1.90)

[2,3-¹³C₂]Pyruvic acid (**7**) was synthesized and administered to cultures of *Pseudomonas syringae* pv *tabaci*. C(2) and C(3) of **7** were incorporated as an intact unit into the β-lactam moiety of tabtoxin (**1**). The result suggests that the biosynthesis of **1** is proceeding in part along the lysine pathway. The labelling pattern in **1** and an incorporation experiment with α,α'-dideuterated (±)-2,6-diaminopimelic acid (**19**) indicate that the branching in the biosynthesis of **1** occurs before lysine is formed.

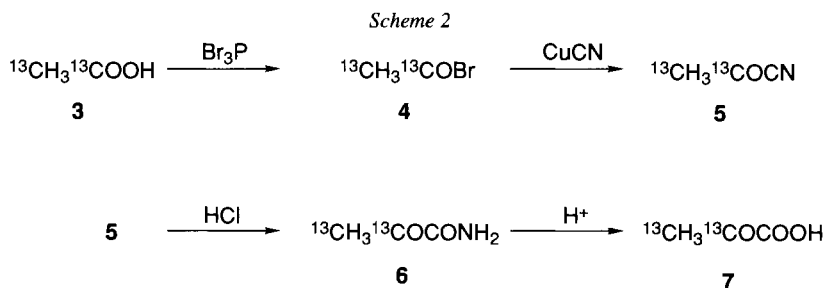
Introduction. – Tabtoxin (Wildfire toxin; **1**) is an extracellular pretoxin produced by the phytopathogenic bacterium *Pseudomonas syringae* pv *tabaci*. The biologically active β-lactam **1** is unstable [1]. It is converted to the stable but biologically inactive δ-lactam isotabtoxin (**2**) by intramolecular transacylation. In our previous investigations on the biosynthesis of tabtoxin, three amino acids were found to be involved in forming the molecule [2]. Incorporation experiments with ¹³C-labelled compounds demonstrated that L-aspartate and L-threonine are precursors of the side chain, whereas the Me group of L-methionine was found to provide the carbonyl C-atom of the β-lactam moiety. Furthermore, it was shown that the remaining C₂ unit of the β-lactam moiety is not derived from acetate as *e.g.* in thienamycin [3], but from an intermediate originating from the C₃ pool (*Scheme 1*). In this paper, we wish to report the synthesis and results of the incorporation experiments of doubly ¹³C-labelled pyruvic acid **7** and α,α'-dideuterated diaminopimelic acid **18** (*cf. Scheme 4*).

Scheme 1



Results and Discussion. – To improve the tabtoxin production, a suspension of *P. tabaci* NCPPB 2730 was applied to its host, the tobacco plant, and re-isolated after *ca.* 5 days [2]. Erlenmeyer flasks containing Woolley's medium [4] were inoculated and incubated on a rotary shaker at 28°. ¹³C- and ²H-Labelled potential precursors were administered by using pulse-feeding in 5 portions. The cultures were harvested after 72 h, yielding 50–100 mg/l isotabtoxin (**2**) as an artefact of the workup. Since the incorporation of the precursor into **2** is representative for the biosynthesis of **1**, no effort was made to isolate **1** in its original form (*Scheme 1*).

After the biogenetic origin of all C-atoms except C(5) and C(8) had been established, we assumed that the remaining C₂ unit in the β-lactam moiety was connected to the metabolism of the C₃ pool. This assumption is supported by the fact that racemic [1,2-¹³C₂]glycerol is incorporated into this C₂ unit, whereas [1,2-¹³C₂]acetic acid is not [2]. To clarify this point, we decided to synthesize doubly labelled pyruvic acid for incorporation experiments. [2,3-¹³C₂]Pyruvic acid (**7**) was prepared using [1,2-¹³C₂]acetic acid (**3**) as starting material (*Scheme 2*). The latter was converted first to [2,3-¹³C₂]acetyl bromide (**4**) [5] which reacted with CuCN to yield [2,3-¹³C₂]pyruvonitrile (**5**) [6]. Addition of HCl gas led to [2,3-¹³C₂]pyruvamide (**6**) from which the desired [2,3-¹³C₂]pyruvic acid (**7**) was obtained by treatment with aq. HCl.



