

Bis(2-carboxyanilinium) sulfate monohydrate

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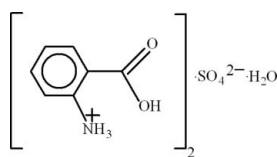
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.046; wR factor = 0.118; data-to-parameter ratio = 13.3.

In the title hydrated molecular salt, $2\text{C}_7\text{H}_8\text{NO}_2^+\cdot\text{SO}_4^{2-}\cdot\text{H}_2\text{O}$, each cation in the asymmetric unit is stabilized by an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond. The O atoms of the sulfate ion are disordered over two sets of sites with an occupancy ratio of 0.541 (13):0.459 (13), which possibly optimizes the acceptance of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds from the cations. The crystal structure also features aromatic $\pi-\pi$ stacking [centroid-centroid separation = $3.842(2)$ Å] and a $\text{C}-\text{H}\cdots\pi$ interaction.

Related literature

For background to the properties and uses of aminobenzoic acids, see: Griss *et al.* (1984); Pedanova *et al.* (1984); Refaat (2010); Rogers & Clark (1973).



Experimental

Crystal data

$2\text{C}_7\text{H}_8\text{NO}_2^+\cdot\text{SO}_4^{2-}\cdot\text{H}_2\text{O}$

$M_r = 390.36$

Monoclinic, $P2_1/n$

$a = 11.260(5)$ Å

$b = 10.542(4)$ Å

$c = 15.358(5)$ Å

$\beta = 109.737(5)^\circ$

$V = 1715.9(12)$ Å³

$Z = 4$

Mo $K\alpha$ radiation

$\mu = 0.24$ mm⁻¹

$T = 296$ K

$0.28 \times 0.25 \times 0.20$ mm

Data collection

Bruker Kappa APEXII CCD diffractometer

Absorption correction: multi-scan (SADABS; Bruker, 2005)

$T_{\min} = 0.934$, $T_{\max} = 0.955$

12530 measured reflections

3748 independent reflections
2528 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.036$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.118$

$S = 1.02$

3748 reflections

282 parameters

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.22$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.40$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$\text{Cg}2$ is the centroid of the C8–C13 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N}1-\text{H}1A\cdots\text{O}6A^i$	0.89	1.83	2.721 (6)	174
$\text{N}1-\text{H}1B\cdots\text{O}2$	0.89	1.99	2.708 (3)	137
$\text{N}1-\text{H}1B\cdots\text{O}4^{ii}$	0.89	2.33	3.041 (3)	137
$\text{N}1-\text{H}1C\cdots\text{O}8A^{iii}$	0.89	1.98	2.860 (11)	168
$\text{N}2-\text{H}2A\cdots\text{O}8A^{iv}$	0.89	1.83	2.698 (12)	166
$\text{N}2-\text{H}2B\cdots\text{O}4$	0.89	1.94	2.689 (3)	140
$\text{N}2-\text{H}2B\cdots\text{O}2^{ii}$	0.89	2.28	2.906 (3)	128
$\text{N}2-\text{H}2C\cdots\text{O}5A^i$	0.89	2.00	2.839 (6)	157
$\text{O}1-\text{H}1\cdots\text{O}9^v$	0.82	1.75	2.557 (3)	168
$\text{O}3-\text{H}3A\cdots\text{O}7A^v$	0.82	1.70	2.512 (13)	167
$\text{O}9-\text{H}9A\cdots\text{O}6A^{vi}$	0.80 (3)	2.46 (3)	3.102 (8)	138 (3)
$\text{O}9-\text{H}9A\cdots\text{O}8A^{vi}$	0.80 (3)	2.38 (4)	3.115 (11)	153 (3)
$\text{O}9-\text{H}9B\cdots\text{O}5A$	0.86 (3)	1.96 (3)	2.792 (7)	161 (3)
$\text{C}4-\text{H}4\cdots\text{Cg}2^{vii}$	0.93	2.75	3.600 (3)	153

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

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and PLATON (Spek, 2009); software used to prepare material for publication: WinGX (Farrugia, 1999) and PLATON.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5357).

