

**(1*R*,2*R*,3*S*,6*aS*,7*R*,8*R*,9*S*,12*aS*)-
1,2,3,7,8,9-Hexahydroxyperhydro-
dipyrido[1,2-a:1',2'-d]pyrazine-6,12-
dione**

**S. F. Jenkinson,^{a*} D. Best,^a F. X. Wilson,^b G. W. J. Fleet^a
and D. J. Watkin^c**

^aDepartment of Organic Chemistry, Chemistry Research Laboratory, University of Oxford, Oxford OX1 3TA, England, ^bSummit PLC, 91 Milton Park, Abingdon, Oxon OX14 4RY, England, and ^cDepartment of Chemical Crystallography, Chemistry Research Laboratory, University of Oxford, Oxford OX1 3TA, England
Correspondence e-mail: sarah.jenkinson@chem.ox.ac.uk

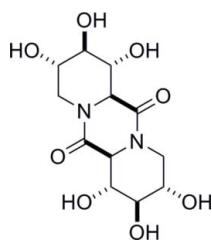
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Key indicators: single-crystal X-ray study; $T = 150\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; R factor = 0.042; wR factor = 0.101; data-to-parameter ratio = 8.4.

The crystal structure of the title compound, $\text{C}_{12}\text{H}_{18}\text{N}_2\text{O}_8$, exists as $\text{O}-\text{H}\cdots\text{O}$ hydrogen-bonded layers of molecules running parallel to the ab plane. Each molecule is a donor and acceptor for six hydrogen bonds. The absolute stereochemistry was determined by the use of D -glucuronolactone as the starting material.

Related literature

For the isolation and biological activity of pipecolic acids, see: Manning *et al.* (1985); di Bello *et al.* (1984). For the synthesis of pipecolic acids, see: Bashyal *et al.* (1986); Bashyal, Chow & Fleet (1987); Bashyal, Chow, Fellows & Fleet (1987).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{18}\text{N}_2\text{O}_8$	$V = 1256.34(5)\text{ \AA}^3$
$M_r = 318.28$	$Z = 4$
Orthorhombic, $P2_12_12_1$	$\text{Mo K}\alpha$ radiation
$a = 7.8711(2)\text{ \AA}$	$\mu = 0.14\text{ mm}^{-1}$
$b = 8.1526(2)\text{ \AA}$	$T = 150\text{ K}$
$c = 19.5783(5)\text{ \AA}$	$0.40 \times 0.10 \times 0.10\text{ mm}$

Data collection

Nonius KappaCCD area-detector diffractometer
Absorption correction: multi-scan (*DENZO/SCALEPACK*;
Otwinowski & Minor, 1997)
 $R_{\text{int}} = 0.061$
 $T_{\min} = 0.85$, $T_{\max} = 0.99$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.101$
 $S = 0.93$
1662 reflections

199 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.35\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.43\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O8—H8i \cdots O17 ⁱ	0.83	1.95	2.756 (4)	162
O22—H22i \cdots O1 ⁱⁱ	0.83	2.22	2.917 (4)	141
O19—H19i \cdots O11 ⁱⁱⁱ	0.82	2.12	2.793 (4)	139
O11—H11i \cdots O13 ^{iv}	0.83	1.86	2.685 (4)	173
O17—H17i \cdots O8 ⁱⁱⁱ	0.80	1.87	2.633 (4)	157
O6—H6i \cdots O19 ^v	0.83	1.97	2.680 (4)	143

Symmetry codes: (i) $x + 1, y, z$; (ii) $x, y - 1, z$; (iii) $x - \frac{1}{2}, -y + \frac{3}{2}, -z + 1$; (iv) $x, y + 1, z$; (v) $x + \frac{1}{2}, -y + \frac{1}{2}, -z + 1$.

Data collection: *COLLECT* (Nonius, 2001); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; 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*.

We would like to thank the Chemical Crystallography department and ALT at the University of Oxford for use of the diffractometers.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5005).

