Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

N-(4-Acetylphenyl)-4-methoxybenzenesulfonamide

Thawanrat Kobkeatthawin,[a] Suchada Chantrapromma,[a]*‡ C. S. Chidan Kumar[b] and Hoong-Kun Fun[b]§

[a]Department of Chemistry, Faculty of Science, Prince of Songkla University, Hat-Yai, Songkhla 90112, Thailand, and [b]X-ray Crystallography Unit, School of Physics, Universiti Sains Malaysia, 11800 USM, Penang, Malaysia
Correspondence e-mail: suchada.c@psu.ac.th

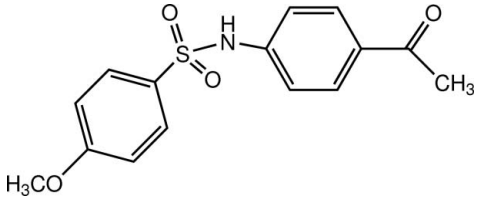
Received 21 October 2013; accepted 31 October 2013

Key indicators: single-crystal X-ray study; T = 298 K; mean σ(C–C) = 0.003 Å; R factor = 0.047; wR factor = 0.135; data-to-parameter ratio = 21.0.

The title compound, $C_{15}H_{15}NO_4S$, was obtained by the condensation of 4-aminoacetophenone and 4-methoxybenzenesulfonyl chloride. The dihedral angle between the benzene rings is 86.56 (9)° and the molecule has an approximate V-shaped conformation. The C atom of the methoxy group is roughly coplanar with its attached ring [deviation = 0.177 (3) Å], as is the methyl C atom of the acetyl group with its ring [deviation = 0.065 (2) Å]. An intramolecular C—H···O interaction generates an S(6) ring. In the crystal, N—H···O and C—H···O hydrogen bonds link the molecules into [010] chains. Weak C—H···π interactions are also observed.

Related literature

For related structures, see: Li *et al.* (2006); Xu *et al.* (2005). For background to and applications of sulfonamides, see: Alsughayer *et al.* (2011); Dragostin *et al.* (2013);



Experimental

Crystal data

$C_{15}H_{15}NO_4S$
M_r = 305.35
Monoclinic, $P2_1/c$
a = 12.8220 (3) Å
b = 8.2709 (2) Å
c = 14.6165 (4) Å
β = 112.841 (1)°
V = 1428.52 (6) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.24 mm^{-1}
T = 298 K
0.48 × 0.44 × 0.33 mm

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
T_{min} = 0.894, T_{max} = 0.924
15571 measured reflections
4120 independent reflections
2593 reflections with $I > 2\sigma(I)$
R_{int} = 0.043

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.047
$wR(F^2)$ = 0.135
S = 1.04
4120 reflections
196 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.20 e Å^{-3}
$\Delta\rho_{min}$ = −0.34 e Å^{-3}

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

*Cg*1 and *Cg*2 are the centroids of the C1–C6 and C7–C12 rings, respectively.

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1*N*1···O3^{i}	0.85 (2)	2.05 (2)	2.896 (2)	172.3 (19)
C8—H8*A*···O2	0.93	2.38	3.030 (2)	127
C9—H9*A*···O1^{ii}	0.93	2.53	3.459 (2)	174
C14—H14*A*···*Cg*1^{i}	0.96	2.83	3.630 (3)	141
C14—H14*C*···*Cg*2^{iii}	0.96	2.83	3.529 (2)	130
C15—H15*C*···*Cg*1^{iv}	0.96	2.99	3.804 (3)	144

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

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*, *PLATON* (Spek, 2009), *Mercury* (Macrae *et al.*, 2006) and *publCIF* (Westrip, 2010).

Financial support from the Thailand Research Fund through the Royal Golden Jubilee PhD Program (grant No. PHD/0137/2554) is gratefully acknowledged. CSCK thanks the Universiti Sains Malaysia for a postdoctoral research fellowship. The authors extend their appreciation to Prince of Songkla University and the Universiti Sains Malaysia for the APEX DE2012 grant No. 1002/PFIZIK/910323.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB7153).

