



The crystal structure of 2-[5-(dimethylamino)naphthalene-1-sulfonamido]-phenyl 5-(dimethylamino)naphthalene-1-sulfonate

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The complete molecule of the title compound, C₃₀H₂₉N₃O₅S₂, is generated by a crystallographic twofold axis: the O atom and NH group attached to the central benzene ring are statistically disordered. The dihedral angle between the naphthalene ring system and the central benzene ring is 52.99 (6)°, while the pendant naphthalene ring systems subtend a dihedral angle of 68.17 (4)°. An intramolecular C—H···O hydrogen bond closes an S(6) ring. In the crystal, the molecules are linked by weak C—H···O hydrogen bonds.

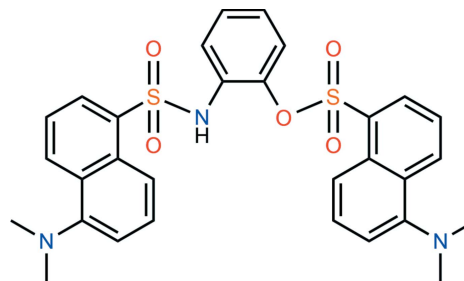
Keywords: crystal structure; dansyl derivatives; disorder; hydrogen bonding; π -stacking.

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1. Related literature

For the use of dansyl tags to monitor biological activity in enzyme systems, see: Brown *et al.* (1970); Liu *et al.* (2010). Dansyl-conjugated liposome has been used to modulate the fluorescence resonance energy transfer (FRET) mechanism, see: Li *et al.* (2006). Dansyl fluorogenic sensors have been used for the recognition and detection of targets such as cationic and anionic species, see: Cao *et al.* (2014); Jisha *et al.* (2009); Bhalla *et al.* (2007). For crystal structures of dansyl derivatives, see: Bhatt *et al.* (2011); Zhang *et al.* (2009) and of metal-

calix[4]arene complexes bearing two dansyl carboxamide units, see: Buie *et al.* (2008).



2. Experimental

2.1. Crystal data

C ₃₀ H ₂₉ N ₃ O ₅ S ₂	$V = 2769.1 (5) \text{ \AA}^3$
$M_r = 575.68$	$Z = 4$
Monoclinic, $C2/c$	Mo $K\alpha$ radiation
$a = 12.7594 (13) \text{ \AA}$	$\mu = 0.24 \text{ mm}^{-1}$
$b = 13.3481 (14) \text{ \AA}$	$T = 296 \text{ K}$
$c = 16.4331 (17) \text{ \AA}$	$0.26 \times 0.22 \times 0.22 \text{ mm}$
$\beta = 98.349 (4)^\circ$	

2.2. Data collection

Bruker D8 QUEST CMOS diffractometer	17644 measured reflections
Absorption correction: multi-scan (SADABS; Bruker, 2014)	3444 independent reflections
$T_{\min} = 0.698$, $T_{\max} = 0.746$	2246 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.041$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$	186 parameters
$wR(F^2) = 0.113$	H-atom parameters constrained
$S = 1.03$	$\Delta\rho_{\text{max}} = 0.26 \text{ e \AA}^{-3}$
3444 reflections	$\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C7—H7···O3	0.93	2.37	3.030 (2)	128
C13—H13···O3 ⁱ	0.93	2.73	3.386 (2)	129

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

Data collection: APEX2 (Bruker, 2014); cell refinement: SAINT (Bruker, 2014); data reduction: SAINT; program(s) used to solve structure: SHELXT (Sheldrick, 2015a); program(s) used to refine structure: SHELXL2014 (Sheldrick, 2015b); molecular graphics: OLEX2 (Dolomanov *et al.*, 2009); software used to prepare material for publication: OLEX2.

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: GW2153).