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supporting information

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(1*R*,2*R*,3*S*,6*aS*,7*R*,8*R*,9*S*,12*aS*)-1,2,3,7,8,9-Hexahydroxyperhydrodipyrido[1,2-a:1',2'-*d*]pyrazine-6,12-dione

S. F. Jenkinson, D. Best, F. X. Wilson, G. W. J. Fleet and D. J. Watkin

S1. Comment

2*S*,3*R*,4*R*,5*S*-Trihydroxypipeolic acid (BR1) **2** (Fig. 1), a sugar mimic of glucuronic acid, has been isolated from the seeds of *Baphia racemosa* (Manning *et al.*, 1985) and shown to inhibit both glucuronidase and iduronidase activity (di Bello *et al.*, 1984). In a modification of the original synthesis of BR1 from D-glucuronolactone (Bashyal *et al.*, 1986, Bashyal, Chow & Fleet, 1987, Bashyal, Chow, Fellows & Fleet, 1987), reduction of the azide **1** afforded a low yield of **2** together with by-products. One of the components of the mixture was crystallized; the structure of this material was determined unequivocally by X-ray crystallographic analysis and shown to be the diketopiperazine **3** (Fig. 2). The absolute stereochemistry was determined by the use of D-glucuronolactone as the starting material. The structure consists of layers of hydrogen bonded molecules running parallel to the *ab* plane (Fig. 3, Fig. 4). Each molecule is a donor and acceptor for 6 hydrogen bonds. Only classical hydrogen bonding was considered.

S2. Experimental

The title compound was recrystallised by diffusion from a mixture of water and acetonitrile: m.p. 511 K decomposed; $[\alpha]_D^{20} + 29.7$ (*c*, 0.35 in H₂O).

S3. Refinement

In the absence of significant anomalous scattering, Friedel pairs were merged and the absolute configuration was assigned from the starting material.

The H atoms were all located in a difference map, but those attached to carbon 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 in the range 0.93–0.98, N—H in the range 0.86–0.89 N—H to 0.86 O—H = 0.82 Å) and *U*_{iso}(H) (in the range 1.2–1.5 times *U*_{eq} of the parent atom), after which the positions were refined with riding constraints.

One outlying reflection was omitted from the refinement as it was thought to be partially occluded by the beam stop.

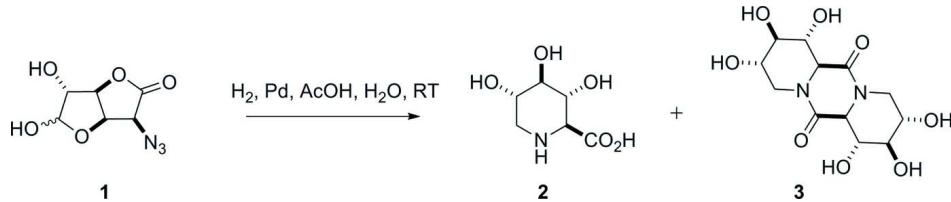
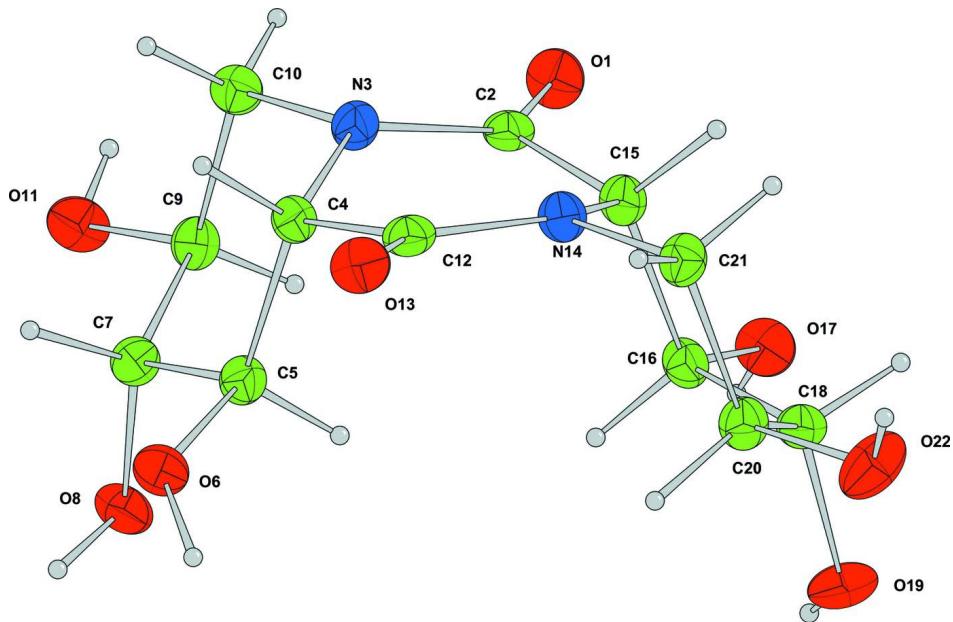


