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## Structure Reports

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# 4-(Diethylamino)salicylaldehyde phenylsulfonylhydrazone

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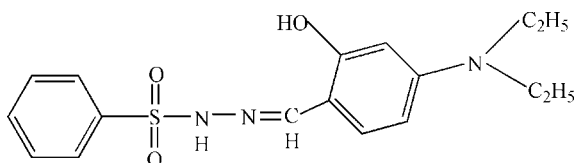
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 Key indicators: single-crystal X-ray study;  $T = 293$  K; mean  $\sigma(\text{C}-\text{C}) = 0.007$  Å; disorder in main residue;  $R$  factor = 0.069;  $wR$  factor = 0.175; data-to-parameter ratio = 11.8.

In the title compound,  $\text{C}_{17}\text{H}_{21}\text{N}_3\text{O}_3\text{S}$ , the dihedral angle between the aromatic ring planes is  $84.2(2)^\circ$ . The pendant ethyl groups of the  $-\text{N}(\text{C}_2\text{H}_5)_2$  group are disordered over two sets of positions in a 0.84(2):0.16(2) ratio. The molecular conformation is stabilized by an intramolecular  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bond, and intermolecular  $\text{N}-\text{H}\cdots\text{O}$  bonds lead to [010] chains in the crystal structure.

## Related literature

 For related literature, see: Tai *et al.* (2008).


## Experimental

### Crystal data

 $\text{C}_{17}\text{H}_{21}\text{N}_3\text{O}_3\text{S}$   
 $M_r = 347.43$ 

 Orthorhombic,  $Pbcn$   
 $a = 29.874(3)$  Å

 $b = 7.5153(12)$  Å  
 $c = 15.4456(19)$  Å  
 $V = 3467.8(8)$  Å<sup>3</sup>  
 $Z = 8$ 

 Mo  $K\alpha$  radiation  
 $\mu = 0.21$  mm<sup>-1</sup>  
 $T = 293(2)$  K  
 $0.43 \times 0.38 \times 0.04$  mm

### Data collection

 Bruker SMART CCD diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 2000)  
 $T_{\min} = 0.916$ ,  $T_{\max} = 0.992$ 

 16321 measured reflections  
 3052 independent reflections  
 2061 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.048$ 

### Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.069$   
 $wR(F^2) = 0.175$   
 $S = 1.08$   
 3052 reflections

 258 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.44$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.30$  e Å<sup>-3</sup>
**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O}3-\text{H}3\cdots\text{N}2$	0.82	1.92	2.637(4)	146
$\text{N}1-\text{H}1\cdots\text{O}1^i$	0.90	2.06	2.944(5)	169

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

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2720).

## References

- Bruker (2000). SMART, SAINT and SADABS. Bruker AXS Inc., Madison, Wisconsin, USA.  
 Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.  
 Tai, X.-S., Feng, Y.-M. & Kong, F.-Y. (2008). *Acta Cryst.* **E64**, o750.

## supporting information

*Acta Cryst.* (2008). E64, o865 [doi:10.1107/S1600536808010118]

## 4-(Diethylamino)salicylaldehyde phenylsulfonylhydrazone

Xi-Shi Tai, Yi-Min Feng and Fan-Yuan Kong

### S1. Comment

As part of our ongoing studies of aroylhydrazones as potential ligands (Tai *et al.*, 2008), we now report the synthesis and structure of the title compound, (I), (Fig. 1).

