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Key indicators

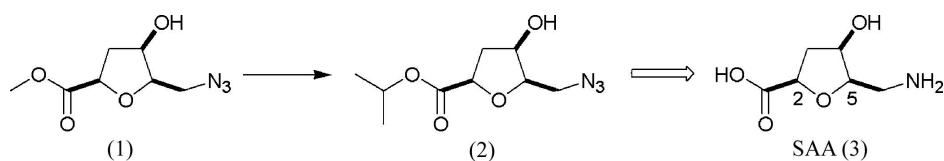
Single-crystal X-ray study
 $T = 190$ K
Mean $\sigma(\text{C}-\text{C}) = 0.003$ Å
 R factor = 0.040
 wR factor = 0.104
Data-to-parameter ratio = 9.1For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Isopropyl 2,5-anhydro-6-azido-3,6-dideoxy-D-xylo-hexonate

Determination of the crystal structure of the title isopropyl azido ester, $\text{C}_9\text{H}_{15}\text{N}_3\text{O}_4$, confirmed its relative stereochemistry and validated further work on the use of a derived sugar amino acid (SAA) as a peptidomimetic.Received 14 December 2005
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Comment

Sugar amino acids (SAAs) are carbohydrates which contain amine and acid groups; SAAs have been the focus of much interest as dipeptide isosteres, foldamers and library scaffolds (Hill *et al.*, 2001; Gruner *et al.*, 2002; Schweizer, 2002; Chakraborty, Srinivasu, Tapadar & Mohan, 2004; Trabocchi *et al.*, 2005). The preference for SAA oligomers (carbopeptoids) to adopt compact conformations as relatively short homooligomers will provide insight into the paradigm of protein folding. SAAs (2,5-*O*-*cis* configuration) structurally related to SAA (3) have a high propensity to adopt repeating β -turn conformations (Hungerford *et al.*, 2000; Smith *et al.*, 2003; Chakraborty, Srinivasu, Sakunthala *et al.*, 2004). In contrast, some 2,5-*O*-*trans* SAAs have been shown to adopt helical conformations (Claridge *et al.*, 1999; Claridge *et al.*, 2005). The conformational complexity of these dipeptide isosteres is being further explored by preparation of structurally related analogues of the original SAA systems *i.e.* SAA (3) and corresponding diastereoisomers (Watterson *et al.*, 2003).



The X-ray crystal structure (Fig. 1) firmly established the relative stereochemistry of the stereogenic centres in the title compound, (2). The absolute configuration of (2) (see scheme) is determined by the use of D-gulono-1,4-lactone as starting material.

The crystal packing consists of chains of molecules linked by hydrogen bonds and lying parallel to the *a* axis (Fig. 2). There are no unusual intermolecular contacts.

Experimental

The title compound, (2), was prepared from the methyl azido ester (1) in good yield by transesterification in acidic propan-2-ol, as described by Watterson *et al.* (2003); subsequent deprotection by hydrolysis and hydrogenation afforded SAA (3). The sample of (2) was crystallized from diethyl ether–hexane.

Crystal data

C₉H₁₅N₃O₄
M_r = 229.24
 Orthorhombic, *P*2₁2₁2₁
a = 5.4778 (7) Å
b = 11.0701 (13) Å
c = 18.2529 (15) Å
V = 1106.9 (2) Å³
Z = 4
D_x = 1.376 Mg m⁻³

Cu Kα radiation
 Cell parameters from 22 reflections
 $\theta = 21\text{--}44^\circ$
 $\mu = 0.92\text{ mm}^{-1}$
T = 190 K
 Block, colourless
 0.60 × 0.40 × 0.40 mm

Data collection

Enraf–Nonius Mach3 diffractometer
 $\omega/2\theta$ scans
 Absorption correction: ψ scan (North *et al.*, 1968)
T_{min} = 0.58, *T_{max}* = 0.69
 1335 measured reflections
 1335 independent reflections

1329 reflections with *I* > 2σ(*I*)
 $\theta_{\text{max}} = 73.9^\circ$
h = 0 → 6
k = 0 → 13
l = 0 → 22
 3 standard reflections
 frequency: 60 min
 intensity decay: 2.2%

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.040
wR(*F*²) = 0.104
S = 0.94
 1335 reflections
 146 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F^2) + (0.07P)^2 + 0.57P]$,
 where $P = [\max(F_o^2, 0) + 2F_c^2]/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.25\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.17\text{ e \AA}^{-3}$
 Extinction correction: Larson (1970), Equation 22
 Extinction coefficient: 159 (14)

Table 1

Selected geometric parameters (Å, °).