References

Alsughayer, A., Elassar, A.-Z. A., Mustafa, S. & Sagheer, F. A. (2011). *J. Biomater. Nanobiotechnol.* **2**, 144–149.
Bruker (2009). *APEX2*, *SAINT* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.
Dragostin, O. M., Lupascu, F., Vasile, C., Mares, M., Nastasa, V., Moraru, R. F., Pieptu, D. & Profire, L. (2013). *Molecules*, **18**, 4140–4157.
Li, X.-M., Zeng, C.-C., Xu, Y.-S., Yan, H., Zheng, D.-W. & Zhong, R. G. (2006). *J. Chem. Crystallogr.* **36**, 357–363.

‡ Thomson Reuters ResearcherID: A-5085-2009.
§ Additional correspondence author, e-mail: hkfun@usm.my. Thomson Reuters ResearcherID: A-3561-2009.

Macrae, C. F., Edgington, P. R., McCabe, P., Pidcock, E., Shields, G. P., Taylor, R., Towler, M. & van de Streek, J. (2006). *J. Appl. Cryst.* **39**, 453–457.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Spek, A. L. (2009). *Acta Cryst*. D**65**, 148–155.
Westrip, S. P. (2010). *J. Appl. Cryst.* **43**, 920–925.
Xu, Y.-S., Zeng, C.-C., Li, X.-M. & Zhong, R.-G. (2005). *Acta Cryst.* E**61**, o3802–o3804.

supplementary materials

Acta Cryst. (2013). E**69**, o1750–o1751 [doi:10.1107/S1600536813029875]

N-(4-Acetylphenyl)-4-methoxybenzenesulfonamide

Thawanrat Kobkeatthawin, Suchada Chantrapromma, C. S. Chidan Kumar and Hoong-Kun Fun

1. Comment

Sulfonamides containing an $–SO_2NH–$ group can be found in many pharmacologically active compounds: recent reports have described antibacterial (Alsughayer *et al.*, 2011) and antioxidant (Dragostin *et al.*, 2013) behaviour. As part of our ongoing research in this field, the title compound (I), a 4-methoxybenzene-sulfonamide derivative, was synthesized for being used as starting material for various syntheses. Herein the crystal structure of (I) is reported.

Figure 1 shows the molecular structure of (I), $C_{15}H_{15}NO_4S$, suggesting a V-shaped conformation (Fig. 2). The benzene rings make the dihedral angle of 86.56 (9)°. The methoxy group is almost co-planar with its attached benzene ring with the deviation of 0.0310 (2) Å for the eight non H atoms (C1–C6/O4/C15) and the torsion angle C15–O4–C4–C5 = -2.2 (3)°. The amide group and acetyl substituent also lie in almost the same plane with the bound benzene ring with the deviation of 0.0220 (2) Å for the ten non H atoms (C7–C14/N1/O3) and the torsion angles of C11–C10–C13–O3 = -177.28 (18)° and C11–C10–C13–C14 = 2.6 (3)°. The dihedral angle between these two planes [C1–C6/O4/C15 and C7–C14/N1/O3] is 88.38 (7)° (Fig. 2). An intramolecular C8—H8A···O2 weak interaction generates an S(6) ring (Fig. 1) Bond distances of (I) are comparable with those in related structures (Li *et al.*, 2006 and Xu *et al.*, 2005).

In the crystal (Fig. 3), the molecules are linked by N—H···O hydrogen bonds and C—H···O weak interactions (Table 1) into chains along [010]. Weak C—H···π interactions are also observed (Table 1).

2. Experimental

The title compound was synthesized by condensation of 4-aminoacetophenone (0.40 g, 3 mmol) and 4-methoxybenzene-sulfonyl chloride in CH_2Cl_2 (30 ml) in the presence of pyridine. The reaction mixture was refluxed for 24 hr at 40 °C and monitored with TLC for the completion of the reaction. Water was then added and the concoction was extracted with CH_2Cl_2. The solvent was evaporated under reduced pressure to yield the resulting solid of the title compound (yield 68%). Yellow blocks of (I) were recrystalized from acetone:CH_3OH solution (1:1 *v*/*v*) by slow evaporation of the solvent at room temperature after several days, Mp. 448–449 K.