Figure 1

Synthetic Scheme.

**Figure 2**

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

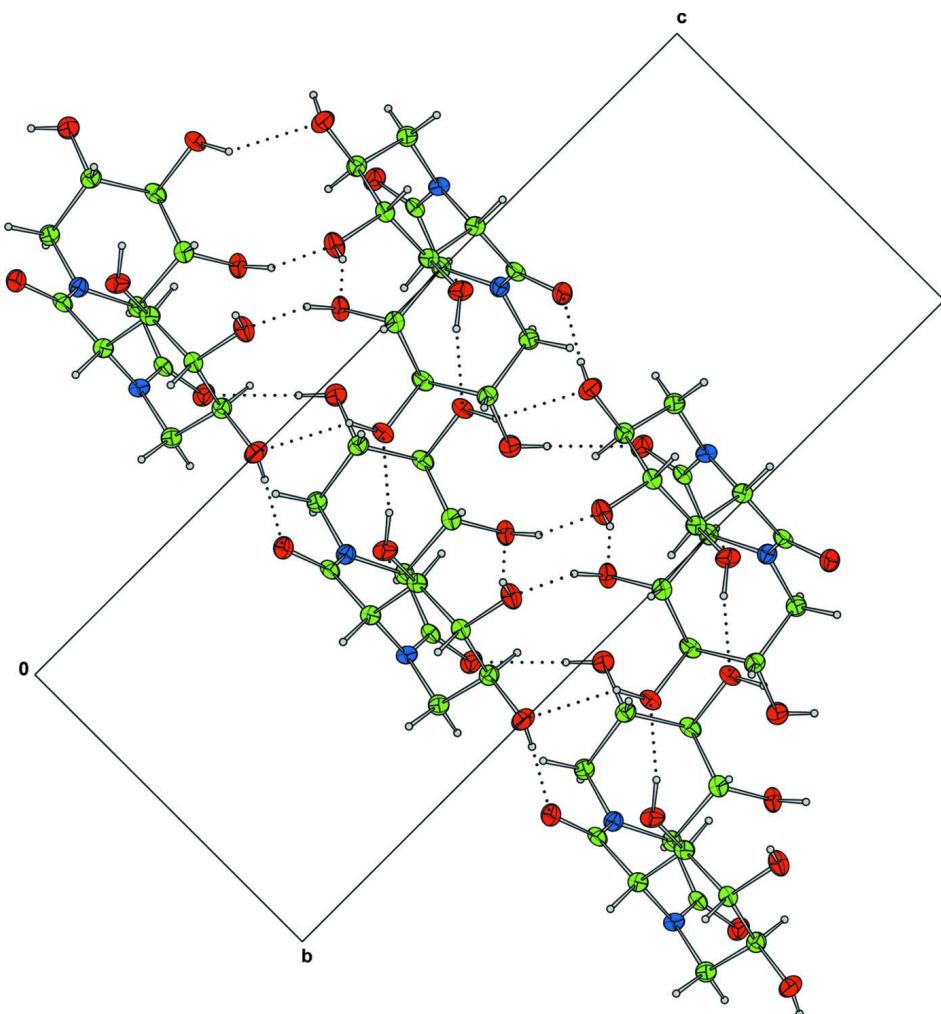
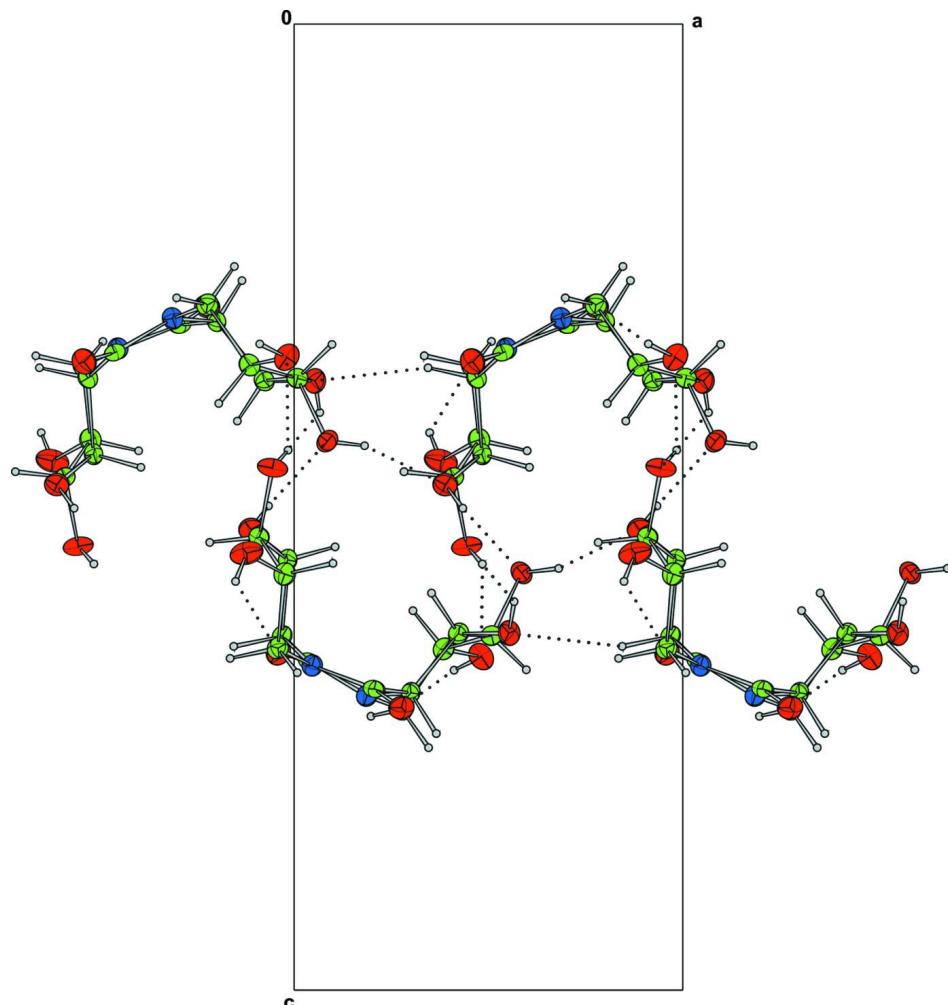


