Ethyl 2-[N-(2-formylphenyl)benzene-sulfonamido]acetate

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In the molecule of the title compound, $C_{17}H_{17}NO_5S$, the two aromatic rings are oriented at an angle of 30.13 (10)°. The ethyl acetate group assumes an extended conformation. Molecules are linked into $C(7)$ chains running along the a axis by intermolecular C—H···O hydrogen bonds, and the chains are crosslinked via C—H···π interactions, with the sulfonyl-bound phenyl ring acting as an acceptor.

Related literature

For the activities of sulfonamides, see: Krishnaiah et al. (1995); Dupont et al. (1978); Sethu Sankar et al. (2002). For related literature, see: Bassindale (1984).
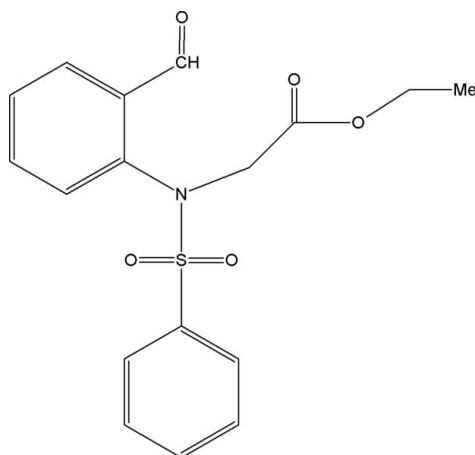
Experimental

Crystal data

$C_{17}H_{17}NO_5S$	$M_r = 347.38$
Orthorhombic, $P2_12_12_1$	$Z = 4$
$a = 11.3512$ (6) Å	Mo $K\alpha$ radiation
$b = 11.7820$ (6) Å	$\mu = 0.22$ mm^{-1}
$c = 12.8045$ (6) Å	$T = 293$ (2) K
$V = 1712.47$ (15) Å^3	$0.25 \times 0.22 \times 0.19$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer	13978 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	5831 independent reflections
$T_{min} = 0.948$, $T_{max} = 0.960$	3738 reflections with $I > 2\sigma(I)$
	$R_{int} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.042$	H-atom parameters constrained
$wR(F^2) = 0.109$	$\Delta\rho_{max} = 0.29$ e Å^{-3}
$S = 1.02$	$\Delta\rho_{min} = -0.25$ e Å^{-3}
5831 reflections	Absolute structure: Flack (1983), with 2533 Friedel pairs
218 parameters	Flack parameter: 0.04 (6)
2 restraints	

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

$Cg1$ is the centroid of the C8–C13 ring.

D—H···A	D—H	H···A	D···A	D—H···A
C5—H5···O4^i	0.93	2.57	3.220 (2)	127
C16—H16B···$Cg1^{ii}$	0.97	2.75	3.615 (2)	150

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); 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 (Farrugia, 1997); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2003).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2739).

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supporting information

Ethyl 2-[*N*-(2-formylphenyl)benzenesulfonamido]acetate

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S1. Comment

The title compound is a potential intermediate for the synthesis of 2-alkylbenzoic acid and exhibits insecticidal, germicidal and antimicrobial activities (Krishnaiah *et al.*, 1995; Dupont *et al.*, 1978). The sulfonamides inhibit the growth of bacterial organism and are also useful for treating urinary and gastrointestinal infections (Sethu Sankar *et al.*, 2002).

Atom S1 has a distorted tetrahedral configuration. The widening of angle O2—S1—O3 [120.46 (10)°] and narrowing of angle C8—S1—N1 [105.97 (8)°] from the ideal tetrahedral value are attributed to the Thorpe-Ingold effect (Bassindale, 1984). The two phenyl rings are oriented at an angle of 30.13 (10)°. The ethylacetate moiety assumes an extended conformation as can be seen from torsion angles C14—C15—O5—C16 of 178.12 (15)° and C15—O5—C16—C17 of 173.12 (19)°.

The molecules are linked into C(7) chains running along the *a* axis by C—H⋯O hydrogen bonding (Table 1). In addition C—H⋯*π* interactions (Table 1) with C8–C13 ring (centroid Cg1) as an acceptor is observed.