O1—C2	1.331 (3)	C6—C11	1.543 (3)
O1—C14	1.462 (3)	C7—N8	1.494 (3)
C2—O3	1.218 (3)	N8—N9	1.227 (3)
C2—C4	1.525 (3)	N9—N10	1.133 (3)
C4—O5	1.421 (3)	C11—C12	1.519 (3)
C4—C12	1.539 (3)	C11—O13	1.424 (3)
O5—C6	1.439 (3)	C14—C15	1.509 (3)
C6—C7	1.505 (3)	C14—C16	1.510 (4)
C2—O1—C14	116.44 (17)	C6—C7—N8	109.04 (19)
O1—C2—O3	124.2 (2)	C7—N8—N9	113.4 (2)
O1—C2—C4	111.63 (18)	N8—N9—N10	173.7 (3)
O3—C2—C4	124.07 (19)	C6—C11—C12	102.52 (17)
C2—C4—O5	109.68 (17)	C6—C11—O13	108.97 (17)
C2—C4—C12	113.67 (18)	C12—C11—O13	110.87 (18)
O5—C4—C12	104.84 (17)	C4—C12—C11	101.61 (18)
C4—O5—C6	109.60 (16)	O1—C14—C15	106.30 (18)
O5—C6—C7	111.26 (19)	O1—C14—C16	108.05 (19)
O5—C6—C11	106.93 (16)	C15—C14—C16	113.3 (2)
C7—C6—C11	112.91 (18)		

Table 2

Hydrogen-bond geometry (Å, °).

<i>D</i> — <i>H</i> ⋯ <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ⋯ <i>A</i>	<i>D</i> ⋯ <i>A</i>	<i>D</i> — <i>H</i> ⋯ <i>A</i>
O13—H13⋯O3 ⁱ	0.82	2.08	2.897 (2)	175

Symmetry code: (i) *x* + 1, *y*, *z*.

Attempted refinement of the Flack (1983) parameter gave an inconclusive result, in the absence of Friedel pairs and the presence of only weak anomalous scattering effects. The absolute configuration

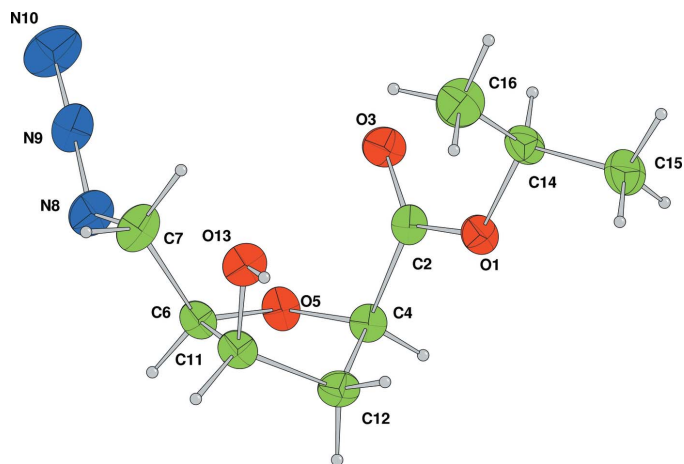


Figure 1

The title compound, with displacement ellipsoids drawn at the 50% probability level. H atoms are shown as spheres of arbitrary radius.

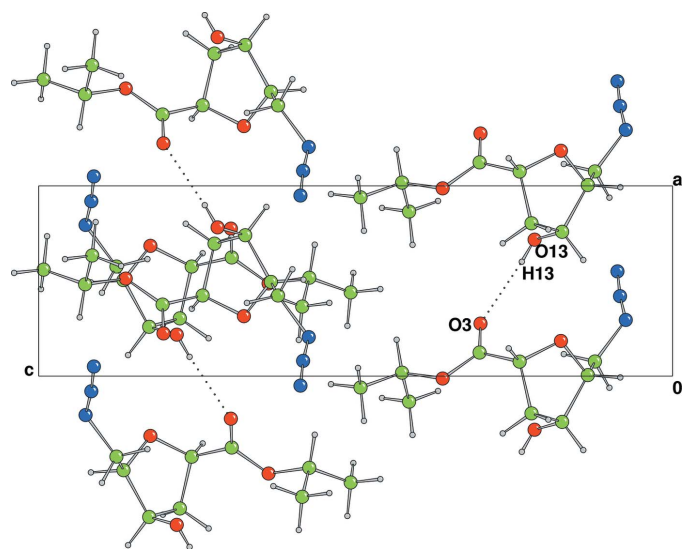


Figure 2

Projection of the title compound down the *b* axis, showing the hydrogen bonds (dashed lines) which link the molecules into columns.

was assigned from the known configuration of the starting material. The H atoms were all located in a difference map, but those attached to C atoms were repositioned geometrically. The H atoms were initially refined with soft restraints on the bond lengths and angles to regularize their geometry (C—H = 0.93–0.98 Å and O—H = 0.82 Å) and *U_{iso}*(H) values (in the range 1.2–1.5 times *U_{eq}* of the parent atom), after which they were refined with riding constraints.

Data collection: *CAD-4 EXPRESS*, (Straver, 1992); cell refinement: *CAD-4 EXPRESS*; data reduction: *RC93* (Watkin *et al.*, 1994); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *CRYSTALS* (Betteridge *et al.*, 2003); molecular graphics: *CAMERON* (Watkin *et al.*, 1996); software used to prepare material for publication: *CRYSTALS*.

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