

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

# Ethyl (2*E*,4*Z*)-5-diethylamino-2-(phenyl-sulfonyl)penta-2,4-dienoate

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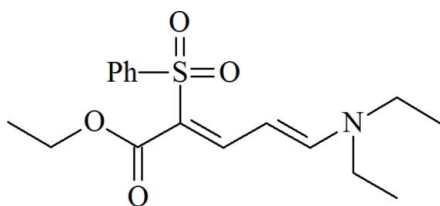
Received 4 April 2012; accepted 23 May 2012

 Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å;  $R$  factor = 0.059;  $wR$  factor = 0.168; data-to-parameter ratio = 19.8.

In the title compound,  $\text{C}_{17}\text{H}_{23}\text{NO}_4\text{S}$ , the pentadiene group adopts a planar conformation, with an r.m.s. deviation of 0.0410 (14) Å. The phenyl ring makes a dihedral angle of 85.73 (11)° with the pentadiene group, while the pentadiene group makes dihedral angles of 11.38 (11) and 14.08 (10)°, respectively, with the amino and ester groups. In the crystal, molecules are linked *via* pairs of C—H···O interactions, forming inversion dimers.

## Related literature

For background information on pentadienoates, see: Sorbetti *et al.* (2007). For structural data of pentadienoates, see: Ceard *et al.* (2002). For details of weak hydrogen-bonding interactions, see: Steiner (2002).



## Experimental

### Crystal data

$\text{C}_{17}\text{H}_{23}\text{NO}_4\text{S}$	$V = 1830.6$ (5) Å <sup>3</sup>
$M_r = 337.42$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 14.489$ (2) Å	$\mu = 0.20$ mm <sup>-1</sup>
$b = 8.2989$ (12) Å	$T = 296$ K
$c = 16.706$ (3) Å	$0.28 \times 0.24 \times 0.20$ mm
$\beta = 114.313$ (3)°	

### Data collection

Bruker APEXII CCD area-detector diffractometer	11618 measured reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 2003)	4183 independent reflections
$T_{\min} = 0.669$ , $T_{\max} = 0.746$	2977 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.031$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.059$	211 parameters
$wR(F^2) = 0.168$	H-atom parameters constrained
$S = 1.04$	$\Delta\rho_{\text{max}} = 0.47$ e Å <sup>-3</sup>
4183 reflections	$\Delta\rho_{\text{min}} = -0.44$ e Å <sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C13}-\text{H13A}\cdots\text{O1}^i$	0.93	2.41	3.277 (3)	155

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

Data collection: APEX2 (Bruker, 2007); cell refinement: APEX2 and SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL and DIAMOND (Brandenburg, 2005); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PK2405).

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## supporting information

*Acta Cryst.* (2012). E68, o1919 [doi:10.1107/S1600536812023483]

**Ethyl (2*E*,4*Z*)-5-diethylamino-2-(phenylsulfonyl)penta-2,4-dienoate****Guo-Qiang Song, Pei-Pei Yu and Xian-Feng Huang****S1. Comment**

