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N-(2-Methoxyphenyl)benzene-sulfonamide

 Aziz-ur-Rehman,^{a*} Muhammad Arif Sajjad,^a Mehmet Akkurt,^{b*} Shahzad Sharif,^a Muhammad Athar Abbasi^a and Islam Ullah Khan^a
^aDepartment of Chemistry, Government College University, Lahore 54000, Pakistan, and ^bDepartment of Physics, Faculty of Arts and Sciences, Erciyes University, 38039 Kayseri, Turkey

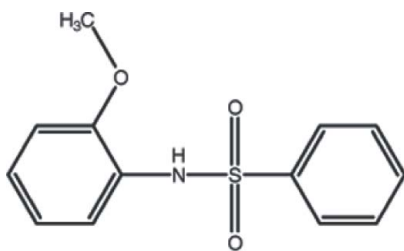
Correspondence e-mail: azizhej@hotmail.com, akkurt@erciyes.edu.tr

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 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.047; wR factor = 0.113; data-to-parameter ratio = 19.0.

The asymmetric unit of the title compound, $\text{C}_{13}\text{H}_{13}\text{NO}_3\text{S}$, contains two crystallographically independent molecules in which the dihedral angles between the phenyl and benzene rings are 88.16 (12) and 44.50 (12)°. One of the molecules features an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond. In the crystal, the molecules are linked into dimers by pairs of $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds. The dimers are further connected by $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\pi$ interactions, forming a three-dimensional network.

Related literature

 For the biological activity of sulfonamides, see: Arshad *et al.* (2008); Gennarti *et al.* (1994); Kayser *et al.* (2004); Rough *et al.* (1998).


Experimental

Crystal data

 $\text{C}_{13}\text{H}_{13}\text{NO}_3\text{S}$
 $M_r = 263.31$

 Monoclinic, $P2_1/n$
 $a = 8.7705$ (2) Å

 $b = 28.1684$ (7) Å
 $c = 10.7256$ (3) Å
 $\beta = 105.968$ (1)°
 $V = 2547.53$ (11) Å³
 $Z = 8$

 Mo $K\alpha$ radiation
 $\mu = 0.25$ mm⁻¹
 $T = 296$ K
 $0.25 \times 0.17 \times 0.07$ mm

Data collection

 Bruker APEXII CCD
 diffractometer
 24823 measured reflections

 6318 independent reflections
 4145 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.043$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.113$
 $S = 1.02$
 6318 reflections
 333 parameters
 2 restraints

 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.29$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.29$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

 Cg4 is the centroid of the $\text{C7}'\text{-C12}'$ phenyl ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}'-\text{H1N}'\cdots\text{O2}$	0.828 (18)	2.310 (17)	3.074 (2)	153.6 (17)
$\text{N1}-\text{H1N}\cdots\text{O1}'$	0.843 (17)	2.129 (17)	2.961 (2)	168.7 (17)
$\text{N1}-\text{H1N}\cdots\text{O3}$	0.843 (17)	2.258 (18)	2.592 (2)	103.8 (14)
$\text{C4}-\text{H4}\cdots\text{O2}^i$	0.93	2.47	3.377 (3)	167
$\text{C4}'-\text{H4}'\cdots\text{Cg4}^{ii}$	0.93	2.85	3.601 (2)	138

 Symmetry codes: (i) $-x + 1, -y + 2, -z + 1$; (ii) $-x, -y + 2, -z$.

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON* (Spek, 2009).

The authors are grateful to the Higher Education Commission of Pakistan for financial support to purchase the diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5501).