3. Refinement

Amide H atoms was located from the difference maps and refined isotropically. The remaining H atoms were positioned geometrically and allowed to ride on their parent atoms, with d(C—H) = 0.93 Å for aromatic and 0.96 for CH_3 atoms. The U_{iso} values were constrained to be $1.5U_{eq}$ of the carrier atom for methyl H atoms and $1.2U_{eq}$ for the remaining H atoms. A rotating group model was used for the methyl groups.

Computing details

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

SHELXTL (Sheldrick, 2008), *PLATON* (Spek, 2009), *Mercury* (Macrae *et al.*, 2006) and *publCIF* (Westrip, 2010).

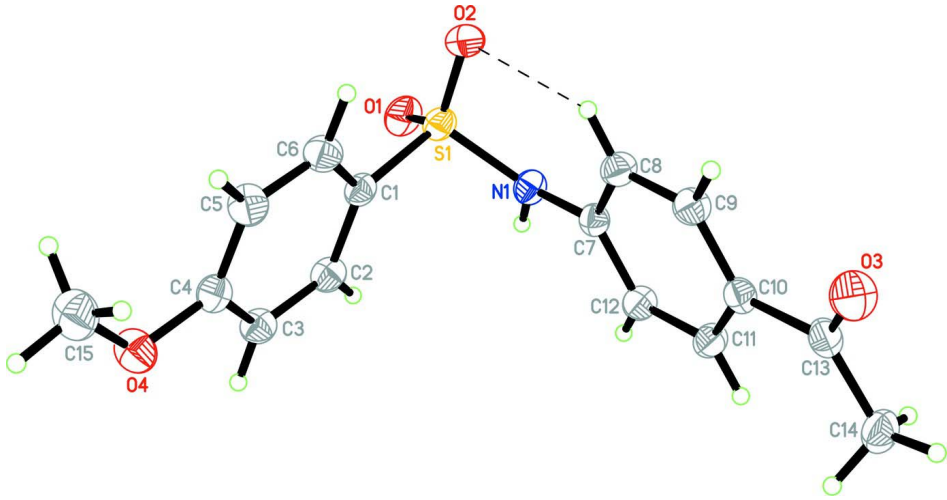


Figure 1

The molecular structure of (I), showing 40% probability displacement ellipsoids. The intramolecular C—H···O hydrogen bond is shown as a dashed line.

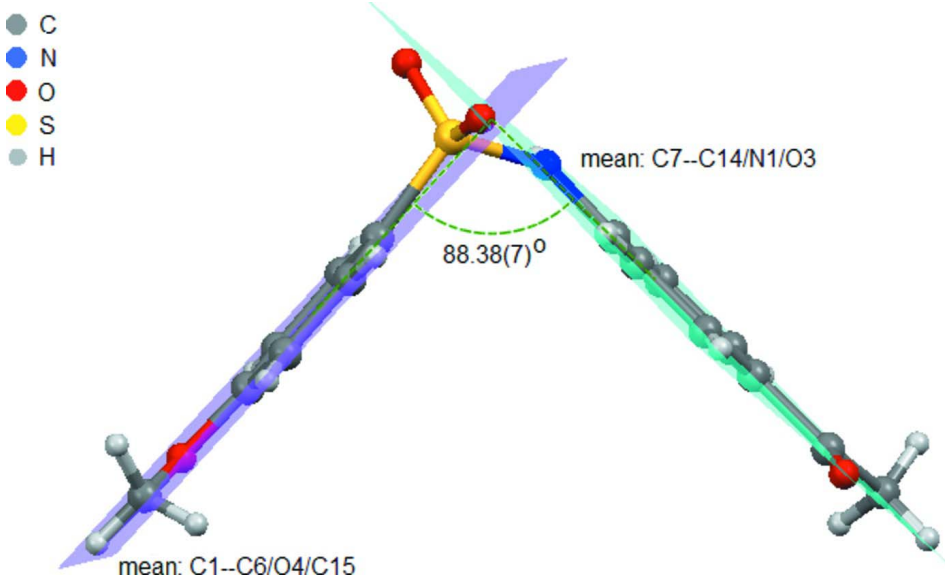


Figure 2

The V-shape conformation of the molecule.