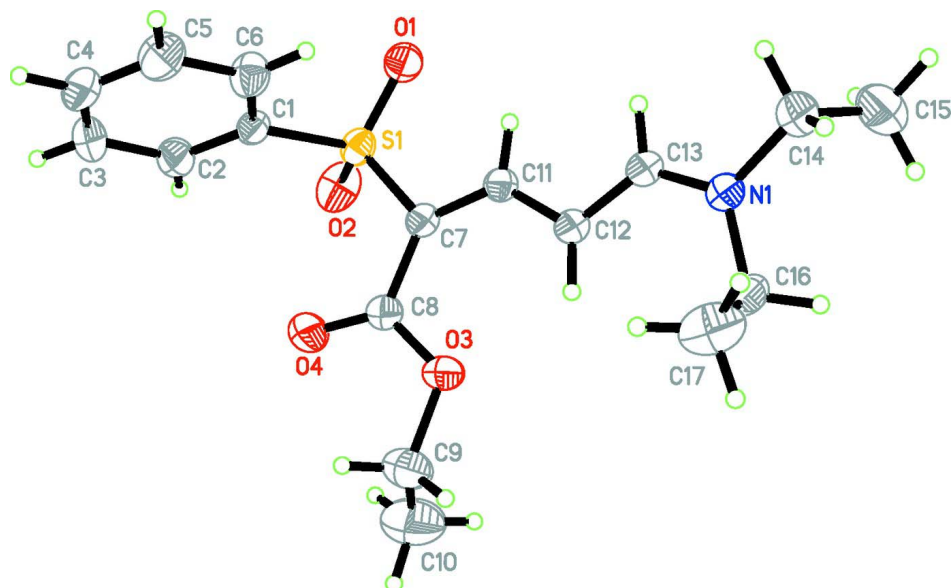
Analogues of pentadienoate have been used widely in perfumes, lubricants, as pharmaceutical intermediates and in other chemical industries (Ceard *et al.*, 2002; Sorbetti *et al.*, 2007). In this work, we report the crystal structure of the title compound, C<sub>17</sub>H<sub>23</sub>NO<sub>4</sub>S, (I). In the crystal (Fig. 1), the pentadiene group adopts a planar conformation with an r.m.s. deviation of 0.0410 (14) Å. The phenyl ring makes a dihedral angle of 85.73 (11)° with the pentadiene group. The dihedral angle of pentadiene with the amino group (atoms N1, C14, C16) is 11.38 (11)° and the dihedral angle between pentadiene and the ester group (atoms C8, O3, O4) is 14.08 (10)°. The crystal structure exhibits weak intermolecular C—H···O hydrogen bonds (Steiner, 2002; Table 1) between pairs of inversion related (-x, 1-y, 1-z) molecules to produce a weakly hydrogen-bonded dimer as shown in Fig. 2.

**S2. Experimental**

The title compound was prepared by stirring a mixture of malonaldehyde bis(dimethyl acetal) (411 mg, 2.5 mmol), ethanamine (366 mg, 2.5 mmol) and acetic acid (300 mg, 5 mmol) under reflux for 1 h. After cooling, ethyl 2-(phenylsulfonyl)acetate (428 mg, 1.87 mmol), DMF (1.5 ml) and 1,8-diazabicyclo[5.4.0]undec-7-ene (DBU, 761 mg, 5 mmol) were added and stirred for 6 h. Then the reaction mixture was added dropwise to ice water (20 ml) to give a yellow solid (259 mg), which was dissolved in 2-propanol. Pale yellow rhomboid-shaped crystals of (I) formed upon evaporation after 7d.