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supporting information

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***N*-(2-Methoxyphenyl)benzenesulfonamide**

Aziz-ur-Rehman, Muhammad Arif Sajjad, Mehmet Akkurt, Shahzad Sharif, Muhammad Athar Abbasi and Islam Ullah Khan

S1. Comment

Sulfonamides are well known for their enormous potential as biologically active molecules (Rough *et al.*, 1998) in areas such as anti-microbial (Kayser *et al.*, 2004), anti-convulsant (Arshad *et al.*, 2008), anti-cancer agents and for the treatment of inflammatory rheumatic and non-rheumatic processes including onsets and traumatologic lesions (Gennarti *et al.*, 1994). In the present paper, the structure of *N*-(2-methoxyphenyl)benzene sulfonamide has been determined as part of a research program involving the synthesis and biological evaluation of sulfur containing compounds.

In the crystal structure of the title compound (I), (Fig. 1), there exist two independent molecules, A (with S1) and B (with S1'). Both independent molecules are bent at their *S* atoms with the C—S—N(H)—C torsion angles of 67.25 (15)° in molecule A and -81.17 (16)° in molecule B. The dihedral angles between the phenyl and benzene rings is 88.16 (12)° in molecule A and 44.50 (12)° in molecule B.

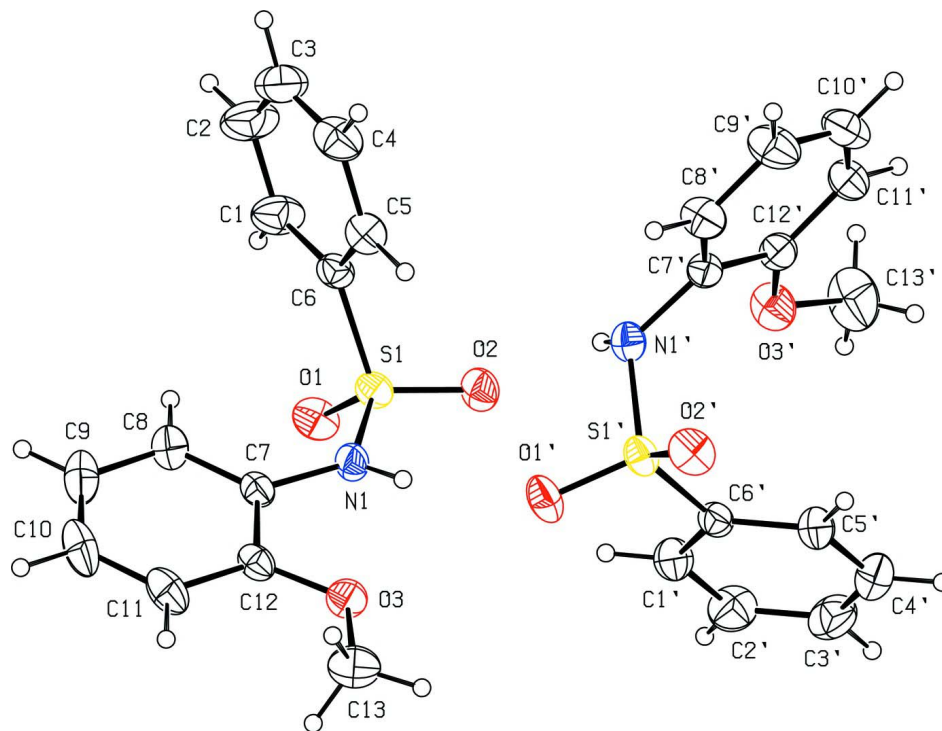
Molecular packing of (I) is stabilized by N—H···O, C—H···O interactions and C—H··· π interactions, forming a three dimensional network (Table 1). Fig. 2 shows N—H···O hydrogen bonds between the molecules A and B in the asymmetric unit.

