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N-Benzyl-5-(dimethylamino)-naphthalene-1-sulfonamide

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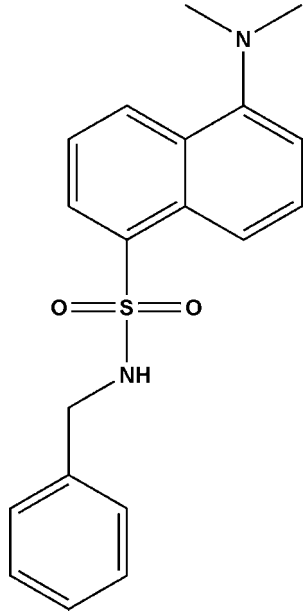
Key indicators: single-crystal X-ray study; T = 173 K; mean σ(C–C) = 0.002 Å; R factor = 0.036; wR factor = 0.096; data-to-parameter ratio = 19.2.

The structure of the title compound, $C_{19}H_{20}N_2O_2S$, displays intermolecular N—H···O hydrogen bonding, which generates inversion dimers. There is no π–π stacking in the crystal structure. The dihedral angle between the phenyl ring and naphthalene ring system is 59.16 (11)°.

Related literature

For the use of dansyl fluorescent analogs as insecticides and synergists, see: Himel *et al.* (1971). Dansyl probes have also been covalently incorporated into a variety of polymeric networks, see: Shea *et al.* (1989). Dansyl chromophoric compounds have been investigated for intramolecular energy transfer in aromatic ring systems, see: Schael *et al.* (1998) and for host–guest interations shown by fluoresence studies of dansyl-labelled calix[6]arene, see: Schonefeld *et al.* (2006). For related structures, see: Illos *et al.* (2005); Hongmei *et al.* (2009); Hong-Wei *et al.* (2009); Chui *et al.* (2010).



Experimental

Crystal data

$C_{19}H_{20}N_2O_2S$
M_r = 340.43
Monoclinic, $C2/c$
a = 16.6635 (5) Å
b = 9.5722 (2) Å
c = 22.8942 (7) Å
β = 108.779 (1)°
V = 3457.38 (16) Å^3
Z = 8
Mo $K\alpha$ radiation
μ = 0.20 mm^{-1}
T = 173 K
0.30 × 0.24 × 0.22 mm

Data collection

Nonius KappaCCD diffractometer
4275 measured reflections
4275 independent reflections
3747 reflections with $I > 2\sigma(I)$

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.036
$wR(F^2)$ = 0.096
S = 1.03
4275 reflections
223 parameters
1 restraint
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.37 e Å^{-3}
$\Delta\rho_{min}$ = −0.40 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O2^{i}	0.86 (2)	2.12 (2)	2.9351 (14)	158 (2)

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

Data collection: *COLLECT* (Nonius, 2000); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *SHELXL97*.

The authors wish to thank Dr Hong Su from the University of the Cape Town for her assistance with the data collection and refinement.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG5080).

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supporting information

Acta Cryst. (2011). E**67**, o2458–o2459 [doi:10.1107/S1600536811033083]

N-Benzyl-5-(dimethylamino)naphthalene-1-sulfonamide

Pralav Bhatt, Thavendran Govender, Hendrik G. Kruger and Glenn E. M. Maguire

S1. Comment

Dansyl fluorescent analogs have been reported as insecticides and synergists (Himel *et al.*,1971*).* Dansyl probes have also been covalently incorporated into a variety of polymeric networks (Shea *et al.*,1989). Dansyl chromophoric compounds were investigated for intramolecular energy transfer in aromatic ring systems (Schael *et al.*,1998). Dansyl labelled calix[6]arene are reported to show host–guest interations using fluoresence studies (Schonefeld *et al.*, 2006).

The title compound is a novel benzylated dansyl derivative (Fig. 1.).