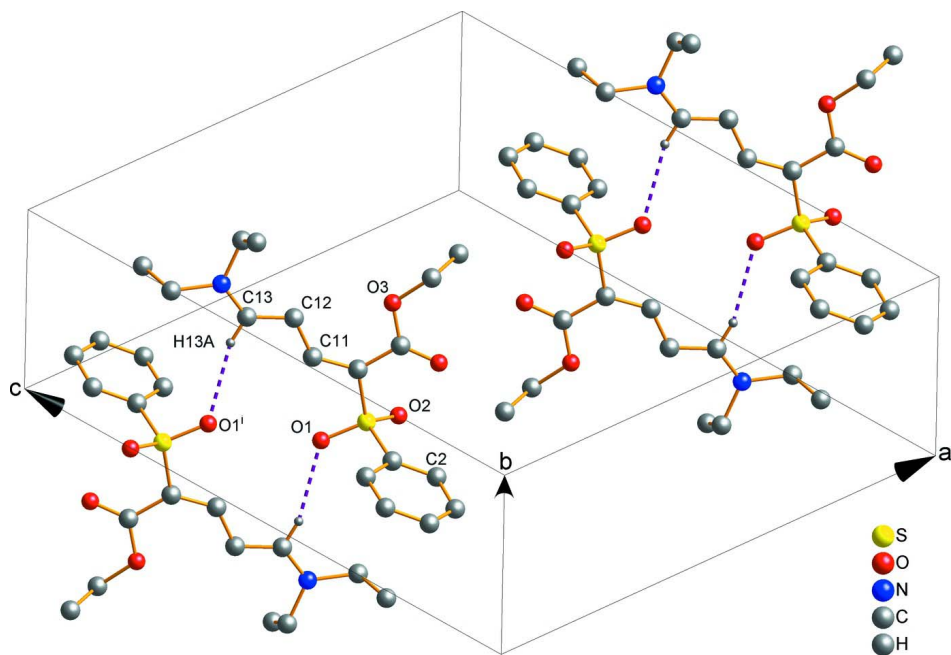
**S3. Refinement**

All the H atoms were positioned geometrically (C—H = 0.93–0.97 Å) and refined as riding, with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{carrier})$  or  $1.5U_{\text{eq}}(\text{methyl C})$ .



**Figure 1**

The molecular structure of (I) showing 30% probability displacement ellipsoids.



**Figure 2**

Crystal packing diagram of (I) with weak hydrogen bonds indicated as dashed lines.

### Ethyl (2*E*,4*Z*)-5-diethylamino-2-(phenylsulfonyl)penta-2,4-dienoate

#### Crystal data

$C_{17}H_{23}NO_4S$

$M_r = 337.42$

Monoclinic,  $P2_1/n$

Hall symbol:  $-P 2_1n$

$a = 14.489 (2) \text{ \AA}$

$b = 8.2989 (12) \text{ \AA}$

$c = 16.706 (3) \text{ \AA}$

$\beta = 114.313 (3)^\circ$

$V = 1830.6 (5) \text{ \AA}^3$   
 $Z = 4$   
 $F(000) = 720$   
 $D_x = 1.224 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$   
 Cell parameters from 3554 reflections

$\theta = 2.7\text{--}25.5^\circ$   
 $\mu = 0.20 \text{ mm}^{-1}$   
 $T = 296 \text{ K}$   
 Block, pale yellow  
 $0.28 \times 0.24 \times 0.20 \text{ mm}$

*Data collection*

Bruker APEXII CCD area-detector  
 diffractometer  
 Radiation source: fine-focus sealed tube  
 Graphite monochromator  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan  
 (SADABS; Sheldrick, 2003)  
 $T_{\min} = 0.669$ ,  $T_{\max} = 0.746$

11618 measured reflections  
 4183 independent reflections  
 2977 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.031$   
 $\theta_{\max} = 27.6^\circ$ ,  $\theta_{\min} = 1.6^\circ$   
 $h = -18 \rightarrow 18$   
 $k = -10 \rightarrow 10$   
 $l = -21 \rightarrow 19$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.059$   
 $wR(F^2) = 0.168$   
 $S = 1.04$   
 4183 reflections  
 211 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0806P)^2 + 0.7143P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.47 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.44 \text{ e \AA}^{-3}$

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

	$x$	$y$	$z$	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.11499 (4)	0.59041 (7)	0.39164 (4)	0.04777 (19)
O1	0.02067 (12)	0.5770 (2)	0.40107 (12)	0.0639 (5)
O2	0.12167 (17)	0.7075 (2)	0.33125 (13)	0.0767 (6)
O3	0.38118 (13)	0.6802 (3)	0.58018 (13)	0.0790 (7)
O4	0.33628 (15)	0.6001 (3)	0.44145 (14)	0.0809 (7)
N1	0.22370 (15)	0.6412 (3)	0.79523 (13)	0.0586 (5)
C1	0.13732 (16)	0.3988 (3)	0.35660 (15)	0.0445 (5)
C2	0.1630 (2)	0.3849 (3)	0.28691 (19)	0.0635 (7)
H2A	0.1709	0.4763	0.2581	0.076*
C3	0.1772 (2)	0.2316 (4)	0.2595 (2)	0.0825 (10)
H3A	0.1942	0.2202	0.2119	0.099*

