

## Ethyl 4-(phenylsulfonyl)piperazine-1-carboxylate

Mohammad T.M. Al-Dajani,<sup>a</sup> Hassan H. Adballah,<sup>b</sup> Nornisah Mohamed,<sup>a</sup> Madhukar Hemamalini<sup>c</sup> and Hoong-Kun Fun<sup>c\*</sup>#

<sup>a</sup>School of Pharmaceutical Sciences, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, <sup>b</sup>School of Chemical Sciences, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia, and <sup>c</sup>X-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia

Correspondence e-mail: hkfun@usm.my

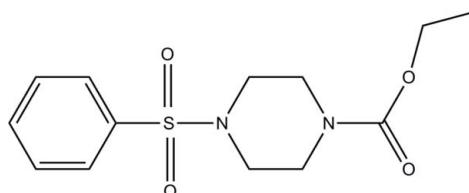
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Key indicators: single-crystal X-ray study;  $T = 296\text{ K}$ ; mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$ ;  $R$  factor = 0.048;  $wR$  factor = 0.158; data-to-parameter ratio = 23.4.

In the title compound,  $\text{C}_{13}\text{H}_{18}\text{N}_2\text{O}_4\text{S}$ , the piperazine ring adopts a chair conformation. The dihedral angle between the least-squares planes through the piperazine and benzene rings is  $73.23(10)^\circ$ . In the crystal, there are no classical hydrogen bonds but stabilization is provided by weak  $\text{C}-\text{H}\cdots\pi$  interactions.

### Related literature

For the biological activity of piperazine derivatives, see: Emami *et al.* (2006); Foroumadi *et al.* (2007). For puckering parameters, see: Cremer & Pople (1975).



### Experimental

#### Crystal data

$\text{C}_{13}\text{H}_{18}\text{N}_2\text{O}_4\text{S}$

$M_r = 298.35$

Monoclinic,  $P2_1/c$

$a = 6.1433(5)\text{ \AA}$

$b = 20.5966(17)\text{ \AA}$

$c = 12.5626(8)\text{ \AA}$

$\beta = 114.026(3)^\circ$

$V = 1451.84(19)\text{ \AA}^3$

$Z = 4$

Mo  $K\alpha$  radiation

$\mu = 0.24\text{ mm}^{-1}$   
 $T = 296\text{ K}$

$0.58 \times 0.38 \times 0.17\text{ mm}$

#### Data collection

Bruker APEXII DUO CCD area-detector diffractometer  
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)  
 $T_{\min} = 0.875$ ,  $T_{\max} = 0.961$

16507 measured reflections  
4255 independent reflections  
3400 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.023$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$   
 $wR(F^2) = 0.158$   
 $S = 1.04$   
4255 reflections

182 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.60\text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30\text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$Cg1$  is the centroid of the C1–C6 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C13—H13A $\cdots$ $Cg1^1$	0.96	2.97	3.900 (4)	165

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: TK2781).

### References

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# supporting information

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### S1. Comment

Piperazine derivatives are used as antibiotic drugs, *e.g.* Norfloxacin, Ciprofloxacin, Enoxacin, Ofloxacin and Levofloxacin (Emami *et al.*, 2006; Foroumadi *et al.*, 2007). Due to the biological importance of piperazine, herein, we present the crystal and molecular structure of the title compound, (I).

The piperazine (N1–N2/C7–C10) ring in (I), Fig. 1, adopts a chair conformation [puckering parameters:  $Q = 0.5682$  (18) Å,  $\theta = 2.56$  (17) ° and  $\varphi = 349$  (4) ° (Cremer & Pople, 1975)] with atoms N1 and C9 deviating by 0.253 (1) and 0.223 (2) Å from the least-squares plane defined by the remaining atoms (N2/C7,C8/C10) in the ring. The dihedral angle between the piperazine (N1–N2/C7–C10) ring and the benzene (C1–C6) ring is 73.23 (10) °.