The ¹³C-NMR spectral lines of the enriched isotabtoxin (**2**) sample, isolated after incorporation of **7**, is shown in *Fig. 1*. The observed ¹³C,¹³C-coupling pattern (*Fig. 2*) is consistent with the assumption that the atoms C(5) and C(8) are derived from pyruvate. A specific incorporation rate of 1.4% (*Table*) is found. The splitting pattern in the aspartate- and the threonine-derived part of the molecule is as expected. Pyruvate may enter the tricarboxylic-acid cycle in different ways. One of them leads from [2,3-¹³C₂]pyruvate via [1,2-¹³C₂]acetate into the cycle. Thus, [1,2-¹³C₂]- or [3,4-¹³C₂]aspartate is formed. In another route, **7** can be converted to [2,3-¹³C₂]oxalacetate or to [2,3-¹³C₂]malate. In this case, [2,3-¹³C₂]aspartate is formed. The specific incorporation rate for C(2)/C(3) of the aspartate moiety of **1** is 1.0%. Since aspartate is a direct precursor of threonine, we find the same coupling pattern in **1** for C(1') to C(4') as for C(1) to C(4).

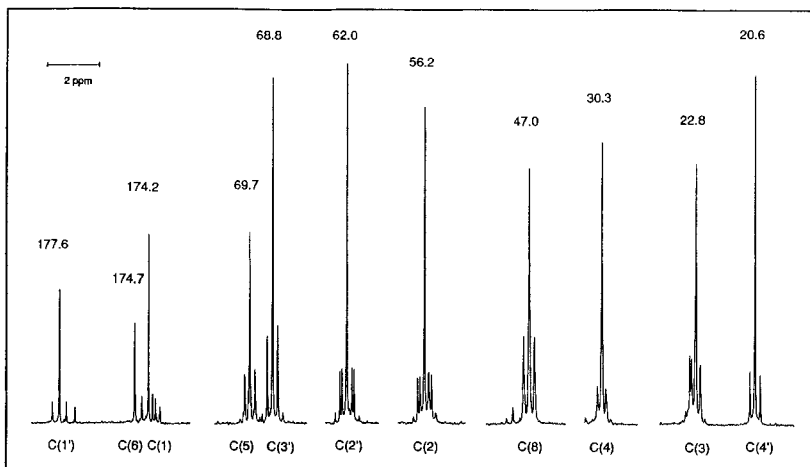


Fig. 1. 400-MHz Proton-noise-decoupled ^{13}C -NMR spectral lines (D_2O) of enriched isotabtoxin (**2**) after incorporation of $[2,3\text{-}^{13}\text{C}_2]\text{pyruvate}$. From left to right: C(1'), C(6), C(1), C(5), C(3'), C(2'), C(2), C(8), C(4), C(3), C(4').

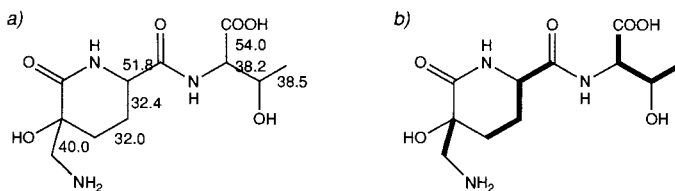


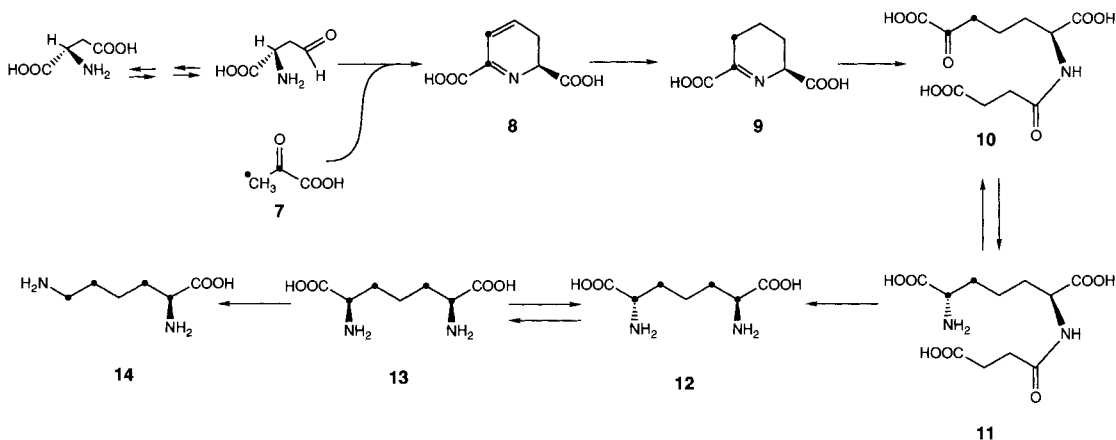
Fig. 2. a) $^{13}\text{C},^{13}\text{C}$ -Coupling constants and b) coupling pattern in isotabtoxin after incorporation of $[2,3\text{-}^{13}\text{C}_2]\text{pyruvate}$