C4	0.1661 (2)	0.0986 (4)	0.3026 (2)	0.0787 (9)
H4A	0.1764	-0.0032	0.2844	0.094*
C5	0.1404 (3)	0.1130 (3)	0.3716 (2)	0.0789 (9)
H5A	0.1338	0.0213	0.4009	0.095*
C6	0.1240 (2)	0.2627 (3)	0.39856 (18)	0.0623 (7)
H6A	0.1041	0.2724	0.4446	0.075*
C7	0.21004 (16)	0.6228 (3)	0.49653 (15)	0.0455 (5)
C8	0.31330 (19)	0.6342 (3)	0.50106 (17)	0.0572 (6)
C9	0.4909 (2)	0.6711 (6)	0.5980 (3)	0.1000 (12)
H9A	0.5311	0.6566	0.6604	0.120*
H9B	0.5027	0.5796	0.5673	0.120*
C10	0.5202 (4)	0.8158 (8)	0.5691 (4)	0.153 (2)
H10A	0.5916	0.8128	0.5836	0.229*
H10B	0.5053	0.9064	0.5975	0.229*
H10C	0.4836	0.8258	0.5066	0.229*
C11	0.18153 (17)	0.6212 (3)	0.56550 (15)	0.0453 (5)
H11A	0.1130	0.6014	0.5493	0.054*
C12	0.23716 (17)	0.6439 (3)	0.65536 (15)	0.0480 (5)
H12A	0.3055	0.6710	0.6780	0.058*
C13	0.18773 (18)	0.6250 (3)	0.70915 (16)	0.0522 (6)
H13A	0.1197	0.5969	0.6813	0.063*
C14	0.1598 (3)	0.6048 (4)	0.8424 (2)	0.0772 (9)
H14A	0.1037	0.5372	0.8056	0.093*
H14B	0.1993	0.5447	0.8953	0.093*
C15	0.1196 (3)	0.7509 (5)	0.8664 (3)	0.1104 (14)
H15A	0.0789	0.7212	0.8971	0.166*
H15B	0.0788	0.8094	0.8141	0.166*
H15C	0.1748	0.8175	0.9035	0.166*
C16	0.32789 (19)	0.6914 (3)	0.84751 (16)	0.0595 (6)
H16A	0.3300	0.7548	0.8970	0.071*
H16B	0.3504	0.7597	0.8119	0.071*
C17	0.3990 (3)	0.5521 (4)	0.8809 (3)	0.0984 (11)
H17A	0.4649	0.5909	0.9193	0.148*
H17B	0.4034	0.4962	0.8323	0.148*
H17C	0.3744	0.4799	0.9126	0.148*

Atomic displacement parameters ( $\text{\AA}^2$ )

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0517 (3)	0.0390 (3)	0.0470 (3)	0.0037 (2)	0.0147 (2)	-0.0007 (2)
O1	0.0429 (9)	0.0721 (12)	0.0661 (11)	0.0088 (8)	0.0118 (8)	-0.0138 (9)
O2	0.1097 (16)	0.0469 (10)	0.0628 (12)	0.0035 (10)	0.0246 (11)	0.0143 (9)
O3	0.0504 (10)	0.1203 (18)	0.0705 (12)	-0.0272 (11)	0.0292 (9)	-0.0313 (12)
O4	0.0694 (12)	0.1177 (18)	0.0694 (12)	-0.0237 (11)	0.0426 (10)	-0.0264 (12)
N1	0.0557 (12)	0.0719 (14)	0.0515 (12)	-0.0186 (10)	0.0256 (10)	-0.0151 (10)
C1	0.0456 (11)	0.0395 (11)	0.0466 (12)	-0.0035 (9)	0.0172 (9)	-0.0058 (9)
C2	0.0724 (17)	0.0638 (16)	0.0657 (16)	-0.0099 (13)	0.0399 (14)	-0.0077 (13)
C3	0.088 (2)	0.092 (2)	0.090 (2)	-0.0117 (18)	0.0583 (19)	-0.0339 (19)

