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## Structure Reports

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## *N,N*-Diethyl-2-(4-methylbenzene-sulfonamido)benzamide

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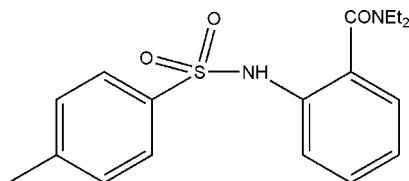
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Key indicators: single-crystal X-ray study;  $T = 150$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.045;  $wR$  factor = 0.118; data-to-parameter ratio = 17.0.

The asymmetric unit of the title compound,  $\text{C}_{18}\text{H}_{22}\text{N}_2\text{O}_3\text{S}$ , contains two molecules, exhibiting similar conformations [ $\text{C}-\text{S}-\text{N}-\text{C}$  torsion angles of  $-82.2$  (2) and  $-70.4$  (2)°, and dihedral angles between the mean planes of the aromatic rings of  $56.6$  (6) and  $51.6$  (6)° in molecules I and II, respectively]. However, the two independent molecules show distinctly different hydrogen-bonding patterns. In the crystal, molecules I form inversion dimers *via* pairs of  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds, whereas for molecules II the  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bond is intramolecular. The hydrogen-bonded dimers of I further propagate along the  $b$ -axis direction through  $\pi-\pi$  interactions [the distance between ring centroids is  $3.8424$  (8) Å].

### Related literature

For the synthesis of the title compound, see: Bakker *et al.* (1997); Kaul *et al.* (2002). For the biological activity of compounds having the sulfonamide  $-\text{SO}_2\text{NH}-$  group, see: Lu & Tucker (2007); Tappe *et al.* (2008); Chegwiddden *et al.* (2000); Purushottamachar *et al.* (2008). For structural and conformational studies of molecules featuring the sulfonamide moiety, see: Parkin *et al.* (2008); Perlovich *et al.* (2009, 2011); Altamura *et al.* (2009); Vega-Hissi *et al.* (2011).



### Experimental

#### Crystal data

$\text{C}_{18}\text{H}_{22}\text{N}_2\text{O}_3\text{S}$   
 $M_r = 346.43$

Triclinic,  $P\bar{1}$   
 $a = 9.4674$  (6) Å

$b = 12.2882$  (9) Å  
 $c = 16.0569$  (12) Å  
 $\alpha = 108.426$  (7)°  
 $\beta = 97.357$  (6)°  
 $\gamma = 100.245$  (6)°  
 $V = 1709.7$  (2) Å<sup>3</sup>

$Z = 4$   
Mo  $K\alpha$  radiation  
 $\mu = 0.21$  mm<sup>-1</sup>  
 $T = 150$  K  
 $0.54 \times 0.43 \times 0.38$  mm

#### Data collection

Oxford Diffraction Xcalibur3 CCD diffractometer  
Absorption correction: multi-scan (ABSPACK in CrysAlis RED; Oxford Diffraction, 2006)  
 $T_{\min} = 0.894$ ,  $T_{\max} = 1.000$   
17890 measured reflections  
7512 independent reflections  
4728 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.025$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$   
 $wR(F^2) = 0.118$   
 $S = 0.96$   
7512 reflections  
441 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.42$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.32$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}'-\text{HN1}'\cdots\text{O3}'$	0.81 (2)	2.15 (2)	2.809 (2)	139 (2)
$\text{N1}-\text{HN1}\cdots\text{O3}$	0.86 (2)	2.15 (2)	2.969 (2)	159 (2)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2006); cell refinement: *CrysAlis CCD*; data reduction: *CrysAlis RED* (Oxford Diffraction, 2006); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*, *WinGX* (Farrugia, 1999) and *PARST* (Nardelli, 1995).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LD2075).

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## supporting information

*Acta Cryst.* (2012). E68, o3144–o3145 [doi:10.1107/S160053681204264X]

***N,N*-Diethyl-2-(4-methylbenzenesulfonamido)benzamide**

**Maria Altamura, Valentina Fedi, Rossano Nannicini, Paola Paoli and Patrizia Rossi**

**S1. Comment**

The sulfonamide moiety is a common pharmacophore in many biologically active compounds, such as HIV inhibitors (Lu & Tucker, 2007), antimicrobial drugs (Tappe *et al.*, 2008), carbonic anhydrase inhibitors (Chegwidden *et al.*, 2000), and anti-tumor agents (Purushottamachar *et al.*, 2008). Because the structural and conformational properties of a compound usually are related to its biological properties, their study would provide useful information to design new effective drugs. In this regard, there are many recent publications reporting structural data on related sulfonamides (Parkin *et al.*, 2008, Altamura *et al.*, 2009, Perlovich *et al.*, 2009, Perlovich *et al.*, 2011, Vega-Hissi *et al.*, 2011).

