organic compounds

4-(4-Diethylamino-2-hydroxybenzylideneammonio)-3-methylbenzenesulfonate dihydrate

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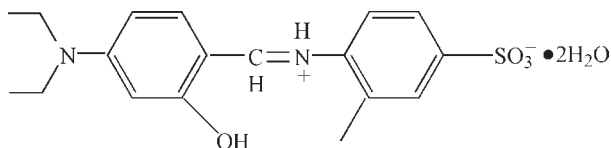Correspondence e-mail: chenyt@qhnu.edu.cn

In the crystal of the title compound, $C_{18}H_{22}N_2O_4S\cdot2H_2O$, molecules are linked into a one-dimensional chain structure by C—H···O, N—H···O, O—H···O and O—H···N hydrogen bonds.

Related literature

For the biological and pharmacological activities of Schiff base compounds, see: Bu *et al.* (2001); Ranford *et al.* (1998).

Experimental

Crystal data

$C_{18}H_{22}N_2O_4S\cdot2H_2O$	V = 3806.9 (9) Å^3
M_r = 398.47	Z = 8
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
a = 21.690 (3) Å	μ = 0.21 mm^{-1}
b = 11.4142 (17) Å	T = 273 K
c = 16.639 (2) Å	$0.19 \times 0.16 \times 0.06$ mm
β = 112.459 (2)°	

Data collection

Bruker SMART CCD area-detector diffractometer	9821 measured reflections
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)	3379 independent reflections
	2910 reflections with $I > 2\sigma(I)$
T_{min} = 0.962, T_{max} = 0.988	R_{int} = 0.026

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.054	246 parameters
$wR(F^2)$ = 0.171	H-atom parameters constrained
S = 1.05	$\Delta\rho_{max}$ = 0.73 e Å^{-3}
3379 reflections	$\Delta\rho_{min}$ = −0.41 e Å^{-3}

Table 1
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O6	0.86	2.10	2.723 (3)	129
N1—H1···O2^i	0.86	2.58	3.160 (3)	125
O6—H6···O2^{ii}	0.82	1.91	2.709 (3)	166
O4—H16···O3^i	0.85	2.00	2.828 (5)	166
O5—H18···N1^{iii}	0.85	2.51	3.35 (3)	174
C2—H2···O3	0.93	2.56	2.915 (4)	103
C5—H5···O1^{iv}	0.93	2.55	3.392 (3)	150
C7—H7B···O2^i	0.96	2.44	3.325 (3)	154
C8—H8···O1^{iv}	0.93	2.43	3.353 (3)	172
C15—H15B···O3^{ii}	0.97	2.50	3.444 (4)	166
C16—H16C···O5^v	0.96	2.51	3.42 (3)	160

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

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: AT2871).

References

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Bu, X. H., Gao, Y. X., Chen, W. & Zhang, R. H. (2001). *J. Rare Earth*, **19**, 70–75.
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supporting information

4-(4-Diethylamino-2-hydroxybenzylideneammonio)-3-methylbenzenesulfonate dihydrate

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S1. Comment

Schiff bases play an important role in the field of bioinorganic chemistry because they have remarkable wide biological and pharmacological activities, such as antitumor, antidiabetic, antitubercular activities [Ranford, *et al.*, 1998; Bu, *et al.*, 2001]. Therefore, investigating the synthesis and proper ties of hydrazone of these compounds seems to be a very interesting problem. as one part of our systematic work, In this paper, we report on the synthesis and crystal structure of the title compound, (I), (Fig. 1).

The dihedral angle between the aromatic ring planes (C1–C6) and (C9–C14) is 5.70 (12)°, showing that the whole compound is not a plane molecule. The bond distances of C8—N1 [1.311 (3) Å], S1—O2 [1.449 (2) Å] and S1—O3 [1.426 (3) Å] are consistent with the carbon-nitrogen and sulfur-oxygen double-bond lengths, respectively. In the crystal packing, the molecules form a one-dimensional chain structure by C—H⋯O, N—H⋯O, O—H⋯O and O—H⋯N hydrogen bonds (Table 1).

S2. Experimental

The solution of 1.0 mmol 4-(diethylamino)salicylaldehyde was added to a solution of 1.0 mmol 3-methyl-benzene-sulfonic acid in 5 ml ethanol at room temperature. The mixture was refluxed for 4 h with stirring, then the resulting precipitate was filtered, washed, and dried *in vacuo* over P_4O_{10} for 48 h. Single crystals suitable for X-ray structural analysis was obtained by slowly evaporating from methanol at room temperature.