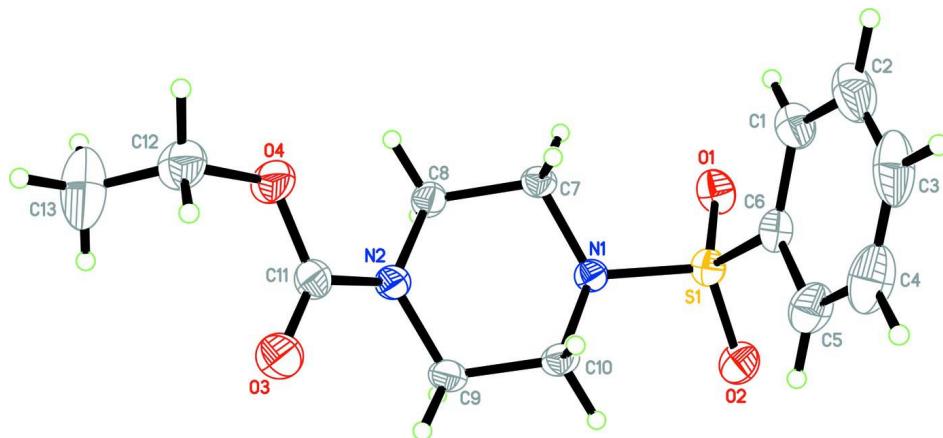
In the crystal structure (Fig. 2), there are no classical hydrogen bonds but stabilization is provided by weak C—H $\cdots$  $\pi$  interactions (Table 1).

### S2. Experimental

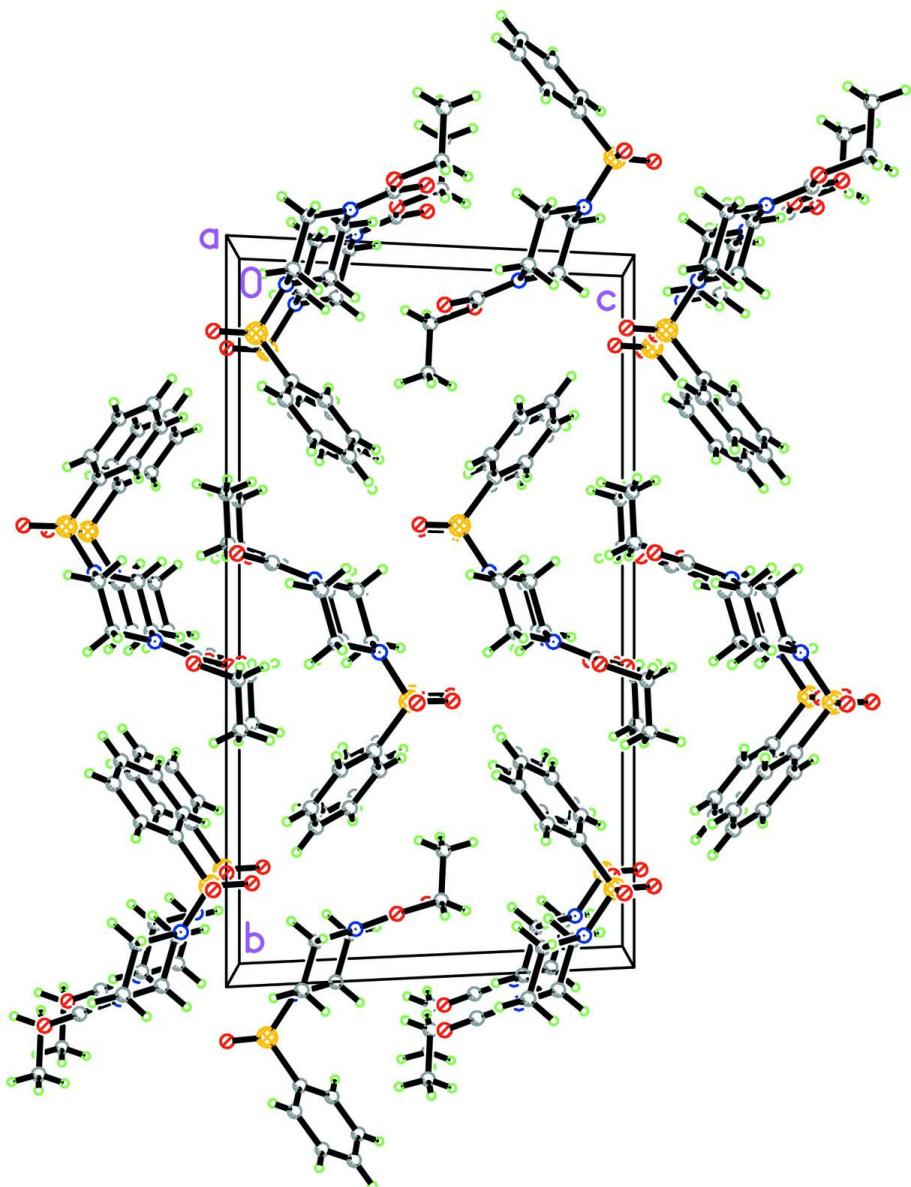
In a round bottom flask, 25ml of toluene was mixed with benzenesulfonyl chloride (0.01 mol, 1.0 g) with stirring. Ethyl-1-piperazine-carboxylate (0.01 mol, 1.7ml) dissolved in toluene was then added drop wise. The reaction mixture was refluxed for 30 min. The yellow precipitate formed was washed with alkaline water. The precipitate was then dissolved in methanol at room temperature. After few days, yellow crystals were formed by slow evaporation.

