organic papers

Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

Shahid Iqbal Awan,^a M. Iqbal Choudhary,^a Atta-ur-Rahman,^a Shazia Anjum,^a* Shamsher Ali^a and Hoong-Kun Fun^b

^aHEJ Research Institute of Chemistry, International Centre for Chemical Sciences, University of Karachi, Karachi 75270, Pakistan, and ^bX-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia

Correspondence e-mail: anjumshazia@yahoo.com

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.003 Å R factor = 0.030 wR factor = 0.078 Data-to-parameter ratio = 6.4

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

© 2005 International Union of Crystallography

Methyl [2,4,5-trihydroxy-6-(hydroxymethyl)perhydropyran-3-yl]carbamate

The title compound, $C_8H_{15}NO_7$, the methyl carbamate of β -D-glucosamine has been synthesized, in moderate yield, from the reaction of β -D-glucosamine hydrochloride and methyl chloroformate. There are two crystallographically independent molecules in the asymmetric unit. In both molecules, the pyranose ring adopts a slightly distorted chair conformation. In the crystal structure, molecules are packed along the *a* axis, with intra- and intermolecular O-H···O, N-H···O and C-H···O hydrogen bonds.

Comment

Amino-sugars have attracted growing interest due to their broad spectrum of application in chemistry, biochemistry, medicinal and pharmaceutical fields (Kirschning et al., 1997; Johnson & Liu, 1998; Elchert et al., 2004). For more than 30 years, non-steroidal anti-inflammatory drugs (NSAID) have been used in the treatment of osteoarthritis (OA). Serious and often life-threatening adverse effects due to these agents are common. Clinical findings have revealed that glucosamine sulfate and chondrotin sulfate are effective and safer alternatives for the alleviation of the symptoms of OA. Experimental evidence indicates that these compounds and their low molecular weight derivatives have a particular tropism for cartilage where they serve as substrates in the biosynthesis of component building blocks of glycosamineglycans (Ferguson 2005). Because joint pain is so debilitating, glucosamine alone is sometimes not enough, and it is important to further improve its biological activity.



Our research group has recently taken an interest in the synthetic manipulations of amino-sugars to develop some efficient pharmacophores to combat OA. The title compound, (II), is the methyl carbamate derivative of β -D-glucosamine (Viscontini & Meier, 1952). However, their procedure did not produce (II) in sufficient yield. Instead, we employed the Boullanger strategy (Boullanger *et al.*, 1987]; this resulted in a moderate yield (57%) from the reaction of β -D-glucosamine hydrochloride, (I), with methyl chloroformate in the presence of triethylamine and sodium methoxide.

The conformation of the pyranose ring is ${}^{4}C_{1}$ in both independent molecules; the puckering parameters are $Q = 0.565 (2)/0.564 (2) \text{ Å}, \theta = 5.1 (2)/6.2 (2)^{\circ}$ and $\varphi = 52 (3)/55 (2)^{\circ}$

Received 9 June 2005 Accepted 27 June 2005 Online 6 July 2005

Printed in Great Britain - all rights reserved



Figure 1

The asymmetric unit of (II), showing 50% probability displacement ellipsoids and the atom-numbering scheme.



Figure 2

Molecular packing of (II), viewed along the *a* axis. Hydrogen bonds are indicated by dashed lines.

(Cremer & Pople, 1975). The pyranose rings are distorted from an ideal chair conformation, allowing the C1-O1-C5 and C1A - O1A - C5A bond angles to widen to 113.0 (2) and 114.3 (2) $^{\circ}$, respectively, while the other internal ring angles remain close to the tetrahedral value. The C-C bond lengths lie in the range 1.516 (3)–1.530 (3) Å. The shortening of the O2-C1 and O2A-C1A bond lengths [1.391 (3) and 1.392 (3) Å, respectively] compared to the O1-C1 and O1A - C1A bond lengths [1.414 (3) & 1.416 (3) Å, respectively] indicates a significant anomeric effect, which is normally found in free sugar derivatives (Berman et al., 1967). There is electron delocalization over O6=C7-O7 and O6A = C7A - O7A, resulting in the shortening of the O7 - C7 and O7A - C7A bond lengths [1.344 (3) and 1.345 (3) Å, respectively]. The torsion angles C5-O1-C1-O2 = $174.2 (17)^{\circ}$ and C5A - O1A - C1A - O2A = 172.6 (17) Åindicate the β -configuration of the substitutent at the C1 and C1A anomeric centers.

Extensive $O-H \cdots O$ type hydrogen bonding is present in (II), accounting for the molecular conformation and stability of the crystal structure. All available O and N atoms are involved in the hydrogen bonding (Fig. 2 and Table 1).

Experimental

 β -D-Glucosamine hydrochloride, (I), (1.0 g, 4.6 mmol) was added to a sodium methoxide solution (4.6 ml methanol and 106 mg sodium metal were mixed and shaken at room temperature for 10 min and then at 273 K for 5 min). Triethylamine (0.3 ml, 4.6 mmol) was added dropwise to the reaction flask, with subsequent addition of methyl chloroformate (0.18 ml, 4.6 mmol) at 273 K. The reaction was completed after 20 min and the solvent was then evaporated. The title compound, (II), was purified by flash column chromatography, and crystallized from dichloromethane and methanol (20:80) in 57% yield (616 mg; m.p. 469 K).