S2. Experimental

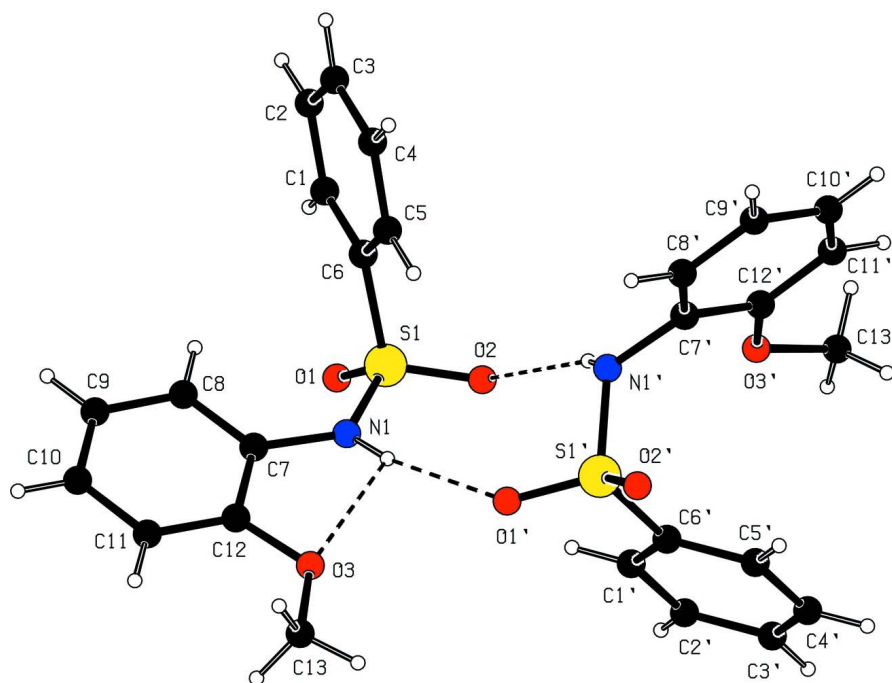
A mixture benzenesulfonyl chloride (10.0 mmol; 1.45 ml), *ortho*-methoxy aniline (*o*-anisidine) (10.0 mmol; 1.12 ml), aqueous sodium carbonate (10%; 15.0 ml) and water (25 ml) was stirred for one hour at room temperature. The crude mixture was washed with water and dried. The product was dissolved in methanol and recrystallized by slow evaporation of the solvent, to generate colourless blocks of (I) in 74% yield.

S3. Refinement

The H atoms of the NH groups were located in a difference Fourier map and refined with the N—H distance restrained to 0.86 (2) %A. The other H atoms were positioned geometrically using a riding model with C—H = 0.93 and 0.96 Å. All H atoms were refined with isotropic displacement parameters with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{aromatic, NH})$ and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{methyl})$.

**Figure 1**

View of the two independent molecules in the asymmetric unit of (10) with displacement ellipsoids drawn at the 30% probability level.

**Figure 2**

View of N—H···O hydrogen bonds shown as dashed lines between the two independent molecules in the asymmetric unit.

***N*-(2-Methoxyphenyl)benzenesulfonamide**

Crystal data

$C_{13}H_{13}NO_3S$

$M_r = 263.31$

Monoclinic, $P2_1/n$

Hall symbol: $-P\ 2_1/n$

$a = 8.7705\ (2)\ \text{\AA}$

$b = 28.1684\ (7)\ \text{\AA}$

$c = 10.7256\ (3)\ \text{\AA}$

$\beta = 105.968\ (1)^\circ$

$V = 2547.53\ (11)\ \text{\AA}^3$

$Z = 8$

$F(000) = 1104$

$D_x = 1.373\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 5117 reflections

$\theta = 2.5\text{--}23.9^\circ$

$\mu = 0.25\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Block, colourless