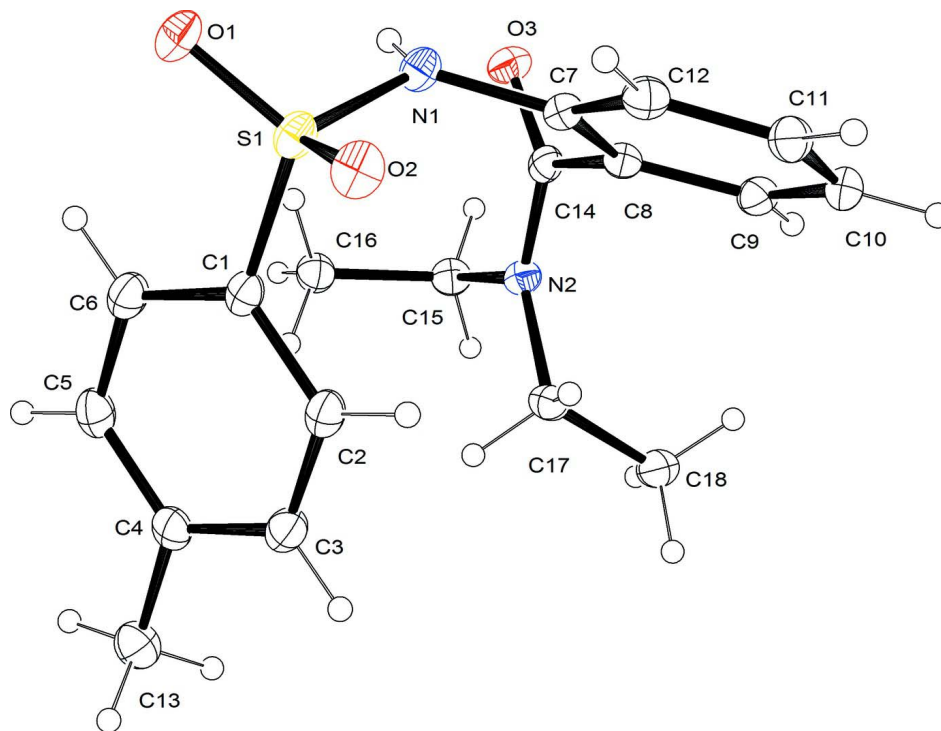
The asymmetric unit of the title compound contains two independent molecules, I and II, which are almost superimposable (Table 1). As expected, a staggered conformation about the N—S bond is adopted, with the N lone pair bisecting the O $\hat{S}$ O angle, and with the p orbital at the *ipso* carbon bisecting the same angle (Table 1, Fig. 1). The sulfonamide nitrogen atom is almost planar-trigonal in molecule I ( $\Sigma\langle N=355$  (1) $^\circ$ ), while in II it is definitely more pyramidal ( $\Sigma\langle N=341$  (1) $^\circ$ ). The conformation of molecule II is stabilized by an intramolecular H-bond involving the H atom of the sulfonamide grouping (HN1') and the oxygen atom O3' of the amide moiety (Table 2). In the crystal packing, molecules I form dimers instead, which are held together by a couple of N—H $\cdots$ O=C hydrogen bonds (Table 2, Fig. 2). Dimers propagate along the *b* axis direction through  $\pi$ - $\pi$  stacking interactions involving two symmetry related C1—C6 rings (centroid-centroid distance 3.8424 (8) Å, symmetry code:  $-x + 2, -y + 1, -z$ ). No further significant intermolecular interactions are present in the crystal structure.

**S2. Experimental**

For the synthesis of the title compound, see: Bakker *et al.* (1997); Kaul *et al.*, (2002). Crystals of *N,N*-diethyl-2-(4-methylphenylsulfonamido)benzamide suitable for single-crystal X-ray diffraction analysis were obtained by slow evaporation of an ethanol/water solution of *N,N*-diethyl-2-(4-methylphenylsulfonamido)benzamide.