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supporting information

Acta Cryst. (2010). E66, o819 [doi:10.1107/S1600536810008913]

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S1. Comment

The salts of aminobenzoic acids have been reported as medicines as well as precursor for in the field of pharmaceutics (Griss *et al.*, 1984). These are also useful for autoxidation of the fats forming free radicals, hence, in comparison to sulfated metal oxides, these can be used for production of biodiesel (Pedanova *et al.*, 1984; Refaat, 2010). The structure of title compound (I, Fig. 1) is being reported here.

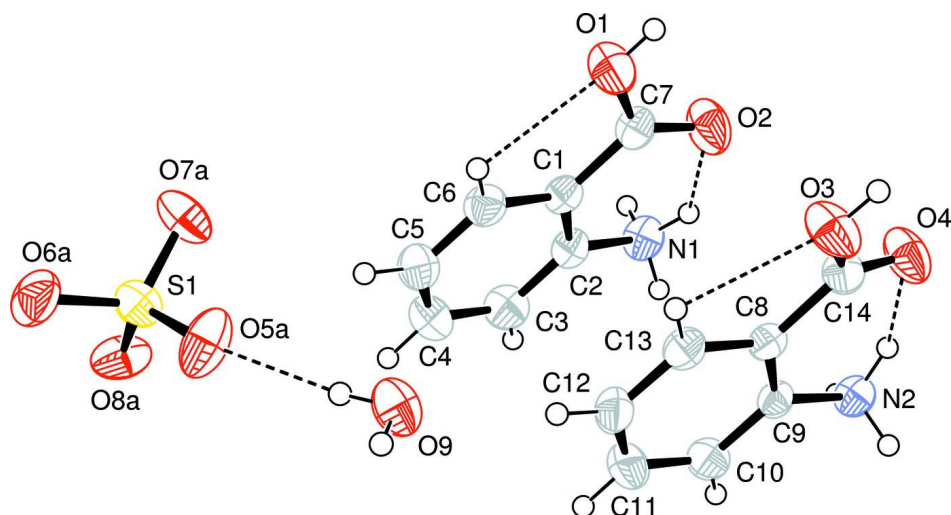
In the title compound (I) benzene rings A (C1–C6) and B (C8–C13) are of course planar with maximum r. m. s. deviations of 0.0025 and 0.0034 Å from the mean squares planes. The dihedral angle between A/B is 7.91 (13)°. The carboxylate groups C (O1/C7/O2) and D (O3/C14/O4) are oriented at dihedral angles of 11.94 (38)° and 10.86 (41)° with benzene rings A and B respectively. The O-atoms of SO₄²⁻ are disordered over two set of sites with occupancy ratio of 0.541 (13):0.459 (13). The molecules are stabilized due to intra as well as inter-molecular H-bondings and C–H⋯π interactions (Table 1, Fig. 2). The π–π interaction between the centroids CgA and CgB of benzene rings A and B respectively, also play a role in the stabilization of molecules. The centroid to centroid distance is 3.842 (2) Å.

S2. Experimental

Ethanol solution of anthranilic acid (0.02 M) was refluxed in the presence of conc. H₂SO₄ (0.01 M) for 30 min. The contents were kept at room temperature for 24 h. The crystalline material formed was washed with n-hexane, ethyl acetate and diethyl ether, respectively and dried. This material was dissolved in ethanol and colorless prisms of (I) were obtained by slow evaporation at room temperature.

S3. Refinement

Although all H-atoms were recognised from the difference fourier map but only coordinates of H-atoms of H₂O were refined. The H-atoms were positioned geometrically (C–H = 0.93, O–H = 0.82, N–H = 0.89 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C, N, O})$, where $x = 1.5$ for NH₃ and hydroxy H atoms, whereas $x = 1.2$ for all other H-atoms.

**Figure 1**

View of (I) with displacement ellipsoids drawn at the 50% probability level. The dotted lines show the hydrogen bonds.