S3. Refinement

All H atoms were included in the refinement at calculated positions, in the riding-model approximation, with C–H distances of 0.93 (ArH), 0.98 (CH_3), 0.97Å (CH_2), N–H = 0.86 Å (NH), O–H = 0.82 Å (OH) and 0.85 Å (H_2O) . The isotropic displacement parameters for all H atoms were set equal to 1.2 or $1.5U_{eq}$ of the carrier atom.
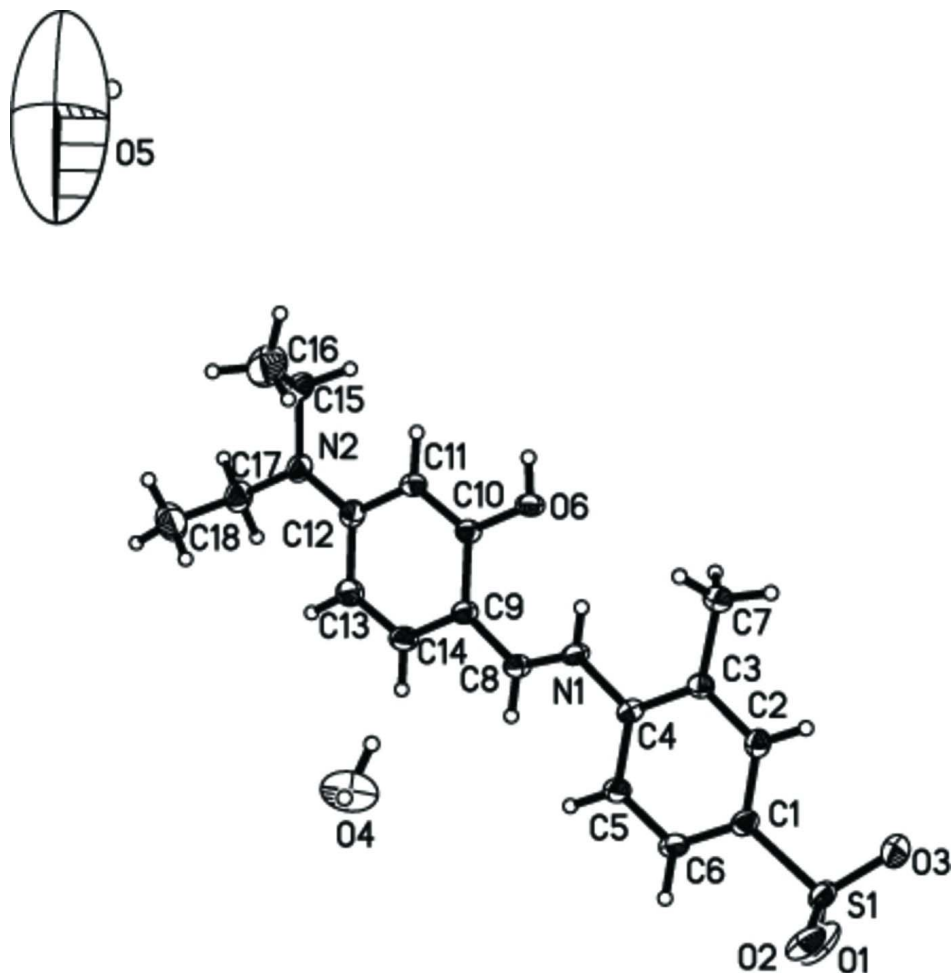
Figure 1

The molecular structure of (I) showing 30% displacement ellipsoids.

4-(4-Diethylamino-2-hydroxybenzylideneammonio)-3-methylbenzenesulfonate dihydrate

Crystal data

$C_{18}H_{22}N_2O_4S\cdot 2H_2O$	$F(000) = 1696$
$M_r = 398.47$	$D_x = 1.390$ Mg m^{-3}
Monoclinic, $C2/c$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: -C 2yc	Cell parameters from 4718 reflections
$a = 21.690$ (3) Å	$\theta = 2.2–28.4°$
$b = 11.4142$ (17) Å	$\mu = 0.21$ mm^{-1}
$c = 16.639$ (2) Å	$T = 273$ K
$\beta = 112.459$ (2)°	Block, colourless
$V = 3806.9$ (9) Å^3	$0.19 \times 0.16 \times 0.06$ mm
$Z = 8$	