The comparison of the specific incorporation rates in this experiment with those obtained earlier after incorporation of $[1,2\text{-}^{13}\text{C}_2]\text{glycerol}$ [2] clearly demonstrates that pyruvate is the more specific precursor (Table). Unkefer *et al.* [7] have recently reached the same conclusion, but on the basis of the incorporation of ^{13}C -labelled glucose.

Table. Specific Incorporation Rates for Some C_2 Units of **2** after Incorporation of $\text{rac}\text{-}[1,2\text{-}^{13}\text{C}_2]\text{Glycerol}$, $[2,3\text{-}^{13}\text{C}_2]\text{Pyruvate}$, $[1,2\text{-}^{13}\text{C}_2]\text{Acetate}$, and $\text{L}\text{-}[1,2\text{-}^{13}\text{C}_2]\text{-}$ and $\text{L}\text{-}[3,4\text{-}^{13}\text{C}_2]\text{Aspartate}$ [8]

Precursor	Specific incorporation rate [%]			Source
	C(2)/C(3)	C(3)/C(4)	C(5)/C(8)	
$\text{rac}\text{-}[1,2\text{-}^{13}\text{C}_2]\text{Glycerol}$	1.1	1.3	0.77	[2]
$[2,3\text{-}^{13}\text{C}_2]\text{Pyruvate}$	1.0	1.2	1.4	this work
$[1,2\text{-}^{13}\text{C}_2]\text{Acetate}$	2.6	3.5	0.0	[2]
$\text{L}\text{-}[1,2\text{-}^{13}\text{C}_2]\text{Aspartate/L}\text{-}[3,4\text{-}^{13}\text{C}_2]\text{Aspartate}$ 1:1	2.6	2.9	0.0	[2]

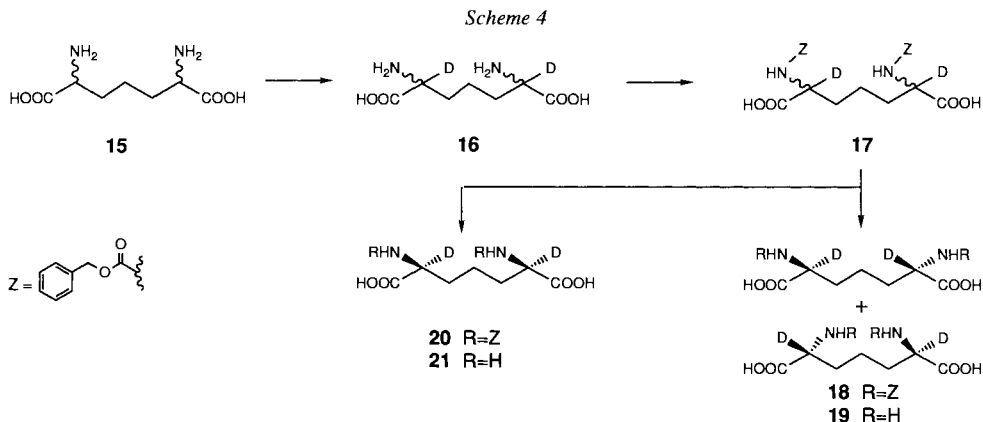
Since the building blocks of the tabtoxinine moiety (*cf.* **1a**) of **1** are the same as in the biosynthesis of lysine (**14**; with exception of the Me group from methionine), the following working hypothesis can be postulated: the tabtoxin biosynthesis is identical with the lysine pathway [9] (*Scheme 3*), until an unknown intermediate is reached. A detailed analysis of the lysine pathway reveals six possible intermediates. Lysine itself may probably be precluded, because, in an earlier feeding experiment, [^{14}C]lysine was incorporated to a much lesser extent than [^{14}C]aspartate [2]. One of the six intermediates **8–13** shows a C_2 axis of symmetry. Thus, the labelling is distributed between two parts of the molecule (one molecule with the labelling in its left-hand side, one molecule with the labelling in its right-hand side). The corresponding compound is (*S,S*)-2,6-diaminopimelic acid ((*S,S*)-2,6-DAP; **12**).