### S3. Refinement

All hydrogen atoms were positioned geometrically [ $C-H = 0.93$ – $0.97$  Å] and were refined using a riding model, with  $U_{iso}(H) = 1.2$  or  $1.5 U_{eq}(C)$ . A rotating group model was applied to the methyl group.

**Figure 1**

The molecular structure of (I), showing the atom labelling scheme and 50% probability displacement ellipsoids.

**Figure 2**

A view of the crystal packing in (I).

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#### Crystal data

$C_{13}H_{18}N_2O_4S$

$M_r = 298.35$

Monoclinic,  $P2_1/c$

Hall symbol: -P 2ybc

$a = 6.1433 (5) \text{ \AA}$

$b = 20.5966 (17) \text{ \AA}$

$c = 12.5626 (8) \text{ \AA}$

$\beta = 114.026 (3)^\circ$

$V = 1451.84 (19) \text{ \AA}^3$

$Z = 4$

$F(000) = 632$

$D_x = 1.365 \text{ Mg m}^{-3}$

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

Cell parameters from 5957 reflections

$\theta = 2.7\text{--}29.7^\circ$

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

$T = 296 \text{ K}$

Block, yellow

$0.58 \times 0.38 \times 0.17 \text{ mm}$

*Data collection*

Bruker APEXII DUO CCD area-detector  
diffractometer  
Radiation source: fine-focus sealed tube  
Graphite monochromator  
 $\varphi$  and  $\omega$  scans  
Absorption correction: multi-scan  
(*SADABS*; Bruker, 2009)  
 $T_{\min} = 0.875$ ,  $T_{\max} = 0.961$

16507 measured reflections  
4255 independent reflections  
3400 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.023$   
 $\theta_{\max} = 30.1^\circ$ ,  $\theta_{\min} = 2.0^\circ$   
 $h = -8 \rightarrow 8$   
 $k = -29 \rightarrow 24$   
 $l = -17 \rightarrow 17$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.048$   
 $wR(F^2) = 0.158$   
 $S = 1.04$   
4255 reflections  
182 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-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.0887P)^2 + 0.3986P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.60 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

*Special details*

**Geometry.** All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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  $F^2 > 2\sigma(F^2)$  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 ( $\text{\AA}^2$ )*