Data collection

Bruker SMART CCD area-detector diffractometer	Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
Radiation source: fine-focus sealed tube	$T_{min} = 0.962$, $T_{max} = 0.988$
Graphite monochromator	9821 measured reflections
φ and ω scans	3379 independent reflections

2910 reflections with $I > 2\sigma(I)$	$h = -11 \rightarrow 25$	
$R_{int} = 0.026$	$k = -13 \rightarrow 13$	
$\theta_{max} = 25.1°$, $\theta_{min} = 2.0°$	$l = -19 \rightarrow 19$	

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	
$R[F^2 > 2\sigma(F^2)] = 0.054$	H-atom parameters constrained
$wR(F^2) = 0.171$	$w = 1/[\sigma^2(F_o^2) + (0.1015P)^2 + 5.082P]$ where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.05$	
3379 reflections	$(\Delta/\sigma)_{max} < 0.001$
246 parameters	$\Delta\rho_{max} = 0.73$ e Å⁻³
0 restraints	$\Delta\rho_{min} = -0.41$ e Å⁻³
Primary atom site location: structure-invariant direct methods	Extinction correction: *SHELXL97* (Sheldrick, 2008), Fc*=kFc[1+0.001xFc²λ³/sin(2θ)]⁻¹ᐟ⁴
Secondary atom site location: difference Fourier map	Extinction coefficient: 0.0019 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

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 (Å²)

	x	y	z	U_{iso}*/U_{eq}
S1	0.18156 (3)	0.09473 (5)	−0.21298 (4)	0.0405 (3)
O1	0.17722 (12)	0.1276 (2)	−0.29916 (14)	0.0750 (8)
O2	0.11837 (9)	0.1127 (2)	−0.20400 (13)	0.0573 (6)
O3	0.20786 (13)	−0.0203 (2)	−0.1892 (2)	0.0980 (11)
O4	0.18994 (19)	0.6795 (4)	0.0958 (2)	0.1365 (14)
H16	0.2252	0.6420	0.1254	0.205*
H15	0.1630	0.6669	0.1209	0.205*
O5	0.276 (2)	0.6202 (16)	1.001 (2)	0.81 (3)
H18	0.3031	0.5664	1.0016	1.213*
H17	0.2582	0.6426	0.9487	1.213*
O6	0.50747 (8)	0.45092 (15)	0.15864 (11)	0.0392 (4)
H6	0.5392	0.4405	0.2047	0.059*
N1	0.39104 (9)	0.41470 (17)	0.02004 (12)	0.0332 (5)
H1	0.4154	0.3891	0.0709	0.040*
N2	0.60480 (10)	0.83477 (18)	0.19161 (14)	0.0404 (5)
C1	0.24039 (11)	0.1925 (2)	−0.13942 (15)	0.0343 (5)
C2	0.28796 (12)	0.1511 (2)	−0.06245 (15)	0.0345 (5)
H2	0.2863	0.0733	−0.0468	0.041*
C3	0.33840 (11)	0.2239 (2)	−0.00783 (14)	0.0306 (5)
C4	0.33861 (11)	0.3409 (2)	−0.03262 (14)	0.0303 (5)

C5	0.28903 (12)	0.3837 (2)	−0.10848 (16)	0.0387 (6)
H5	0.2886	0.4624	−0.1230	0.046*
C6	0.24063 (12)	0.3090 (2)	−0.16183 (16)	0.0412 (6)
H6A	0.2081	0.3371	−0.2129	0.049*
C7	0.39164 (13)	0.1746 (2)	0.07302 (16)	0.0442 (6)
H7A	0.3837	0.0925	0.0774	0.066*
H7B	0.3905	0.2145	0.1232	0.066*
H7C	0.4346	0.1853	0.0700	0.066*
C8	0.40592 (11)	0.5183 (2)	−0.00150 (15)	0.0338 (5)
H8	0.3790	0.5459	−0.0563	0.041*
C9	0.45786 (11)	0.59191 (19)	0.04893 (15)	0.0322 (5)
C10	0.50910 (11)	0.5615 (2)	0.13088 (14)	0.0304 (5)
C11	0.55669 (11)	0.6412 (2)	0.17671 (15)	0.0343 (5)
H11	0.5891	0.6194	0.2299	0.041*
C12	0.55782 (11)	0.7562 (2)	0.14517 (16)	0.0355 (5)
C13	0.50789 (13)	0.7859 (2)	0.06268 (17)	0.0417 (6)
H13	0.5073	0.8605	0.0400	0.050*
C14	0.46131 (13)	0.7060 (2)	0.01705 (16)	0.0412 (6)
H14	0.4303	0.7270	−0.0374	0.049*
C15	0.65369 (14)	0.8065 (3)	0.27868 (18)	0.0514 (7)
H15A	0.6912	0.8598	0.2929	0.062*
H15B	0.6702	0.7277	0.2782	0.062*
C16	0.6252 (3)	0.8148 (4)	0.3481 (2)	0.0935 (13)
H16A	0.5824	0.7771	0.3282	0.140*
H16B	0.6203	0.8957	0.3603	0.140*
H16C	0.6547	0.7768	0.4000	0.140*
C17	0.60749 (14)	0.9541 (2)	0.1600 (2)	0.0490 (7)
H17A	0.5944	0.9516	0.0973	0.059*
H17B	0.6532	0.9818	0.1850	0.059*
C18	0.56346 (19)	1.0404 (3)	0.1815 (3)	0.0720 (10)
H18A	0.5791	1.0499	0.2434	0.108*
H18B	0.5185	1.0116	0.1598	0.108*
H18C	0.5648	1.1145	0.1549	0.108*