$0.25 \times 0.17 \times 0.07\ \text{mm}$

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: sealed tube

Graphite monochromator

φ and ω scans

24823 measured reflections

6318 independent reflections

4145 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.043$

$\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 3.3^\circ$

$h = -11 \rightarrow 11$

$k = -37 \rightarrow 37$

$l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.113$
 $S = 1.02$
 6318 reflections
 333 parameters
 2 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
 $w = 1/[\sigma^2(F_o^2) + (0.0488P)^2 + 0.3381P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.29 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.29 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.33115 (5)	0.81887 (2)	0.56961 (5)	0.0412 (2)
O1	0.29814 (16)	0.77142 (5)	0.59898 (15)	0.0561 (5)
O2	0.21225 (14)	0.84480 (5)	0.47617 (14)	0.0518 (5)
O3	0.68975 (15)	0.82034 (5)	0.38159 (14)	0.0507 (5)
N1	0.48440 (18)	0.81979 (6)	0.51516 (16)	0.0430 (5)
C1	0.3503 (3)	0.83266 (9)	0.8243 (2)	0.0681 (9)
C2	0.3878 (4)	0.85873 (11)	0.9379 (3)	0.0865 (11)
C3	0.4523 (3)	0.90282 (11)	0.9397 (3)	0.0764 (10)
C4	0.4848 (3)	0.92123 (8)	0.8326 (3)	0.0663 (9)
C5	0.4496 (2)	0.89536 (7)	0.7189 (2)	0.0511 (7)
C6	0.3823 (2)	0.85108 (7)	0.71581 (19)	0.0424 (6)
C7	0.6322 (2)	0.79744 (6)	0.57144 (18)	0.0388 (6)
C8	0.6717 (3)	0.77602 (7)	0.6914 (2)	0.0536 (7)
C9	0.8209 (3)	0.75621 (9)	0.7392 (3)	0.0681 (9)
C10	0.9276 (3)	0.75748 (9)	0.6684 (3)	0.0728 (9)
C11	0.8897 (2)	0.77854 (8)	0.5478 (3)	0.0596 (8)
C12	0.7411 (2)	0.79839 (6)	0.4983 (2)	0.0417 (6)
C13	0.7966 (3)	0.82674 (9)	0.3050 (2)	0.0653 (9)
S1'	0.31232 (5)	0.92918 (2)	0.25688 (5)	0.0434 (2)
O1'	0.41671 (16)	0.89021 (5)	0.30204 (16)	0.0599 (5)
O2'	0.37153 (17)	0.97058 (5)	0.21000 (15)	0.0587 (5)
O3'	-0.07087 (16)	0.95549 (5)	0.29180 (16)	0.0619 (5)
N1'	0.24881 (18)	0.94574 (6)	0.37937 (16)	0.0432 (6)
C1'	0.1023 (3)	0.86096 (7)	0.1393 (2)	0.0540 (7)