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$M_r = 390.36$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1n$

$a = 11.260\ (5)\ \text{\AA}$

$b = 10.542\ (4)\ \text{\AA}$

$c = 15.358\ (5)\ \text{\AA}$

$\beta = 109.737\ (5)^\circ$

$V = 1715.9\ (12)\ \text{\AA}^3$

$Z = 4$

$F(000) = 816$

$D_x = 1.511\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 3752 reflections

$\theta = 2.0\text{--}27.1^\circ$

$\mu = 0.24\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Prisms, colorless

$0.28 \times 0.25 \times 0.20\ \text{mm}$

Data collection

Bruker Kappa APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube
Graphite monochromator

Detector resolution: $7.50\ \text{pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2005)

$T_{\min} = 0.934$, $T_{\max} = 0.955$

12530 measured reflections

3748 independent reflections

2528 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.036$

$\theta_{\max} = 27.1^\circ$, $\theta_{\min} = 2.0^\circ$

$h = -14 \rightarrow 14$

$k = -12 \rightarrow 13$

$l = -19 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.118$

$S = 1.02$

3748 reflections

282 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H atoms treated by a mixture of independent
and constrained refinement

$$w = 1/[\sigma^2(F_o^2) + (0.0551P)^2 + 0.3168P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$$\Delta\rho_{\max} = 0.22 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.40 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell esds are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.45161 (14)	0.45846 (15)	0.61664 (12)	0.0529 (6)	
O2	0.24999 (14)	0.41658 (17)	0.58921 (13)	0.0613 (7)	
N1	0.15921 (15)	0.19068 (17)	0.51077 (12)	0.0408 (6)	
C1	0.36494 (17)	0.28683 (19)	0.51932 (13)	0.0326 (6)	
C2	0.27346 (18)	0.19261 (19)	0.48532 (14)	0.0346 (6)	
C3	0.2899 (2)	0.0970 (2)	0.43001 (16)	0.0459 (8)	
C4	0.3988 (2)	0.0928 (2)	0.40722 (16)	0.0518 (8)	
C5	0.4909 (2)	0.1840 (2)	0.44025 (15)	0.0478 (8)	
C6	0.47351 (18)	0.2800 (2)	0.49500 (14)	0.0404 (7)	
C7	0.34884 (18)	0.3926 (2)	0.57799 (14)	0.0357 (6)	
O3	0.20126 (14)	0.69772 (16)	0.43653 (12)	0.0569 (6)	
O4	0.01902 (14)	0.62640 (15)	0.44221 (12)	0.0557 (6)	
N2	-0.09794 (14)	0.43896 (16)	0.32614 (12)	0.0364 (6)	
C8	0.12219 (17)	0.51165 (19)	0.35454 (13)	0.0307 (6)	
C9	0.02378 (18)	0.42708 (18)	0.31295 (13)	0.0308 (6)	
C10	0.0387 (2)	0.3301 (2)	0.25830 (15)	0.0412 (7)	
C11	0.1524 (2)	0.3134 (2)	0.24375 (16)	0.0488 (8)	
C12	0.2513 (2)	0.3945 (2)	0.28460 (15)	0.0442 (7)	
C13	0.23563 (18)	0.4927 (2)	0.33872 (14)	0.0376 (6)	
C14	0.10804 (19)	0.6174 (2)	0.41481 (14)	0.0367 (7)	
S1	0.82551 (5)	0.10652 (5)	0.37522 (4)	0.0404 (2)	
O5A	0.7633 (7)	0.2145 (4)	0.3274 (4)	0.0608 (16)	0.541 (13)
O6A	0.9595 (4)	0.1055 (9)	0.3666 (3)	0.0576 (19)	0.541 (13)
O7A	0.8109 (9)	0.1137 (12)	0.4669 (9)	0.053 (2)	0.541 (13)
O8A	0.7732 (9)	-0.0123 (10)	0.3402 (8)	0.051 (2)	0.541 (13)
O5B	0.6994 (8)	0.1600 (10)	0.3072 (4)	0.078 (3)	0.459 (13)
O6B	0.9227 (7)	0.1737 (8)	0.3650 (3)	0.0500 (19)	0.459 (13)
O7B	0.8590 (9)	0.1102 (15)	0.4772 (10)	0.049 (2)	0.459 (13)
O8B	0.8163 (12)	-0.0309 (13)	0.3482 (10)	0.057 (3)	0.459 (13)
O9	0.55554 (17)	0.37630 (18)	0.26338 (13)	0.0535 (7)	
H1	0.43834	0.51227	0.65081	0.0793*	
H1A	0.09356	0.16783	0.46196	0.0612*	
H1B	0.14569	0.26764	0.52934	0.0612*	