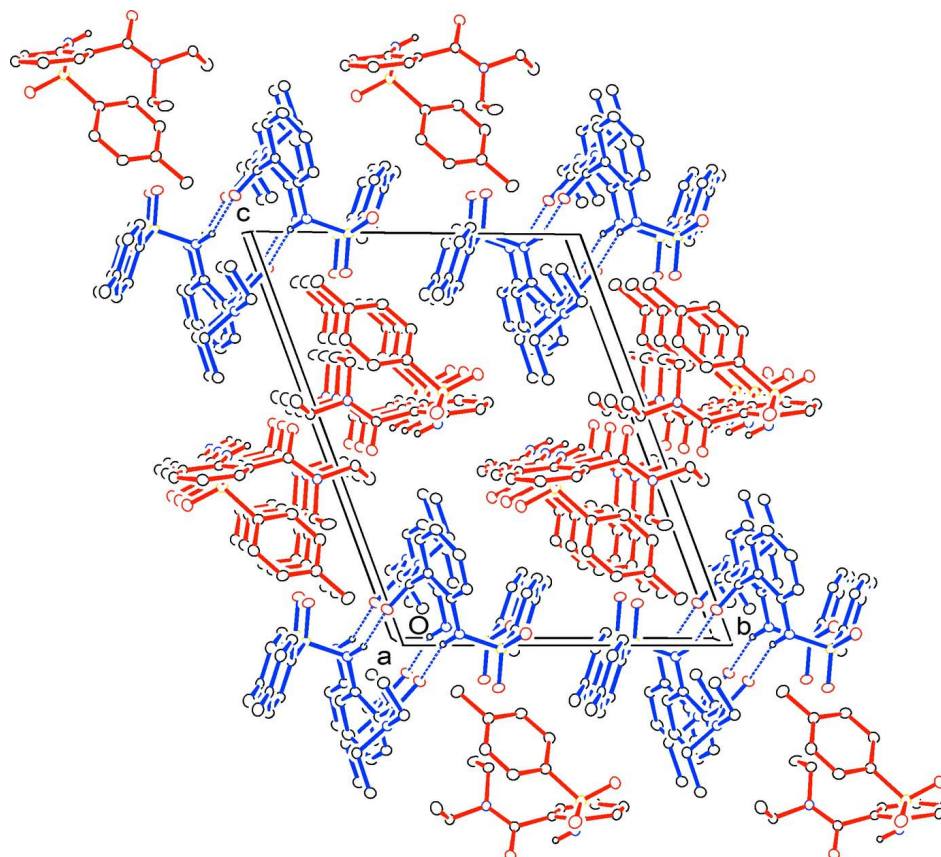
**S3. Refinement**

The N—H H atoms were located in the Fourier difference map and their coordinates were refined with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$ . All other H atoms were positioned using idealized geometry, and refined using a riding model with  $U_{\text{iso}}(\text{H})$  1.2 times  $U_{\text{eq}}(\text{C})$  (1.5 for methyl H atoms).



**Figure 1**

The symmetrically independent molecule I (molecule II has a similar shape and the same labelling scheme). Displacement ellipsoids are drawn at the 30% probability level.

**Figure 2**

Crystal structure of the title compound as viewed along the *a*-axis (molecules I are shown in blue and molecules II - in red). Intermolecular NH $\cdots$ O hydrogen bonding is shown as dashed lines.

### *N,N*-Diethyl-2-(4-methylbenzenesulfonamido)benzamide

#### Crystal data

$C_{18}H_{22}N_2O_3S$

$M_r = 346.43$

Triclinic,  $P\bar{1}$

$a = 9.4674$  (6) Å

$b = 12.2882$  (9) Å

$c = 16.0569$  (12) Å

$\alpha = 108.426$  (7)°

$\beta = 97.357$  (6)°

$\gamma = 100.245$  (6)°

$V = 1709.7$  (2) Å<sup>3</sup>

$Z = 4$

$F(000) = 736$

$D_x = 1.346$  Mg m<sup>-3</sup>

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

$\theta = 4.1$ – $28.6$ °

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

$T = 150$  K

Parallelepiped, colourless

$0.54 \times 0.43 \times 0.38$  mm

#### Data collection

Oxford Diffraction Xcalibur3 CCD  
diffractometer

Radiation source: Enhance (Mo) X-ray Source

Graphite monochromator

Detector resolution: 16.4547 pixels mm<sup>-1</sup>

$\omega$  scans

Absorption correction: multi-scan

(*ABSPACK* in *CrysAlis RED*; Oxford

Diffraction, 2006)

$T_{\min} = 0.894$ ,  $T_{\max} = 1.000$

17890 measured reflections

7512 independent reflections

4728 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$   
 $\theta_{\text{max}} = 28.7^\circ$ ,  $\theta_{\text{min}} = 4.1^\circ$   
 $h = -12 \rightarrow 11$

$k = -15 \rightarrow 16$   
 $l = -21 \rightarrow 21$

### Refinement

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.045$   
 $wR(F^2) = 0.118$   
 $S = 0.96$   
 7512 reflections  
 441 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 atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0677P)^2]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} = 0.001$   
 $\Delta\rho_{\text{max}} = 0.42 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\text{min}} = -0.32 \text{ e } \text{\AA}^{-3}$

