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N-(Phenylsulfonyl)naphtho[2,1-b]furan-1-carboxamide

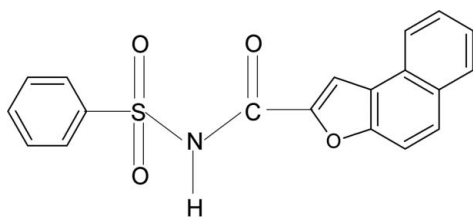
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 Key indicators: single-crystal X-ray study; $T = 299$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.041; wR factor = 0.133; data-to-parameter ratio = 12.9.

In the title compound, $\text{C}_{19}\text{H}_{13}\text{NO}_4\text{S}$, the molecule is twisted at the S atom with a C—S—N—C torsion angle of -65.2 (2)° between the benzene ring and the $-\text{SO}_2-\text{NH}-\text{C}=\text{O}$ segment. The dihedral angle between the benzene and the naphthofuran ring system is 83.3 (1)°. In the crystal, molecules are linked by N—H...O hydrogen bonds into chains running along the c axis. An intramolecular N—H...O(furan) interaction is also observed.

Related literature

 For related structures, see: Gowda *et al.* (2009, 2010).


Experimental

Crystal data

 $\text{C}_{19}\text{H}_{13}\text{NO}_4\text{S}$
 $M_r = 351.36$
 Monoclinic, $P2_1/c$
 $a = 13.8504$ (10) Å

 $b = 12.2166$ (8) Å
 $c = 9.7164$ (6) Å
 $\beta = 101.248$ (2)°
 $V = 1612.48$ (19) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.23$ mm⁻¹
 $T = 299$ K
 $0.35 \times 0.3 \times 0.25$ mm

Data collection

 Bruker APEXII CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.924$, $T_{\max} = 0.945$
 15289 measured reflections
 2986 independent reflections
 2394 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.133$
 $S = 0.85$
 2986 reflections
 231 parameters
 1 restraint

 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.25$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.38$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
N1—H1N...O4	0.79 (3)	2.33 (3)	2.653 (2)	106 (2)
N1—H1N...O2 ⁱ	0.79 (3)	2.66 (1)	3.057 (2)	113 (2)

 Symmetry code: (i) $x, -y + \frac{1}{2}, z + \frac{1}{2}$.

Data collection: APEX2 (Bruker, 2004); cell refinement: APEX2 and SAINT-Plus (Bruker, 2004); data reduction: SAINT-Plus and XPREP (Bruker, 2004); 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.

The authors acknowledge Dr K. Gunasekaran and Jagadeesan CAS in Crystallography and Biophysics, University of Madras, and Dr H. C. Devarajegowda, Yuvaraja's College, Mysore, for useful discussions. The Department of Chemistry, IIT Madras, is acknowledged for the data collection.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DS2146).

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supporting information

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***N*-(Phenylsulfonyl)naphtho[2,1-*b*]furan-1-carboxamide**

M. Shetprakash, P. A. Suchetan, B. S. Palakshamurthy, K. M. Mahadevan and V. P. Vaidya

S1. Comment

Aryl Acyl sulfonamides are known as a potent antitumor agent against a broad spectrum of human tumor xenografts in nude mice. Further, the title compound exhibits antibacterial and antifungal activities (our unpublished results). In order to study the effect of the ring substituents on the solid-state structures of *N*-naphthofuroyl-sulfonamides, in the present work the structure of *N*-(Naphthofuroyl)benzenesulfonamide has been determined. The title compound (I) crystallizes in Monoclinic $P2_1/c$ space group compared to *N*-(benzoyl)benzenesulfonamide (II) (Gowda *et al.*, 2009) and *N*-(Phenylsulfonyl)acetamide (III) (Gowda *et al.*, 2010) which crystallizes in Triclinic P-1 and Tetragonal $P4_3$ space groups respectively. In III, the packing of molecules is linked by N—H \cdots O(C) hydrogen bonds and in II by N—H \cdots O(S) bonds, whereas in I, the molecules are linked by intermolecular N—H \cdots O(S) hydrogen bonds. Intramolecular C—H \cdots O(S) and N—H \cdots O(furan) interactions are also observed. The molecules are twisted at S atoms with the C—S(O2)—NH—C(O) torsion angle of -65.2 (2) $^\circ$, compared to the values of -66.9 (3) $^\circ$ in (II) and -58.8 (4) $^\circ$ in (III). The dihedral angle between the benzene ring and the naphthofuran ring in (I) is 83.3 (1) $^\circ$, compared to 80.3 (1) $^\circ$ observed between the two benzene rings in (II) and 38.7 (0) $^\circ$ in (III) between the benzene ring and the mean plane of CH3 fragment respectively. The packing of the molecules *via* intermolecular N—H \cdots O(S) hydrogen bonds is shown in Fig. 2.