There are a number of examples in literature where amino-sulfonamides dansyl structures have shown intermolecular hydrogen bonding. These arrangments can be described in two broad categories. First, where hydrogen bonds occur between the sulfonyl oxygen and the nitrogen of two adjacent molecules in a alternating chain arrangement (Hongmei *et al.*, 2009, Chui *et al.*,2010). Second, where they interact with an adjacent molecule in a head to tail manner, (Illos *et al.*, 2005, Hong-Wei *et al.*, 2009). Our system falls into the latter category. Our structure thus displays N1—H1···O2, 2.9351 (14) Å intermolecular hydrogen bonding, generating inversion dimers (Fig. 2). There is no π–π stacking in the crystal.

S2. Experimental

To a dry THF 5 ml benzyl amine (107 mg, 1 m*M*) was added triethyl amine (303 mg, 3 m*M*). Dansyl chloride (269 mg, 1 m*M*) was then added and the resulting solution was stirred until the reaction was completed (TLC R_f = 0.27 in 60% ethyl acetate/hexane). The reaction contents were filtered. The filtrate was evaporated under reduced pressure yielding a yellow oil. To this residue was added 20 ml of dichloromethane and then the organic layer was washed with water and then separated. After drying over anhydrous magnesium sulfate the solvent was evaporated once again under reduced pressure to yield a yellow crystalline solid (240 mg, 71%). *M*.p. = 408 K.

Crystals suitable for X-ray analysis were grown in ethyl acetate/hexane at room temperature.

S3. Refinement

All non-hydrogen atoms were refined anisotropically. All hydrogen atoms, except H1 on N1, were placed in idealized positions in a riding model and refined with U_{iso} set at 1.2 or 1.5 times those of their parent atoms. The position of H1 was located in the difference electron density map and refined with bond length constraint d(N—H) = 0.88 (2) Å.

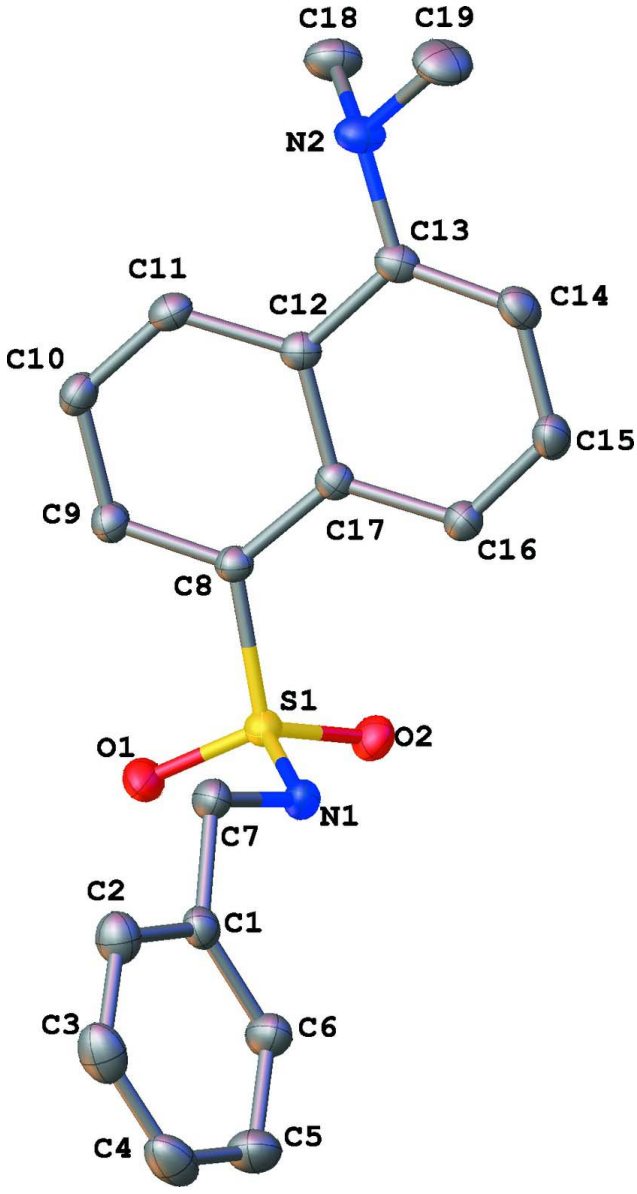

Figure 1

The molecular structure of the title compound with atomic numbering scheme. The H atoms have been omitted for clarity. Displacement elipsoids are drawn at 40% probability.