Atomic displacement parameters (Å^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0310 (4)	0.0404 (4)	0.0444 (4)	−0.0065 (2)	0.0080 (3)	−0.0132 (3)
O1	0.0836 (16)	0.1016 (18)	0.0462 (12)	−0.0473 (14)	0.0318 (12)	−0.0338 (12)
O2	0.0341 (10)	0.0851 (15)	0.0515 (11)	−0.0193 (9)	0.0150 (8)	−0.0287 (10)
O3	0.0664 (15)	0.0437 (13)	0.128 (2)	0.0007 (11)	−0.0253 (15)	−0.0222 (14)
O4	0.102 (3)	0.148 (3)	0.120 (3)	0.026 (2)	−0.002 (2)	0.010 (2)
O5	1.22 (7)	0.36 (2)	1.12 (5)	−0.11 (3)	0.75 (6)	−0.22 (3)
O6	0.0317 (9)	0.0359 (9)	0.0378 (9)	−0.0044 (7)	−0.0004 (7)	0.0086 (7)
N1	0.0283 (10)	0.0372 (11)	0.0262 (10)	−0.0048 (8)	0.0018 (8)	0.0015 (8)
N2	0.0344 (11)	0.0346 (11)	0.0480 (12)	−0.0076 (9)	0.0112 (9)	−0.0070 (9)
C1	0.0255 (11)	0.0394 (13)	0.0366 (12)	−0.0036 (9)	0.0101 (10)	−0.0069 (10)
C2	0.0353 (12)	0.0305 (12)	0.0386 (13)	−0.0035 (9)	0.0150 (10)	−0.0023 (9)

C3	0.0296 (11)	0.0340 (12)	0.0279 (11)	0.0010 (9)	0.0106 (9)	0.0014 (9)
C4	0.0271 (11)	0.0359 (12)	0.0266 (11)	−0.0046 (9)	0.0087 (9)	−0.0020 (9)
C5	0.0355 (13)	0.0324 (12)	0.0380 (13)	−0.0030 (10)	0.0027 (10)	0.0054 (10)
C6	0.0332 (13)	0.0428 (14)	0.0363 (13)	−0.0028 (10)	0.0005 (10)	0.0015 (10)
C7	0.0459 (15)	0.0393 (14)	0.0376 (13)	0.0019 (11)	0.0049 (11)	0.0055 (10)
C8	0.0305 (12)	0.0382 (13)	0.0284 (11)	−0.0007 (9)	0.0066 (9)	0.0026 (9)
C9	0.0278 (12)	0.0349 (12)	0.0304 (12)	−0.0024 (9)	0.0073 (9)	0.0012 (9)
C10	0.0269 (11)	0.0323 (11)	0.0304 (11)	−0.0003 (9)	0.0091 (9)	0.0022 (9)
C11	0.0279 (11)	0.0361 (12)	0.0332 (12)	0.0001 (9)	0.0053 (9)	0.0001 (9)
C12	0.0309 (12)	0.0356 (13)	0.0417 (13)	−0.0022 (10)	0.0157 (10)	−0.0053 (10)
C13	0.0430 (14)	0.0343 (13)	0.0427 (14)	−0.0049 (10)	0.0105 (11)	0.0065 (10)
C14	0.0394 (14)	0.0411 (14)	0.0354 (13)	−0.0014 (10)	0.0056 (11)	0.0078 (10)
C15	0.0443 (16)	0.0477 (16)	0.0514 (16)	−0.0131 (12)	0.0061 (13)	−0.0089 (12)
C16	0.137 (4)	0.091 (3)	0.053 (2)	0.004 (3)	0.038 (2)	−0.004 (2)
C17	0.0440 (15)	0.0393 (14)	0.0621 (17)	−0.0096 (12)	0.0185 (13)	−0.0068 (13)
C18	0.072 (2)	0.0533 (19)	0.089 (3)	0.0109 (17)	0.0290 (19)	−0.0022 (17)