S2. Experimental

The title compound was prepared by refluxing a mixture of naphthofuran-2-carboxylic acid (10 mmol), benzenesulfonamide (10 mmol) and phosphorous oxychloride for 1 h on a water bath. The resultant mixture was cooled and poured into ice cold water. The solid, *N*-(Naphthofuroyl)benzenesulfonamide obtained was filtered, washed thoroughly with water and then dissolved in sodium bicarbonate solution. The compound was later reprecipitated by acidifying the filtered solution with dilute hydrochloric acid. The filtered and dried compound was recrystallized to constant melting point. The compound was characterized by its characteristic carbonyl C=O stretching (1698.2 cm $^{-1}$), N—H stretching (3233.1 cm $^{-1}$), symmetric SO₂ (1173.3 cm $^{-1}$) and asymmetric SO₂ (1326.2 cm $^{-1}$) infrared absorption frequencies. Single crystals suitable for *x*-ray diffraction were grown from a slow evaporation of its ethanolic solution at room temperature.

S3. Refinement

The H atom of the NH group was located in a difference map and later restrained to N—H = 0.86 (1) $\%$ A. The other H atoms were positioned with idealized geometry using a riding model with C—H = 0.93 Å. All H atoms were refined with isotropic displacement parameters (set to 1.2 times of the U_{eq} of the parent atom).

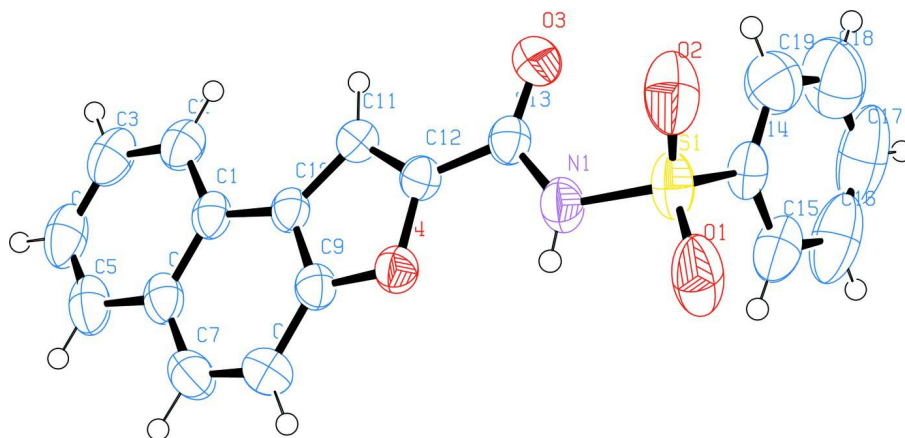


Figure 1

Molecular structure of the title compound, showing the atom-labelling scheme.