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The crystal structure of 2-[5-(dimethylamino)naphthalene-1-sulfonamido]phenyl 5-(dimethylamino)naphthalene-1-sulfonate

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S1. Introduction

Dansyl derivatives can be widely used as fluorescence probes in biological and environmental systems. Dansyl tags have been increasingly used to monitor biological activities in the enzyme system for providing the accurate information (Brown *et al.*, 1970; Liu *et al.*, 2010). An example is dansyl-conjugated liposome for modulating fluorescence resonance energy transfer (FRET) mechanism (Li *et al.* 2006). Furthermore, dansyl fluorogenic sensors were prepared for recognition and detection of many targets such as cationic and anionic species (Cao *et al.*, 2014; Jisha *et al.*, 2009; Bhalla *et al.*, 2007). Crystal structures of dansyl derivatives (Bhatt *et al.*, 2011; Zhang *et al.*, 2009) and metal complexes of calix[4]arene bearing two dansyl carboxamide units have been reported (Buie *et al.*, 2008).

S2. Synthesis and crystallization

The title compound was synthesized by condensation of 2-aminophenol (0.55 g, 5.04 mmol) and dansyl chloride (2.72 g, 10.08 mmol) using potassium carbonate (17.27 g, 12.50 mmol) as a base in acetonitrile (30 ml). The solution was heated and stirred under N₂ atmosphere for 24 h. The solvent was then removed by a rotary evaporator. Water (10 ml) was added to the residue and the organic phase was extracted with dichloromethane (3 x 20 ml). The organic layer was dried with Na₂SO₄. The product was purified by column chromatography using dichloromethane as an eluent. The solvent was evaporated to afford a yellow crystalline solid in 55% yield. Single crystals suitable for X-ray measurements were obtained by recrystallization using the mixture solution of dichloromethane and hexane (1:1, v/v) at room temperature.

S3. Refinement

Atom O1 and the N1H1 group attached to the central benzene ring are statistically disordered and were refined with the occupancies of the N, H and O atoms fixed at 0.5. All H atoms were positioned geometrically and refined using a riding model, with C—H = 0.93 Å for aryl and 0.96 Å for methyl H atoms, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aryl and $1.5U_{\text{eq}}(\text{C})$ for methyl H atoms. The N-bound H-atom was refined with N—H = 0.86 Å, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$.

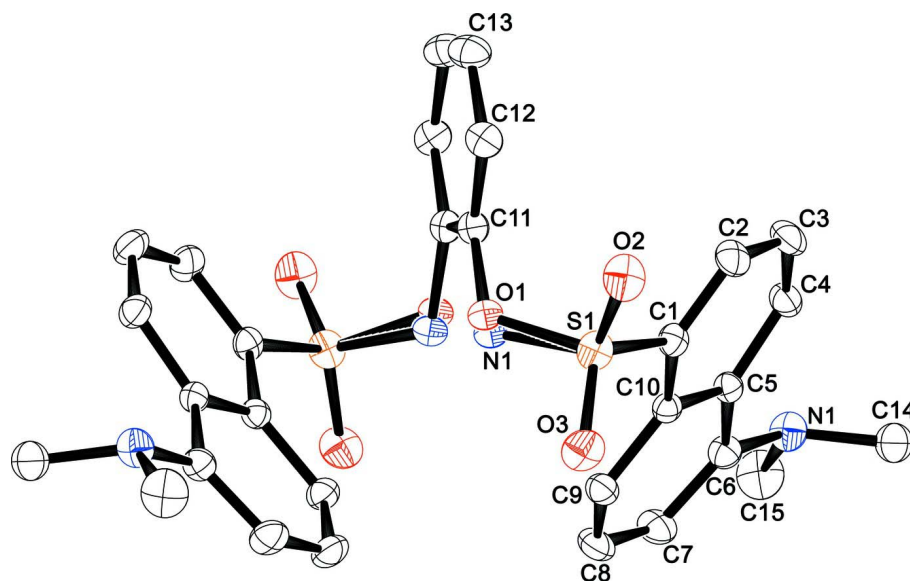


Figure 1

The molecular structure of the title compound with 30% probability ellipsoids and atom numbering. Hydrogen atoms are omitted for clarity.