C4	0.0771 (19)	0.0543 (16)	0.112 (3)	-0.0126 (14)	0.0456 (19)	-0.0323 (17)
C5	0.095 (2)	0.0443 (15)	0.104 (2)	-0.0059 (14)	0.047 (2)	-0.0028 (15)
C6	0.0828 (19)	0.0460 (13)	0.0643 (16)	-0.0011 (12)	0.0366 (14)	0.0006 (12)
C7	0.0447 (11)	0.0422 (11)	0.0499 (12)	-0.0043 (9)	0.0197 (10)	-0.0079 (9)
C8	0.0533 (13)	0.0644 (15)	0.0590 (15)	-0.0143 (12)	0.0284 (12)	-0.0131 (12)
C9	0.0607 (18)	0.154 (4)	0.091 (2)	-0.019 (2)	0.0365 (18)	-0.023 (3)
C10	0.114 (4)	0.185 (6)	0.178 (5)	-0.061 (4)	0.079 (4)	-0.030 (4)
C11	0.0413 (11)	0.0406 (11)	0.0544 (13)	-0.0010 (9)	0.0203 (10)	-0.0077 (10)
C12	0.0435 (11)	0.0493 (12)	0.0522 (13)	-0.0061 (10)	0.0205 (10)	-0.0104 (10)
C13	0.0470 (12)	0.0545 (14)	0.0549 (13)	-0.0104 (10)	0.0208 (10)	-0.0143 (11)
C14	0.082 (2)	0.092 (2)	0.0699 (18)	-0.0248 (17)	0.0428 (16)	-0.0180 (16)
C15	0.102 (3)	0.121 (3)	0.137 (3)	-0.019 (2)	0.079 (3)	-0.033 (3)
C16	0.0630 (15)	0.0623 (16)	0.0482 (13)	-0.0171 (12)	0.0177 (11)	-0.0118 (12)
C17	0.080 (2)	0.078 (2)	0.115 (3)	-0.0047 (18)	0.018 (2)	0.005 (2)

*Geometric parameters (Å, °)*

S1—O2	1.4324 (19)	C9—C10	1.423 (7)
S1—O1	1.4429 (18)	C9—H9A	0.9700
S1—C7	1.747 (2)	C9—H9B	0.9700
S1—C1	1.770 (2)	C10—H10A	0.9600
O3—C8	1.338 (3)	C10—H10B	0.9600
O3—C9	1.494 (4)	C10—H10C	0.9600
O4—C8	1.206 (3)	C11—C12	1.394 (3)
N1—C13	1.319 (3)	C11—H11A	0.9300
N1—C16	1.459 (3)	C12—C13	1.369 (3)
N1—C14	1.473 (3)	C12—H12A	0.9300
C1—C2	1.365 (3)	C13—H13A	0.9300
C1—C6	1.384 (3)	C14—C15	1.471 (5)
C2—C3	1.396 (4)	C14—H14A	0.9700
C2—H2A	0.9300	C14—H14B	0.9700
C3—C4	1.363 (5)	C15—H15A	0.9600
C3—H3A	0.9300	C15—H15B	0.9600
C4—C5	1.355 (5)	C15—H15C	0.9600
C4—H4A	0.9300	C16—C17	1.495 (4)
C5—C6	1.375 (4)	C16—H16A	0.9700
C5—H5A	0.9300	C16—H16B	0.9700
C6—H6A	0.9300	C17—H17A	0.9600
C7—C11	1.375 (3)	C17—H17B	0.9600
C7—C8	1.470 (3)	C17—H17C	0.9600
O2—S1—O1	118.10 (12)	C9—C10—H10A	109.5
O2—S1—C7	110.37 (12)	C9—C10—H10B	109.5
O1—S1—C7	107.05 (11)	H10A—C10—H10B	109.5
O2—S1—C1	107.58 (11)	C9—C10—H10C	109.5
O1—S1—C1	106.07 (11)	H10A—C10—H10C	109.5
C7—S1—C1	107.10 (10)	H10B—C10—H10C	109.5
C8—O3—C9	118.1 (2)	C7—C11—C12	131.4 (2)