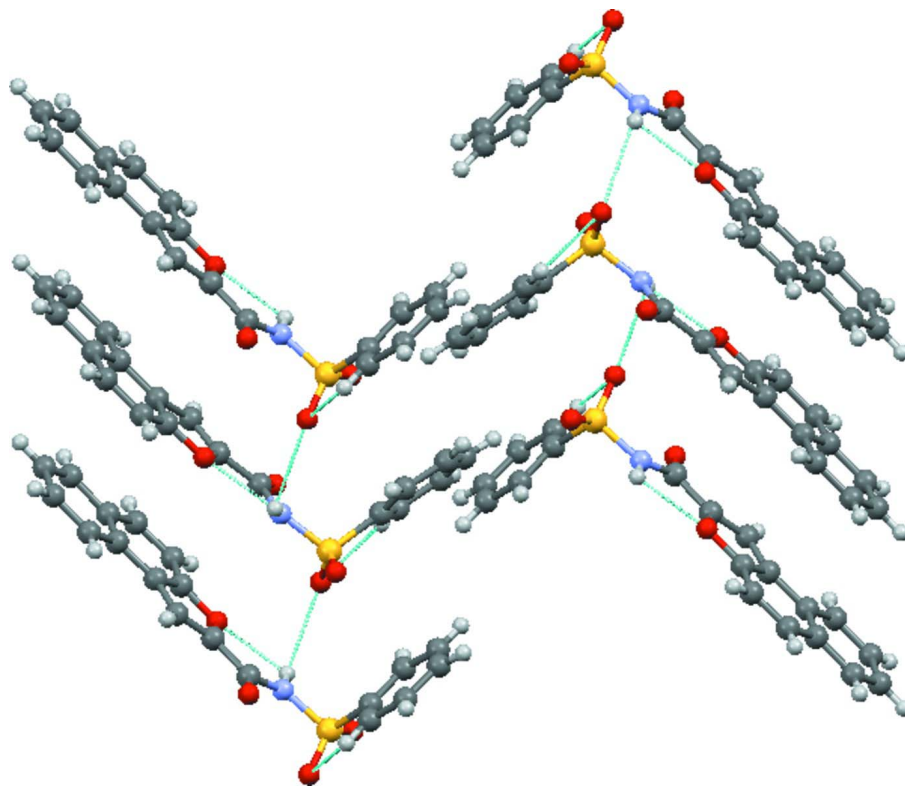


Figure 2

Molecular packing in the title compound. Hydrogen bonds are shown as dashed lines.

N-(Phenylsulfonyl)naphtho[2,1-*b*]furan-1-carboxamide

Crystal data

$C_{19}H_{13}NO_4S$

$M_r = 351.36$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 13.8504$ (10) Å

$b = 12.2166$ (8) Å

$c = 9.7164$ (6) Å

$\beta = 101.248$ (2)°

$V = 1612.48$ (19) Å³

$Z = 4$

$F(000) = 728$
 1.447 Mg m⁻³
 $D_x = 1.447 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 2986 reflections

$\theta = 2.2^\circ$
 $\mu = 0.23 \text{ mm}^{-1}$
 $T = 299 \text{ K}$
 Prism, colourless
 $0.35 \times 0.3 \times 0.25 \text{ mm}$

Data collection

Bruker APEXII CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 phi and scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.924$, $T_{\max} = 0.945$

15289 measured reflections
 2986 independent reflections
 2394 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.027$
 $\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 2.9^\circ$
 $h = -16 \rightarrow 16$
 $k = -14 \rightarrow 14$
 $l = -11 \rightarrow 11$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.133$
 $S = 0.85$
 2986 reflections
 231 parameters
 1 restraint
 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
 $w = 1/[\sigma^2(F_o^2) + (0.0827P)^2 + 1.1854P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.38 \text{ e \AA}^{-3}$
 Extinction correction: SHELXL
 Extinction coefficient: 0.0054 (12)

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
H1N	0.8221 (19)	0.1816 (19)	0.346 (3)	0.053 (8)*
S1	0.72588 (4)	0.25754 (5)	0.18450 (6)	0.0566 (2)
N1	0.82025 (14)	0.24177 (18)	0.3158 (2)	0.0527 (5)
O1	0.69636 (15)	0.14945 (17)	0.1467 (2)	0.0910 (7)
O2	0.75502 (14)	0.3299 (2)	0.08507 (18)	0.0853 (7)
O3	0.85990 (12)	0.42070 (13)	0.35980 (18)	0.0626 (5)
O4	0.96215 (10)	0.17651 (11)	0.52390 (14)	0.0448 (4)
C1	1.14610 (14)	0.26862 (18)	0.8155 (2)	0.0430 (5)
C2	1.18623 (16)	0.3662 (2)	0.8774 (2)	0.0545 (6)
H2	1.1644	0.4332	0.8376	0.065*