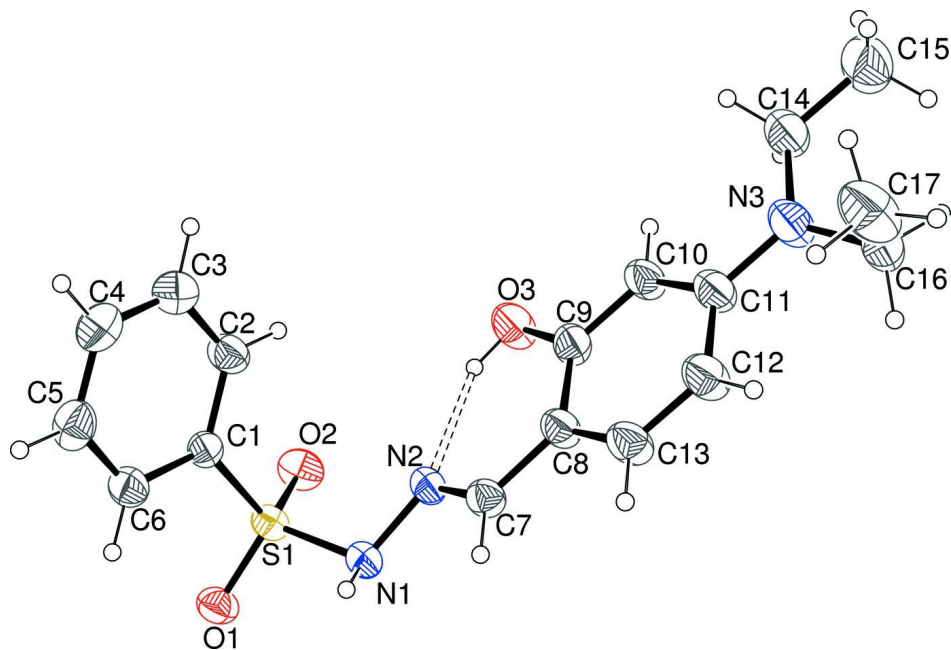
The dihedral angle between the aromatic ring planes is  $84.2(2)^\circ$ . The pendant ethyl groups of the  $-\text{N}(\text{C}_2\text{H}_5)_2$  grouping are disordered over two sets of positions in a 0.84(2):0.16(2) ratio. The molecular conformation is stabilised by an intramolecular O—H $\cdots$ N hydrogen bond and intermolecular N—H $\cdots$ O bonds lead to [010] chains in the crystal (Table 1).

### S2. Experimental

3 mmol of *p*-(diethylamino)salicylaldehyde (3 mmol) was added to a solution of benzenesulfonyl hydrazide (3 mmol) in 10 ml of 95% ethanol. The mixture was continuously stirred for 4 h at refluxing temperature, evaporating some ethanol, then, upon cooling, the solid product was collected by filtration and dried *in vacuo* (yield 67%). Colourless plates of (I) were obtained by evaporation from a methanol solution after several days.

### S3. Refinement

The H atoms were placed geometrically (C—H = 0.93–0.96 Å, N—H = 0.86 Å, O—H = 0.82 Å) and refined as riding with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$  or  $1.5U_{\text{eq}}(\text{methyl C}, \text{O})$ .

**Figure 1**

The molecular structure of (I) showing 30% displacement ellipsoids for the non-hydrogen atoms. Only the major disorder component is shown and the hydrogen bond is indicated by a double-dashed line.

#### 4-(Diethylamino)salicylaldehyde phenylsulfonylhydrazone

##### Crystal data

$C_{17}H_{21}N_3O_3S$

$M_r = 347.43$

Orthorhombic, *Pbcn*

$a = 29.874$  (3) Å

$b = 7.5153$  (12) Å

$c = 15.4456$  (19) Å

$V = 3467.8$  (8) Å<sup>3</sup>

$Z = 8$

$F(000) = 1472$

$D_x = 1.331$  Mg m<sup>-3</sup>

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

Cell parameters from 3546 reflections

$\theta = 2.7\text{--}21.4^\circ$

$\mu = 0.21$  mm<sup>-1</sup>

$T = 293$  K

Plate, colourless

$0.43 \times 0.38 \times 0.04$  mm

##### Data collection

Bruker SMART CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

$\omega$  scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2000)

$T_{\min} = 0.916$ ,  $T_{\max} = 0.992$

16321 measured reflections

3052 independent reflections

2061 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 25.0^\circ$ ,  $\theta_{\min} = 2.6^\circ$

$h = -35 \rightarrow 34$

$k = -8 \rightarrow 7$

$l = -18 \rightarrow 14$

##### Refinement

Refinement on  $F^2$

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.069$

$wR(F^2) = 0.175$

$S = 1.08$

3052 reflections

258 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-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0453P)^2 + 7.4306P]$$