### Special details

**Geometry.** All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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  $F^2 > 2\sigma(F^2)$  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	1.22768 (5)	0.28575 (4)	-0.00356 (3)	0.02970 (14)
O1	1.18234 (15)	0.23441 (12)	-0.09887 (8)	0.0365 (3)
O2	1.36507 (14)	0.36978 (12)	0.03457 (9)	0.0365 (3)
O3	1.03929 (14)	-0.01020 (12)	0.08734 (9)	0.0329 (3)
N1	1.23284 (17)	0.17480 (15)	0.03138 (10)	0.0278 (4)
HN1	1.158 (2)	0.1167 (18)	0.0058 (13)	0.033*
N2	0.98318 (16)	0.13823 (13)	0.19374 (10)	0.0257 (4)
C1	1.0900 (2)	0.35179 (16)	0.03962 (12)	0.0268 (4)
C2	1.1191 (2)	0.42734 (17)	0.12797 (13)	0.0311 (5)
H2	1.2108	0.4429	0.1638	0.037*
C3	1.0114 (2)	0.47931 (17)	0.16265 (13)	0.0331 (5)
H3	1.0315	0.5296	0.2220	0.040*
C4	0.8731 (2)	0.45770 (17)	0.11021 (13)	0.0308 (5)
C5	0.8479 (2)	0.38376 (18)	0.02171 (14)	0.0348 (5)
H5	0.7570	0.3698	-0.0146	0.042*
C6	0.9537 (2)	0.33011 (18)	-0.01428 (13)	0.0334 (5)
H6	0.9339	0.2802	-0.0737	0.040*
C7	1.30951 (19)	0.18780 (16)	0.11830 (12)	0.0262 (4)
C8	1.23673 (19)	0.14737 (16)	0.17723 (12)	0.0257 (4)
C9	1.3163 (2)	0.15919 (17)	0.26008 (12)	0.0304 (5)
H9	1.2691	0.1312	0.2991	0.037*

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C10	1.4645 (2)	0.21191 (18)	0.28524 (13)	0.0343 (5)
H10	1.5162	0.2206	0.3411	0.041*
C11	1.5345 (2)	0.25127 (18)	0.22645 (13)	0.0332 (5)
H11	1.6340	0.2868	0.2431	0.040*
C12	1.4591 (2)	0.23881 (17)	0.14310 (13)	0.0305 (5)
H12	1.5082	0.2644	0.1037	0.037*
C13	0.7557 (2)	0.51226 (19)	0.14887 (15)	0.0397 (5)
H13A	0.6650	0.4798	0.1062	0.060*
H13B	0.7826	0.5960	0.1626	0.060*
H13C	0.7444	0.4959	0.2026	0.060*
C14	1.0785 (2)	0.08570 (17)	0.14925 (12)	0.0250 (4)
C15	0.82675 (19)	0.07980 (17)	0.16405 (13)	0.0293 (4)
H15A	0.7766	0.1054	0.2133	0.035*
H15B	0.8154	-0.0048	0.1472	0.035*
C16	0.7572 (2)	0.10750 (18)	0.08525 (13)	0.0326 (5)
H16A	0.6554	0.0680	0.0677	0.049*
H16B	0.8054	0.0810	0.0360	0.049*
H16C	0.7666	0.1911	0.1020	0.049*
C17	1.0220 (2)	0.25858 (16)	0.25954 (12)	0.0291 (4)
H17A	0.9527	0.3020	0.2443	0.035*
H17B	1.1183	0.2979	0.2561	0.035*
C18	1.0228 (2)	0.26194 (18)	0.35513 (13)	0.0373 (5)
H18A	1.0486	0.3424	0.3951	0.056*
H18B	1.0930	0.2208	0.3712	0.056*
H18C	0.9272	0.2247	0.3594	0.056*
S1'	0.36428 (5)	0.62169 (4)	0.38496 (3)	0.02947 (14)
O1'	0.25700 (14)	0.66660 (13)	0.43239 (9)	0.0389 (4)
O2'	0.34408 (14)	0.49807 (12)	0.34048 (9)	0.0356 (3)
O3'	0.71780 (14)	0.88703 (11)	0.52660 (9)	0.0346 (3)
N1'	0.51694 (17)	0.66700 (15)	0.46038 (11)	0.0279 (4)
HN1'	0.534 (2)	0.7376 (18)	0.4848 (14)	0.033*
N2'	0.81608 (16)	0.90225 (13)	0.40820 (10)	0.0263 (4)
C1'	0.39078 (18)	0.69322 (17)	0.30717 (12)	0.0273 (4)
C2'	0.45199 (19)	0.64421 (18)	0.23389 (13)	0.0297 (4)
H2'	0.4787	0.5727	0.2251	0.036*
C3'	0.4727 (2)	0.70303 (18)	0.17430 (13)	0.0316 (5)
H3'	0.5144	0.6706	0.1256	0.038*
C4'	0.43286 (19)	0.80900 (18)	0.18556 (13)	0.0314 (5)
C5'	0.3715 (2)	0.85577 (19)	0.25892 (14)	0.0369 (5)
H5'	0.3431	0.9265	0.2671	0.044*
C6'	0.3517 (2)	0.79973 (18)	0.32016 (13)	0.0345 (5)
H6'	0.3124	0.8333	0.3697	0.041*
C7'	0.64714 (19)	0.63062 (17)	0.43794 (12)	0.0246 (4)
C8'	0.76801 (19)	0.71269 (16)	0.43306 (12)	0.0251 (4)
C9'	0.8955 (2)	0.67471 (17)	0.41748 (12)	0.0279 (4)
H9'	0.9765	0.7277	0.4145	0.034*
C10'	0.9044 (2)	0.56046 (18)	0.40639 (13)	0.0319 (5)
H10'	0.9906	0.5369	0.3963	0.038*