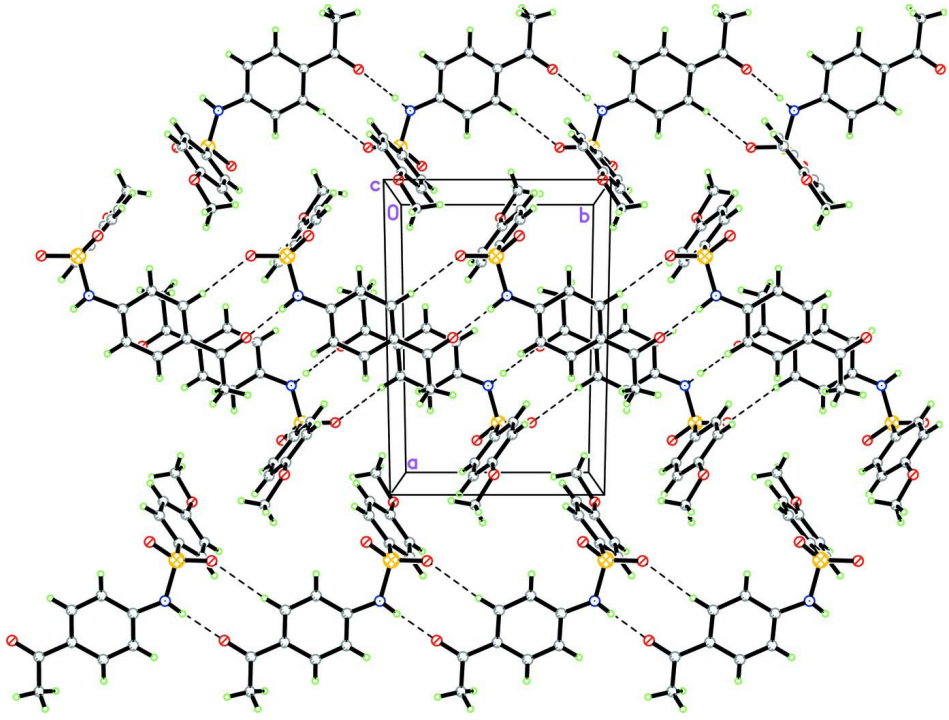


Figure 3

The crystal packing of the title compound viewed along the *c* axis. Hydrogen bonds were shown as dashed lines.

N-(4-Acetylphenyl)-4-methoxybenzenesulfonamide

Crystal data

$C_{15}H_{15}NO_4S$
$M_r = 305.35$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
$a = 12.8220\ (3)$ Å
$b = 8.2709\ (2)$ Å
$c = 14.6165\ (4)$ Å
$\beta = 112.841\ (1)°$
$V = 1428.52\ (6)$ Å^3
$Z = 4$
$F(000) = 640$
$D_x = 1.420$ Mg m^{-3}
Melting point = 448–449 K
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 4120 reflections
$\theta = 1.7$–$29.9°$
$\mu = 0.24$ mm^{-1}
$T = 298$ K
Block, yellow
0.48 × 0.44 × 0.33 mm

Data collection

Bruker APEXII CCD diffractometer
Radiation source: sealed tube
Graphite monochromator
φ and ω scans
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
$T_{min} = 0.894$, $T_{max} = 0.924$
15571 measured reflections
4120 independent reflections
2593 reflections with $I > 2\sigma(I)$
$R_{int} = 0.043$
$\theta_{max} = 29.9°$, $\theta_{min} = 1.7°$
$h = -17 \rightarrow 17$
$k = -11 \rightarrow 11$
$l = -20 \rightarrow 19$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.047$
$wR(F^2) = 0.135$
$S = 1.04$
4120 reflections
196 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
$w = 1/[\sigma^2(F_o^2) + (0.0625P)^2 + 0.0913P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} = 0.001$
$\Delta\rho_{max} = 0.20$ e Å^{-3}
$\Delta\rho_{min} = -0.34$ e Å^{-3}