where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
N1	0.31274 (11)	0.2572 (5)	0.9493 (2)	0.0502 (9)	
H1	0.2949	0.3499	0.9363	0.060*	
N2	0.35710 (11)	0.3011 (5)	0.9253 (2)	0.0512 (9)	
N3	0.54548 (13)	0.6991 (6)	0.8710 (3)	0.0794 (14)	
O1	0.24926 (10)	0.0597 (4)	0.93219 (19)	0.0596 (8)	
O2	0.32557 (10)	-0.0581 (4)	0.9110 (2)	0.0643 (9)	
O3	0.44009 (10)	0.2276 (4)	0.8794 (2)	0.0773 (11)	
H3	0.4137	0.2060	0.8893	0.116*	
S1	0.29375 (3)	0.08000 (14)	0.89918 (7)	0.0481 (3)	
C1	0.29054 (13)	0.1295 (5)	0.7887 (3)	0.0457 (10)	
C2	0.32787 (15)	0.1092 (6)	0.7367 (3)	0.0602 (12)	
H2	0.3547	0.0683	0.7601	0.072*	
C3	0.32493 (18)	0.1505 (7)	0.6499 (3)	0.0727 (14)	
H3A	0.3498	0.1363	0.6142	0.087*	
C4	0.28604 (19)	0.2114 (7)	0.6164 (3)	0.0700 (14)	
H4	0.2846	0.2396	0.5578	0.084*	
C5	0.24886 (19)	0.2323 (7)	0.6664 (3)	0.0693 (14)	
H5	0.2224	0.2744	0.6420	0.083*	
C6	0.25055 (15)	0.1908 (6)	0.7535 (3)	0.0586 (12)	
H6	0.2253	0.2037	0.7882	0.070*	
C7	0.36859 (14)	0.4638 (6)	0.9325 (3)	0.0517 (11)	
H7	0.3473	0.5476	0.9489	0.062*	
C8	0.41362 (13)	0.5202 (6)	0.9160 (3)	0.0545 (11)	
C9	0.44774 (14)	0.4048 (6)	0.8899 (3)	0.0572 (12)	
C10	0.49056 (14)	0.4634 (6)	0.8754 (3)	0.0630 (13)	
H10	0.5124	0.3820	0.8588	0.076*	
C11	0.50227 (15)	0.6425 (6)	0.8849 (3)	0.0661 (13)	
C12	0.46791 (15)	0.7606 (7)	0.9108 (4)	0.0725 (15)	
H12	0.4742	0.8809	0.9179	0.087*	
C13	0.42560 (15)	0.6996 (6)	0.9256 (3)	0.0670 (14)	

H13	0.4038	0.7804	0.9427	0.080*	
C14	0.5780 (8)	0.571 (4)	0.8341 (15)	0.095 (6)	0.84 (2)
H14A	0.5787	0.4677	0.8715	0.114*	0.84 (2)
H14B	0.5666	0.5327	0.7784	0.114*	0.84 (2)
C15	0.6231 (3)	0.6286 (18)	0.8217 (10)	0.114 (4)	0.84 (2)
H15A	0.6242	0.7137	0.7754	0.171*	0.84 (2)
H15B	0.6414	0.5280	0.8074	0.171*	0.84 (2)
H15C	0.6339	0.6826	0.8740	0.171*	0.84 (2)
C16	0.5597 (2)	0.8789 (15)	0.8999 (8)	0.076 (3)	0.84 (2)
H16A	0.5912	0.8770	0.9147	0.092*	0.84 (2)
H16B	0.5430	0.9123	0.9513	0.092*	0.84 (2)
C17	0.5517 (3)	1.012 (2)	0.8301 (10)	0.107 (4)	0.84 (2)
H17A	0.5649	0.9712	0.7772	0.161*	0.84 (2)
H17B	0.5650	1.1239	0.8463	0.161*	0.84 (2)
H17C	0.5201	1.0281	0.8220	0.161*	0.84 (2)
C14'	0.585 (4)	0.574 (18)	0.870 (7)	0.08 (2)	0.16 (2)
H14C	0.5748	0.4534	0.8813	0.098*	0.16 (2)
H14D	0.6063	0.6079	0.9134	0.098*	0.16 (2)
C15'	0.606 (2)	0.586 (8)	0.779 (5)	0.11 (2)	0.16 (2)
H15D	0.5939	0.6879	0.7496	0.170*	0.16 (2)
H15E	0.5985	0.4801	0.7472	0.170*	0.16 (2)
H15F	0.6375	0.5966	0.7846	0.170*	0.16 (2)
C16'	0.5486 (13)	0.875 (8)	0.826 (4)	0.082 (18)	0.16 (2)
H16C	0.5207	0.9021	0.7969	0.099*	0.16 (2)
H16D	0.5724	0.8723	0.7830	0.099*	0.16 (2)
C17'	0.5588 (15)	1.015 (10)	0.896 (4)	0.089 (19)	0.16 (2)
H17D	0.5320	1.0407	0.9280	0.133*	0.16 (2)
H17E	0.5694	1.1224	0.8689	0.133*	0.16 (2)
H17F	0.5813	0.9706	0.9346	0.133*	0.16 (2)