C11'	0.7839 (2)	0.48063 (17)	0.41038 (13)	0.0316 (5)
H11'	0.7894	0.4033	0.4025	0.038*
C12'	0.6555 (2)	0.51581 (17)	0.42602 (13)	0.0293 (4)
H12'	0.5750	0.4620	0.4285	0.035*
C13'	0.4544 (2)	0.8733 (2)	0.12051 (14)	0.0437 (6)
H13D	0.5286	0.8482	0.0885	0.066*
H13E	0.4842	0.9567	0.1528	0.066*
H13F	0.3642	0.8559	0.0789	0.066*
C14'	0.76384 (19)	0.84015 (17)	0.45779 (12)	0.0263 (4)
C15'	0.8238 (2)	1.02962 (17)	0.44075 (13)	0.0318 (5)
H15C	0.9033	1.0690	0.4203	0.038*
H15D	0.8465	1.0589	0.5057	0.038*
C16'	0.6845 (2)	1.0613 (2)	0.40996 (15)	0.0457 (6)
H16D	0.6968	1.1454	0.4334	0.069*
H16E	0.6056	1.0244	0.4312	0.069*
H16F	0.6624	1.0344	0.3457	0.069*
C17'	0.8356 (2)	0.85147 (18)	0.31525 (12)	0.0305 (5)
H17C	0.7752	0.8800	0.2772	0.037*
H17D	0.8015	0.7664	0.2952	0.037*
C18'	0.9930 (2)	0.8811 (2)	0.30377 (14)	0.0443 (6)
H18D	0.9980	0.8455	0.2420	0.066*
H18E	1.0534	0.8515	0.3400	0.066*
H18F	1.0270	0.9651	0.3220	0.066*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0316 (3)	0.0309 (3)	0.0249 (3)	0.0009 (2)	0.0054 (2)	0.0110 (2)
O1	0.0445 (8)	0.0397 (9)	0.0227 (7)	0.0037 (7)	0.0059 (6)	0.0109 (7)
O2	0.0314 (7)	0.0366 (8)	0.0390 (8)	-0.0025 (6)	0.0063 (6)	0.0153 (7)
O3	0.0342 (7)	0.0305 (8)	0.0278 (7)	0.0052 (6)	0.0064 (6)	0.0029 (7)
N1	0.0266 (8)	0.0268 (10)	0.0249 (9)	-0.0003 (7)	0.0016 (7)	0.0071 (8)
N2	0.0283 (8)	0.0233 (9)	0.0245 (8)	0.0048 (7)	0.0064 (7)	0.0069 (7)
C1	0.0323 (10)	0.0229 (10)	0.0241 (10)	-0.0005 (8)	0.0027 (8)	0.0113 (9)
C2	0.0320 (11)	0.0308 (12)	0.0274 (11)	0.0020 (9)	-0.0003 (9)	0.0111 (9)
C3	0.0401 (12)	0.0313 (12)	0.0259 (11)	0.0053 (9)	0.0046 (9)	0.0097 (10)
C4	0.0377 (11)	0.0225 (11)	0.0340 (12)	0.0039 (9)	0.0046 (9)	0.0150 (10)
C5	0.0301 (11)	0.0346 (12)	0.0383 (12)	0.0048 (9)	-0.0028 (9)	0.0159 (10)
C6	0.0372 (11)	0.0312 (12)	0.0268 (11)	0.0021 (9)	-0.0005 (9)	0.0090 (10)
C7	0.0280 (10)	0.0261 (11)	0.0225 (10)	0.0079 (8)	0.0044 (8)	0.0052 (9)
C8	0.0284 (10)	0.0219 (10)	0.0263 (10)	0.0073 (8)	0.0053 (8)	0.0069 (9)
C9	0.0367 (11)	0.0311 (12)	0.0254 (10)	0.0115 (9)	0.0082 (9)	0.0096 (9)
C10	0.0358 (11)	0.0357 (12)	0.0276 (11)	0.0132 (9)	-0.0023 (9)	0.0063 (10)
C11	0.0238 (10)	0.0357 (12)	0.0338 (11)	0.0071 (9)	0.0004 (9)	0.0055 (10)
C12	0.0294 (10)	0.0304 (11)	0.0317 (11)	0.0075 (9)	0.0089 (9)	0.0093 (10)
C13	0.0403 (12)	0.0358 (13)	0.0450 (13)	0.0087 (10)	0.0081 (10)	0.0168 (11)
C14	0.0305 (10)	0.0246 (11)	0.0216 (10)	0.0052 (8)	0.0050 (8)	0.0108 (9)
C15	0.0265 (10)	0.0307 (11)	0.0315 (11)	0.0053 (8)	0.0098 (9)	0.0108 (9)