S2. Experimental

2-(Benzenesulfonylamino)benzaldehyde (2 mmol) was added with ethyl bromoacetate (2.2 mmol) in the presence of potassium carbonate (4.7 mmol) and dimethyl acetamide (15 ml). The mixture was stirred at room temperature for 6 h. The reaction mass was poured into crushed ice (50 g) containing 4 to 5 drops of concentrated HCl and extracted with ethyl acetate. The product was obtained by column chromatography (hexane-ethyl acetate 9:1). The removal of the solvent followed by column chromatography of the residue (ethyl acetate) afforded white crystalline solid (yield 25%, m.p. 381-383 K). Single crystals suitable for the X-ray diffraction were obtained by slow evaporation of an ethyl acetate solution of the title compound at room temperature.

S3. Refinement

H atoms were positioned geometrically (C-H = 0.93-0.97 Å) and were treated as riding on their parent C atoms with $U_{iso}(H) = 1.2$-$1.5U_{eq}(C)$. The U^{ij} components of atoms C1, C2 and C6 in the direction of the bond between them were restrained to be equal within an effective standard deviation of 0.001.
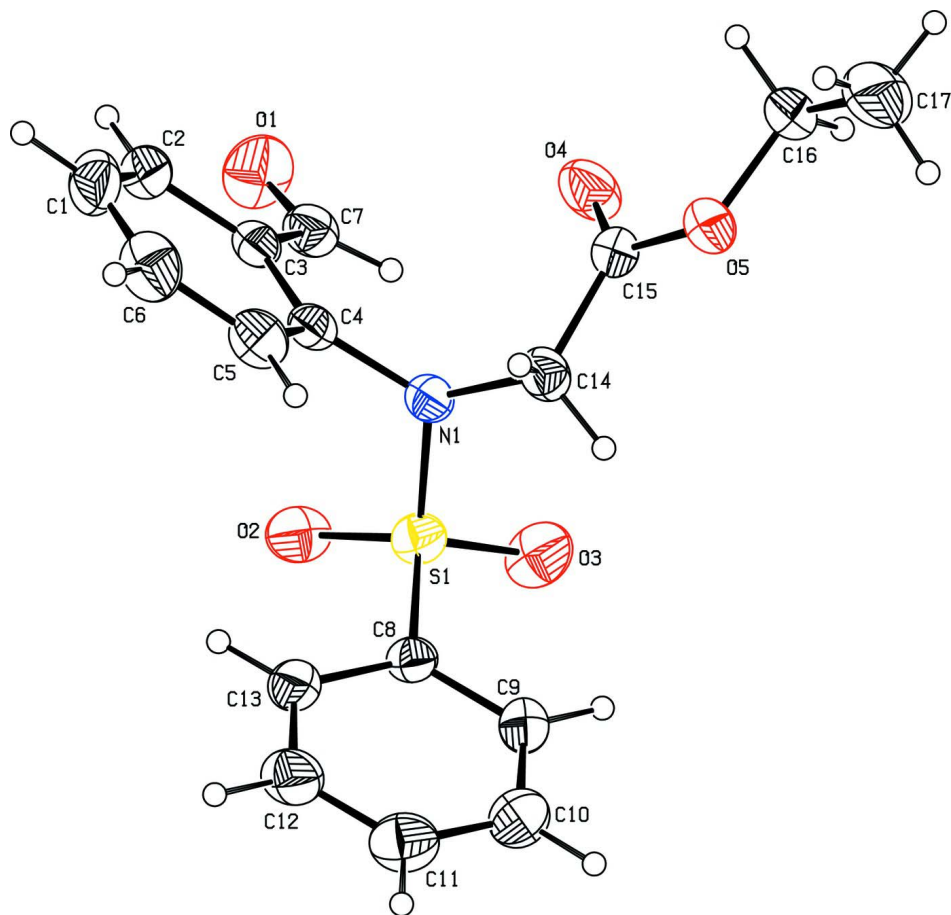
Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 30% probability level.
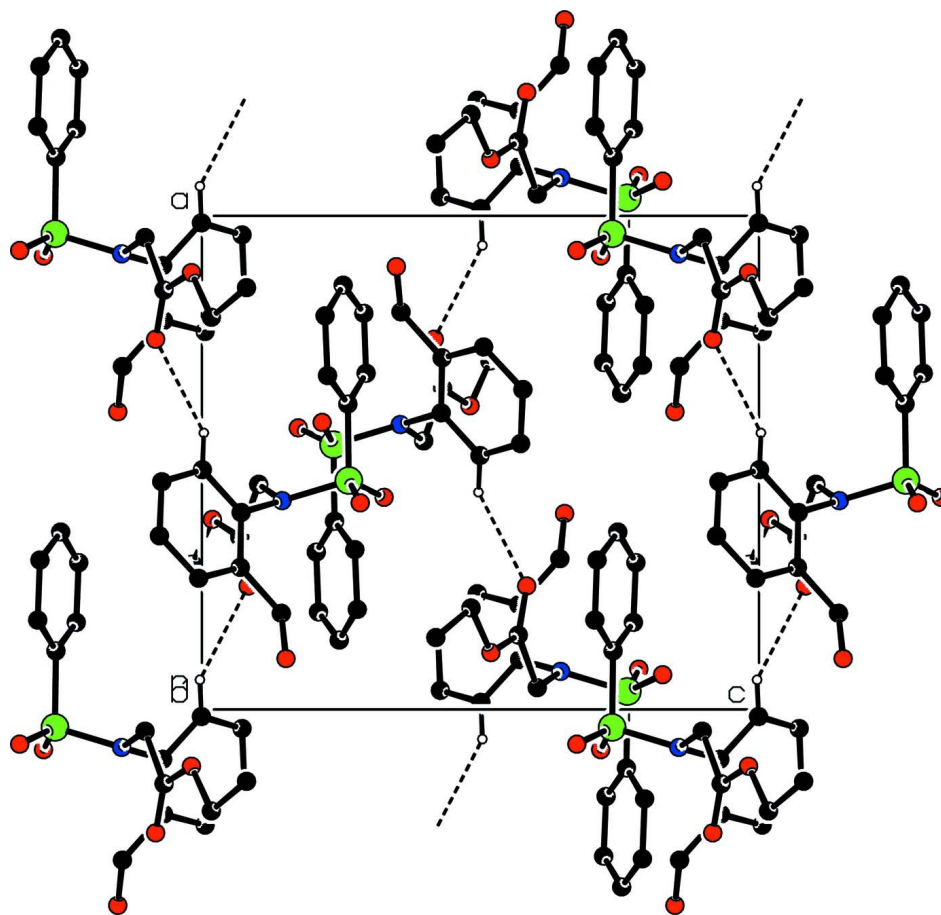
Figure 2