C2'	-0.0250 (3)	0.84437 (9)	0.0434 (3)	0.0675 (9)
C3'	-0.1041 (3)	0.87391 (10)	-0.0551 (2)	0.0720 (10)
C4'	-0.0567 (3)	0.92019 (9)	-0.0586 (2)	0.0687 (9)
C5'	0.0710 (3)	0.93721 (8)	0.0355 (2)	0.0542 (7)
C6'	0.1504 (2)	0.90740 (7)	0.13445 (18)	0.0412 (6)
C7'	0.1746 (2)	0.99118 (7)	0.37841 (18)	0.0415 (6)
C8'	0.2668 (3)	1.02998 (7)	0.4245 (2)	0.0570 (8)
C9'	0.1992 (3)	1.07399 (8)	0.4279 (3)	0.0711 (10)
C10'	0.0390 (3)	1.07863 (9)	0.3853 (3)	0.0694 (10)
C11'	-0.0561 (3)	1.04027 (8)	0.3392 (2)	0.0593 (8)
C12'	0.0103 (2)	0.99586 (7)	0.33485 (19)	0.0453 (7)
C13'	-0.2368 (3)	0.95970 (11)	0.2309 (3)	0.0931 (13)
H1	0.30370	0.80290	0.82150	0.0820*
H1N	0.476 (2)	0.8381 (6)	0.4514 (17)	0.0520*
H2	0.36910	0.84620	1.01260	0.1040*
H3	0.47460	0.92060	1.01560	0.0910*
H4	0.53060	0.95110	0.83600	0.0800*
H5	0.47120	0.90770	0.64510	0.0610*
H8	0.59870	0.77490	0.74000	0.0640*
H9	0.84830	0.74190	0.82050	0.0820*
H10	1.02730	0.74400	0.70170	0.0870*
H11	0.96360	0.77940	0.50010	0.0710*
H13A	0.88270	0.84670	0.35050	0.0980*
H13B	0.74210	0.84140	0.22420	0.0980*
H13C	0.83730	0.79650	0.28840	0.0980*
H1'	0.15550	0.84110	0.20650	0.0650*
H1N'	0.210 (2)	0.9225 (6)	0.4068 (19)	0.0520*
H2'	-0.05770	0.81300	0.04530	0.0810*
H3'	-0.19020	0.86250	-0.11970	0.0860*
H4'	-0.11130	0.94010	-0.12520	0.0820*
H5'	0.10370	0.96850	0.03280	0.0650*
H8'	0.37640	1.02660	0.45380	0.0680*
H9'	0.26260	1.10020	0.45910	0.0850*
H10'	-0.00680	1.10830	0.38740	0.0830*
H11'	-0.16560	1.04410	0.31080	0.0710*
H13D	-0.28950	0.97050	0.29310	0.1400*
H13E	-0.27860	0.92930	0.19770	0.1400*
H13F	-0.25390	0.98210	0.16090	0.1400*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0350 (2)	0.0431 (3)	0.0485 (3)	0.0004 (2)	0.0165 (2)	-0.0006 (2)
O1	0.0577 (9)	0.0437 (8)	0.0721 (11)	-0.0080 (7)	0.0265 (8)	-0.0027 (7)
O2	0.0329 (7)	0.0658 (9)	0.0574 (9)	0.0070 (6)	0.0137 (6)	0.0053 (7)
O3	0.0403 (7)	0.0654 (9)	0.0508 (8)	0.0048 (7)	0.0200 (6)	0.0008 (7)
N1	0.0351 (8)	0.0546 (10)	0.0419 (9)	0.0105 (7)	0.0148 (7)	0.0079 (7)
C1	0.0940 (18)	0.0595 (15)	0.0663 (16)	-0.0063 (13)	0.0479 (14)	-0.0035 (12)