C16	0.0302 (10)	0.0376 (12)	0.0325 (11)	0.0104 (9)	0.0082 (9)	0.0137 (10)
C17	0.0333 (11)	0.0253 (11)	0.0313 (11)	0.0103 (9)	0.0095 (9)	0.0101 (9)
C18	0.0493 (13)	0.0328 (12)	0.0299 (11)	0.0142 (10)	0.0081 (10)	0.0086 (10)
S1'	0.0235 (2)	0.0323 (3)	0.0314 (3)	0.0011 (2)	0.0062 (2)	0.0119 (2)
O1'	0.0278 (7)	0.0482 (9)	0.0444 (9)	0.0060 (7)	0.0164 (7)	0.0190 (8)
O2'	0.0335 (7)	0.0288 (8)	0.0369 (8)	-0.0040 (6)	0.0007 (6)	0.0093 (7)
O3'	0.0431 (8)	0.0284 (8)	0.0290 (8)	0.0027 (6)	0.0121 (7)	0.0067 (7)
N1'	0.0286 (9)	0.0247 (9)	0.0289 (9)	0.0024 (8)	0.0072 (7)	0.0090 (8)
N2'	0.0269 (8)	0.0237 (9)	0.0255 (9)	-0.0003 (7)	0.0012 (7)	0.0096 (7)
C1'	0.0190 (9)	0.0322 (11)	0.0278 (10)	0.0023 (8)	0.0008 (8)	0.0100 (9)
C2'	0.0238 (10)	0.0305 (11)	0.0316 (11)	0.0048 (8)	0.0037 (9)	0.0082 (10)
C3'	0.0273 (10)	0.0388 (13)	0.0276 (11)	0.0041 (9)	0.0049 (9)	0.0119 (10)
C4'	0.0240 (10)	0.0355 (12)	0.0316 (11)	-0.0006 (9)	-0.0020 (9)	0.0144 (10)
C5'	0.0406 (12)	0.0336 (12)	0.0381 (12)	0.0125 (10)	0.0042 (10)	0.0138 (11)
C6'	0.0345 (11)	0.0387 (13)	0.0312 (11)	0.0134 (10)	0.0085 (9)	0.0100 (10)
C7'	0.0242 (9)	0.0288 (11)	0.0199 (9)	0.0034 (8)	0.0026 (8)	0.0095 (9)
C8'	0.0266 (10)	0.0249 (11)	0.0200 (10)	-0.0006 (8)	0.0006 (8)	0.0079 (9)
C9'	0.0262 (10)	0.0291 (11)	0.0264 (10)	0.0008 (8)	0.0035 (8)	0.0105 (9)
C10'	0.0272 (10)	0.0349 (12)	0.0312 (11)	0.0062 (9)	0.0034 (9)	0.0097 (10)
C11'	0.0366 (11)	0.0260 (11)	0.0304 (11)	0.0069 (9)	0.0023 (9)	0.0090 (9)
C12'	0.0298 (10)	0.0268 (11)	0.0299 (11)	-0.0005 (8)	0.0024 (9)	0.0133 (9)
C13'	0.0447 (13)	0.0475 (14)	0.0386 (13)	0.0027 (11)	0.0016 (11)	0.0216 (11)
C14'	0.0241 (10)	0.0251 (11)	0.0256 (10)	0.0005 (8)	0.0008 (8)	0.0076 (9)
C15'	0.0361 (11)	0.0240 (11)	0.0303 (11)	-0.0016 (9)	0.0020 (9)	0.0092 (9)
C16'	0.0511 (14)	0.0364 (13)	0.0467 (14)	0.0125 (11)	0.0002 (11)	0.0128 (11)
C17'	0.0304 (10)	0.0321 (12)	0.0246 (10)	-0.0013 (9)	0.0012 (9)	0.0102 (9)
C18'	0.0385 (12)	0.0611 (16)	0.0281 (11)	0.0002 (11)	0.0080 (10)	0.0141 (11)

*Geometric parameters (Å, °)*

S1—O1	1.4315 (13)	S1'—O2'	1.4248 (14)
S1—O2	1.4324 (13)	S1'—O1'	1.4311 (14)
S1—N1	1.6358 (17)	S1'—N1'	1.6446 (17)
S1—C1	1.758 (2)	S1'—C1'	1.758 (2)
O3—C14	1.235 (2)	O3'—C14'	1.246 (2)
N1—C7	1.435 (2)	N1'—C7'	1.434 (2)
N1—HN1	0.86 (2)	N1'—HN1'	0.81 (2)
N2—C14	1.350 (2)	N2'—C14'	1.346 (2)
N2—C17	1.469 (2)	N2'—C15'	1.470 (2)
N2—C15	1.470 (2)	N2'—C17'	1.475 (2)
C1—C2	1.388 (3)	C1'—C6'	1.383 (3)
C1—C6	1.390 (3)	C1'—C2'	1.391 (3)
C2—C3	1.381 (3)	C2'—C3'	1.382 (3)
C2—H2	0.9300	C2'—H2'	0.9300
C3—C4	1.395 (3)	C3'—C4'	1.384 (3)
C3—H3	0.9300	C3'—H3'	0.9300
C4—C5	1.386 (3)	C4'—C5'	1.386 (3)
C4—C13	1.500 (3)	C4'—C13'	1.509 (3)