Figure 3

Packing diagram for the title compound projected along the a -axis. Hydrogen bonds are shown by dotted lines.

**Figure 4**

Packing diagram for the title compound projected along the b axis. Hydrogen bonds are shown by dotted lines.

(1*R*,2*R*,3*S*,6*aS*,7*R*,8*R*,9*S*,12*aS*)-1,2,3,7,8,9-Hexahydroxyperhydrodipyrido[1,2-*a*:1',2'-*d*]pyrazine-6,12-dione

Crystal data

$C_{12}H_{18}N_2O_8$
 $M_r = 318.28$
Orthorhombic, $P2_12_12_1$
Hall symbol: P 2ac 2ab
 $a = 7.8711 (2) \text{ \AA}$
 $b = 8.1526 (2) \text{ \AA}$
 $c = 19.5783 (5) \text{ \AA}$
 $V = 1256.34 (5) \text{ \AA}^3$
 $Z = 4$

$F(000) = 672$
 $D_x = 1.683 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 1646 reflections
 $\theta = 5-27^\circ$
 $\mu = 0.14 \text{ mm}^{-1}$
 $T = 150 \text{ K}$
Plate, colourless
 $0.40 \times 0.10 \times 0.10 \text{ mm}$

Data collection

Nonius KappaCCD area-detector
diffractometer
Graphite monochromator
 ω scans

Absorption correction: multi-scan
(DENZO/SCALEPACK; Otwinowski & Minor,
1997)
 $T_{\min} = 0.85$, $T_{\max} = 0.99$

12748 measured reflections
 1663 independent reflections
 1348 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.061$

$\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 5.1^\circ$
 $h = -10 \rightarrow 10$
 $k = -10 \rightarrow 10$
 $l = -25 \rightarrow 25$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.042$
 $wR(F^2) = 0.101$
 $S = 0.93$
 1662 reflections
 199 parameters
 0 restraints
 Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained
 Method = Modified Sheldrick $w = 1/\sigma^2(F^2) + (0.06P)^2 + 0.76P$, where $P = [\max(F_o^2, 0) + 2F_c^2]/3$
 $(\Delta/\sigma)_{\max} = 0.000342$
 $\Delta\rho_{\max} = 0.35 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.43 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.4593 (3)	0.7922 (2)	0.34949 (10)	0.0245
C2	0.5380 (4)	0.6635 (3)	0.34006 (13)	0.0181
N3	0.6870 (3)	0.6583 (3)	0.30476 (12)	0.0186
C4	0.8001 (3)	0.5165 (3)	0.30769 (13)	0.0178
C5	0.9227 (3)	0.5430 (3)	0.36900 (14)	0.0187
O6	1.0531 (3)	0.4227 (2)	0.36945 (10)	0.0233
C7	1.0081 (4)	0.7115 (3)	0.36672 (14)	0.0191
O8	1.0852 (2)	0.7438 (2)	0.43187 (9)	0.0238
C9	0.8851 (4)	0.8502 (3)	0.35308 (14)	0.0196
C10	0.7767 (4)	0.8115 (4)	0.29066 (15)	0.0202
O11	0.9812 (3)	0.9966 (2)	0.34507 (10)	0.0247
C12	0.7057 (4)	0.3552 (3)	0.31154 (14)	0.0192
O13	0.7761 (3)	0.2272 (2)	0.29231 (10)	0.0234
N14	0.5463 (3)	0.3566 (3)	0.33597 (12)	0.0186
C15	0.4689 (3)	0.5025 (3)	0.36625 (13)	0.0188
C16	0.4834 (3)	0.4874 (3)	0.44528 (13)	0.0187
O17	0.3887 (3)	0.6143 (2)	0.47677 (10)	0.0240
C18	0.4098 (4)	0.3245 (3)	0.46900 (13)	0.0203
O19	0.4442 (3)	0.3016 (3)	0.54059 (9)	0.0269
C20	0.4757 (4)	0.1754 (3)	0.43053 (13)	0.0201
C21	0.4581 (4)	0.2045 (3)	0.35390 (13)	0.0199
O22	0.3776 (3)	0.0390 (2)	0.45221 (10)	0.0310
H41	0.8655	0.5150	0.2656	0.0187*
H51	0.8535	0.5336	0.4109	0.0227*
H71	1.0974	0.7116	0.3316	0.0230*
H91	0.8058	0.8605	0.3927	0.0225*
H102	0.8459	0.7964	0.2509	0.0230*
H101	0.6975	0.8990	0.2835	0.0223*
H151	0.3443	0.5003	0.3557	0.0220*
H161	0.6054	0.4942	0.4588	0.0216*