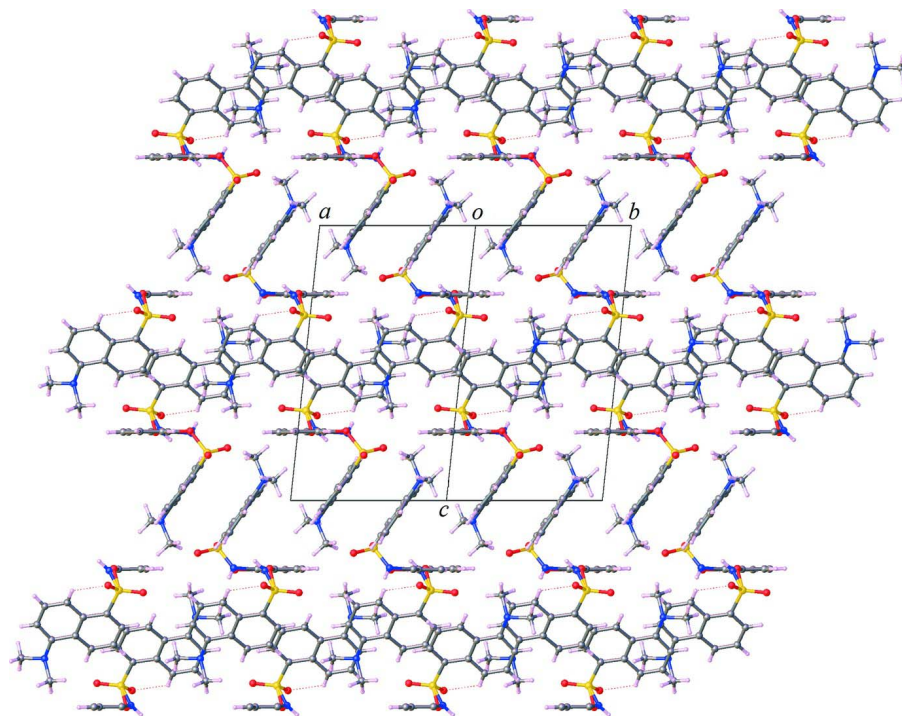


Figure 2

The crystal packing of the title compound, viewed along the [110] direction.

2-[5-(Dimethylamino)naphthalene-1-sulfonamido]phenyl 5-(dimethylamino)naphthalene-1-sulfonate

Crystal data

$C_{30}H_{29}N_3O_5S_2$

$M_r = 575.68$

Monoclinic, $C2/c$

$a = 12.7594$ (13) Å

$b = 13.3481$ (14) Å

$c = 16.4331$ (17) Å

$\beta = 98.349$ (4)°

$V = 2769.1$ (5) Å³

$Z = 4$

$F(000) = 1208$

$D_x = 1.381$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4957 reflections

$\theta = 3.1$ – 25.7 °

$\mu = 0.24$ mm⁻¹

$T = 296$ K

Block, light green

$0.26 \times 0.22 \times 0.22$ mm

Data collection

Bruker D8 QUEST CMOS
diffractometer

Radiation source: microfocus sealed x-ray tube,
Incoatec μ us

GraphiteDouble Bounce Multilayer Mirror
monochromator

Detector resolution: 10.5 pixels mm⁻¹

ω and φ scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2014)

$T_{\min} = 0.698$, $T_{\max} = 0.746$

17644 measured reflections

3444 independent reflections

2246 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.041$

$\theta_{\max} = 28.3$ °, $\theta_{\min} = 3.1$ °

$h = -15 \rightarrow 16$

$k = -17 \rightarrow 17$

$l = -21 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.113$

$S = 1.03$

3444 reflections

186 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0488P)^2 + 1.1669P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.26$ e Å⁻³

$\Delta\rho_{\min} = -0.23$ 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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
S1	0.69732 (3)	0.73580 (4)	0.32645 (3)	0.05431 (17)	
N1	0.603 (2)	0.709 (2)	0.2569 (14)	0.0418 (18)	0.5
H1	0.5801	0.7537	0.2215	0.050*	0.5
O2	0.77825 (10)	0.66139 (12)	0.33521 (10)	0.0736 (5)	
O3	0.72349 (11)	0.83601 (11)	0.30878 (10)	0.0711 (4)	
N2	0.35617 (13)	0.82675 (13)	0.57539 (10)	0.0609 (5)	
C11	0.55450 (13)	0.61136 (13)	0.25005 (10)	0.0433 (4)	