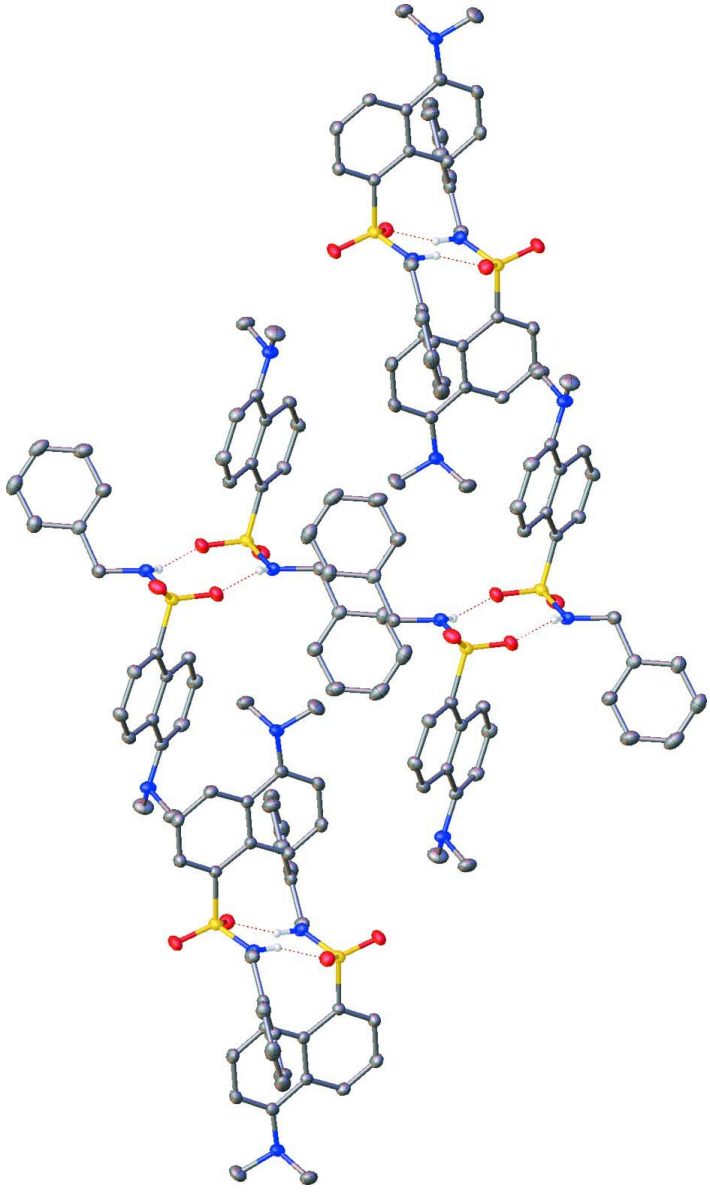

Figure 2

The hydrogen bonding interactions of the title compound along the [110] axis. All H atoms except those involved in hydrogen bonding interactions have been omitted for clarity.