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
N1	0.0406 (19)	0.055 (2)	0.055 (2)	0.0017 (16)	-0.0002 (16)	-0.0013 (18)
N2	0.0378 (18)	0.057 (2)	0.059 (2)	0.0001 (16)	-0.0014 (16)	-0.0042 (18)
N3	0.053 (2)	0.065 (3)	0.120 (4)	-0.008 (2)	0.015 (2)	-0.019 (3)
O1	0.0506 (16)	0.061 (2)	0.067 (2)	-0.0052 (15)	0.0125 (15)	-0.0017 (16)
O2	0.0634 (19)	0.0535 (19)	0.076 (2)	0.0149 (15)	0.0025 (16)	0.0112 (17)
O3	0.0547 (19)	0.053 (2)	0.125 (3)	-0.0024 (16)	0.0098 (19)	-0.022 (2)
S1	0.0427 (6)	0.0449 (6)	0.0566 (7)	0.0016 (5)	0.0036 (5)	0.0027 (5)
C1	0.044 (2)	0.038 (2)	0.054 (3)	0.0016 (18)	-0.001 (2)	-0.0076 (19)
C2	0.047 (3)	0.069 (3)	0.065 (3)	0.007 (2)	0.002 (2)	0.002 (3)
C3	0.068 (3)	0.085 (4)	0.065 (4)	-0.003 (3)	0.006 (3)	-0.001 (3)
C4	0.087 (4)	0.066 (3)	0.057 (3)	-0.001 (3)	-0.006 (3)	0.000 (3)
C5	0.074 (3)	0.063 (3)	0.071 (3)	0.017 (3)	-0.020 (3)	-0.005 (3)
C6	0.049 (2)	0.062 (3)	0.065 (3)	0.009 (2)	-0.003 (2)	-0.007 (2)
C7	0.046 (2)	0.048 (3)	0.061 (3)	0.005 (2)	0.001 (2)	-0.003 (2)
C8	0.042 (2)	0.053 (3)	0.069 (3)	0.002 (2)	0.003 (2)	-0.003 (2)

C9	0.048 (2)	0.051 (3)	0.073 (3)	0.001 (2)	-0.002 (2)	-0.011 (2)
C10	0.043 (2)	0.055 (3)	0.090 (4)	0.003 (2)	0.008 (2)	-0.018 (3)
C11	0.045 (2)	0.068 (3)	0.085 (4)	-0.003 (2)	0.008 (2)	-0.009 (3)
C12	0.056 (3)	0.053 (3)	0.109 (4)	-0.003 (2)	0.014 (3)	-0.011 (3)
C13	0.050 (3)	0.054 (3)	0.098 (4)	0.005 (2)	0.011 (3)	-0.009 (3)
C14	0.060 (9)	0.096 (8)	0.129 (17)	-0.014 (7)	0.018 (10)	-0.045 (13)
C15	0.067 (6)	0.131 (9)	0.145 (11)	0.011 (6)	-0.007 (6)	-0.040 (7)
C16	0.052 (4)	0.075 (7)	0.101 (8)	-0.006 (3)	0.001 (4)	-0.019 (5)
C17	0.072 (6)	0.097 (10)	0.152 (11)	0.004 (5)	0.011 (7)	0.011 (9)
C14'	0.06 (4)	0.08 (4)	0.11 (7)	-0.01 (3)	0.00 (4)	-0.02 (5)
C15'	0.09 (4)	0.11 (4)	0.13 (6)	-0.01 (3)	0.00 (4)	-0.03 (4)
C16'	0.06 (2)	0.08 (4)	0.11 (5)	0.00 (2)	0.01 (2)	-0.02 (3)
C17'	0.06 (2)	0.08 (4)	0.13 (5)	0.00 (2)	0.00 (3)	-0.02 (4)