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	0.21179 (4)	1.00630 (5)	1.08283 (4)	0.04186 (15)
O1	0.21200 (12)	1.17413 (16)	1.10664 (11)	0.0546 (4)
O2	0.14687 (11)	0.89569 (17)	1.11408 (11)	0.0529 (4)
O3	0.48523 (14)	0.21499 (16)	1.11408 (12)	0.0624 (4)
O4	0.05688 (12)	0.98389 (16)	0.64850 (11)	0.0540 (4)
N1	0.34483 (13)	0.95337 (19)	1.13298 (12)	0.0408 (4)
C1	0.16988 (14)	0.9867 (2)	0.95362 (14)	0.0378 (4)
C2	0.22355 (15)	1.0787 (2)	0.90451 (14)	0.0418 (4)
H2A	0.2849	1.1439	0.9404	0.050*
C3	0.18492 (15)	1.0722 (2)	0.80272 (15)	0.0432 (4)
H3A	0.2209	1.1321	0.7697	0.052*
C4	0.09193 (16)	0.9760 (2)	0.74878 (14)	0.0415 (4)
C5	0.04067 (17)	0.8825 (2)	0.79829 (15)	0.0487 (5)
H5A	−0.0201	0.8161	0.7627	0.058*
C6	0.07979 (16)	0.8882 (2)	0.90017 (15)	0.0466 (5)
H6A	0.0454	0.8254	0.9333	0.056*
C7	0.38891 (14)	0.7983 (2)	1.12776 (12)	0.0353 (4)
C8	0.32855 (16)	0.6554 (2)	1.12191 (14)	0.0422 (4)
H8A	0.2549	0.6591	1.1191	0.051*
C9	0.37892 (16)	0.5091 (2)	1.12029 (15)	0.0427 (4)
H9A	0.3381	0.4144	1.1157	0.051*
C10	0.48954 (16)	0.4993 (2)	1.12535 (13)	0.0381 (4)
C11	0.54871 (16)	0.6432 (2)	1.13233 (14)	0.0422 (4)
H11A	0.6230	0.6394	1.1368	0.051*
C12	0.49943 (15)	0.7902 (2)	1.13275 (14)	0.0413 (4)

H12A	0.5400	0.8849	1.1364	0.050*
C13	0.54079 (17)	0.3380 (2)	1.12393 (14)	0.0427 (4)
C14	0.66124 (18)	0.3279 (3)	1.13447 (15)	0.0542 (5)
H14A	0.6814	0.2168	1.1316	0.081*
H14B	0.7094	0.3736	1.1970	0.081*
H14C	0.6704	0.3870	1.0815	0.081*
C15	−0.0427 (2)	0.8943 (3)	0.59083 (17)	0.0666 (6)
H15A	−0.0602	0.9117	0.5215	0.100*
H15B	−0.1050	0.9300	0.6067	0.100*
H15C	−0.0297	0.7812	0.6057	0.100*
H1N1	0.3889 (18)	1.031 (2)	1.1334 (15)	0.047 (6)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0414 (2)	0.0396 (3)	0.0475 (3)	0.00720 (19)	0.0204 (2)	−0.00056 (19)
O1	0.0597 (9)	0.0422 (8)	0.0628 (9)	0.0125 (6)	0.0248 (7)	−0.0084 (6)
O2	0.0485 (8)	0.0588 (9)	0.0603 (9)	0.0041 (6)	0.0309 (7)	0.0070 (7)
O3	0.0769 (10)	0.0329 (7)	0.0842 (11)	−0.0013 (7)	0.0386 (9)	0.0008 (7)
O4	0.0561 (9)	0.0552 (9)	0.0464 (9)	−0.0140 (7)	0.0152 (7)	−0.0038 (6)
N1	0.0411 (8)	0.0332 (8)	0.0468 (10)	0.0015 (7)	0.0156 (7)	−0.0022 (7)
C1	0.0337 (8)	0.0337 (9)	0.0464 (10)	0.0046 (7)	0.0158 (8)	0.0028 (7)
C2	0.0365 (9)	0.0369 (10)	0.0508 (12)	−0.0058 (7)	0.0156 (8)	−0.0021 (8)
C3	0.0404 (10)	0.0378 (10)	0.0528 (12)	−0.0039 (8)	0.0196 (9)	0.0016 (8)
C4	0.0412 (9)	0.0374 (10)	0.0438 (11)	0.0011 (8)	0.0143 (8)	−0.0015 (8)
C5	0.0444 (10)	0.0437 (11)	0.0526 (12)	−0.0126 (8)	0.0130 (9)	−0.0018 (9)
C6	0.0435 (10)	0.0425 (10)	0.0552 (12)	−0.0073 (8)	0.0209 (9)	0.0039 (9)
C7	0.0381 (8)	0.0344 (9)	0.0321 (9)	0.0022 (7)	0.0121 (7)	−0.0007 (7)
C8	0.0385 (9)	0.0405 (10)	0.0496 (11)	−0.0001 (8)	0.0193 (8)	0.0004 (8)
C9	0.0455 (10)	0.0337 (9)	0.0513 (11)	−0.0035 (8)	0.0213 (9)	0.0010 (8)
C10	0.0440 (9)	0.0346 (9)	0.0356 (9)	0.0016 (7)	0.0153 (8)	0.0026 (7)
C11	0.0373 (9)	0.0411 (10)	0.0494 (11)	0.0022 (8)	0.0179 (8)	0.0020 (8)
C12	0.0391 (9)	0.0346 (9)	0.0506 (11)	−0.0030 (7)	0.0180 (8)	0.0004 (8)
C13	0.0552 (11)	0.0377 (10)	0.0356 (10)	0.0069 (8)	0.0180 (9)	0.0038 (7)
C14	0.0590 (12)	0.0497 (12)	0.0530 (13)	0.0166 (10)	0.0208 (10)	−0.0003 (9)
C15	0.0587 (14)	0.0790 (16)	0.0519 (14)	−0.0175 (12)	0.0103 (11)	−0.0074 (12)