N-Benzyl-5-(dimethylamino)naphthalene-1-sulfonamide

Crystal data

$C_{19}H_{20}N_2O_2S$
$M_r = 340.43$
Monoclinic, $C2/c$
Hall symbol: -C 2yc
$a = 16.6635\ (5)$ Å
$b = 9.5722\ (2)$ Å
$c = 22.8942\ (7)$ Å
$\beta = 108.779\ (1)°$
$V = 3457.38\ (16)$ Å^3
$Z = 8$
$F(000) = 1440$
$D_x = 1.308$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 4275 reflections
$\theta = 2.5–28.3°$
$\mu = 0.20$ mm^{-1}

T = 173 K
Block, colourless

0.30 × 0.24 × 0.22 mm

Data collection

Nonius KappaCCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
1.2° φ scans and ω scans
4275 measured reflections
4275 independent reflections

3747 reflections with $I > 2\sigma(I)$
R_{int} = 0.000
θ_{max} = 28.3°, θ_{min} = 2.5°
h = 0→22
k = 0→12
l = −30→28

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)]$ = 0.036
$wR(F^2)$ = 0.096
S = 1.03
4275 reflections
223 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
$w = 1/[\sigma^2(F_o^2) + (0.0451P)^2 + 3.0274P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} < 0.001$
$\Delta\rho_{max}$ = 0.37 e Å^{-3}
$\Delta\rho_{min}$ = −0.40 e Å^{-3}

Special details

Experimental. Half sphere of data collected using *COLLECT* strategy (Nonius, 2000). Crystal to detector distance = 40 mm; combination of φ and ω scans of 1.0°, 60 s per °, 2 iterations.

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 (Å^2)

	x	y	z	U_{iso}*/U_{eq}
S1	0.20046 (2)	0.47743 (3)	0.473136 (13)	0.02444 (9)
O1	0.18380 (7)	0.60223 (10)	0.50185 (4)	0.0342 (2)
O2	0.16204 (6)	0.34869 (10)	0.48342 (4)	0.0307 (2)
N1	0.30155 (7)	0.45093 (11)	0.49739 (5)	0.0276 (2)
H1	0.3140 (12)	0.3655 (16)	0.4932 (8)	0.050 (5)*
N2	0.12696 (7)	0.39094 (13)	0.17747 (5)	0.0300 (2)
C1	0.43601 (8)	0.58023 (13)	0.54635 (6)	0.0268 (3)
C2	0.49467 (9)	0.68229 (15)	0.54407 (7)	0.0348 (3)
H2	0.4855	0.7356	0.5075	0.042*
C3	0.56607 (9)	0.70685 (17)	0.59432 (8)	0.0424 (4)
H3	0.6051	0.7775	0.5924	0.051*
C4	0.58056 (10)	0.62813 (18)	0.64761 (8)	0.0435 (4)