Geometric parameters (Å, º)

S1—O3	1.426 (2)	C7—H7A	0.9600
S1—O2	1.449 (2)	C7—H7B	0.9600
S1—O1	1.450 (2)	C7—H7C	0.9600
S1—C1	1.783 (2)	C8—C9	1.399 (3)
O4—H16	0.8502	C8—H8	0.9300
O4—H15	0.8505	C9—C14	1.419 (3)
O5—H18	0.8491	C9—C10	1.434 (3)
O5—H17	0.8505	C10—C11	1.369 (3)
O6—C10	1.349 (3)	C11—C12	1.417 (3)
O6—H6	0.8200	C11—H11	0.9300
N1—C8	1.311 (3)	C12—C13	1.427 (3)
N1—C4	1.417 (3)	C13—C14	1.357 (3)
N1—H1	0.8600	C13—H13	0.9300
N2—C12	1.356 (3)	C14—H14	0.9300
N2—C15	1.467 (3)	C15—C16	1.507 (5)
N2—C17	1.470 (3)	C15—H15A	0.9700
C1—C6	1.382 (4)	C15—H15B	0.9700
C1—C2	1.385 (3)	C16—H16A	0.9600
C2—C3	1.398 (3)	C16—H16B	0.9600
C2—H2	0.9300	C16—H16C	0.9600
C3—C4	1.399 (3)	C17—C18	1.506 (4)
C3—C7	1.508 (3)	C17—H17A	0.9700
C4—C5	1.397 (3)	C17—H17B	0.9700
C5—C6	1.380 (3)	C18—H18A	0.9600
C5—H5	0.9300	C18—H18B	0.9600
C6—H6A	0.9300	C18—H18C	0.9600
O3—S1—O2	113.29 (17)	C8—C9—C14	118.2 (2)
O3—S1—O1	112.40 (19)	C8—C9—C10	125.4 (2)

O2—S1—O1	111.02 (14)	C14—C9—C10	116.4 (2)
O3—S1—C1	106.26 (12)	O6—C10—C11	123.0 (2)
O2—S1—C1	107.32 (11)	O6—C10—C9	116.08 (19)
O1—S1—C1	106.02 (12)	C11—C10—C9	121.0 (2)
H16—O4—H15	105.3	C10—C11—C12	121.7 (2)
H18—O5—H17	105.4	C10—C11—H11	119.1
C10—O6—H6	109.5	C12—C11—H11	119.1
C8—N1—C4	126.16 (19)	N2—C12—C11	121.1 (2)
C8—N1—H1	116.9	N2—C12—C13	121.4 (2)
C4—N1—H1	116.9	C11—C12—C13	117.5 (2)
C12—N2—C15	121.3 (2)	C14—C13—C12	120.5 (2)
C12—N2—C17	122.2 (2)	C14—C13—H13	119.7
C15—N2—C17	116.4 (2)	C12—C13—H13	119.7
C6—C1—C2	120.0 (2)	C13—C14—C9	122.8 (2)
C6—C1—S1	119.52 (18)	C13—C14—H14	118.6
C2—C1—S1	120.39 (18)	C9—C14—H14	118.6
C1—C2—C3	121.3 (2)	N2—C15—C16	112.9 (3)
C1—C2—H2	119.3	N2—C15—H15A	109.0
C3—C2—H2	119.3	C16—C15—H15A	109.0
C2—C3—C4	117.8 (2)	N2—C15—H15B	109.0
C2—C3—C7	120.1 (2)	C16—C15—H15B	109.0
C4—C3—C7	122.1 (2)	H15A—C15—H15B	107.8
C5—C4—C3	120.8 (2)	C15—C16—H16A	109.5
C5—C4—N1	120.7 (2)	C15—C16—H16B	109.5
C3—C4—N1	118.52 (19)	H16A—C16—H16B	109.5
C6—C5—C4	119.9 (2)	C15—C16—H16C	109.5
C6—C5—H5	120.0	H16A—C16—H16C	109.5
C4—C5—H5	120.0	H16B—C16—H16C	109.5
C5—C6—C1	120.1 (2)	N2—C17—C18	113.9 (3)
C5—C6—H6A	120.0	N2—C17—H17A	108.8
C1—C6—H6A	120.0	C18—C17—H17A	108.8
C3—C7—H7A	109.5	N2—C17—H17B	108.8
C3—C7—H7B	109.5	C18—C17—H17B	108.8
H7A—C7—H7B	109.5	H17A—C17—H17B	107.7
C3—C7—H7C	109.5	C17—C18—H18A	109.5
H7A—C7—H7C	109.5	C17—C18—H18B	109.5
H7B—C7—H7C	109.5	H18A—C18—H18B	109.5
N1—C8—C9	127.1 (2)	C17—C18—H18C	109.5
N1—C8—H8	116.4	H18A—C18—H18C	109.5
C9—C8—H8	116.4	H18B—C18—H18C	109.5
O3—S1—C1—C6	159.2 (2)	N1—C8—C9—C10	7.0 (4)
O2—S1—C1—C6	−79.3 (2)	C8—C9—C10—O6	2.7 (3)
O1—S1—C1—C6	39.4 (2)	C14—C9—C10—O6	−176.7 (2)
O3—S1—C1—C2	−17.3 (3)	C8—C9—C10—C11	−177.9 (2)
O2—S1—C1—C2	104.2 (2)	C14—C9—C10—C11	2.7 (3)
O1—S1—C1—C2	−137.1 (2)	O6—C10—C11—C12	178.9 (2)
C6—C1—C2—C3	−2.6 (4)	C9—C10—C11—C12	−0.6 (4)