Scheme 3

If our working hypothesis is correct, the incorporation experiment with **7** as well as the earlier experiments with doubly labelled aspartic acid, glycerol and AcOH [2] should provide further information. If the biosynthesis of tabtoxinine (**1a**) is branching off after the formation of **12**, then the distribution of labelled C_2 units in the chain C(2)–C(3)–C(4)–C(5)–C(8) of **2** should be symmetrical with respect to C(4). In other words, a coupling between the pairs C(3)/C(4) and C(4)/C(5) should appear at the same intensity. If we do not find this symmetry, we can assume that the branching is attained, before **12** is formed. This consideration is only valid on the assumption that **12** is not enzyme-bound, but is moving unhindered in the cytoplasm. Only in this case, free rotation around the C_2 axis with resulting distribution of the label is possible. With this restriction, our experiments indicate that the branching might occur before the formation of **12**, since we do not observe a symmetrical distribution of the label. In particular, a coupling between C(4) and C(5) has never been observed, whereas the C_2 unit C(3)–C(4) was labelled in all incorporation experiments mentioned above (*Table*).

Since the cyclic intermediates **8** and **9** on the lysine pathway are known to be unstable [10], they could not be synthesized and administered in a labelled form. We, therefore,

decided to perform incorporation experiments with (\pm) -[2,6- $^2\text{H}_2$]diaminopimelic acid ((\pm) -[2,6- $^2\text{H}_2$]DAP; **19**). For this purpose, 2,6-diaminopimelic acid (**15**) was deuterated by stirring with $\text{D}_2\text{O}/\text{KOH}$ [11]. A mixture of diastereoisomers **16** is formed. To remove the *meso*-compound from the racemate, the isomers were converted into the benzyloxycarbonyl derivatives **17**, and the latter were separated by fractional crystallization (*cf.* Wade *et al.* [12]). Subsequent catalytic hydrogenolysis with Pd yielded pure (\pm) -[2,6- $^2\text{H}_2$]-DAP (**19**) and *meso*-[2,6- $^2\text{H}_2$]DAP (**21**).



A sample of isotabtoxin (**2**) isolated, after administration of **19** was investigated by ^2H -NMR spectroscopy. With the technique used, an incorporation of $> 0.25\%$ would have been detected, but none was found at C(2) or at C(8). The stability of the C- ^2H bond of the precursor under the usual pH conditions had already been tested. This result is consistent with the assumption that the biosynthesis of tabtoxin (**1**) is branching off from the lysine pathway before the formation of **12**. However, other reasons could as well explain the lack of incorporation of ^2H . Thus, **12** might not pass the bacterial cell membrane or might be metabolized in the primary metabolism prior to its possible use in tabtoxin biosynthesis.

The support of these investigations by the Swiss National Science Foundation is gratefully acknowledged.

Experimental Part

General. $^{13}\text{CH}_3^{13}\text{COOH}$ was purchased from ICN Stable Isotopes, Innerberg, Switzerland. *P. tabaci* strain NCPPB 2730 was from the National Collection of Plant Pathogenic Bacteria, England. Stock cultures of *P. tabaci* were maintained on agar slants containing NBY medium [13] and stored at 4° . The tobacco plants (*Nicotiana glauca*) were supplied by Prof. F. Meins (Friedrich-Miescher Institute, Ciba-Geigy AG, Basel). We thank Prof. F. Meins for this gift. The org. extracts were dried (Na_2SO_4) and evaporated under reduced pressure below 45° . TLC: silica gel 60 F_{254} (Merck); detection with UV or ninhydrin soln. Column chromatography: silica gel 60 (63–200 μm , Merck); cellulose microcrystallin Avicel pH 101 (50 μm , Fluka); solvent ratios in *v/v*. HPLC: Nucleosil- C_{18} (5 μm , 4.0×250 mm, Macherey-Nagel) with 0.1% CF_3COOH in H_2O or 0.1% CH_3COOH in H_2O . M.p.: Kofler Block, corrected. IR: Perkin-Elmer-781 spectrometer. NMR:

Varian-EM-360 (^1H , 60 MHz), Bruker-WH-90 with Fourier transform (^1H , 90 MHz; ^{13}C , 22.63 MHz), and Varian VXR-400 spectrometer with Fourier transform (^1H , 400 MHz; ^{13}C , 101 MHz); chemical shifts in ppm relative to internal TMS or TSP (1.7 ppm). MS: VG-70-250 instrument.

Production of Tabtoxin (= (S)-N-[4-(3-Hydroxy-2-oxo-3-azetidiny)-L-2-aminobutanoyl]-L-threonine; 1) and Isolation of Isotabtoxin (= (2S-trans)-N-[[5-(Aminomethyl)-5-hydroxy-6-oxo-2-piperidiny]carbonyl]-L-threonine; 2). Activation of bacteria on tobacco plants, growing of cultures, pulse feeding (150 mg/l of **7**, 250 mg/l of **18**), and isolation of **2** were carried out as described in [2]. $^1\text{H-NMR}$ (400 MHz, D_2O): 1.20 (*d*, $J = 6.5$, $\text{CH}_2(4')$); 1.96 (*m*, $\text{CH}_2(4)$); 2.07, 2.36 (*m*, *AB*, $\text{CH}_2(3)$); 3.13, 3.33 (*AB*, $J = 13.5$, $\text{CH}_2(8)$); 4.18 (*d*, $J = 4.5$, $\text{H-C}(2')$); 4.26 (*m*, $\text{H-C}(3')$); 4.36 (*t*, $J = 11.0$, $\text{H-C}(2)$). $^{13}\text{C-NMR}$ (101 MHz, D_2O): 20.6 ($\text{C}(4')$); 22.8 ($\text{C}(3)$); 30.3 ($\text{C}(4)$); 47.0 ($\text{C}(8)$); 56.2 ($\text{C}(2)$); 62.0 ($\text{C}(2')$); 68.8 ($\text{C}(3')$); 69.7 ($\text{C}(5)$); 174.2 ($\text{C}(1)$); 174.7 ($\text{C}(6)$); 177.6 ($\text{C}(1')$).

[1,2- $^{13}\text{C}_2$]Acetyl Bromide (**4**) [5]. To 1.12 ml (11.8 mmol) of Br_3P , 1.9 ml (33 mmol) of [1,2- $^{13}\text{C}_2$]acetic acid (90% ^{13}C) were added under stirring within 10 min. After stirring at 40–50° for 1 h, distillation yielded 1.55 g (12.6 mmol) of **4**. B.p. 69–73°.

[2,3- $^{13}\text{C}_2$]Pyruvonitrile (**5**) [6]. To 1.55 g (12.6 mmol) of **4**, 1.12 g (12.6 mmol) of CuCN were added within 20 min. After subsequent stirring at 80° for 2 h, 608 mg (8.8 mmol) of **5** were obtained by distillation. B.p. 86–90°. IR (CH_2Cl_2): 2200, 1728, 1360, 1170. $^1\text{H-NMR}$ (400 MHz, CDCl_3): 2.55 (*s*, CH_3).

[2,3- $^{13}\text{C}_2$]Pyruvamide (**6**) [6]. A soln. of 608 mg (8.8 mmol) of **5** in 35 ml of abs. Et_2O was saturated with dried HCl gas at 0°. After slow addition of 0.16 ml (8.8 mmol) of H_2O , the soln. was further treated with HCl gas for 10 min. The precipitate was washed with $\text{Et}_2\text{O}/\text{cyclohexane}$ 5:1 and dried to yield 410 mg (4.7 mmol) of **6**. M.p. 120–123°. IR (CH_2Cl_2): 3510, 3390, 1710, 1570, 1350, 1170. $^1\text{H-NMR}$ (400 MHz, D_2O , DMSO): 2.32 (*s*, CH_3); 7.61 (NH); 7.86 (NH). $^{13}\text{C-NMR}$ (101 MHz, D_2O , DMSO): 24.6 ($\text{C}(3)$); 163.2 ($\text{C}(1)$); 197.7 ($\text{C}(2)$). EI-MS: 89 (M^+), 45 ($[\text{M} - ^{13}\text{CH}_3\text{CO}]^+$, $[\text{M} - \text{CH}_3^{13}\text{CO}]^+$), 44 ($[\text{M} - ^{13}\text{CH}_3^{13}\text{CO}]^+$).