Molecular packing in the title compound, viewed down the *b* axis. Dashed lines represent hydrogen bonds.

Ethyl 2-[*N*-(2-formylphenyl)benzenesulfonamido]acetate

Crystal data

$C_{17}H_{17}NO_5S$	$F(000) = 728$
$M_r = 347.38$	$D_x = 1.347$ Mg m^{-3}
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Hall symbol: P 2ac 2ab	Cell parameters from 5831 reflections
$a = 11.3512$ (6) Å	$\theta = 2.4$–$31.9°$
$b = 11.7820$ (6) Å	$\mu = 0.22$ mm^{-1}
$c = 12.8045$ (6) Å	$T = 293$ K
$V = 1712.47$ (15) Å^3	Block, white
$Z = 4$	$0.25 \times 0.22 \times 0.19$ mm

Data collection

Bruker APEXII CCD area-detector diffractometer	13978 measured reflections
	5831 independent reflections
Radiation source: fine-focus sealed tube	3738 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{int} = 0.024$
ω and φ scans	$\theta_{max} = 31.9°$, $\theta_{min} = 2.4°$
Absorption correction: multi-scan	$h = -16 \rightarrow 16$
(*SADABS*; Sheldrick, 1996)	$k = -17 \rightarrow 14$
$T_{min} = 0.948$, $T_{max} = 0.960$	$l = -19 \rightarrow 17$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.042$	$w = 1/[\sigma^2(F_o^2) + (0.0543P)^2 + 0.0353P]$
$wR(F^2) = 0.109$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.02$	$(\Delta/\sigma)_{max} = 0.001$
5831 reflections	$\Delta\rho_{max} = 0.29$ e Å⁻³
218 parameters	$\Delta\rho_{min} = -0.25$ e Å⁻³
2 restraints	Absolute structure: Flack (1983), 2533 Friedel pairs
Primary atom site location: structure-invariant direct methods	Absolute structure parameter: 0.04 (6)
Secondary atom site location: difference Fourier map	