*Geometric parameters (Å, °)*

N1—N2	1.415 (4)	C10—C11	1.398 (6)
N1—S1	1.642 (4)	C10—H10	0.9300
N1—H1	0.9000	C11—C12	1.415 (6)
N2—C7	1.275 (5)	C12—C13	1.363 (6)
N3—C11	1.376 (6)	C12—H12	0.9300
N3—C14	1.48 (3)	C13—H13	0.9300
N3—C16	1.485 (11)	C14—C15	1.42 (3)
N3—C16'	1.50 (6)	C14—H14A	0.9700
N3—C14'	1.50 (15)	C14—H14B	0.9700
O1—S1	1.432 (3)	C15—H15A	0.9600
O2—S1	1.419 (3)	C15—H15B	0.9600
O3—C9	1.361 (5)	C15—H15C	0.9600
O3—H3	0.8200	C16—C17	1.49 (2)
S1—C1	1.750 (4)	C16—H16A	0.9700
C1—C2	1.382 (6)	C16—H16B	0.9700
C1—C6	1.391 (6)	C17—H17A	0.9600
C2—C3	1.380 (7)	C17—H17B	0.9600
C2—H2	0.9300	C17—H17C	0.9600
C3—C4	1.351 (7)	C14'—C15'	1.53 (16)
C3—H3A	0.9300	C14'—H14C	0.9700
C4—C5	1.362 (7)	C14'—H14D	0.9700
C4—H4	0.9300	C15'—H15D	0.9600
C5—C6	1.383 (6)	C15'—H15E	0.9600
C5—H5	0.9300	C15'—H15F	0.9600
C6—H6	0.9300	C16'—C17'	1.54 (10)
C7—C8	1.433 (6)	C16'—H16C	0.9700
C7—H7	0.9300	C16'—H16D	0.9700
C8—C9	1.398 (6)	C17'—H17D	0.9600
C8—C13	1.403 (6)	C17'—H17E	0.9600
C9—C10	1.371 (6)	C17'—H17F	0.9600
N2—N1—S1	112.9 (3)	C9—C10—C11	121.7 (4)

N2—N1—H1	108.5	C9—C10—H10	119.1
S1—N1—H1	108.5	C11—C10—H10	119.1
C7—N2—N1	116.9 (4)	N3—C11—C10	121.1 (4)
C11—N3—C14	118.4 (10)	N3—C11—C12	122.0 (4)
C11—N3—C16	120.2 (5)	C10—C11—C12	116.9 (4)
C14—N3—C16	121.2 (10)	C13—C12—C11	120.6 (5)
C11—N3—C16'	113.9 (16)	C13—C12—H12	119.7
C14—N3—C16'	111 (2)	C11—C12—H12	119.7
C16—N3—C16'	47 (2)	C12—C13—C8	122.8 (4)
C11—N3—C14'	123 (5)	C12—C13—H13	118.6
C14—N3—C14'	23 (4)	C8—C13—H13	118.6
C16—N3—C14'	111 (4)	C15—C14—N3	118.4 (18)
C16'—N3—C14'	120 (6)	C15—C14—H14A	107.7
C9—O3—H3	109.5	N3—C14—H14A	107.7
O2—S1—O1	119.90 (19)	C15—C14—H14B	107.7
O2—S1—N1	107.52 (19)	N3—C14—H14B	107.7
O1—S1—N1	103.83 (18)	H14A—C14—H14B	107.1
O2—S1—C1	108.52 (19)	N3—C16—C17	110.4 (12)
O1—S1—C1	108.61 (19)	N3—C16—H16A	109.6
N1—S1—C1	107.86 (19)	C17—C16—H16A	109.6
C2—C1—C6	120.2 (4)	N3—C16—H16B	109.6
C2—C1—S1	119.9 (3)	C17—C16—H16B	109.6
C6—C1—S1	119.9 (3)	H16A—C16—H16B	108.1
C3—C2—C1	119.2 (4)	N3—C14'—C15'	107 (8)
C3—C2—H2	120.4	N3—C14'—H14C	110.3
C1—C2—H2	120.4	C15'—C14'—H14C	110.3
C4—C3—C2	120.2 (5)	N3—C14'—H14D	110.3
C4—C3—H3A	119.9	C15'—C14'—H14D	110.3
C2—C3—H3A	119.9	H14C—C14'—H14D	108.5
C3—C4—C5	121.6 (5)	C14'—C15'—H15D	109.5
C3—C4—H4	119.2	C14'—C15'—H15E	109.5
C5—C4—H4	119.2	H15D—C15'—H15E	109.5
C4—C5—C6	119.7 (5)	C14'—C15'—H15F	109.5
C4—C5—H5	120.1	H15D—C15'—H15F	109.5
C6—C5—H5	120.1	H15E—C15'—H15F	109.5
C5—C6—C1	119.1 (5)	N3—C16'—C17'	107 (6)
C5—C6—H6	120.5	N3—C16'—H16C	110.4
C1—C6—H6	120.5	C17'—C16'—H16C	110.4
N2—C7—C8	121.4 (4)	N3—C16'—H16D	110.4
N2—C7—H7	119.3	C17'—C16'—H16D	110.4
C8—C7—H7	119.3	H16C—C16'—H16D	108.6
C9—C8—C13	116.2 (4)	C16'—C17'—H17D	109.5
C9—C8—C7	123.5 (4)	C16'—C17'—H17E	109.5
C13—C8—C7	120.3 (4)	H17D—C17'—H17E	109.5
O3—C9—C10	116.9 (4)	C16'—C17'—H17F	109.5
O3—C9—C8	121.3 (4)	H17D—C17'—H17F	109.5
C10—C9—C8	121.9 (4)	H17E—C17'—H17F	109.5