H4	0.6299	0.6438	0.6820	0.052*
C5	0.52285 (10)	0.52681 (18)	0.65038 (7)	0.0426 (4)
H5	0.5326	0.4730	0.6869	0.051*
C6	0.45021 (9)	0.50297 (15)	0.59985 (7)	0.0348 (3)
H6	0.4106	0.4337	0.6022	0.042*
C7	0.35869 (8)	0.56091 (13)	0.49002 (6)	0.0286 (3)
H7A	0.3271	0.6502	0.4809	0.034*
H7B	0.3774	0.5378	0.4543	0.034*
C8	0.16952 (7)	0.51250 (12)	0.39278 (5)	0.0221 (2)
C9	0.14023 (8)	0.64467 (13)	0.37456 (6)	0.0262 (2)
H9	0.1360	0.7110	0.4043	0.031*
C10	0.11642 (8)	0.68209 (13)	0.31179 (6)	0.0286 (3)
H10	0.0963	0.7738	0.2993	0.034*
C11	0.12215 (8)	0.58726 (13)	0.26895 (6)	0.0263 (3)
H11	0.1070	0.6146	0.2269	0.032*
C12	0.15032 (7)	0.44820 (13)	0.28577 (5)	0.0226 (2)
C13	0.15527 (7)	0.34835 (14)	0.24015 (5)	0.0250 (2)
C14	0.18974 (8)	0.21871 (14)	0.25901 (6)	0.0295 (3)
H14	0.1961	0.1540	0.2293	0.035*
C15	0.21569 (8)	0.18099 (14)	0.32193 (6)	0.0301 (3)
H15	0.2390	0.0908	0.3339	0.036*
C16	0.20798 (8)	0.27152 (13)	0.36610 (6)	0.0261 (2)
H16	0.2239	0.2427	0.4080	0.031*
C17	0.17616 (7)	0.40859 (12)	0.34938 (5)	0.0217 (2)
C18	0.03465 (9)	0.40632 (18)	0.15059 (7)	0.0391 (3)
H18A	0.0086	0.3139	0.1402	0.059*
H18B	0.0214	0.4634	0.1132	0.059*
H18C	0.0124	0.4519	0.1805	0.059*
C19	0.16041 (11)	0.31121 (19)	0.13632 (7)	0.0440 (4)
H19A	0.2222	0.3041	0.1543	0.066*
H19B	0.1460	0.3585	0.0963	0.066*
H19C	0.1356	0.2174	0.1306	0.066*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.03185 (16)	0.02196 (15)	0.01989 (15)	0.00216 (11)	0.00886 (11)	0.00062 (10)
O1	0.0511 (6)	0.0282 (5)	0.0261 (5)	0.0076 (4)	0.0164 (4)	−0.0023 (4)
O2	0.0370 (5)	0.0270 (5)	0.0301 (5)	−0.0001 (4)	0.0139 (4)	0.0051 (4)
N1	0.0310 (5)	0.0205 (5)	0.0262 (5)	0.0004 (4)	0.0021 (4)	0.0006 (4)
N2	0.0290 (5)	0.0407 (6)	0.0206 (5)	0.0027 (5)	0.0082 (4)	−0.0020 (4)
C1	0.0280 (6)	0.0226 (6)	0.0286 (6)	0.0030 (5)	0.0076 (5)	−0.0041 (5)
C2	0.0314 (7)	0.0320 (7)	0.0420 (8)	0.0005 (5)	0.0130 (6)	−0.0016 (6)
C3	0.0302 (7)	0.0378 (8)	0.0578 (10)	−0.0039 (6)	0.0122 (7)	−0.0103 (7)
C4	0.0307 (7)	0.0461 (9)	0.0453 (9)	0.0016 (6)	0.0003 (6)	−0.0161 (7)
C5	0.0422 (8)	0.0462 (9)	0.0314 (7)	0.0025 (7)	0.0006 (6)	−0.0014 (6)
C6	0.0356 (7)	0.0328 (7)	0.0310 (7)	−0.0022 (6)	0.0039 (6)	0.0001 (5)
C7	0.0338 (6)	0.0233 (6)	0.0262 (6)	−0.0009 (5)	0.0063 (5)	0.0012 (5)

C8	0.0232 (5)	0.0235 (6)	0.0197 (5)	0.0009 (4)	0.0068 (4)	0.0004 (4)
C9	0.0303 (6)	0.0235 (6)	0.0259 (6)	0.0029 (5)	0.0106 (5)	−0.0003 (5)
C10	0.0317 (6)	0.0243 (6)	0.0290 (6)	0.0056 (5)	0.0086 (5)	0.0048 (5)
C11	0.0267 (6)	0.0293 (6)	0.0222 (5)	0.0013 (5)	0.0068 (5)	0.0047 (5)
C12	0.0198 (5)	0.0263 (6)	0.0221 (5)	0.0002 (4)	0.0073 (4)	0.0000 (4)
C13	0.0211 (5)	0.0317 (6)	0.0227 (5)	−0.0004 (5)	0.0076 (4)	−0.0027 (5)
C14	0.0296 (6)	0.0301 (7)	0.0288 (6)	0.0025 (5)	0.0093 (5)	−0.0078 (5)
C15	0.0310 (6)	0.0237 (6)	0.0332 (7)	0.0049 (5)	0.0068 (5)	−0.0023 (5)
C16	0.0270 (6)	0.0240 (6)	0.0252 (6)	0.0018 (5)	0.0055 (5)	0.0003 (5)
C17	0.0193 (5)	0.0232 (6)	0.0224 (5)	−0.0002 (4)	0.0064 (4)	−0.0007 (4)
C18	0.0320 (7)	0.0515 (9)	0.0283 (7)	0.0022 (6)	0.0020 (5)	−0.0016 (6)
C19	0.0523 (9)	0.0567 (10)	0.0288 (7)	0.0067 (8)	0.0212 (7)	−0.0047 (7)