C2	0.114 (2)	0.098 (2)	0.0639 (18)	0.0013 (19)	0.0522 (17)	-0.0102 (16)
C3	0.0722 (17)	0.093 (2)	0.0642 (17)	0.0062 (15)	0.0193 (14)	-0.0275 (15)
C4	0.0516 (13)	0.0576 (15)	0.0858 (19)	-0.0042 (11)	0.0126 (13)	-0.0189 (13)
C5	0.0466 (11)	0.0494 (12)	0.0588 (14)	-0.0019 (10)	0.0169 (10)	0.0001 (10)
C6	0.0410 (10)	0.0436 (11)	0.0469 (11)	0.0050 (8)	0.0195 (9)	0.0004 (9)
C7	0.0333 (9)	0.0354 (10)	0.0463 (11)	0.0035 (8)	0.0084 (8)	-0.0031 (8)
C8	0.0521 (12)	0.0505 (13)	0.0560 (13)	0.0049 (10)	0.0114 (10)	0.0089 (10)
C9	0.0635 (15)	0.0601 (15)	0.0686 (16)	0.0082 (12)	-0.0022 (13)	0.0190 (12)
C10	0.0448 (12)	0.0618 (16)	0.101 (2)	0.0167 (11)	0.0017 (13)	0.0162 (14)
C11	0.0365 (11)	0.0537 (13)	0.0886 (18)	0.0076 (10)	0.0174 (11)	-0.0007 (12)
C12	0.0328 (9)	0.0362 (10)	0.0543 (12)	0.0000 (8)	0.0091 (8)	-0.0062 (9)
C13	0.0596 (13)	0.0753 (16)	0.0730 (16)	-0.0073 (12)	0.0383 (13)	-0.0062 (13)
S1'	0.0351 (2)	0.0419 (3)	0.0543 (3)	0.0050 (2)	0.0143 (2)	0.0066 (2)
O1'	0.0452 (8)	0.0579 (9)	0.0768 (11)	0.0199 (7)	0.0172 (7)	0.0121 (8)
O2'	0.0542 (8)	0.0521 (9)	0.0770 (11)	-0.0079 (7)	0.0304 (8)	0.0062 (8)
O3'	0.0368 (7)	0.0664 (10)	0.0780 (11)	-0.0051 (7)	0.0085 (7)	-0.0066 (8)
N1'	0.0405 (9)	0.0425 (10)	0.0463 (10)	0.0013 (7)	0.0113 (7)	0.0067 (7)
C1'	0.0641 (13)	0.0433 (12)	0.0555 (13)	0.0007 (10)	0.0178 (11)	0.0076 (10)
C2'	0.0769 (16)	0.0555 (15)	0.0686 (16)	-0.0142 (12)	0.0174 (14)	-0.0068 (12)
C3'	0.0709 (16)	0.0831 (19)	0.0549 (15)	-0.0082 (14)	0.0053 (12)	-0.0136 (13)
C4'	0.0729 (16)	0.0741 (17)	0.0503 (14)	0.0100 (13)	0.0023 (12)	0.0064 (12)
C5'	0.0635 (13)	0.0466 (12)	0.0515 (13)	0.0081 (10)	0.0141 (11)	0.0089 (10)
C6'	0.0431 (10)	0.0406 (11)	0.0429 (11)	0.0061 (8)	0.0168 (9)	0.0027 (8)
C7'	0.0403 (10)	0.0456 (11)	0.0391 (10)	0.0035 (8)	0.0120 (8)	0.0001 (8)
C8'	0.0460 (11)	0.0542 (13)	0.0696 (15)	-0.0039 (10)	0.0141 (11)	-0.0114 (11)
C9'	0.0695 (16)	0.0507 (14)	0.094 (2)	-0.0067 (12)	0.0242 (14)	-0.0202 (13)
C10'	0.0766 (17)	0.0563 (15)	0.0807 (18)	0.0155 (13)	0.0306 (14)	-0.0106 (13)
C11'	0.0474 (12)	0.0714 (16)	0.0611 (15)	0.0170 (11)	0.0183 (11)	-0.0006 (12)
C12'	0.0389 (10)	0.0543 (13)	0.0433 (11)	0.0013 (9)	0.0122 (9)	-0.0005 (9)
C13'	0.0414 (13)	0.098 (2)	0.123 (3)	-0.0113 (13)	-0.0060 (14)	-0.0054 (18)

Geometric parameters (Å, °)

S1—O1	1.4212 (15)	C5—H5	0.9300
S1—O2	1.4319 (15)	C8—H8	0.9300
S1—N1	1.6063 (17)	C9—H9	0.9300
S1—C6	1.760 (2)	C10—H10	0.9300
S1'—C6'	1.7582 (19)	C11—H11	0.9300
S1'—N1'	1.6299 (17)	C13—H13A	0.9600
S1'—O1'	1.4260 (15)	C13—H13C	0.9600
S1'—O2'	1.4235 (15)	C13—H13B	0.9600
O3—C12	1.357 (2)	C1'—C2'	1.375 (4)
O3—C13	1.417 (3)	C1'—C6'	1.380 (3)
O3'—C12'	1.354 (2)	C2'—C3'	1.374 (4)
O3'—C13'	1.427 (3)	C3'—C4'	1.372 (4)
N1—C7	1.418 (2)	C4'—C5'	1.372 (3)
N1—H1N	0.843 (17)	C5'—C6'	1.382 (3)
N1'—C7'	1.435 (3)	C7'—C12'	1.394 (3)