S1—N1—N2—C7	153.6 (3)	C16'—N3—C11—C10	141 (3)
N2—N1—S1—O2	52.0 (3)	C14'—N3—C11—C10	-18 (4)
N2—N1—S1—O1	-179.9 (3)	C14—N3—C11—C12	-172.9 (12)
N2—N1—S1—C1	-64.8 (3)	C16—N3—C11—C12	12.5 (9)
O2—S1—C1—C2	-31.0 (4)	C16'—N3—C11—C12	-40 (3)
O1—S1—C1—C2	-162.8 (3)	C14'—N3—C11—C12	161 (4)
N1—S1—C1—C2	85.2 (4)	C9—C10—C11—N3	179.3 (5)
O2—S1—C1—C6	149.7 (3)	C9—C10—C11—C12	0.4 (8)
O1—S1—C1—C6	17.8 (4)	N3—C11—C12—C13	-178.8 (5)
N1—S1—C1—C6	-94.1 (4)	C10—C11—C12—C13	0.1 (8)
C6—C1—C2—C3	0.0 (7)	C11—C12—C13—C8	-0.2 (9)
S1—C1—C2—C3	-179.4 (4)	C9—C8—C13—C12	-0.1 (8)
C1—C2—C3—C4	0.6 (8)	C7—C8—C13—C12	179.9 (5)
C2—C3—C4—C5	-0.6 (8)	C11—N3—C14—C15	-178.5 (15)
C3—C4—C5—C6	0.0 (8)	C16—N3—C14—C15	-4 (2)
C4—C5—C6—C1	0.5 (7)	C16'—N3—C14—C15	48 (3)
C2—C1—C6—C5	-0.5 (7)	C14'—N3—C14—C15	-71 (15)
S1—C1—C6—C5	178.8 (4)	C11—N3—C16—C17	-89.7 (8)
N1—N2—C7—C8	175.4 (4)	C14—N3—C16—C17	95.8 (13)
N2—C7—C8—C9	1.0 (7)	C16'—N3—C16—C17	6 (2)
N2—C7—C8—C13	-179.0 (5)	C14'—N3—C16—C17	118 (5)
C13—C8—C9—O3	179.3 (5)	C11—N3—C14'—C15'	118 (7)
C7—C8—C9—O3	-0.8 (7)	C14—N3—C14'—C15'	32 (11)
C13—C8—C9—C10	0.6 (7)	C16—N3—C14'—C15'	-90 (6)
C7—C8—C9—C10	-179.4 (5)	C16'—N3—C14'—C15'	-39 (8)
O3—C9—C10—C11	-179.5 (5)	C11—N3—C16'—C17'	101 (3)
C8—C9—C10—C11	-0.8 (8)	C14—N3—C16'—C17'	-123 (2)
C14—N3—C11—C10	8.3 (13)	C16—N3—C16'—C17'	-9 (2)
C16—N3—C11—C10	-166.3 (7)	C14'—N3—C16'—C17'	-100 (4)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
O3—H3...N2	0.82	1.92	2.637 (4)	146
N1—H1...O1 <sup>i</sup>	0.90	2.06	2.944 (5)	169

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