C5	0.49120 (13)	0.77874 (12)	0.49182 (10)	0.0427 (4)	
C6	0.54736 (13)	0.79731 (12)	0.42414 (10)	0.0413 (4)	
C1	0.62854 (13)	0.72779 (13)	0.41220 (11)	0.0450 (4)	
C10	0.40779 (14)	0.84536 (14)	0.50673 (12)	0.0495 (4)	
C12	0.60785 (15)	0.52183 (15)	0.24889 (12)	0.0543 (5)	
H12	0.6806	0.5215	0.2481	0.065*	
C7	0.51636 (15)	0.87928 (13)	0.37174 (11)	0.0500 (4)	
H7	0.5533	0.8940	0.3285	0.060*	
C4	0.51413 (16)	0.69084 (14)	0.53896 (12)	0.0531 (5)	
H4	0.4748	0.6766	0.5810	0.064*	
C2	0.65034 (16)	0.64565 (15)	0.46148 (12)	0.0582 (5)	
H2	0.7046	0.6022	0.4527	0.070*	
C9	0.37880 (17)	0.92039 (15)	0.45129 (13)	0.0612 (5)	
H9	0.3220	0.9615	0.4583	0.073*	
C8	0.43259 (17)	0.93625 (15)	0.38477 (13)	0.0611 (5)	
H8	0.4104	0.9876	0.3480	0.073*	
C13	0.55357 (17)	0.43329 (16)	0.24897 (14)	0.0665 (6)	
H13	0.5895	0.3728	0.2476	0.080*	
C3	0.59130 (17)	0.62698 (15)	0.52481 (13)	0.0635 (6)	
H3	0.6052	0.5702	0.5574	0.076*	
C14	0.41854 (19)	0.83896 (17)	0.65611 (13)	0.0707 (6)	
H14A	0.4905	0.8197	0.6537	0.106*	
H14B	0.3898	0.7975	0.6951	0.106*	
H14C	0.4165	0.9078	0.6727	0.106*	
C15	0.25143 (19)	0.8715 (2)	0.57308 (17)	0.0878 (8)	
H15A	0.2587	0.9419	0.5846	0.132*	
H15B	0.2153	0.8401	0.6136	0.132*	
H15C	0.2114	0.8620	0.5195	0.132*	
O1	0.6121 (16)	0.7010 (16)	0.2444 (11)	0.0418 (18)	0.5

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0322 (2)	0.0701 (4)	0.0622 (3)	-0.0032 (2)	0.0121 (2)	-0.0040 (3)
N1	0.035 (3)	0.051 (3)	0.043 (5)	0.0008 (19)	0.017 (2)	-0.002 (3)
O2	0.0366 (7)	0.1019 (12)	0.0833 (10)	0.0177 (7)	0.0120 (7)	-0.0060 (9)
O3	0.0528 (8)	0.0783 (10)	0.0848 (10)	-0.0260 (7)	0.0188 (7)	-0.0017 (8)
N2	0.0564 (10)	0.0708 (12)	0.0595 (10)	0.0057 (8)	0.0213 (8)	-0.0053 (8)
C11	0.0430 (9)	0.0499 (10)	0.0401 (9)	-0.0004 (8)	0.0161 (8)	0.0007 (8)
C5	0.0422 (9)	0.0435 (10)	0.0411 (9)	0.0018 (7)	0.0023 (7)	-0.0002 (7)
C6	0.0387 (9)	0.0416 (9)	0.0426 (9)	-0.0021 (7)	0.0023 (7)	-0.0026 (7)
C1	0.0367 (9)	0.0517 (10)	0.0454 (10)	0.0024 (8)	0.0017 (7)	-0.0020 (8)
C10	0.0474 (10)	0.0508 (11)	0.0509 (11)	0.0029 (8)	0.0088 (8)	-0.0037 (8)
C12	0.0515 (11)	0.0591 (13)	0.0566 (11)	0.0084 (9)	0.0225 (9)	0.0023 (9)
C7	0.0572 (11)	0.0458 (10)	0.0478 (10)	0.0015 (9)	0.0101 (9)	0.0054 (8)
C4	0.0616 (12)	0.0523 (11)	0.0459 (10)	0.0048 (9)	0.0098 (9)	0.0076 (9)
C2	0.0543 (11)	0.0605 (12)	0.0585 (12)	0.0223 (9)	0.0033 (9)	0.0026 (10)
C9	0.0606 (12)	0.0543 (12)	0.0702 (13)	0.0212 (10)	0.0144 (10)	0.0039 (10)