H1C	0.16899	0.13539	0.55646	0.0612*
H3	0.22814	0.03516	0.40785	0.0551*
H4	0.40991	0.02812	0.36949	0.0622*
H5	0.56438	0.18049	0.42545	0.0573*
H6	0.53543	0.34189	0.51640	0.0484*
H2A	-0.15481	0.46829	0.27447	0.0546*
H2B	-0.09109	0.49252	0.37237	0.0546*
H2C	-0.12228	0.36332	0.33962	0.0546*
H3A	0.18855	0.75376	0.46936	0.0854*
H10	-0.02800	0.27507	0.23078	0.0495*
H11	0.16195	0.24753	0.20641	0.0586*
H12	0.32830	0.38297	0.27570	0.0531*
H13	0.30250	0.54790	0.36542	0.0451*
H9A	0.576 (3)	0.407 (3)	0.2229 (19)	0.0642*
H9B	0.607 (2)	0.313 (3)	0.2786 (18)	0.0642*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0400 (8)	0.0520 (11)	0.0720 (12)	-0.0148 (8)	0.0259 (8)	-0.0236 (9)
O2	0.0370 (9)	0.0661 (12)	0.0872 (13)	-0.0092 (8)	0.0294 (9)	-0.0292 (10)
N1	0.0334 (9)	0.0396 (11)	0.0503 (11)	-0.0066 (8)	0.0153 (8)	0.0028 (9)
C1	0.0296 (10)	0.0350 (12)	0.0325 (10)	-0.0007 (9)	0.0097 (8)	0.0052 (9)
C2	0.0318 (10)	0.0353 (12)	0.0369 (11)	0.0001 (9)	0.0117 (9)	0.0064 (10)
C3	0.0463 (12)	0.0388 (13)	0.0513 (14)	-0.0073 (11)	0.0147 (11)	-0.0023 (11)
C4	0.0571 (15)	0.0498 (15)	0.0506 (14)	0.0039 (12)	0.0209 (12)	-0.0097 (12)
C5	0.0390 (12)	0.0606 (16)	0.0478 (13)	0.0031 (12)	0.0200 (10)	-0.0015 (12)
C6	0.0316 (10)	0.0486 (14)	0.0401 (11)	-0.0039 (10)	0.0111 (9)	-0.0012 (11)
C7	0.0306 (10)	0.0365 (12)	0.0403 (11)	-0.0015 (9)	0.0125 (9)	0.0048 (10)
O3	0.0507 (9)	0.0519 (10)	0.0768 (13)	-0.0231 (8)	0.0328 (9)	-0.0311 (9)
O4	0.0425 (9)	0.0538 (10)	0.0792 (12)	-0.0122 (8)	0.0314 (9)	-0.0287 (9)
N2	0.0299 (9)	0.0365 (10)	0.0410 (10)	-0.0058 (8)	0.0095 (8)	-0.0005 (8)
C8	0.0290 (9)	0.0297 (11)	0.0302 (10)	-0.0005 (9)	0.0059 (8)	0.0015 (9)
C9	0.0304 (10)	0.0279 (11)	0.0315 (10)	0.0023 (9)	0.0070 (8)	0.0038 (9)
C10	0.0428 (12)	0.0328 (12)	0.0439 (12)	-0.0038 (10)	0.0091 (10)	-0.0067 (10)
C11	0.0585 (15)	0.0395 (13)	0.0495 (14)	0.0089 (12)	0.0196 (12)	-0.0070 (11)
C12	0.0386 (12)	0.0509 (14)	0.0453 (12)	0.0107 (11)	0.0169 (10)	0.0033 (12)
C13	0.0302 (10)	0.0409 (12)	0.0389 (11)	-0.0025 (9)	0.0082 (9)	0.0007 (10)
C14	0.0322 (10)	0.0341 (12)	0.0425 (12)	-0.0052 (9)	0.0111 (9)	-0.0047 (10)
S1	0.0493 (3)	0.0288 (3)	0.0455 (3)	-0.0044 (3)	0.0192 (3)	-0.0041 (3)
O5A	0.045 (3)	0.033 (2)	0.096 (3)	0.004 (2)	0.013 (2)	0.020 (2)
O6A	0.0306 (19)	0.080 (5)	0.061 (2)	-0.002 (2)	0.0141 (16)	0.007 (2)
O7A	0.075 (5)	0.039 (3)	0.058 (4)	-0.011 (5)	0.041 (5)	-0.014 (3)
O8A	0.065 (4)	0.026 (4)	0.048 (2)	-0.013 (3)	-0.001 (3)	-0.006 (2)
O5B	0.046 (4)	0.084 (5)	0.093 (4)	0.018 (4)	0.009 (3)	0.019 (3)
O6B	0.045 (3)	0.046 (4)	0.064 (3)	-0.015 (3)	0.025 (2)	0.005 (2)
O7B	0.061 (5)	0.043 (3)	0.047 (3)	-0.006 (5)	0.025 (5)	-0.009 (2)
O8B	0.099 (8)	0.028 (3)	0.057 (6)	-0.003 (5)	0.045 (6)	-0.004 (3)