C3	1.25770 (18)	0.3628 (3)	0.9968 (2)	0.0663 (7)
H3	1.2841	0.4278	1.0377	0.08*
C4	1.29114 (18)	0.2631 (3)	1.0575 (3)	0.0703 (8)
H4	1.3395	0.2621	1.1387	0.084*
C5	1.25403 (17)	0.1678 (3)	0.9995 (2)	0.0636 (7)
H5	1.2771	0.102	1.0416	0.076*
C6	1.18038 (15)	0.16646 (19)	0.8755 (2)	0.0488 (5)
C7	1.14042 (17)	0.0666 (2)	0.8153 (2)	0.0566 (6)
H7	1.1644	0.0011	0.8573	0.068*
C8	1.06823 (17)	0.06330 (19)	0.6984 (2)	0.0550 (6)
H8	1.0426	-0.0025	0.6593	0.066*
C9	1.03480 (14)	0.16388 (17)	0.6404 (2)	0.0425 (5)
C10	1.06976 (14)	0.26419 (16)	0.6922 (2)	0.0395 (4)
C11	1.01546 (14)	0.34374 (17)	0.6020 (2)	0.0412 (4)
H11	1.0225	0.4194	0.609	0.049*
C12	0.95188 (13)	0.28806 (17)	0.5044 (2)	0.0404 (4)
C13	0.87507 (14)	0.32539 (18)	0.3889 (2)	0.0430 (5)
C14	0.63289 (15)	0.3220 (2)	0.2518 (2)	0.0502 (5)
C16	0.4993 (2)	0.3091 (5)	0.3712 (4)	0.1040 (13)
H16	0.4601	0.2673	0.4186	0.125*
C15	0.57551 (19)	0.2612 (3)	0.3241 (3)	0.0725 (8)
H15	0.5888	0.1872	0.3409	0.087*
C19	0.6169 (2)	0.4313 (3)	0.2335 (4)	0.0876 (9)
H19	0.6567	0.4741	0.1882	0.105*
C17	0.4807 (3)	0.4141 (5)	0.3499 (5)	0.1213 (18)
H17	0.4271	0.445	0.3801	0.146*
C18	0.5377 (3)	0.4776 (4)	0.2854 (5)	0.1210 (15)
H18	0.5249	0.5522	0.275	0.145*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0409 (3)	0.0788 (5)	0.0460 (3)	0.0105 (3)	-0.0012 (2)	-0.0159 (3)
N1	0.0427 (10)	0.0551 (13)	0.0552 (11)	0.0046 (8)	-0.0028 (8)	-0.0053 (9)
O1	0.0699 (12)	0.0851 (14)	0.1006 (15)	0.0202 (10)	-0.0264 (11)	-0.0464 (12)
O2	0.0582 (10)	0.154 (2)	0.0440 (9)	0.0078 (12)	0.0119 (8)	0.0091 (11)
O3	0.0571 (9)	0.0564 (10)	0.0672 (10)	-0.0011 (8)	-0.0051 (8)	0.0136 (8)
O4	0.0406 (7)	0.0441 (8)	0.0477 (8)	0.0006 (6)	0.0034 (6)	-0.0032 (6)
C1	0.0329 (10)	0.0591 (13)	0.0383 (10)	-0.0002 (8)	0.0104 (8)	0.0039 (9)
C2	0.0465 (12)	0.0679 (15)	0.0478 (12)	-0.0094 (10)	0.0056 (9)	0.0003 (10)
C3	0.0515 (13)	0.092 (2)	0.0534 (13)	-0.0177 (13)	0.0049 (11)	-0.0072 (13)
C4	0.0443 (13)	0.118 (2)	0.0443 (13)	-0.0070 (14)	-0.0007 (10)	0.0093 (14)
C5	0.0421 (12)	0.098 (2)	0.0498 (13)	0.0085 (13)	0.0078 (10)	0.0199 (13)
C6	0.0367 (10)	0.0675 (14)	0.0442 (11)	0.0060 (9)	0.0127 (9)	0.0108 (10)
C7	0.0541 (13)	0.0563 (14)	0.0599 (13)	0.0134 (11)	0.0124 (11)	0.0145 (11)
C8	0.0574 (13)	0.0458 (13)	0.0615 (13)	0.0057 (10)	0.0104 (11)	0.0033 (10)
C9	0.0364 (10)	0.0466 (11)	0.0444 (10)	0.0023 (8)	0.0082 (8)	0.0011 (9)
C10	0.0319 (9)	0.0480 (11)	0.0392 (10)	0.0002 (8)	0.0084 (8)	0.0028 (8)