C8	0.0715 (14)	0.0484 (11)	0.0626 (12)	0.0162 (10)	0.0074 (10)	0.0128 (9)
C13	0.0791 (14)	0.0509 (12)	0.0752 (14)	0.0108 (10)	0.0306 (13)	0.0021 (11)
C3	0.0775 (14)	0.0554 (12)	0.0570 (12)	0.0212 (11)	0.0078 (11)	0.0144 (10)
C14	0.0904 (17)	0.0706 (15)	0.0549 (13)	0.0006 (12)	0.0233 (12)	-0.0004 (11)
C15	0.0640 (14)	0.111 (2)	0.0949 (19)	0.0165 (14)	0.0343 (14)	-0.0055 (15)
O1	0.035 (3)	0.051 (3)	0.043 (5)	0.0008 (19)	0.017 (2)	-0.002 (3)

Geometric parameters (Å, °)

S1—N1	1.58 (3)	C12—H12	0.9300
S1—O2	1.4248 (14)	C12—C13	1.370 (3)
S1—O3	1.4189 (15)	C7—H7	0.9300
S1—C1	1.7681 (18)	C7—C8	1.354 (3)
S1—O1	1.67 (2)	C4—H4	0.9300
N1—H1	0.8600	C4—C3	1.348 (3)
N1—C11	1.44 (3)	C2—H2	0.9300
N2—C10	1.409 (2)	C2—C3	1.393 (3)
N2—C14	1.454 (3)	C9—H9	0.9300
N2—C15	1.459 (3)	C9—C8	1.389 (3)
C11—C11 ⁱ	1.391 (3)	C8—H8	0.9300
C11—C12	1.377 (2)	C13—C13 ⁱ	1.372 (4)
C11—O1	1.41 (2)	C13—H13	0.9300
C5—C6	1.429 (2)	C3—H3	0.9300
C5—C10	1.435 (2)	C14—H14A	0.9600
C5—C4	1.413 (2)	C14—H14B	0.9600
C6—C1	1.425 (2)	C14—H14C	0.9600
C6—C7	1.413 (2)	C15—H15A	0.9600
C1—C2	1.367 (3)	C15—H15B	0.9600
C10—C9	1.368 (3)	C15—H15C	0.9600
N1—S1—C1	98.6 (8)	C6—C7—H7	120.1
O2—S1—N1	112.1 (9)	C8—C7—C6	119.74 (17)
O2—S1—C1	108.28 (9)	C8—C7—H7	120.1
O2—S1—O1	105.4 (6)	C5—C4—H4	119.0
O3—S1—N1	104.3 (10)	C3—C4—C5	121.93 (18)
O3—S1—O2	119.29 (9)	C3—C4—H4	119.0
O3—S1—C1	112.27 (9)	C1—C2—H2	120.0
O3—S1—O1	104.0 (7)	C1—C2—C3	120.02 (18)
O1—S1—C1	106.6 (6)	C3—C2—H2	120.0
S1—N1—H1	118.6	C10—C9—H9	119.4
C11—N1—S1	122.8 (18)	C10—C9—C8	121.27 (18)
C11—N1—H1	118.6	C8—C9—H9	119.4
C10—N2—C14	116.97 (16)	C7—C8—C9	122.05 (18)
C10—N2—C15	116.09 (18)	C7—C8—H8	119.0
C14—N2—C15	110.84 (18)	C9—C8—H8	119.0
C11 ⁱ —C11—N1	115.0 (10)	C12—C13—C13 ⁱ	120.37 (12)
C11 ⁱ —C11—O1	121.9 (8)	C12—C13—H13	119.8
C12—C11—N1	125.1 (10)	C13 ⁱ —C13—H13	119.8