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.36299 (7)	-0.120155 (19)	0.94030 (3)	0.03858 (14)
O1	0.3060 (3)	-0.12052 (6)	1.04000 (10)	0.0517 (3)
O2	0.6074 (2)	-0.12032 (7)	0.95617 (12)	0.0547 (3)
O3	-0.1090 (3)	0.07756 (8)	0.52229 (12)	0.0659 (4)
O4	-0.3840 (2)	0.07205 (8)	0.59831 (12)	0.0605 (4)
N1	0.2424 (2)	-0.05482 (6)	0.86538 (10)	0.0346 (3)
N2	-0.0094 (2)	0.04239 (7)	0.70682 (12)	0.0412 (3)
C1	0.0247 (4)	-0.21553 (9)	0.86231 (18)	0.0544 (4)
H1A	-0.0323	-0.1999	0.9156	0.065*
C2	-0.0839 (4)	-0.26783 (11)	0.7902 (2)	0.0749 (7)
H2A	-0.2149	-0.2877	0.7957	0.090*
C3	-0.0001 (5)	-0.29053 (11)	0.7108 (2)	0.0813 (8)
H3A	-0.0749	-0.3255	0.6630	0.098*
C4	0.1924 (5)	-0.26197 (12)	0.7021 (2)	0.0785 (7)
H4A	0.2477	-0.2776	0.6482	0.094*
C5	0.3063 (4)	-0.20992 (11)	0.77261 (18)	0.0595 (5)

H5A	0.4375	-0.1905	0.7666	0.071*
C6	0.2208 (3)	-0.18732 (8)	0.85247 (14)	0.0421 (3)
C7	-0.0086 (3)	-0.04191 (8)	0.84341 (13)	0.0376 (3)
H7A	-0.0340	-0.0501	0.9135	0.045*
H7B	-0.1126	-0.0704	0.7824	0.045*
C8	-0.0652 (3)	0.02844 (8)	0.80650 (14)	0.0406 (3)
H8A	-0.2328	0.0367	0.7862	0.049*
H8B	0.0271	0.0567	0.8708	0.049*
C9	0.2364 (3)	0.02842 (9)	0.72652 (16)	0.0472 (4)
H9A	0.3431	0.0565	0.7873	0.057*
H9B	0.2594	0.0367	0.6558	0.057*
C10	0.2931 (3)	-0.04207 (9)	0.76225 (14)	0.0423 (3)
H10A	0.1969	-0.0703	0.6987	0.051*
H10B	0.4597	-0.0508	0.7804	0.051*
C11	-0.1625 (3)	0.06530 (8)	0.60275 (14)	0.0436 (4)
C12	-0.5669 (4)	0.09306 (12)	0.48616 (18)	0.0637 (5)
H12A	-0.5300	0.0763	0.4233	0.076*
H12C	-0.7206	0.0759	0.4769	0.076*
C13	-0.5775 (7)	0.16308 (16)	0.4804 (3)	0.1174 (13)
H13A	-0.6971	0.1763	0.4064	0.176*
H13B	-0.4254	0.1799	0.4891	0.176*
H13C	-0.6170	0.1795	0.5418	0.176*