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}
C1	0.6483 (2)	0.49883 (17)	0.57621 (17)	0.0725 (5)
H1	0.6643	0.5535	0.6266	0.087*
C2	0.7343 (2)	0.46784 (15)	0.50573 (16)	0.0599 (4)
H2	0.8076	0.5030	0.5079	0.072*
C3	0.71222 (15)	0.38413 (13)	0.43117 (13)	0.0455 (4)
C4	0.60095 (14)	0.33360 (13)	0.42953 (13)	0.0439 (4)
C5	0.51443 (18)	0.36705 (17)	0.49915 (16)	0.0601 (5)
H5	0.4401	0.3341	0.4965	0.072*
C6	0.5388 (2)	0.44896 (19)	0.57199 (17)	0.0749 (5)
H6	0.4807	0.4711	0.6190	0.090*
C7	0.80521 (16)	0.35124 (16)	0.35744 (16)	0.0541 (4)
H7	0.7922	0.2885	0.3149	0.065*
C8	0.38129 (14)	0.27847 (14)	0.23805 (13)	0.0441 (3)
C9	0.31891 (18)	0.18498 (16)	0.20531 (16)	0.0567 (5)
H9	0.3582	0.1211	0.1807	0.068*
C10	0.1974 (2)	0.18686 (19)	0.20932 (19)	0.0706 (6)
H10	0.1544	0.1243	0.1867	0.085*
C11	0.14047 (18)	0.2805 (2)	0.24649 (18)	0.0712 (6)
H11	0.0586	0.2811	0.2499	0.085*
C12	0.20264 (19)	0.37386 (19)	0.27891 (17)	0.0684 (6)
H12	0.1627	0.4371	0.3041	0.082*
C13	0.32368 (18)	0.37453 (15)	0.27450 (15)	0.0554 (4)
H13	0.3661	0.4381	0.2955	0.066*

C14	0.54399 (17)	0.13373 (14)	0.39808 (16)	0.0540 (4)
H14A	0.5015	0.0911	0.3453	0.065*
H14B	0.4919	0.1444	0.4573	0.065*
C15	0.65056 (17)	0.06720 (14)	0.43210 (14)	0.0493 (4)
C16	0.7056 (2)	−0.10263 (15)	0.51699 (17)	0.0665 (6)
H16A	0.7630	−0.0624	0.5592	0.080*
H16B	0.7459	−0.1367	0.4580	0.080*
C17	0.6456 (3)	−0.1912 (2)	0.5801 (3)	0.1116 (11)
H17A	0.6075	−0.1565	0.6389	0.167*
H17B	0.7026	−0.2453	0.6044	0.167*
H17C	0.5879	−0.2292	0.5378	0.167*
N1	0.57707 (12)	0.24388 (11)	0.35598 (11)	0.0472 (3)
O1	0.89698 (13)	0.40067 (14)	0.34903 (16)	0.0853 (5)
O2	0.58147 (12)	0.38549 (11)	0.21580 (11)	0.0643 (4)
O3	0.57026 (13)	0.18131 (14)	0.17302 (12)	0.0758 (4)
O4	0.75012 (13)	0.09354 (12)	0.41701 (15)	0.0768 (5)
O5	0.61509 (11)	−0.02585 (10)	0.48124 (11)	0.0592 (3)
S1	0.53632 (4)	0.27483 (4)	0.23629 (3)	0.04953 (12)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.1104 (14)	0.0522 (10)	0.0549 (11)	0.0040 (10)	−0.0043 (10)	−0.0089 (9)
C2	0.0728 (13)	0.0477 (9)	0.0593 (10)	−0.0020 (9)	−0.0145 (8)	0.0032 (8)
C3	0.0469 (9)	0.0410 (8)	0.0484 (8)	0.0041 (7)	−0.0058 (7)	0.0073 (7)
C4	0.0445 (9)	0.0421 (8)	0.0451 (8)	0.0043 (7)	−0.0039 (7)	0.0069 (7)
C5	0.0544 (11)	0.0629 (11)	0.0629 (11)	0.0072 (9)	0.0080 (9)	0.0063 (9)
C6	0.0906 (13)	0.0731 (12)	0.0611 (11)	0.0163 (12)	0.0180 (13)	−0.0021 (10)
C7	0.0456 (10)	0.0544 (10)	0.0624 (11)	0.0023 (9)	−0.0015 (8)	0.0070 (8)
C8	0.0436 (8)	0.0493 (8)	0.0395 (7)	0.0012 (7)	−0.0038 (6)	−0.0011 (8)
C9	0.0585 (12)	0.0463 (9)	0.0653 (12)	0.0026 (9)	−0.0056 (9)	−0.0044 (8)
C10	0.0562 (12)	0.0676 (12)	0.0879 (15)	−0.0114 (11)	−0.0087 (11)	0.0002 (11)
C11	0.0472 (10)	0.0903 (16)	0.0763 (14)	0.0025 (11)	0.0026 (10)	−0.0014 (13)
C12	0.0615 (12)	0.0784 (14)	0.0653 (12)	0.0238 (11)	−0.0038 (10)	−0.0146 (11)
C13	0.0579 (11)	0.0552 (10)	0.0532 (9)	0.0078 (8)	−0.0074 (8)	−0.0093 (8)
C14	0.0459 (10)	0.0443 (8)	0.0719 (10)	−0.0015 (8)	−0.0038 (9)	0.0079 (8)
C15	0.0518 (10)	0.0414 (8)	0.0546 (9)	−0.0014 (8)	−0.0071 (8)	−0.0024 (8)
C16	0.0828 (15)	0.0493 (9)	0.0674 (12)	0.0114 (10)	−0.0237 (11)	0.0029 (9)
C17	0.133 (3)	0.0727 (16)	0.129 (3)	−0.0176 (16)	−0.048 (2)	0.0442 (17)
N1	0.0434 (7)	0.0416 (7)	0.0567 (8)	−0.0004 (6)	−0.0075 (6)	0.0048 (6)
O1	0.0493 (8)	0.0925 (11)	0.1142 (13)	−0.0147 (8)	0.0139 (8)	−0.0077 (10)
O2	0.0613 (8)	0.0729 (9)	0.0587 (7)	−0.0149 (7)	−0.0028 (6)	0.0174 (7)
O3	0.0589 (9)	0.0942 (11)	0.0743 (9)	0.0085 (8)	0.0091 (7)	−0.0305 (8)
O4	0.0472 (8)	0.0625 (8)	0.1207 (13)	0.0017 (7)	−0.0119 (8)	0.0208 (9)
O5	0.0664 (8)	0.0460 (6)	0.0652 (8)	0.0016 (6)	−0.0078 (6)	0.0095 (6)
S1	0.0434 (2)	0.0569 (2)	0.0483 (2)	0.0004 (2)	0.00160 (19)	−0.00257 (19)