H181	0.2825	0.3283	0.4634	0.0238*
H201	0.5996	0.1598	0.4417	0.0244*
H212	0.5077	0.1118	0.3286	0.0232*
H211	0.3349	0.2163	0.3433	0.0228*
H81	1.1809	0.7027	0.4365	0.0340*
H221	0.3488	-0.0299	0.4231	0.0459*
H191	0.4842	0.3832	0.5589	0.0407*
H111	0.9114	1.0645	0.3311	0.0377*
H171	0.4338	0.6804	0.5013	0.0351*
H61	1.0660	0.3593	0.4021	0.0354*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0253 (11)	0.0201 (10)	0.0280 (11)	0.0043 (9)	0.0024 (9)	-0.0012 (8)
C2	0.0189 (14)	0.0226 (14)	0.0128 (13)	0.0006 (13)	-0.0029 (11)	-0.0002 (10)
N3	0.0185 (12)	0.0171 (11)	0.0203 (13)	0.0000 (10)	0.0012 (10)	0.0030 (9)
C4	0.0153 (12)	0.0194 (13)	0.0186 (13)	0.0024 (12)	0.0012 (10)	0.0011 (11)
C5	0.0174 (13)	0.0198 (13)	0.0188 (13)	0.0017 (11)	0.0017 (11)	0.0014 (10)
O6	0.0206 (10)	0.0224 (10)	0.0270 (10)	0.0069 (9)	-0.0006 (9)	0.0050 (9)
C7	0.0193 (14)	0.0209 (13)	0.0170 (13)	0.0003 (11)	0.0006 (11)	-0.0014 (11)
O8	0.0191 (10)	0.0316 (11)	0.0208 (10)	0.0048 (9)	-0.0041 (8)	-0.0056 (8)
C9	0.0191 (14)	0.0192 (14)	0.0205 (14)	-0.0001 (12)	0.0020 (11)	-0.0008 (10)
C10	0.0200 (15)	0.0205 (13)	0.0203 (15)	0.0011 (12)	-0.0010 (11)	0.0004 (11)
O11	0.0265 (10)	0.0169 (10)	0.0307 (10)	0.0000 (10)	-0.0056 (8)	0.0012 (8)
C12	0.0212 (14)	0.0235 (15)	0.0129 (14)	0.0016 (12)	-0.0019 (11)	0.0007 (11)
O13	0.0274 (12)	0.0206 (10)	0.0223 (11)	0.0049 (9)	0.0016 (9)	-0.0027 (8)
N14	0.0182 (12)	0.0180 (12)	0.0197 (12)	-0.0009 (11)	0.0002 (10)	-0.0004 (9)
C15	0.0168 (12)	0.0208 (14)	0.0186 (12)	0.0026 (13)	0.0021 (10)	-0.0015 (12)
C16	0.0151 (12)	0.0214 (14)	0.0197 (12)	0.0014 (12)	0.0006 (10)	-0.0017 (11)
O17	0.0259 (11)	0.0240 (10)	0.0220 (10)	0.0037 (9)	0.0005 (9)	-0.0075 (8)
C18	0.0182 (14)	0.0270 (14)	0.0158 (14)	0.0028 (12)	-0.0007 (11)	0.0015 (11)
O19	0.0392 (12)	0.0261 (10)	0.0153 (9)	0.0003 (10)	-0.0045 (9)	0.0013 (8)
C20	0.0200 (13)	0.0183 (13)	0.0220 (14)	-0.0013 (12)	0.0010 (11)	0.0007 (11)
C21	0.0195 (14)	0.0200 (14)	0.0201 (14)	-0.0034 (12)	0.0015 (12)	-0.0005 (11)
O22	0.0468 (13)	0.0227 (11)	0.0235 (10)	-0.0087 (10)	0.0064 (10)	-0.0005 (8)

Geometric parameters (\AA , ^\circ)

O1—C2	1.232 (3)	O11—H111	0.826
C2—N3	1.362 (4)	C12—O13	1.240 (3)
C2—C15	1.510 (4)	C12—N14	1.343 (4)
N3—C4	1.461 (3)	N14—C15	1.462 (3)
N3—C10	1.461 (4)	N14—C21	1.464 (3)
C4—C5	1.555 (4)	C15—C16	1.556 (3)
C4—C12	1.513 (4)	C15—H151	1.002
C4—H41	0.972	C16—O17	1.416 (3)
C5—O6	1.420 (3)	C16—C18	1.521 (4)