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0397 (2)	0.0421 (2)	0.0286 (2)	0.00210 (13)	0.00853 (15)	0.00111 (13)
O1	0.0710 (9)	0.0526 (7)	0.0297 (6)	0.0009 (6)	0.0187 (6)	0.0025 (5)
O2	0.0368 (6)	0.0628 (8)	0.0521 (8)	0.0056 (5)	0.0053 (5)	0.0070 (6)
O3	0.0668 (9)	0.0926 (11)	0.0443 (7)	0.0176 (8)	0.0288 (6)	0.0217 (7)
O4	0.0449 (7)	0.0863 (10)	0.0481 (7)	0.0147 (6)	0.0167 (6)	0.0204 (7)
N1	0.0342 (6)	0.0394 (6)	0.0305 (6)	0.0018 (5)	0.0136 (5)	0.0023 (5)
N2	0.0382 (6)	0.0502 (7)	0.0376 (6)	0.0057 (5)	0.0179 (5)	0.0107 (6)
C1	0.0568 (10)	0.0465 (9)	0.0550 (10)	-0.0022 (8)	0.0178 (8)	0.0017 (8)
C2	0.0690 (14)	0.0462 (11)	0.0853 (17)	-0.0098 (9)	0.0065 (12)	0.0024 (11)
C3	0.0979 (19)	0.0399 (10)	0.0691 (15)	0.0061 (11)	-0.0039 (13)	-0.0143 (10)
C4	0.1010 (19)	0.0625 (14)	0.0608 (13)	0.0193 (13)	0.0213 (13)	-0.0212 (11)
C5	0.0699 (12)	0.0580 (11)	0.0521 (10)	0.0106 (9)	0.0262 (9)	-0.0092 (9)
C6	0.0470 (8)	0.0366 (8)	0.0374 (7)	0.0065 (6)	0.0119 (6)	0.0000 (6)
C7	0.0366 (7)	0.0439 (8)	0.0364 (7)	0.0003 (6)	0.0192 (6)	0.0037 (6)
C8	0.0435 (7)	0.0449 (8)	0.0373 (7)	0.0074 (6)	0.0205 (6)	0.0040 (6)
C9	0.0376 (7)	0.0587 (10)	0.0494 (9)	0.0020 (7)	0.0220 (7)	0.0164 (8)
C10	0.0380 (7)	0.0556 (9)	0.0386 (8)	0.0075 (6)	0.0210 (6)	0.0076 (7)
C11	0.0455 (8)	0.0477 (9)	0.0375 (8)	0.0038 (6)	0.0169 (6)	0.0053 (6)
C12	0.0510 (10)	0.0811 (15)	0.0494 (10)	0.0032 (10)	0.0104 (8)	0.0043 (10)
C13	0.131 (3)	0.077 (2)	0.101 (2)	0.0254 (18)	0.003 (2)	0.0136 (17)

Geometric parameters ( $\text{\AA}$ ,  $\text{^{\circ}}$ )

S1—O1	1.4312 (13)	C4—H4A	0.9300
S1—O2	1.4320 (14)	C5—C6	1.389 (2)
S1—N1	1.6365 (13)	C5—H5A	0.9300
S1—C6	1.7634 (17)	C7—C8	1.518 (2)
O3—C11	1.210 (2)	C7—H7A	0.9700
O4—C11	1.346 (2)	C7—H7B	0.9700
O4—C12	1.465 (2)	C8—H8A	0.9700
N1—C10	1.4745 (18)	C8—H8B	0.9700
N1—C7	1.4755 (18)	C9—C10	1.518 (2)
N2—C11	1.347 (2)	C9—H9A	0.9700
N2—C8	1.4554 (19)	C9—H9B	0.9700
N2—C9	1.455 (2)	C10—H10A	0.9700
C1—C6	1.387 (3)	C10—H10B	0.9700
C1—C2	1.391 (3)	C12—C13	1.444 (4)
C1—H1A	0.9300	C12—H12A	0.9700
C2—C3	1.377 (4)	C12—H12C	0.9700
C2—H2A	0.9300	C13—H13A	0.9600
C3—C4	1.365 (4)	C13—H13B	0.9600
C3—H3A	0.9300	C13—H13C	0.9600
C4—C5	1.385 (3)		
O1—S1—O2	119.62 (9)	C8—C7—H7B	109.9
O1—S1—N1	107.02 (7)	H7A—C7—H7B	108.3
O2—S1—N1	106.60 (7)	N2—C8—C7	110.26 (12)
O1—S1—C6	107.94 (8)	N2—C8—H8A	109.6
O2—S1—C6	108.04 (8)	C7—C8—H8A	109.6
N1—S1—C6	107.00 (7)	N2—C8—H8B	109.6
C11—O4—C12	115.91 (15)	C7—C8—H8B	109.6
C10—N1—C7	112.47 (11)	H8A—C8—H8B	108.1
C10—N1—S1	116.33 (10)	N2—C9—C10	109.69 (13)
C7—N1—S1	116.73 (10)	N2—C9—H9A	109.7
C11—N2—C8	126.01 (13)	C10—C9—H9A	109.7
C11—N2—C9	120.02 (13)	N2—C9—H9B	109.7
C8—N2—C9	113.96 (12)	C10—C9—H9B	109.7
C6—C1—C2	118.1 (2)	H9A—C9—H9B	108.2
C6—C1—H1A	121.0	N1—C10—C9	108.96 (13)
C2—C1—H1A	121.0	N1—C10—H10A	109.9
C3—C2—C1	120.9 (2)	C9—C10—H10A	109.9
C3—C2—H2A	119.6	N1—C10—H10B	109.9
C1—C2—H2A	119.6	C9—C10—H10B	109.9
C4—C3—C2	120.2 (2)	H10A—C10—H10B	108.3
C4—C3—H3A	119.9	O3—C11—O4	123.90 (15)
C2—C3—H3A	119.9	O3—C11—N2	124.31 (16)
C3—C4—C5	120.7 (2)	O4—C11—N2	111.78 (14)
C3—C4—H4A	119.6	C13—C12—O4	110.2 (2)
C5—C4—H4A	119.6	C13—C12—H12A	109.6