[2,3- $^{13}\text{C}_2$]Pyruvic Acid (**7**). A soln. of 410 mg (4.7 mmol) of **6** in 4.8 ml of 1N HCl and 5 ml of H_2O was refluxed for 2 h at 105°. After adding 1 g of NaCl , the product was extracted with Et_2O in a Kutscher-Steudel apparatus for 48 h. A subsequent bulb-to-bulb distillation (15 mm Hg, 110°) yielded 313 mg (3.5 mmol) of **7**. IR (CH_2Cl_2): 3400, 1785, 1730. $^1\text{H-NMR}$ (400 MHz, CDCl_3): 2.60 (*s*, $^{12}\text{CH}_3^{12}\text{CO}$; *d*, $J = 130.0$, $^{13}\text{CH}_3^{12}\text{CO}$; *d*, $J = 7.0$, $^{12}\text{CH}_3^{13}\text{CO}$; *dd*, $J = 130.0$, 7.0, $^{13}\text{CH}_3^{13}\text{CO}$); 8.68 (*s*, OH). $^{13}\text{C-NMR}$ (101 MHz, CDCl_3): 24.7 (*s*, *d*; $J = 45.0$ $\text{C}(3)$); 159.9 ($\text{C}(1)$); 193.5 (*s*, *d*; $J = 45.0$ $\text{C}(2)$). EI-MS: 90 (M^+), 45 (100).

2,6-Diamino[2,6- $^2\text{H}_2$]pimelic Acid (**16**) [11]. As described in [11], 2,6-diaminopimelic acid (**15**; a mixture of (*R,R*)-, (*S,S*)-, and (*R,S*)-isomers; 4.9 g, 25.7 mmol), KOH (5.85 g), and pyridoxal hydrochloride (525 mg; 2.57 mmol) were dissolved in 24.5 ml of D_2O . After stirring for 9.5 h, the solvent was removed by freeze-drying. The residue was dissolved in D_2O (35 ml), heated to reflux for 2 h, cooled, and the pH value adjusted to 5.0 by addition of 4N HCl . EtOH (100 ml) was added, until permanent cloudiness was observed. Cooling to 0° and filtration gave 3.47 g (21.2 mmol) of **16**. $^1\text{H-NMR}$ (400 MHz, D_2O): 1.48 (*m*, $\text{CH}_2(4)$); 1.91 (*m*, $\text{CH}_2(3)$, $\text{CH}_2(5)$); 3.74 (*t*, $J = 5.8$, $\text{H-C}(2)$, $\text{H-C}(6)$ of (*R,S*)-isomer); 3.78 (*t*, $J = 6.0$, $\text{H-C}(2)$, $\text{H-C}(6)$ of (*R,R*)- and (*S,S*)-isomer, resp.). FAB-MS: 193 (100, $[\text{M} + 3]^+$, $[\text{2,6-}^2\text{H}_2]\text{DAP}$), 192 ($[\text{M} + 2]^+$, $[\text{2,6-}^2\text{H}_1]\text{DAP}$), 191 ($[\text{MH}]^+$, 2,6-DAP).

2,6-Bis[(benzyloxy)carbonylamino][2,6- $^2\text{H}_2$]pimelic Acid (**17**) [12]. As described in [12], 1.56 g (8 mmol) of **16** were dissolved in 20.5 ml of 1N NaOH . After dropwise addition of 3.2 ml (22.6 mmol) of benzyl chloroformate within 30 min at 0°, the mixture was stirred for 2 h at r.t. The soln. was extracted with AcOEt , then the aq. layer was acidified (4N HCl) to pH 1.7.