C5—C6	1.384 (3)	C5'—C6'	1.380 (3)
C5—H5	0.9300	C5'—H5'	0.9300
C6—H6	0.9300	C6'—H6'	0.9300
C7—C12	1.393 (3)	C7'—C12'	1.381 (3)
C7—C8	1.400 (3)	C7'—C8'	1.413 (2)
C8—C9	1.393 (3)	C8'—C9'	1.393 (3)
C8—C14	1.493 (2)	C8'—C14'	1.498 (3)
C9—C10	1.385 (3)	C9'—C10'	1.377 (3)
C9—H9	0.9300	C9'—H9'	0.9300
C10—C11	1.378 (3)	C10'—C11'	1.388 (3)
C10—H10	0.9300	C10'—H10'	0.9300
C11—C12	1.382 (3)	C11'—C12'	1.388 (3)
C11—H11	0.9300	C11'—H11'	0.9300
C12—H12	0.9300	C12'—H12'	0.9300
C13—H13A	0.9600	C13'—H13D	0.9600
C13—H13B	0.9600	C13'—H13E	0.9600
C13—H13C	0.9600	C13'—H13F	0.9600
C15—C16	1.513 (3)	C15'—C16'	1.508 (3)
C15—H15A	0.9700	C15'—H15C	0.9700
C15—H15B	0.9700	C15'—H15D	0.9700
C16—H16A	0.9600	C16'—H16D	0.9600
C16—H16B	0.9600	C16'—H16E	0.9600
C16—H16C	0.9600	C16'—H16F	0.9600
C17—C18	1.521 (3)	C17'—C18'	1.518 (3)
C17—H17A	0.9700	C17'—H17C	0.9700
C17—H17B	0.9700	C17'—H17D	0.9700
C18—H18A	0.9600	C18'—H18D	0.9600
C18—H18B	0.9600	C18'—H18E	0.9600
C18—H18C	0.9600	C18'—H18F	0.9600
O1—S1—O2	119.57 (8)	O2'—S1'—O1'	120.27 (8)
O1—S1—N1	105.49 (8)	O2'—S1'—N1'	107.60 (8)
O2—S1—N1	107.66 (8)	O1'—S1'—N1'	104.98 (8)
O1—S1—C1	108.16 (9)	O2'—S1'—C1'	108.68 (9)
O2—S1—C1	107.99 (9)	O1'—S1'—C1'	107.76 (9)
N1—S1—C1	107.40 (8)	N1'—S1'—C1'	106.80 (8)
C7—N1—S1	123.27 (13)	C7'—N1'—S1'	120.56 (13)
C7—N1—HN1	118.4 (13)	C7'—N1'—HN1'	110.6 (14)
S1—N1—HN1	113.0 (13)	S1'—N1'—HN1'	110.0 (15)
C14—N2—C17	124.10 (15)	C14'—N2'—C15'	117.36 (15)
C14—N2—C15	117.84 (15)	C14'—N2'—C17'	125.41 (16)
C17—N2—C15	117.24 (14)	C15'—N2'—C17'	115.80 (15)
C2—C1—C6	120.03 (18)	C6'—C1'—C2'	120.19 (18)
C2—C1—S1	119.44 (14)	C6'—C1'—S1'	119.29 (14)
C6—C1—S1	120.52 (15)	C2'—C1'—S1'	120.51 (15)
C3—C2—C1	119.89 (18)	C3'—C2'—C1'	119.20 (19)
C3—C2—H2	120.1	C3'—C2'—H2'	120.4
C1—C2—H2	120.1	C1'—C2'—H2'	120.4