C4—C5—C6	118.7 (2)	O4—C12—H12A	109.6
C4—C5—H5A	120.7	C13—C12—H12C	109.6
C6—C5—H5A	120.7	O4—C12—H12C	109.6
C1—C6—C5	121.44 (18)	H12A—C12—H12C	108.1
C1—C6—S1	119.99 (14)	C12—C13—H13A	109.5
C5—C6—S1	118.55 (15)	C12—C13—H13B	109.5
N1—C7—C8	108.73 (12)	H13A—C13—H13B	109.5
N1—C7—H7A	109.9	C12—C13—H13C	109.5
C8—C7—H7A	109.9	H13A—C13—H13C	109.5
N1—C7—H7B	109.9	H13B—C13—H13C	109.5
O1—S1—N1—C10	175.82 (11)	N1—S1—C6—C5	85.65 (15)
O2—S1—N1—C10	46.71 (13)	C10—N1—C7—C8	-58.52 (17)
C6—S1—N1—C10	-68.69 (13)	S1—N1—C7—C8	163.26 (10)
O1—S1—N1—C7	-47.57 (12)	C11—N2—C8—C7	122.68 (18)
O2—S1—N1—C7	-176.67 (11)	C9—N2—C8—C7	-56.43 (18)
C6—S1—N1—C7	67.92 (12)	N1—C7—C8—N2	55.05 (16)
C6—C1—C2—C3	-0.5 (3)	C11—N2—C9—C10	-122.59 (17)
C1—C2—C3—C4	0.2 (4)	C8—N2—C9—C10	56.58 (19)
C2—C3—C4—C5	0.0 (4)	C7—N1—C10—C9	59.10 (17)
C3—C4—C5—C6	0.0 (3)	S1—N1—C10—C9	-162.49 (11)
C2—C1—C6—C5	0.5 (3)	N2—C9—C10—N1	-55.79 (18)
C2—C1—C6—S1	179.12 (15)	C12—O4—C11—O3	2.6 (3)
C4—C5—C6—C1	-0.3 (3)	C12—O4—C11—N2	-176.51 (17)
C4—C5—C6—S1	-178.91 (16)	C8—N2—C11—O3	178.53 (18)
O1—S1—C6—C1	21.89 (16)	C9—N2—C11—O3	-2.4 (3)
O2—S1—C6—C1	152.57 (14)	C8—N2—C11—O4	-2.4 (2)
N1—S1—C6—C1	-92.99 (15)	C9—N2—C11—O4	176.66 (15)
O1—S1—C6—C5	-159.47 (14)	C11—O4—C12—C13	-88.6 (3)
O2—S1—C6—C5	-28.79 (17)		

*Hydrogen-bond geometry (Å, °)*

Cg1 is the centroid of the C1—C6 ring.

D—H···A	D—H	H···A	D···A	D—H···A
C13—H13A···Cg1 <sup>i</sup>	0.96	2.97	3.900 (4)	165

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