C13—N1—C16	121.9 (2)	C7—C11—H11A	114.3
C13—N1—C14	120.6 (2)	C12—C11—H11A	114.3
C16—N1—C14	117.5 (2)	C13—C12—C11	117.7 (2)
C2—C1—C6	120.5 (2)	C13—C12—H12A	121.2
C2—C1—S1	120.57 (19)	C11—C12—H12A	121.2
C6—C1—S1	118.88 (18)	N1—C13—C12	128.7 (2)
C1—C2—C3	119.0 (3)	N1—C13—H13A	115.7
C1—C2—H2A	120.5	C12—C13—H13A	115.7
C3—C2—H2A	120.5	C15—C14—N1	112.5 (3)
C4—C3—C2	120.0 (3)	C15—C14—H14A	109.1
C4—C3—H3A	120.0	N1—C14—H14A	109.1
C2—C3—H3A	120.0	C15—C14—H14B	109.1
C5—C4—C3	120.7 (3)	N1—C14—H14B	109.1
C5—C4—H4A	119.6	H14A—C14—H14B	107.8
C3—C4—H4A	119.6	C14—C15—H15A	109.5
C4—C5—C6	120.2 (3)	C14—C15—H15B	109.5
C4—C5—H5A	119.9	H15A—C15—H15B	109.5
C6—C5—H5A	119.9	C14—C15—H15C	109.5
C5—C6—C1	119.5 (3)	H15A—C15—H15C	109.5
C5—C6—H6A	120.3	H15B—C15—H15C	109.5
C1—C6—H6A	120.3	N1—C16—C17	112.8 (2)
C11—C7—C8	127.5 (2)	N1—C16—H16A	109.0
C11—C7—S1	117.03 (17)	C17—C16—H16A	109.0
C8—C7—S1	115.27 (17)	N1—C16—H16B	109.0
O4—C8—O3	122.8 (2)	C17—C16—H16B	109.0
O4—C8—C7	124.3 (2)	H16A—C16—H16B	107.8
O3—C8—C7	112.8 (2)	C16—C17—H17A	109.5
C10—C9—O3	109.3 (4)	C16—C17—H17B	109.5
C10—C9—H9A	109.8	H17A—C17—H17B	109.5
O3—C9—H9A	109.8	C16—C17—H17C	109.5
C10—C9—H9B	109.8	H17A—C17—H17C	109.5
O3—C9—H9B	109.8	H17B—C17—H17C	109.5
H9A—C9—H9B	108.3		
O2—S1—C1—C2	4.3 (2)	C1—S1—C7—C8	64.6 (2)
O1—S1—C1—C2	131.5 (2)	C9—O3—C8—O4	-7.0 (5)
C7—S1—C1—C2	-114.4 (2)	C9—O3—C8—C7	170.3 (3)
O2—S1—C1—C6	-172.9 (2)	C11—C7—C8—O4	163.0 (3)
O1—S1—C1—C6	-45.7 (2)	S1—C7—C8—O4	-11.1 (4)
C7—S1—C1—C6	68.4 (2)	C11—C7—C8—O3	-14.3 (4)
C6—C1—C2—C3	-1.0 (4)	S1—C7—C8—O3	171.59 (19)
S1—C1—C2—C3	-178.2 (2)	C8—O3—C9—C10	86.6 (4)
C1—C2—C3—C4	-0.5 (5)	C8—C7—C11—C12	6.7 (4)
C2—C3—C4—C5	0.7 (5)	S1—C7—C11—C12	-179.3 (2)
C3—C4—C5—C6	0.7 (5)	C7—C11—C12—C13	-176.0 (2)
C4—C5—C6—C1	-2.2 (5)	C16—N1—C13—C12	2.6 (4)
C2—C1—C6—C5	2.4 (4)	C14—N1—C13—C12	-175.8 (3)
S1—C1—C6—C5	179.6 (2)	C11—C12—C13—N1	-179.4 (2)

O2—S1—C7—C11	132.99 (19)	C13—N1—C14—C15	-101.9 (3)
O1—S1—C7—C11	3.2 (2)	C16—N1—C14—C15	79.6 (4)
C1—S1—C7—C11	-110.18 (18)	C13—N1—C16—C17	-91.9 (3)
O2—S1—C7—C8	-52.3 (2)	C14—N1—C16—C17	86.6 (3)
O1—S1—C7—C8	177.97 (18)		

*Hydrogen-bond geometry (Å, °)*

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
C13—H13A...O1 <sup>i</sup>	0.93	2.41	3.277 (3)	155

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