O9	0.0570 (11)	0.0506 (11)	0.0635 (12)	-0.0020 (8)	0.0343 (9)	-0.0038 (9)
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Geometric parameters (Å, °)

S1—O7A	1.474 (13)	N2—H2B	0.8900
S1—O8A	1.411 (11)	C1—C2	1.399 (3)
S1—O5B	1.557 (8)	C1—C7	1.483 (3)
S1—O6B	1.356 (8)	C1—C6	1.396 (3)
S1—O7B	1.483 (14)	C2—C3	1.370 (3)
S1—O5A	1.407 (5)	C3—C4	1.385 (3)
S1—O6A	1.559 (5)	C4—C5	1.379 (3)
S1—O8B	1.501 (14)	C5—C6	1.371 (3)
O1—C7	1.308 (3)	C3—H3	0.9300
O2—C7	1.209 (3)	C4—H4	0.9300
O1—H1	0.8200	C5—H5	0.9300
O3—C14	1.301 (3)	C6—H6	0.9300
O4—C14	1.215 (3)	C8—C13	1.393 (3)
O3—H3A	0.8200	C8—C14	1.492 (3)
O9—H9A	0.80 (3)	C8—C9	1.397 (3)
O9—H9B	0.86 (3)	C9—C10	1.369 (3)
N1—C2	1.466 (3)	C10—C11	1.383 (3)
N1—H1A	0.8900	C11—C12	1.376 (3)
N1—H1C	0.8900	C12—C13	1.376 (3)
N1—H1B	0.8900	C10—H10	0.9300
N2—C9	1.457 (3)	C11—H11	0.9300
N2—H2A	0.8900	C12—H12	0.9300
N2—H2C	0.8900	C13—H13	0.9300
O5B—S1—O7B	123.3 (6)	C4—C5—C6	119.7 (2)
O5B—S1—O8B	101.6 (7)	C1—C6—C5	121.4 (2)
O6B—S1—O7B	100.4 (5)	O1—C7—C1	113.84 (18)
O6B—S1—O8B	117.1 (6)	O1—C7—O2	122.7 (2)
O7B—S1—O8B	106.7 (8)	O2—C7—C1	123.5 (2)
O5A—S1—O8A	116.7 (5)	C4—C3—H3	120.00
O6A—S1—O7A	120.3 (5)	C2—C3—H3	120.00
O6A—S1—O8A	104.7 (6)	C3—C4—H4	120.00
O7A—S1—O8A	104.1 (7)	C5—C4—H4	120.00
O5B—S1—O6B	108.8 (5)	C4—C5—H5	120.00
O5A—S1—O6A	106.7 (4)	C6—C5—H5	120.00
O5A—S1—O7A	105.1 (6)	C5—C6—H6	119.00
C7—O1—H1	109.00	C1—C6—H6	119.00
C14—O3—H3A	109.00	C9—C8—C13	117.46 (18)
H9A—O9—H9B	100 (3)	C9—C8—C14	121.75 (18)
C2—N1—H1B	109.00	C13—C8—C14	120.79 (18)
H1B—N1—H1C	109.00	N2—C9—C10	117.75 (18)
H1A—N1—H1B	109.00	N2—C9—C8	121.31 (17)
C2—N1—H1A	109.00	C8—C9—C10	121.0 (2)
H1A—N1—H1C	109.00	C9—C10—C11	120.4 (2)