Geometric parameters (Å, º)

S1—O1	1.4331 (9)	C8—C9	1.3718 (17)
S1—O2	1.4425 (9)	C8—C17	1.4351 (16)
S1—N1	1.6150 (11)	C9—C10	1.4084 (17)
S1—C8	1.7754 (12)	C9—H9	0.9500
N1—C7	1.4651 (17)	C10—C11	1.3622 (18)
N1—H1	0.857 (14)	C10—H10	0.9500
N2—C13	1.4184 (16)	C11—C12	1.4227 (17)
N2—C19	1.4558 (17)	C11—H11	0.9500
N2—C18	1.4687 (17)	C12—C17	1.4306 (16)
C1—C6	1.3843 (19)	C12—C13	1.4375 (16)
C1—C2	1.3948 (19)	C13—C14	1.3771 (19)
C1—C7	1.5125 (17)	C14—C15	1.4113 (18)
C2—C3	1.383 (2)	C14—H14	0.9500
C2—H2	0.9500	C15—C16	1.3687 (18)
C3—C4	1.387 (2)	C15—H15	0.9500
C3—H3	0.9500	C16—C17	1.4207 (17)
C4—C5	1.382 (2)	C16—H16	0.9500
C4—H4	0.9500	C18—H18A	0.9800
C5—C6	1.398 (2)	C18—H18B	0.9800
C5—H5	0.9500	C18—H18C	0.9800
C6—H6	0.9500	C19—H19A	0.9800
C7—H7A	0.9900	C19—H19B	0.9800
C7—H7B	0.9900	C19—H19C	0.9800
O1—S1—O2	118.40 (6)	C8—C9—C10	120.04 (11)
O1—S1—N1	107.93 (6)	C8—C9—H9	120.0
O2—S1—N1	106.16 (6)	C10—C9—H9	120.0
O1—S1—C8	106.45 (6)	C11—C10—C9	120.21 (12)
O2—S1—C8	109.54 (6)	C11—C10—H10	119.9
N1—S1—C8	107.99 (6)	C9—C10—H10	119.9
C7—N1—S1	119.48 (9)	C10—C11—C12	121.47 (11)
C7—N1—H1	118.9 (13)	C10—C11—H11	119.3
S1—N1—H1	112.0 (13)	C12—C11—H11	119.3