C2—C3—C4	121.13 (18)	C2'—C3'—C4'	121.49 (18)
C2—C3—H3	119.4	C2'—C3'—H3'	119.3
C4—C3—H3	119.4	C4'—C3'—H3'	119.3
C5—C4—C3	117.83 (18)	C3'—C4'—C5'	118.20 (18)
C5—C4—C13	121.32 (18)	C3'—C4'—C13'	121.76 (18)
C3—C4—C13	120.85 (18)	C5'—C4'—C13'	120.04 (19)
C6—C5—C4	122.02 (18)	C6'—C5'—C4'	121.47 (19)
C6—C5—H5	119.0	C6'—C5'—H5'	119.3
C4—C5—H5	119.0	C4'—C5'—H5'	119.3
C5—C6—C1	119.07 (18)	C5'—C6'—C1'	119.44 (18)
C5—C6—H6	120.5	C5'—C6'—H6'	120.3
C1—C6—H6	120.5	C1'—C6'—H6'	120.3
C12—C7—C8	119.84 (17)	C12'—C7'—C8'	120.49 (17)
C12—C7—N1	119.24 (17)	C12'—C7'—N1'	118.96 (16)
C8—C7—N1	120.90 (16)	C8'—C7'—N1'	120.45 (16)
C9—C8—C7	118.94 (17)	C9'—C8'—C7'	118.05 (17)
C9—C8—C14	120.59 (17)	C9'—C8'—C14'	121.72 (16)
C7—C8—C14	120.36 (16)	C7'—C8'—C14'	119.60 (16)
C10—C9—C8	121.10 (18)	C10'—C9'—C8'	121.49 (17)
C10—C9—H9	119.4	C10'—C9'—H9'	119.3
C8—C9—H9	119.4	C8'—C9'—H9'	119.3
C11—C10—C9	119.21 (19)	C9'—C10'—C11'	119.66 (18)
C11—C10—H10	120.4	C9'—C10'—H10'	120.2
C9—C10—H10	120.4	C11'—C10'—H10'	120.2
C10—C11—C12	121.06 (18)	C12'—C11'—C10'	120.24 (18)
C10—C11—H11	119.5	C12'—C11'—H11'	119.9
C12—C11—H11	119.5	C10'—C11'—H11'	119.9
C11—C12—C7	119.83 (18)	C7'—C12'—C11'	120.06 (17)
C11—C12—H12	120.1	C7'—C12'—H12'	120.0
C7—C12—H12	120.1	C11'—C12'—H12'	120.0
C4—C13—H13A	109.5	C4'—C13'—H13D	109.5
C4—C13—H13B	109.5	C4'—C13'—H13E	109.5
H13A—C13—H13B	109.5	H13D—C13'—H13E	109.5
C4—C13—H13C	109.5	C4'—C13'—H13F	109.5
H13A—C13—H13C	109.5	H13D—C13'—H13F	109.5
H13B—C13—H13C	109.5	H13E—C13'—H13F	109.5
O3—C14—N2	122.48 (16)	O3'—C14'—N2'	121.77 (17)
O3—C14—C8	119.72 (16)	O3'—C14'—C8'	118.54 (16)
N2—C14—C8	117.80 (16)	N2'—C14'—C8'	119.61 (16)
N2—C15—C16	111.74 (15)	N2'—C15'—C16'	113.69 (16)
N2—C15—H15A	109.3	N2'—C15'—H15C	108.8
C16—C15—H15A	109.3	C16'—C15'—H15C	108.8
N2—C15—H15B	109.3	N2'—C15'—H15D	108.8
C16—C15—H15B	109.3	C16'—C15'—H15D	108.8
H15A—C15—H15B	107.9	H15C—C15'—H15D	107.7
C15—C16—H16A	109.5	C15'—C16'—H16D	109.5
C15—C16—H16B	109.5	C15'—C16'—H16E	109.5
H16A—C16—H16B	109.5	H16D—C16'—H16E	109.5

C15—C16—H16C	109.5	C15'—C16'—H16F	109.5
H16A—C16—H16C	109.5	H16D—C16'—H16F	109.5
H16B—C16—H16C	109.5	H16E—C16'—H16F	109.5
N2—C17—C18	113.09 (16)	N2'—C17'—C18'	113.74 (15)
N2—C17—H17A	109.0	N2'—C17'—H17C	108.8
C18—C17—H17A	109.0	C18'—C17'—H17C	108.8
N2—C17—H17B	109.0	N2'—C17'—H17D	108.8
C18—C17—H17B	109.0	C18'—C17'—H17D	108.8
H17A—C17—H17B	107.8	H17C—C17'—H17D	107.7
C17—C18—H18A	109.5	C17'—C18'—H18D	109.5
C17—C18—H18B	109.5	C17'—C18'—H18E	109.5
H18A—C18—H18B	109.5	H18D—C18'—H18E	109.5
C17—C18—H18C	109.5	C17'—C18'—H18F	109.5
H18A—C18—H18C	109.5	H18D—C18'—H18F	109.5
H18B—C18—H18C	109.5	H18E—C18'—H18F	109.5
C1—S1—N1—C7	-82.2 (2)	C1'—S1'—N1'—C7'	-70.4 (2)
HN1—N1—S1—O1	-44 (1)	HN1'—N1'—S1'—O1'	-54 (2)
C7—N1—S1—O2	33.9 (2)	C7'—N1'—S1'—O2'	46.1 (2)
C6—C1—S1—O1	11.7 (2)	C6'—C1'—S1'—O1'	20.8 (2)

*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>
N1'—HN1'...O3'	0.81 (2)	2.15 (2)	2.809 (2)	139 (2)
N1—HN1...O3 <sup>i</sup>	0.86 (2)	2.15 (2)	2.969 (2)	159 (2)

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