C11	0.0387 (10)	0.0414 (11)	0.0436 (10)	-0.0016 (8)	0.0082 (8)	0.0009 (8)
C12	0.0350 (9)	0.0430 (11)	0.0441 (10)	0.0011 (8)	0.0096 (8)	0.0012 (8)
C13	0.0360 (10)	0.0519 (13)	0.0413 (10)	0.0003 (9)	0.0084 (8)	0.0004 (9)
C14	0.0348 (10)	0.0639 (14)	0.0491 (11)	0.0011 (9)	0.0009 (9)	-0.0112 (10)
C16	0.0497 (17)	0.190 (4)	0.075 (2)	-0.017 (2)	0.0200 (15)	-0.028 (3)
C15	0.0501 (14)	0.102 (2)	0.0630 (15)	-0.0128 (13)	0.0048 (12)	-0.0044 (14)
C19	0.0656 (17)	0.072 (2)	0.124 (3)	0.0113 (14)	0.0167 (17)	-0.0006 (18)
C17	0.0517 (18)	0.186 (5)	0.124 (3)	0.007 (3)	0.011 (2)	-0.077 (3)
C18	0.091 (3)	0.094 (3)	0.170 (4)	0.035 (2)	0.007 (3)	-0.037 (3)

Geometric parameters (Å, °)

S1—O1	1.410 (2)	C7—C8	1.360 (3)
S1—O2	1.425 (2)	C7—H7	0.93
S1—N1	1.650 (2)	C8—C9	1.393 (3)
S1—C14	1.742 (2)	C8—H8	0.93
N1—C13	1.383 (3)	C9—C10	1.376 (3)
N1—H1N	0.79 (2)	C10—C11	1.422 (3)
O3—C13	1.207 (2)	C11—C12	1.346 (3)
O4—C9	1.369 (2)	C11—H11	0.93
O4—C12	1.379 (2)	C12—C13	1.461 (3)
C1—C2	1.400 (3)	C14—C19	1.360 (4)
C1—C6	1.419 (3)	C14—C15	1.377 (3)
C1—C10	1.436 (3)	C16—C17	1.316 (6)
C2—C3	1.371 (3)	C16—C15	1.362 (5)
C2—H2	0.93	C16—H16	0.93
C3—C4	1.393 (4)	C15—H15	0.93
C3—H3	0.93	C19—C18	1.412 (5)
C4—C5	1.350 (4)	C19—H19	0.93
C4—H4	0.93	C17—C18	1.346 (6)
C5—C6	1.419 (3)	C17—H17	0.93
C5—H5	0.93	C18—H18	0.93
C6—C7	1.418 (3)		
O1—S1—O2	120.71 (14)	O4—C9—C10	110.56 (17)
O1—S1—N1	103.77 (12)	O4—C9—C8	124.53 (19)
O2—S1—N1	108.08 (11)	C10—C9—C8	124.91 (19)
O1—S1—C14	108.85 (12)	C9—C10—C11	106.10 (17)
O2—S1—C14	107.57 (12)	C9—C10—C1	119.19 (18)
N1—S1—C14	107.12 (10)	C11—C10—C1	134.70 (19)
C13—N1—S1	125.68 (18)	C12—C11—C10	106.48 (18)
C13—N1—H1N	121.3 (19)	C12—C11—H11	126.8
S1—N1—H1N	111.1 (19)	C10—C11—H11	126.8
C9—O4—C12	105.34 (15)	C11—C12—O4	111.50 (17)
C2—C1—C6	119.94 (19)	C11—C12—C13	131.45 (19)
C2—C1—C10	123.80 (19)	O4—C12—C13	117.01 (17)
C6—C1—C10	116.25 (19)	O3—C13—N1	122.6 (2)
C3—C2—C1	119.9 (2)	O3—C13—C12	123.26 (19)