Geometric parameters (Å, º)

C1—C6	1.376 (3)	C11—C12	1.371 (3)
C1—C2	1.378 (3)	C11—H11	0.93
C1—H1	0.93	C12—C13	1.375 (3)
C2—C3	1.395 (3)	C12—H12	0.93
C2—H2	0.93	C13—H13	0.93
C3—C4	1.396 (2)	C14—N1	1.455 (2)
C3—C7	1.468 (2)	C14—C15	1.506 (3)
C4—C5	1.384 (2)	C14—H14A	0.97
C4—N1	1.442 (2)	C14—H14B	0.97
C5—C6	1.370 (3)	C15—O4	1.188 (2)
C5—H5	0.93	C15—O5	1.327 (2)
C6—H6	0.93	C16—O5	1.443 (2)
C7—O1	1.198 (2)	C16—C17	1.486 (3)
C7—H7	0.93	C16—H16A	0.97
C8—C9	1.375 (2)	C16—H16B	0.97
C8—C13	1.388 (2)	C17—H17A	0.96
C8—S1	1.7605 (16)	C17—H17B	0.96
C9—C10	1.380 (3)	C17—H17C	0.96
C9—H9	0.93	N1—S1	1.6419 (14)
C10—C11	1.365 (3)	O2—S1	1.4253 (13)
C10—H10	0.93	O3—S1	1.4209 (15)
C6—C1—C2	120.05 (19)	C13—C12—H12	119.8
C6—C1—H1	120.0	C12—C13—C8	118.67 (18)
C2—C1—H1	120.0	C12—C13—H13	120.7
C1—C2—C3	120.5 (2)	C8—C13—H13	120.7
C1—C2—H2	119.7	N1—C14—C15	111.36 (15)
C3—C2—H2	119.7	N1—C14—H14A	109.4
C2—C3—C4	118.28 (17)	C15—C14—H14A	109.4
C2—C3—C7	119.84 (17)	N1—C14—H14B	109.4
C4—C3—C7	121.88 (16)	C15—C14—H14B	109.4
C5—C4—C3	120.73 (16)	H14A—C14—H14B	108.0
C5—C4—N1	119.76 (16)	O4—C15—O5	125.58 (17)
C3—C4—N1	119.50 (15)	O4—C15—C14	125.54 (16)
C6—C5—C4	119.74 (19)	O5—C15—C14	108.88 (15)
C6—C5—H5	120.1	O5—C16—C17	106.6 (2)
C4—C5—H5	120.1	O5—C16—H16A	110.4
C5—C6—C1	120.6 (2)	C17—C16—H16A	110.4
C5—C6—H6	119.7	O5—C16—H16B	110.4
C1—C6—H6	119.7	C17—C16—H16B	110.4
O1—C7—C3	123.7 (2)	H16A—C16—H16B	108.6
O1—C7—H7	118.2	C16—C17—H17A	109.5
C3—C7—H7	118.2	C16—C17—H17B	109.5
C9—C8—C13	120.88 (16)	H17A—C17—H17B	109.5
C9—C8—S1	119.43 (13)	C16—C17—H17C	109.5
C13—C8—S1	119.68 (14)	H17A—C17—H17C	109.5