Crystal data

$D_x = 1.526 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation
Cell parameters fro
reflections
$\theta = 2.3 - 25.0^{\circ}$
$\mu = 0.14 \text{ mm}^{-1}$
T = 293 (2) K
Block, colorless
$0.64\times0.56\times0.14$

Data collection

Siemens SMART CCD area detector diffractometer ω scans Absorption correction: multi-scan (SADABS: Sheldrick, 1996) $T_{\min} = 0.919, \ T_{\max} = 0.981$ 5804 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.030$ $wR(F^2) = 0.078$ S = 1.082082 reflections 324 parameters H atoms treated by a mixture of independent and constrained refinement

om 5705 mm

2082 independent reflections 2033 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.023$ $\theta_{\rm max} = 26.0^{\circ}$ $h = -8 \rightarrow 7$ $k = -21 \rightarrow 15$ $l = -10 \rightarrow 11$

 $w = 1/[\sigma^2(F_0^2) + (0.0456P)^2]$ + 0.1323P] where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta \rho_{\rm max} = 0.25 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$ Extinction correction: SHELXL97 Extinction coefficient: 0.016 (3)

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1A - H1AA \cdots O2^{i}$	0.86	2.19	3.007 (3)	159
$N1 - H1A \cdots O2A^{ii}$	0.86	2.22	3.016 (3)	155
$O2-H2A\cdots O4^{iii}$	0.81 (4)	1.93 (4)	2.731 (2)	171 (3)
$O3-H3A\cdots O6^{iv}$	0.84 (3)	1.98 (3)	2.800 (3)	168 (3)
$O4-H4A\cdots O5A^{v}$	0.87 (4)	1.89 (3)	2.750 (3)	166 (3)
$O5-H5A\cdots O1A^{iii}$	0.88 (7)	2.18 (6)	2.883 (3)	137 (5)
$O2A - H2AA \cdots O4A^{iv}$	0.93 (3)	1.84 (4)	2.755 (3)	171 (3)
$O3A - H3AA \cdots O6A^{iii}$	0.78 (4)	2.01(4)	2.793 (3)	173 (4)
$O4A - H4AA \cdots O3^{vi}$	0.78 (4)	1.99 (4)	2.766 (3)	171 (4)
$O5A - H5AA \cdots O5^{iv}$	0.81 (5)	2.06 (5)	2.865 (3)	169 (4)
$C2-H2B\cdots O6^{v}$	0.98	2.47	2.843 (3)	102
$C2A - H2AB \cdots O6A^{v}$	0.98	2.46	2.821 (3)	101
$C4-H4B\cdots O5^{v}$	0.98	2.58	2.931 (3)	101
$C4-H4B\cdots O6A^{vii}$	0.98	2.51	3.388 (3)	149
$C4A - H4AB \cdots O6^{vi}$	0.98	2.45	3.432 (3)	175
$C5-H5B\cdots O7A^{ii}$	0.98	2.55	3.446 (3)	152
$C5A - H5AB \cdots O7^{i}$	0.98	2.56	3.477 (3)	157
$C6-H6B\cdots O5A^{v}$	0.97	2.56	3.395 (3)	144
$C6A - H6AA \cdots O4A^{v}$	0.97	2.59	2.969 (3)	103
$C8-H8A\cdotsO1^{viii}$	0.96	2.57	3.229 (4)	126

 $\overline{ \text{Symmetry codes: (i)} -x, y + \frac{1}{2}, -z + 2; (ii) -x + 1, y - \frac{1}{2}, -z + 2; (iii) x - 1, y, z; (iv) x + 1, y, z; (v) x, y, z; (v) -x, y + \frac{1}{2}, -z + 1; (vii) -x + 1, y - \frac{1}{2}, -z + 1; (vii) -x + 1; (vii) -x + 1, y - \frac{1}{2}, -z + 1; (vii) -x +$ $-x-1, y-\frac{1}{2}, -z+2$

All C- and N-bound H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H = 0.97-0.98 Å and N-H = 0.86 Å, and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H and $1.2U_{eq}(C)$ for others. A rotating group model was used for the methyl group. The hydroxyl H atoms were located in a difference map and their parameters were freely refined; the O-H distances lie in the range 0.78 (4)-0.93 (4) Å. In the absence of significant anomalous dispersion effects, Friedel pairs were averaged before the final refinement and the absolute configuration was assigned on the basis of the starting material.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2003).

SA thanks the Higher Education Commission for the research grant R&D/Acad/03/1064 to conduct research on synthetic manipulations of amino-sugars. AUR, SA and HKF

also thank the Malaysian Goverment and Universiti Sains Malaysia for the Scientific Advancement Grant Allocation (SAGA) grant No. 304/PFIZIK/653003/A118.

References

- Berman, H. M., Chu, S. S. C. & Jeffrey, G. A. (1967). Science, 157, 1576–1577.
- Boullanger, P., Banoub, J. & Descotes, G. (1987). Can. J. Chem. 65, 1343–1348.
- Cremer, D. & Pople, J. A. (1975). J. Am. Chem. Soc. 97, 1354-1358.
- Elchert, B., Li, J., Wang, J., Hui, Y., Rai, R., Ptak, R., Ward, P., Takemoto, J. Y., Bensaci, M. & Chang, C.-W. T. (2004). J. Org. Chem. 69, 1513–1523.
- Ferguson, A. (2005). *http://www.rowingact.Org.au/SDO/Masters/Glucosamine.html*.
- Johnson, D. A. & Liu, H.-W. (1998). Curr. Opin. Chem. Biol. 2, 642-649.
- Kirschning, A., Bechthold, A. F.-W. & Rohr, J. (1997). Top. Curr. Chem. 188, 1–84.
- Sheldrick, G. M. (1996). SADABS. University of Göttingen, Germany.
- Sheldrick, G. M. (1997). SHELXTL. Version 5.1. Bruker AXS Inc., Madison, Wisconsin, USA.
- Siemens (1996). SMART and SAINT. Siemens Analytical X-Ray Instruments Inc., Madison, Wisconsin, USA.
- Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.
- Viscontini, M. & Meier, J. (1952). Helv. Chim. Acta, 35, 807-12.