C3—C2—H2	120.1	N1—C13—C12	114.10 (19)
C1—C2—H2	120.1	C19—C14—C15	120.1 (2)
C2—C3—C4	120.7 (2)	C19—C14—S1	120.5 (2)
C2—C3—H3	119.7	C15—C14—S1	119.4 (2)
C4—C3—H3	119.7	C17—C16—C15	120.4 (4)
C5—C4—C3	120.6 (2)	C17—C16—H16	119.8
C5—C4—H4	119.7	C15—C16—H16	119.8
C3—C4—H4	119.7	C16—C15—C14	120.1 (4)
C4—C5—C6	121.1 (2)	C16—C15—H15	119.9
C4—C5—H5	119.4	C14—C15—H15	119.9
C6—C5—H5	119.4	C14—C19—C18	117.7 (3)
C1—C6—C5	117.8 (2)	C14—C19—H19	121.2
C1—C6—C7	120.97 (19)	C18—C19—H19	121.2
C5—C6—C7	121.2 (2)	C16—C17—C18	121.4 (4)
C8—C7—C6	122.3 (2)	C16—C17—H17	119.3
C8—C7—H7	118.8	C18—C17—H17	119.3
C6—C7—H7	118.8	C17—C18—C19	120.2 (4)
C7—C8—C9	116.4 (2)	C17—C18—H18	119.9
C7—C8—H8	121.8	C19—C18—H18	119.9
C9—C8—H8	121.8		
O1—S1—N1—C13	179.8 (2)	C6—C1—C10—C11	179.5 (2)
O2—S1—N1—C13	50.5 (2)	C9—C10—C11—C12	0.6 (2)
C14—S1—N1—C13	-65.2 (2)	C1—C10—C11—C12	-178.9 (2)
C6—C1—C2—C3	-0.8 (3)	C10—C11—C12—O4	-0.9 (2)
C10—C1—C2—C3	178.7 (2)	C10—C11—C12—C13	177.01 (19)
C1—C2—C3—C4	0.1 (4)	C9—O4—C12—C11	0.8 (2)
C2—C3—C4—C5	0.2 (4)	C9—O4—C12—C13	-177.41 (15)
C3—C4—C5—C6	0.2 (4)	S1—N1—C13—O3	-2.0 (3)
C2—C1—C6—C5	1.1 (3)	S1—N1—C13—C12	177.46 (15)
C10—C1—C6—C5	-178.44 (17)	C11—C12—C13—O3	3.4 (3)
C2—C1—C6—C7	179.6 (2)	O4—C12—C13—O3	-178.81 (18)
C10—C1—C6—C7	0.0 (3)	C11—C12—C13—N1	-176.1 (2)
C4—C5—C6—C1	-0.8 (3)	O4—C12—C13—N1	1.7 (2)
C4—C5—C6—C7	-179.3 (2)	O1—S1—C14—C19	-147.8 (2)
C1—C6—C7—C8	0.1 (3)	O2—S1—C14—C19	-15.4 (2)
C5—C6—C7—C8	178.5 (2)	N1—S1—C14—C19	100.6 (2)
C6—C7—C8—C9	-0.3 (3)	O1—S1—C14—C15	32.2 (2)
C12—O4—C9—C10	-0.4 (2)	O2—S1—C14—C15	164.60 (19)
C12—O4—C9—C8	179.38 (19)	N1—S1—C14—C15	-79.4 (2)
C7—C8—C9—O4	-179.39 (19)	C17—C16—C15—C14	-1.1 (5)
C7—C8—C9—C10	0.4 (3)	C19—C14—C15—C16	3.4 (4)
O4—C9—C10—C11	-0.1 (2)	S1—C14—C15—C16	-176.6 (2)
C8—C9—C10—C11	-179.9 (2)	C15—C14—C19—C18	-2.7 (4)
O4—C9—C10—C1	179.47 (15)	S1—C14—C19—C18	177.3 (3)
C8—C9—C10—C1	-0.3 (3)	C15—C16—C17—C18	-2.0 (6)
C2—C1—C10—C9	-179.43 (19)	C16—C17—C18—C19	2.7 (7)
C6—C1—C10—C9	0.1 (3)	C14—C19—C18—C17	-0.3 (6)

C2—C1—C10—C11 0.0 (3)

Hydrogen-bond geometry (Å, °)

<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—H1N \cdots O4	0.79 (3)	2.33 (3)	2.653 (2)	106 (2)
C19—H19 \cdots O2	0.93	2.55	2.890 (4)	102
N1—H1N \cdots O2 ⁱ	0.79 (3)	2.66 (1)	3.057 (2)	113 (2)

Symmetry code: (i) $x, -y+1/2, z+1/2$.