C8—C9—C10	119.37 (18)	H17B—C17—H17C	109.5
C8—C9—H9	120.3	C4—N1—C14	117.44 (14)
C10—C9—H9	120.3	C4—N1—S1	119.99 (10)
C11—C10—C9	119.9 (2)	C14—N1—S1	118.12 (12)
C11—C10—H10	120.0	C15—O5—C16	116.89 (15)
C9—C10—H10	120.0	O3—S1—O2	120.46 (10)
C10—C11—C12	120.7 (2)	O3—S1—N1	106.47 (9)
C10—C11—H11	119.7	O2—S1—N1	105.88 (8)
C12—C11—H11	119.7	O3—S1—C8	107.29 (9)
C11—C12—C13	120.43 (19)	O2—S1—C8	109.85 (9)
C11—C12—H12	119.8	N1—S1—C8	105.97 (8)
C6—C1—C2—C3	−1.4 (3)	N1—C14—C15—O5	172.45 (14)
C1—C2—C3—C4	0.5 (3)	C5—C4—N1—C14	−59.0 (2)
C1—C2—C3—C7	−179.31 (18)	C3—C4—N1—C14	119.59 (16)
C2—C3—C4—C5	0.9 (2)	C5—C4—N1—S1	96.88 (17)
C7—C3—C4—C5	−179.28 (16)	C3—C4—N1—S1	−84.51 (17)
C2—C3—C4—N1	−177.66 (14)	C15—C14—N1—C4	−81.22 (19)
C7—C3—C4—N1	2.1 (2)	C15—C14—N1—S1	122.42 (14)
C3—C4—C5—C6	−1.4 (3)	O4—C15—O5—C16	−1.6 (3)
N1—C4—C5—C6	177.22 (17)	C14—C15—O5—C16	178.12 (15)
C4—C5—C6—C1	0.4 (3)	C17—C16—O5—C15	173.12 (19)
C2—C1—C6—C5	1.0 (3)	C4—N1—S1—O3	154.32 (14)
C2—C3—C7—O1	−8.5 (3)	C14—N1—S1—O3	−49.95 (15)
C4—C3—C7—O1	171.70 (19)	C4—N1—S1—O2	24.99 (16)
C13—C8—C9—C10	−0.3 (3)	C14—N1—S1—O2	−179.28 (13)
S1—C8—C9—C10	178.31 (17)	C4—N1—S1—C8	−91.66 (14)
C8—C9—C10—C11	−0.6 (3)	C14—N1—S1—C8	64.07 (15)
C9—C10—C11—C12	0.8 (3)	C9—C8—S1—O3	15.45 (17)
C10—C11—C12—C13	0.0 (3)	C13—C8—S1—O3	−165.94 (15)
C11—C12—C13—C8	−0.9 (3)	C9—C8—S1—O2	148.05 (15)
C9—C8—C13—C12	1.0 (3)	C13—C8—S1—O2	−33.34 (17)
S1—C8—C13—C12	−177.55 (16)	C9—C8—S1—N1	−98.00 (15)
N1—C14—C15—O4	−7.8 (3)	C13—C8—S1—N1	80.61 (16)

Hydrogen-bond geometry (Å, º)

D—H···A	D—H	H···A	D···A	D—H···A
C5—H5···O4[i]	0.93	2.57	3.220 (2)	127
C16—H16B···Cg1[ii]	0.97	2.75	3.615 (2)	150

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