

Note

Crystal and molecular structure of ethyl 6,7,8-tri-*O*-acetyl-5,9-anhydro-2,3,4-trideoxy-2-diphenylmethyleamino-4-nitro-D-*threo*-D-*altro*-nononate

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In an attempt to prepare the C-glycosyl amino acid analogue of *O*-(β -D-xylopyranosyl)serine, the mixture of the two epimeric addition products of β -D-xylopyranosylnitromethane (1,5-anhydro-6-deoxy-6-nitro-L-*gluco*-hexitol) to one equivalent of formaldehyde was acetylated and treated under basic conditions with *N*-diphenylmethyleneglycine ethyl ester. This reaction, which should proceed via a Michael-type addition to an intermediate nitro-olefin, yielded a mixture of four diastereomers, from which a major isomer (**1**), after partial fractionation by chromatography, could be crystallised in an overall yield of 9% ¹. Because of ambiguities in the configurational assignments of isomers by ¹H- and ¹³C-NMR spectroscopy, compound **1** was subjected to X-ray analysis at 300 K.

The relevant crystallographic data of **1** are given in Table I. The structure was solved by direct methods in the usual way with the help of the programs SHELXS90 ² and SHELX76 ³; **1** crystallized with two independent molecules in the asymmetric unit.

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TABLE I

Crystallographic data for **1**^a

Formula	C ₃₀ H ₃₄ O ₁₁ N ₂	Density (calcd) (g·cm ⁻³)	1.195
Mol wt	598.60	λ (Cu-K α) (pm)	154.051
Mp (degrees)	150	μ (cm ⁻¹)	7.3
Crystal size (mm)	0.4×0.3×0.2	2 θ range (degrees)	4.5–153
Space group	<i>P</i> 2 ₁	Reflections measured	7355
Cell parameters (pm, degrees)		Symmetry independent reflections	6463
<i>a</i>	845.4(1)	Symmetry independent reflections	
<i>b</i>	4383.3(3)	with $F_o > 3\sigma(F_o)$	5973
<i>c</i>	989.5(1)	Number of refined parameters	775
β	114.82(1)	Final residual factors	
Volume (pm ³)	3328.0(6)×10 ⁶	<i>R</i>	0.065
<i>Z</i>	4	<i>R_w</i>	0.062
<i>F</i> (000)	1264	Diffractometer	Enraf-Nonius CAD4

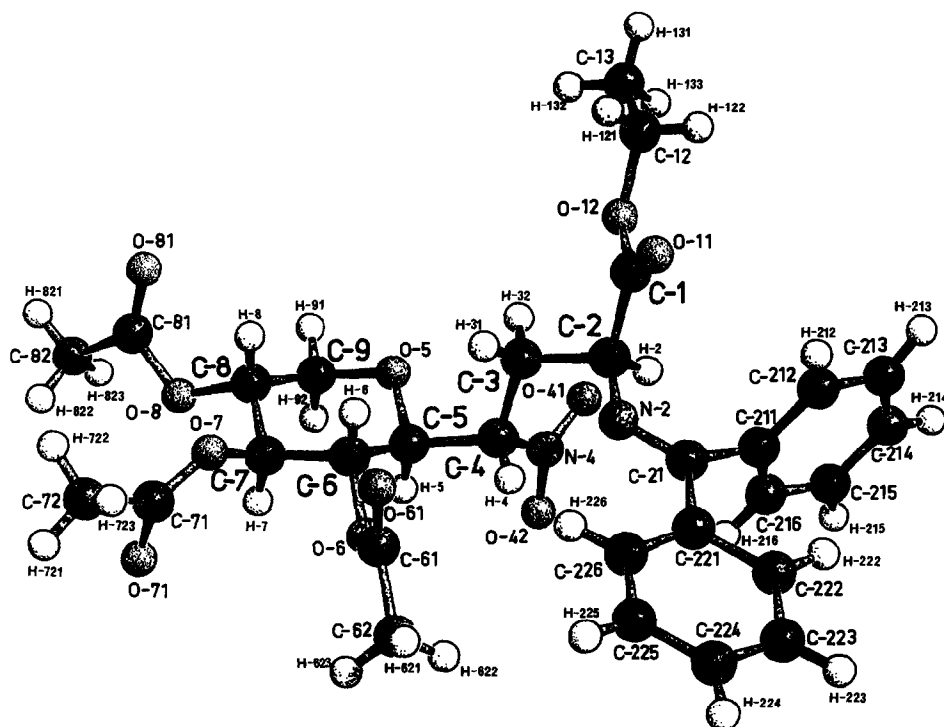
^a Standard deviations in parentheses.Fig. 1. SCHAKAL88⁴ plot of **1** (molecule I) showing atom numbering.