C12—C11—C11 ⁱ	119.76 (11)	C4—C3—C2	120.24 (18)
C12—C11—O1	118.1 (8)	C4—C3—H3	119.9
C6—C5—C10	119.56 (15)	C2—C3—H3	119.9
C4—C5—C6	118.90 (16)	N2—C14—H14A	109.5
C4—C5—C10	121.37 (17)	N2—C14—H14B	109.5
C1—C6—C5	116.78 (15)	N2—C14—H14C	109.5
C7—C6—C5	118.75 (16)	H14A—C14—H14B	109.5
C7—C6—C1	124.40 (16)	H14A—C14—H14C	109.5
C6—C1—S1	121.67 (13)	H14B—C14—H14C	109.5
C2—C1—S1	116.02 (14)	N2—C15—H15A	109.5
C2—C1—C6	122.00 (17)	N2—C15—H15B	109.5
N2—C10—C5	118.17 (16)	N2—C15—H15C	109.5
C9—C10—N2	123.36 (17)	H15A—C15—H15B	109.5
C9—C10—C5	118.38 (17)	H15A—C15—H15C	109.5
C11—C12—H12	120.1	H15B—C15—H15C	109.5
C13—C12—C11	119.85 (18)	C11—O1—S1	117.7 (13)
C13—C12—H12	120.1		
S1—N1—C11—C11 ⁱ	124.2 (13)	C6—C5—C4—C3	3.7 (3)
S1—N1—C11—C12	-51.6 (18)	C6—C1—C2—C3	1.1 (3)
S1—C1—C2—C3	-172.60 (16)	C6—C7—C8—C9	3.7 (3)
N1—S1—C1—C6	-67.6 (10)	C1—S1—N1—C11	-66.4 (15)
N1—S1—C1—C2	106.1 (10)	C1—S1—O1—C11	-48.7 (12)
N1—C11—C12—C13	174.6 (11)	C1—C6—C7—C8	174.71 (18)
O2—S1—N1—C11	47.5 (16)	C1—C2—C3—C4	-1.5 (3)
O2—S1—C1—C6	175.57 (14)	C10—C5—C6—C1	-179.37 (15)
O2—S1—C1—C2	-10.74 (18)	C10—C5—C6—C7	-2.3 (3)
O2—S1—O1—C11	66.2 (11)	C10—C5—C4—C3	179.07 (19)
O3—S1—N1—C11	177.9 (12)	C10—C9—C8—C7	-0.5 (3)
O3—S1—C1—C6	41.79 (17)	C12—C11—O1—S1	-72.7 (11)
O3—S1—C1—C2	-144.52 (15)	C7—C6—C1—S1	-2.0 (2)
O3—S1—O1—C11	-167.5 (9)	C7—C6—C1—C2	-175.27 (19)
N2—C10—C9—C8	179.6 (2)	C4—C5—C6—C1	-4.0 (2)
C11 ⁱ —C11—C12—C13	-1.0 (3)	C4—C5—C6—C7	173.16 (17)
C11 ⁱ —C11—O1—S1	112.6 (11)	C4—C5—C10—N2	6.6 (3)
C11—C12—C13—C13 ⁱ	-0.6 (4)	C4—C5—C10—C9	-170.03 (19)
C5—C6—C1—S1	174.98 (12)	C14—N2—C10—C5	66.8 (2)
C5—C6—C1—C2	1.7 (3)	C14—N2—C10—C9	-116.8 (2)
C5—C6—C7—C8	-2.2 (3)	C15—N2—C10—C5	-159.33 (19)
C5—C10—C9—C8	-4.0 (3)	C15—N2—C10—C9	17.1 (3)
C5—C4—C3—C2	-1.0 (3)	O1—S1—C1—C6	-71.5 (8)
C6—C5—C10—N2	-178.08 (16)	O1—S1—C1—C2	102.2 (8)
C6—C5—C10—C9	5.3 (3)	O1—C11—C12—C13	-175.8 (9)

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

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>
C7—H7 \cdots O3	0.93	2.37	3.030 (2)	128
C13—H13 \cdots O3 ⁱⁱ	0.93	2.73	3.386 (2)	129

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