C5—C7	1.530 (4)	C16—H161	0.998
C5—H51	0.988	O17—H171	0.804
O6—H61	0.829	C18—O19	1.440 (3)
C7—O8	1.437 (3)	C18—C20	1.521 (4)
C7—C9	1.513 (4)	C18—H181	1.008
C7—H71	0.983	O19—H191	0.819
O8—H81	0.830	C20—C21	1.525 (4)
C9—C10	1.523 (4)	C20—O22	1.419 (3)
C9—O11	1.422 (3)	C20—H201	1.008
C9—H91	1.000	C21—H212	0.984
C10—H102	0.958	C21—H211	0.996
C10—H101	0.957	O22—H221	0.832
O1—C2—N3	122.3 (3)	C4—C12—O13	119.8 (3)
O1—C2—C15	120.5 (2)	C4—C12—N14	118.0 (2)
N3—C2—C15	117.1 (2)	O13—C12—N14	122.2 (3)
C2—N3—C4	122.0 (2)	C12—N14—C15	122.7 (2)
C2—N3—C10	119.1 (2)	C12—N14—C21	121.4 (2)
C4—N3—C10	112.9 (2)	C15—N14—C21	113.2 (2)
N3—C4—C5	107.4 (2)	C2—C15—N14	114.8 (2)
N3—C4—C12	113.0 (2)	C2—C15—C16	112.3 (2)
C5—C4—C12	112.8 (2)	N14—C15—C16	108.0 (2)
N3—C4—H41	107.4	C2—C15—H151	107.3
C5—C4—H41	109.1	N14—C15—H151	108.0
C12—C4—H41	106.9	C16—C15—H151	105.9
C4—C5—O6	110.9 (2)	C15—C16—O17	109.7 (2)
C4—C5—C7	112.0 (2)	C15—C16—C18	110.2 (2)
O6—C5—C7	107.6 (2)	O17—C16—C18	107.7 (2)
C4—C5—H51	106.8	C15—C16—H161	109.3
O6—C5—H51	109.8	O17—C16—H161	110.5
C7—C5—H51	109.7	C18—C16—H161	109.5
C5—O6—H61	121.6	C16—O17—H171	121.1
C5—C7—O8	108.9 (2)	C16—C18—O19	109.8 (2)
C5—C7—C9	113.3 (2)	C16—C18—C20	114.6 (2)
O8—C7—C9	106.9 (2)	O19—C18—C20	108.3 (2)
C5—C7—H71	109.6	C16—C18—H181	108.6
O8—C7—H71	108.6	O19—C18—H181	107.2
C9—C7—H71	109.5	C20—C18—H181	108.0
C7—O8—H81	114.0	C18—O19—H191	113.2
C7—C9—C10	110.2 (2)	C18—C20—C21	109.4 (2)
C7—C9—O11	107.8 (2)	C18—C20—O22	107.0 (2)
C10—C9—O11	112.5 (2)	C21—C20—O22	111.5 (2)
C7—C9—H91	109.0	C18—C20—H201	108.8
C10—C9—H91	106.9	C21—C20—H201	108.8
O11—C9—H91	110.3	O22—C20—H201	111.2
C9—C10—N3	107.2 (2)	C20—C21—N14	108.9 (2)
C9—C10—H102	111.1	C20—C21—H212	109.8
N3—C10—H102	108.5	N14—C21—H212	110.0

C9—C10—H101	109.1	C20—C21—H211	107.9
N3—C10—H101	110.5	N14—C21—H211	109.3
H102—C10—H101	110.3	H212—C21—H211	110.9
C9—O11—H111	104.2	C20—O22—H221	118.2

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C4—H41···O11 ⁱ	0.97	2.48	3.455 (4)	176
C15—H151···O6 ⁱⁱ	1.00	2.39	3.337 (4)	157
O8—H81···O17 ⁱⁱⁱ	0.83	1.95	2.756 (4)	162
O22—H221···O1 ^{iv}	0.83	2.22	2.917 (4)	141
O19—H191···O11 ^v	0.82	2.12	2.793 (4)	139
O11—H111···O13 ^{vi}	0.83	1.86	2.685 (4)	173
O17—H171···O8 ^v	0.80	1.87	2.633 (4)	157
O6—H61···O19 ^{vii}	0.83	1.97	2.680 (4)	143

Symmetry codes: (i) $-x+2, y-1/2, -z+1/2$; (ii) $x-1, y, z$; (iii) $x+1, y, z$; (iv) $x, y-1, z$; (v) $x-1/2, -y+3/2, -z+1$; (vi) $x, y+1, z$; (vii) $x+1/2, -y+1/2, -z+1$.