(\pm)-Bis[(benzyloxy)carbonylamino][2,6- $^2\text{H}_2$]pimelic Acid (**18**) [12]. The precipitated oil **17** was extracted with AcOEt , dried, evaporated to 8 ml, and stored at 4° for 8 days. The precipitated solid was recrystallized from AcOEt to yield 1.40 g (3.0 mmol) of **18**. M.p. 152–154°. $^1\text{H-NMR}$ (400 MHz, D_2O , acetone): 1.62 (*m*, $\text{CH}_2(4)$); 1.85 (*m*, $\text{CH}_2(3)$, $\text{CH}_2(5)$); 4.23 (*m*, $\text{H-C}(2)$, $\text{H-C}(6)$, signals from undeuterated sites); 5.08 (*s*, $\text{CH}_2(2')$, $\text{CH}_2(2'')$); 6.5 (br., 2 COOH); 7.36 (*m*, 10 arom. H). FAB-MS: 461 ($[\text{MH}]^+$), 460, 417, 91.

(\pm)-2,6-Diamino[2,6- $^2\text{H}_2$]pimelic Acid (**19**) [12]. According to [12] 1.39 g (3.0 mmol) of **18** were dissolved in 30 ml of $\text{AcOH}/10$ ml of H_2O and hydrogenated in presence of 70 mg Pd black. The catalyst was filtered off and the solvent evaporated *in vacuo*. The residue was taken up in 12 ml of H_2O and evaporated again. The residue was recrystallized from 35% EtOH to obtain 537 mg (3.0 mmol) of **19**. The material consisted of 30% of doubly-deuterated, 52% of single-deuterated, and 18% of undeuterated product. $^1\text{H-NMR}$ (400 MHz, D_2O): 1.47 (*m*, $\text{CH}_2(4)$); 1.91 (*m*, $\text{CH}_2(3)$, $\text{CH}_2(5)$); 3.78 (*t*, $J = 5.9$, $\text{H-C}(2)$, $\text{H-C}(6)$). FAB-MS: 193 ($[\text{M} + 3]^+$, $[\text{2,6-}^2\text{H}_2]\text{DAP}$), 192 (100, $[\text{M} + 2]^+$, $[\text{2,6-}^2\text{H}_1]\text{DAP}$), 191 ($[\text{MH}]^+$, 2,6-DAP).

REFERENCES

- [1] W. J. Stewart, *Nature (London)* **1971**, 229, 174.
- [2] B. Müller, Dissertation, Universität Basel, 1987; B. Müller, A. Hädener, Ch. Tamm, *Helv. Chim. Acta* **1987**, 70, 412.
- [3] J. M. Williamson, *CRC Crit. Rev. Biotechnol.* **1986**, 4, 111.
- [4] D. W. Woolley, R. B. Pringle, A. C. Brown, *J. Biol. Chem.* **1952**, 197, 409.
- [5] M. Calvin, R. M. Lemmon, *J. Am. Chem. Soc.* **1947**, 69, 1232.
- [6] U. Hollstein, D. L. Mock, R. R. Sibbitt, *J. Lab. Compd.* **1980**, 17, 289.
- [7] C. J. Unkefer, R. E. London, R. D. Durbin, T. F. Uchytel, P. J. Langston–Unkefer, *J. Biol. Chem.* **1987**, 262, 4994.
- [8] A. Ian Scott, C. A. Townsend, K. Okada, M. Kajiwara, R. J. Cushley, P. J. Whitman, *J. Am. Chem. Soc.* **1974**, 96, 8069.
- [9] H. E. Umbarger, *Ann. Rev. Biochem.* **1978**, 47, 533.
- [10] a) Y. Yugari, Ch. Gilvarg, *J. Biol. Chem.* **1965**, 240, 4710; b) W. Farkas, Ch. Gilvarg, *ibid.* **1965**, 240, 4717.
- [11] J. G. Kelland, M. M. Palcic, M. A. Pickard, J. C. Vederas, *Biochemistry* **1985**, 24, 3263.
- [12] R. Wade, S. M. Birnbaum, M. Winitz, R. J. Koegel, J. P. Greenstein, *J. Am. Chem. Soc.* **1957**, 79, 648.
- [13] M. J. Gasson, *Appl. Environ. Microbiol.* **1980**, 39, 25.