S1—C1—C2—C3	173.88 (17)	C15—N2—C12—C11	−3.9 (4)
C1—C2—C3—C4	1.2 (3)	C17—N2—C12—C11	179.4 (2)
C1—C2—C3—C7	−176.6 (2)	C15—N2—C12—C13	176.3 (2)
C2—C3—C4—C5	1.4 (3)	C17—N2—C12—C13	−0.4 (4)
C7—C3—C4—C5	179.2 (2)	C10—C11—C12—N2	179.3 (2)
C2—C3—C4—N1	−177.30 (19)	C10—C11—C12—C13	−1.0 (3)
C7—C3—C4—N1	0.5 (3)	N2—C12—C13—C14	179.9 (2)
C8—N1—C4—C5	−12.5 (4)	C11—C12—C13—C14	0.2 (4)
C8—N1—C4—C3	166.3 (2)	C12—C13—C14—C9	2.2 (4)
C3—C4—C5—C6	−2.7 (4)	C8—C9—C14—C13	177.0 (2)
N1—C4—C5—C6	176.0 (2)	C10—C9—C14—C13	−3.6 (4)
C4—C5—C6—C1	1.3 (4)	C12—N2—C15—C16	−78.0 (3)
C2—C1—C6—C5	1.3 (4)	C17—N2—C15—C16	98.9 (3)
S1—C1—C6—C5	−175.2 (2)	C12—N2—C17—C18	85.9 (3)
C4—N1—C8—C9	−179.1 (2)	C15—N2—C17—C18	−91.0 (3)
N1—C8—C9—C14	−173.7 (2)		

Hydrogen-bond geometry (Å, º)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O6	0.86	2.10	2.723 (3)	129
N1—H1···O2ⁱ	0.86	2.58	3.160 (3)	125
O6—H6···O2ⁱⁱ	0.82	1.91	2.709 (3)	166
O4—H16···O3ⁱ	0.85	2.00	2.828 (5)	166
O5—H18···N1ⁱⁱⁱ	0.85	2.51	3.35 (3)	174
C2—H2···O3	0.93	2.56	2.915 (4)	103
C5—H5···O1ⁱᵛ	0.93	2.55	3.392 (3)	150
C7—H7*B*···O2ⁱ	0.96	2.44	3.325 (3)	154
C8—H8···O1ⁱᵛ	0.93	2.43	3.353 (3)	172
C15—H15*B*···O3ⁱⁱ	0.97	2.50	3.444 (4)	166
C16—H16*C*···O5ᵛ	0.96	2.51	3.42 (3)	160

Symmetry codes: (i) −*x*+1/2, −*y*+1/2, −*z*; (ii) *x*+1/2, −*y*+1/2, *z*+1/2; (iii) *x*, *y*, *z*+1; (iv) −*x*+1/2, *y*+1/2, −*z*−1/2; (v) −*x*+1, *y*, −*z*+3/2.