C13—N2—C19	115.60 (11)	C11—C12—C17	119.24 (11)
C13—N2—C18	114.53 (11)	C11—C12—C13	121.10 (11)
C19—N2—C18	110.42 (11)	C17—C12—C13	119.64 (11)
C6—C1—C2	119.02 (13)	C14—C13—N2	123.05 (11)
C6—C1—C7	123.03 (12)	C14—C13—C12	119.15 (11)
C2—C1—C7	117.94 (12)	N2—C13—C12	117.75 (11)
C3—C2—C1	120.89 (14)	C13—C14—C15	120.69 (11)
C3—C2—H2	119.6	C13—C14—H14	119.7
C1—C2—H2	119.6	C15—C14—H14	119.7
C2—C3—C4	119.91 (14)	C16—C15—C14	121.42 (12)
C2—C3—H3	120.0	C16—C15—H15	119.3
C4—C3—H3	120.0	C14—C15—H15	119.3
C5—C4—C3	119.67 (14)	C15—C16—C17	120.09 (11)
C5—C4—H4	120.2	C15—C16—H16	120.0
C3—C4—H4	120.2	C17—C16—H16	120.0
C4—C5—C6	120.44 (15)	C16—C17—C12	118.87 (11)
C4—C5—H5	119.8	C16—C17—C8	123.98 (11)
C6—C5—H5	119.8	C12—C17—C8	117.14 (11)
C1—C6—C5	120.06 (14)	N2—C18—H18A	109.5
C1—C6—H6	120.0	N2—C18—H18B	109.5
C5—C6—H6	120.0	H18A—C18—H18B	109.5
N1—C7—C1	113.36 (10)	N2—C18—H18C	109.5
N1—C7—H7A	108.9	H18A—C18—H18C	109.5
C1—C7—H7A	108.9	H18B—C18—H18C	109.5
N1—C7—H7B	108.9	N2—C19—H19A	109.5
C1—C7—H7B	108.9	N2—C19—H19B	109.5
H7A—C7—H7B	107.7	H19A—C19—H19B	109.5
C9—C8—C17	121.86 (11)	N2—C19—H19C	109.5
C9—C8—S1	116.47 (9)	H19A—C19—H19C	109.5
C17—C8—S1	121.66 (9)	H19B—C19—H19C	109.5

O1—S1—N1—C7	−55.53 (11)	C10—C11—C12—C17	−2.33 (18)
O2—S1—N1—C7	176.58 (9)	C10—C11—C12—C13	179.29 (11)
C8—S1—N1—C7	59.17 (11)	C19—N2—C13—C14	−19.34 (19)
C6—C1—C2—C3	−0.1 (2)	C18—N2—C13—C14	110.75 (14)
C7—C1—C2—C3	178.98 (13)	C19—N2—C13—C12	158.21 (12)
C1—C2—C3—C4	0.8 (2)	C18—N2—C13—C12	−71.70 (15)
C2—C3—C4—C5	−0.9 (2)	C11—C12—C13—C14	174.43 (12)
C3—C4—C5—C6	0.2 (2)	C17—C12—C13—C14	−3.94 (17)
C2—C1—C6—C5	−0.6 (2)	C11—C12—C13—N2	−3.22 (17)
C7—C1—C6—C5	−179.62 (13)	C17—C12—C13—N2	178.41 (10)
C4—C5—C6—C1	0.5 (2)	N2—C13—C14—C15	−179.01 (12)
S1—N1—C7—C1	137.24 (10)	C12—C13—C14—C15	3.47 (19)
C6—C1—C7—N1	−1.11 (18)	C13—C14—C15—C16	−0.4 (2)
C2—C1—C7—N1	179.83 (11)	C14—C15—C16—C17	−2.3 (2)
O1—S1—C8—C9	2.04 (12)	C15—C16—C17—C12	1.77 (18)
O2—S1—C8—C9	131.16 (10)	C15—C16—C17—C8	−177.02 (12)
N1—S1—C8—C9	−113.64 (10)	C11—C12—C17—C16	−177.06 (11)

O1—S1—C8—C17	−178.94 (10)	C13—C12—C17—C16	1.34 (16)
O2—S1—C8—C17	−49.82 (11)	C11—C12—C17—C8	1.81 (16)
N1—S1—C8—C17	65.38 (11)	C13—C12—C17—C8	−179.79 (10)
C17—C8—C9—C10	−0.62 (19)	C9—C8—C17—C16	178.42 (12)
S1—C8—C9—C10	178.40 (10)	S1—C8—C17—C16	−0.55 (16)
C8—C9—C10—C11	0.16 (19)	C9—C8—C17—C12	−0.39 (17)
C9—C10—C11—C12	1.33 (19)	S1—C8—C17—C12	−179.35 (9)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1···O2^{i}	0.86 (2)	2.12 (2)	2.9351 (14)	158 (2)

Symmetry code: (i) −*x*+1/2, −*y*+1/2, −*z*+1.