TABLE II

Fractional positional parameters of C, N, and O atoms ($\times 10^4$)^a for **1**

Atom	Molecule I			Molecule II		
	x	y	z	x	y	z
O-5	7228(4)	3621	378(3)	948(4)	1635(1)	7664(3)
O-6	6544(4)	4328(1)	– 1709(3)	2434(4)	917(1)	9633(3)
O-7	6288(5)	4514(1)	1080(4)	– 558(5)	744(1)	7000(4)
O-8	8636(4)	4158(1)	3486(4)	– 768(4)	1115(1)	4491(3)
O-11	532(5)	3294(1)	– 5186(5)	– 271(6)	1930(1)	13174(5)
O-12	2258(4)	3028(1)	– 3215(4)	– 365(5)	2218(1)	11347(4)
O-41	7298(6)	3164(1)	2270(5)	3714(6)	2090(1)	10341(5)
O-42	8990(6)	3522(1)	– 2248(5)	5338(6)	1734(1)	10210(6)
O-61	3787(7)	4481(2)	– 2529(7)	754(8)	808(1)	10774(6)
O-71	8085(9)	4866(1)	897(9)	913(8)	435(1)	6178(8)
O-81	7033(6)	3977(1)	4600(4)	– 3472(5)	1296(1)	3425(4)
N-2	3207(5)	3647(1)	– 5117(4)	2430(5)	1585(1)	13154(4)
N-4	7634(7)	3429(1)	– 2255(5)	3999(6)	1828(1)	10276(5)
C-1	1942(7)	3245(1)	– 4230(5)	265(6)	1992(1)	12311(5)
C-2	3614(6)	3398(1)	– 4063(5)	1801(6)	1840(1)	12130(5)
C-3	4603(6)	3516(1)	– 2474(5)	1232(6)	1725(1)	10517(5)
C-4	6353(6)	3666(1)	– 2229(5)	2704(6)	1587(1)	10230(5)
C-5	7238(6)	3830(1)	– 721(5)	2085(6)	1427(1)	8724(5)
C-6	6366(6)	4121(1)	– 629(5)	1194(6)	1131(1)	8635(5)
C-7	7311(6)	4261(1)	922(5)	519(6)	994(1)	7083(5)
C-8	7481(6)	4029(1)	2080(5)	– 523(6)	1233(1)	5949(5)
C-9	8259(7)	3741(1)	1850(5)	450(7)	1525(1)	6190(5)
C-12	810(8)	2844(2)	– 3249			

TABLE III

Selected torsion angles (°) in **1**^a (molecules I and II)

<i>Angles in the carbon chain</i>			<i>Angles between vicinal oxygens</i>		
	I	II		I	II
C-1–C-2–C-3–C-4	178.3(4)	176.8(4)	O-5–C-5–C-6–O-6	176.0(3)	174.9(4)
C-2–C-3–C-4–C-5	171.2(4)	170.4(4)	O-6–C-6–C-7–O-7	73.7(5)	70.9(5)
C-3–C-4–C-5–C-6	–72.6(5)	–73.2(5)	O-7–C-7–C-8–O-8	–70.9(5)	–74.0(5)
C-4–C-5–C-6–C-7	178.4(4)	176.6(4)	O-8–C-8–C-9–O-5	–173.9(4)	–172.2(4)
C-5–C-6–C-7–C-8	–52.3(6)	–50.5(6)			
C-6–C-7–C-8–C-9	51.9(6)	49.5(6)			
<i>Angles around the amino and nitro group</i>			<i>Angles between vicinal hydrogens</i>		
	I	II		I	II
O-12–C-1–C-2–N-2	–175.9(4)	–177.4(4)	H-2–C-2–C-3–H-31	176.0(4)	179.1(4)
C-1–C-2–N-2–C-21	–97.1(5)	–95.3(5)	H-2–C-2–C-3–H-32	–60.8(6)	–61.5(6)
N-2–C-2–C-3–C-4	–61.2(5)	–62.5(5)	H-31–C-3–C-4–H-4	–66.0(5)	–67.1(5)
N-4–C-4–C-5–O-5	–72.7(5)	–74.2(5)	H-32–C-3–C-4–H-4	173.1(4)	171.1(4)
N-4–C-4–C-5–C-6	165.9(4)	163.8(4)	H-4–C-4–C-5–H-5	–72.6(6)	–77.0(5)
N-4–C-4–C-3–C-2	–69.9(5)	–68.7(5)	H-5–C-5–C-6–H-6	176.9(4)	177.3(4)
C-3–C-4–C-5–O-5	48.8(5)	48.8(5)	H-6–C-6–C-7–H-7	–168.2(4)	–169.2(4)
			H-7–C-7–C-8–H-8	170.3(5)	166.4(4)
			H-8–C-8–C-9–H-91	–55.1(6)	–56.8(6)
			H-8–C-8–C-9–H-92	–176.3(4)	–175.9(4)

^a Standard deviations in parentheses.

All atoms (hydrogens included in theoretical positions) were refined. The final co-ordinates of C, N, and O are listed in Table II*. Selected torsion angles are given in Table III, and a perspective view (SCHAKAL88⁴ plot) of one of the independent molecules, which are only slightly different, is presented in Fig. 1, which clearly shows that the structure is as given in the title. By reactions already described¹, **1** can be transformed into the above-mentioned C-xylosyl amino acid with the correct “L” configuration of the amino acid part.

Therefore, we conclude that both topologies compete, and it remains to be demonstrated which is the more favoured.

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