Geometric parameters (Å, º)

S1—O2	1.4261 (14)	C7—C12	1.392 (2)
S1—O1	1.4308 (13)	C7—C8	1.397 (2)
S1—N1	1.6335 (16)	C8—C9	1.376 (2)
S1—C1	1.7590 (19)	C8—H8A	0.9300
O3—C13	1.218 (2)	C9—C10	1.394 (3)
O4—C4	1.358 (2)	C9—H9A	0.9300
O4—C15	1.433 (3)	C10—C11	1.394 (2)
N1—C7	1.416 (2)	C10—C13	1.491 (2)
N1—H1N1	0.85 (2)	C11—C12	1.371 (2)
C1—C6	1.381 (3)	C11—H11A	0.9300
C1—C2	1.397 (3)	C12—H12A	0.9300

C2—C3	1.375 (3)	C13—C14	1.494 (3)
C2—H2A	0.9300	C14—H14A	0.9600
C3—C4	1.395 (3)	C14—H14B	0.9600
C3—H3A	0.9300	C14—H14C	0.9600
C4—C5	1.387 (3)	C15—H15A	0.9600
C5—C6	1.375 (3)	C15—H15B	0.9600
C5—H5A	0.9300	C15—H15C	0.9600
C6—H6A	0.9300		

O2—S1—O1	119.39 (9)	C9—C8—C7	119.50 (17)
O2—S1—N1	108.79 (8)	C9—C8—H8A	120.3
O1—S1—N1	104.35 (9)	C7—C8—H8A	120.3
O2—S1—C1	108.17 (9)	C8—C9—C10	121.67 (17)
O1—S1—C1	108.75 (8)	C8—C9—H9A	119.2
N1—S1—C1	106.72 (8)	C10—C9—H9A	119.2
C4—O4—C15	117.12 (16)	C11—C10—C9	117.90 (16)
C7—N1—S1	126.00 (13)	C11—C10—C13	122.34 (17)
C7—N1—H1N1	113.8 (14)	C9—C10—C13	119.76 (16)
S1—N1—H1N1	112.0 (14)	C12—C11—C10	121.26 (17)
C6—C1—C2	120.09 (18)	C12—C11—H11A	119.4
C6—C1—S1	120.10 (15)	C10—C11—H11A	119.4
C2—C1—S1	119.69 (14)	C11—C12—C7	120.25 (16)
C3—C2—C1	119.49 (17)	C11—C12—H12A	119.9
C3—C2—H2A	120.3	C7—C12—H12A	119.9
C1—C2—H2A	120.3	O3—C13—C10	120.55 (18)
C2—C3—C4	120.26 (18)	O3—C13—C14	119.98 (17)
C2—C3—H3A	119.9	C10—C13—C14	119.47 (17)
C4—C3—H3A	119.9	C13—C14—H14A	109.5
O4—C4—C5	124.41 (17)	C13—C14—H14B	109.5
O4—C4—C3	115.76 (17)	H14A—C14—H14B	109.5
C5—C4—C3	119.82 (18)	C13—C14—H14C	109.5
C6—C5—C4	119.90 (18)	H14A—C14—H14C	109.5
C6—C5—H5A	120.0	H14B—C14—H14C	109.5