C2—N1—H1C	109.00	C10—C11—C12	119.9 (2)
C9—N2—H2C	109.00	C11—C12—C13	119.6 (2)
C9—N2—H2B	109.00	C8—C13—C12	121.7 (2)
C9—N2—H2A	109.00	O4—C14—C8	123.2 (2)
H2A—N2—H2C	109.00	O3—C14—C8	113.54 (19)
H2A—N2—H2B	109.00	O3—C14—O4	123.3 (2)
H2B—N2—H2C	109.00	C11—C10—H10	120.00
C2—C1—C7	122.35 (18)	C9—C10—H10	120.00
C2—C1—C6	117.69 (18)	C10—C11—H11	120.00
C6—C1—C7	119.96 (18)	C12—C11—H11	120.00
N1—C2—C3	118.01 (19)	C13—C12—H12	120.00
N1—C2—C1	120.82 (18)	C11—C12—H12	120.00
C1—C2—C3	121.1 (2)	C12—C13—H13	119.00
C2—C3—C4	119.8 (2)	C8—C13—H13	119.00
C3—C4—C5	120.3 (2)		
C6—C1—C2—N1	177.63 (18)	C13—C8—C9—N2	179.56 (18)
C6—C1—C2—C3	0.0 (3)	C13—C8—C9—C10	-0.5 (3)
C7—C1—C2—N1	-3.4 (3)	C14—C8—C9—N2	0.4 (3)
C7—C1—C2—C3	179.0 (2)	C14—C8—C9—C10	-179.60 (19)
C2—C1—C6—C5	-0.5 (3)	C9—C8—C13—C12	-0.2 (3)
C7—C1—C6—C5	-179.48 (19)	C14—C8—C13—C12	178.92 (19)
C2—C1—C7—O1	168.59 (19)	C9—C8—C14—O3	-170.35 (18)
C2—C1—C7—O2	-11.6 (3)	C9—C8—C14—O4	10.8 (3)
C6—C1—C7—O1	-12.4 (3)	C13—C8—C14—O3	10.5 (3)
C6—C1—C7—O2	167.4 (2)	C13—C8—C14—O4	-168.3 (2)
N1—C2—C3—C4	-177.61 (19)	N2—C9—C10—C11	-179.52 (19)
C1—C2—C3—C4	0.1 (3)	C8—C9—C10—C11	0.5 (3)
C2—C3—C4—C5	0.3 (3)	C9—C10—C11—C12	0.2 (3)
C3—C4—C5—C6	-0.8 (3)	C10—C11—C12—C13	-0.8 (3)
C4—C5—C6—C1	0.9 (3)	C11—C12—C13—C8	0.9 (3)

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

Cg2 is the centroid of of the C8—C13 ring.

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1A \cdots O6A ⁱ	0.89	1.83	2.721 (6)	174
N1—H1B \cdots O2	0.89	1.99	2.708 (3)	137
N1—H1B \cdots O4 ⁱⁱ	0.89	2.33	3.041 (3)	137
N1—H1C \cdots O8A ⁱⁱⁱ	0.89	1.98	2.860 (11)	168
N2—H2A \cdots O8A ^{iv}	0.89	1.83	2.698 (12)	166
N2—H2B \cdots O4	0.89	1.94	2.689 (3)	140
N2—H2B \cdots O2 ⁱⁱ	0.89	2.28	2.906 (3)	128
N2—H2C \cdots O5A ⁱ	0.89	2.00	2.839 (6)	157
O1—H1 \cdots O9 ^v	0.82	1.75	2.557 (3)	168
O3—H3A \cdots O7A ^v	0.82	1.70	2.512 (13)	167
O9—H9A \cdots O6A ^{vi}	0.80 (3)	2.46 (3)	3.102 (8)	138 (3)
O9—H9A \cdots O8A ^{vi}	0.80 (3)	2.38 (4)	3.115 (11)	153 (3)

O9—H9B···O5A	0.86 (3)	1.96 (3)	2.792 (7)	161 (3)
C4—H4···Cg2 ^{viii}	0.93	2.75	3.600 (3)	153

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