C4—C5—H5A	120.0	O4—C15—H15A	109.5
C5—C6—C1	120.40 (18)	O4—C15—H15B	109.5
C5—C6—H6A	119.8	H15A—C15—H15B	109.5
C1—C6—H6A	119.8	O4—C15—H15C	109.5
C12—C7—C8	119.41 (16)	H15A—C15—H15C	109.5
C12—C7—N1	117.40 (16)	H15B—C15—H15C	109.5
C8—C7—N1	123.14 (17)		

O2—S1—N1—C7	53.22 (18)	C2—C1—C6—C5	−1.4 (3)
O1—S1—N1—C7	−178.31 (15)	S1—C1—C6—C5	174.65 (15)
C1—S1—N1—C7	−63.27 (17)	S1—N1—C7—C12	151.37 (15)
O2—S1—C1—C6	5.98 (17)	S1—N1—C7—C8	−31.2 (3)
O1—S1—C1—C6	−125.08 (15)	C12—C7—C8—C9	−0.6 (3)
N1—S1—C1—C6	122.88 (15)	N1—C7—C8—C9	−177.92 (17)
O2—S1—C1—C2	−177.94 (13)	C7—C8—C9—C10	0.6 (3)
O1—S1—C1—C2	51.00 (16)	C8—C9—C10—C11	0.1 (3)

N1—S1—C1—C2	−61.03 (15)	C8—C9—C10—C13	179.65 (17)
C6—C1—C2—C3	1.0 (3)	C9—C10—C11—C12	−1.0 (3)
S1—C1—C2—C3	−175.13 (14)	C13—C10—C11—C12	179.50 (17)
C1—C2—C3—C4	0.8 (3)	C10—C11—C12—C7	1.0 (3)
C15—O4—C4—C5	2.7 (3)	C8—C7—C12—C11	−0.3 (3)
C15—O4—C4—C3	−176.22 (18)	N1—C7—C12—C11	177.24 (17)
C2—C3—C4—O4	176.85 (16)	C11—C10—C13—O3	−177.28 (18)
C2—C3—C4—C5	−2.2 (3)	C9—C10—C13—O3	3.2 (3)
O4—C4—C5—C6	−177.21 (18)	C11—C10—C13—C14	2.6 (3)
C3—C4—C5—C6	1.7 (3)	C9—C10—C13—C14	−176.96 (18)
C4—C5—C6—C1	0.1 (3)		

Hydrogen-bond geometry (Å, º)

Cg1 and Cg2 are the centroids of the C1–C6 and C7–C12 rings, respectively.

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1*N*1···O3^{i}	0.85 (2)	2.05 (2)	2.896 (2)	172.3 (19)
C8—H8*A*···O2	0.93	2.38	3.030 (2)	127
C9—H9*A*···O1^{ii}	0.93	2.53	3.459 (2)	174
C14—H14*A*···*Cg*1^{i}	0.96	2.83	3.630 (3)	141
C14—H14*C*···*Cg*2^{iii}	0.96	2.83	3.529 (2)	130
C15—H15*C*···*Cg*1^{iv}	0.96	2.99	3.804 (3)	144

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