N1'—H1N'	0.828 (18)	C7'—C8'	1.369 (3)
C1—C2	1.383 (4)	C8'—C9'	1.379 (3)
C1—C6	1.372 (3)	C9'—C10'	1.359 (4)
C2—C3	1.363 (4)	C10'—C11'	1.371 (4)
C3—C4	1.360 (4)	C11'—C12'	1.386 (3)
C4—C5	1.381 (4)	C1'—H1'	0.9300
C5—C6	1.376 (3)	C2'—H2'	0.9300
C7—C12	1.393 (3)	C3'—H3'	0.9300
C7—C8	1.376 (3)	C4'—H4'	0.9300
C8—C9	1.385 (4)	C5'—H5'	0.9300
C9—C10	1.358 (4)	C8'—H8'	0.9300
C10—C11	1.378 (4)	C9'—H9'	0.9300
C11—C12	1.383 (3)	C10'—H10'	0.9300
C1—H1	0.9300	C11'—H11'	0.9300
C2—H2	0.9300	C13'—H13D	0.9600
C3—H3	0.9300	C13'—H13E	0.9600
C4—H4	0.9300	C13'—H13F	0.9600
O1—S1—O2	118.77 (9)	C9—C10—H10	120.00
O1—S1—N1	109.73 (9)	C11—C10—H10	120.00
O1—S1—C6	107.75 (9)	C10—C11—H11	120.00
O2—S1—N1	104.98 (9)	C12—C11—H11	120.00
O2—S1—C6	108.59 (9)	O3—C13—H13A	109.00
N1—S1—C6	106.38 (9)	H13A—C13—H13C	109.00
O1'—S1'—O2'	119.21 (9)	H13B—C13—H13C	109.00
O1'—S1'—N1'	106.05 (9)	H13A—C13—H13B	109.00
O1'—S1'—C6'	107.24 (9)	O3—C13—H13B	109.00
O2'—S1'—N1'	106.80 (9)	O3—C13—H13C	110.00
O2'—S1'—C6'	108.68 (9)	C2'—C1'—C6'	119.2 (2)
N1'—S1'—C6'	108.48 (9)	C1'—C2'—C3'	120.3 (2)
C12—O3—C13	119.21 (16)	C2'—C3'—C4'	120.3 (2)
C12'—O3'—C13'	117.44 (18)	C3'—C4'—C5'	120.3 (2)
S1—N1—C7	126.62 (14)	C4'—C5'—C6'	119.3 (2)
C7—N1—H1N	118.8 (12)	S1'—C6'—C5'	119.63 (16)
S1—N1—H1N	114.2 (12)	C1'—C6'—C5'	120.68 (19)
S1'—N1'—C7'	120.30 (13)	S1'—C6'—C1'	119.68 (15)
C7'—N1'—H1N'	118.5 (13)	C8'—C7'—C12'	119.96 (19)
S1'—N1'—H1N'	109.0 (13)	N1'—C7'—C8'	119.22 (18)
C2—C1—C6	119.5 (2)	N1'—C7'—C12'	120.80 (17)
C1—C2—C3	119.6 (3)	C7'—C8'—C9'	120.8 (2)
C2—C3—C4	121.2 (3)	C8'—C9'—C10'	119.3 (2)
C3—C4—C5	119.8 (2)	C9'—C10'—C11'	121.0 (2)
C4—C5—C6	119.3 (2)	C10'—C11'—C12'	120.3 (2)
C1—C6—C5	120.58 (19)	C7'—C12'—C11'	118.65 (19)
S1—C6—C1	119.96 (16)	O3'—C12'—C7'	115.67 (17)
S1—C6—C5	119.42 (15)	O3'—C12'—C11'	125.68 (18)
C8—C7—C12	119.97 (19)	C2'—C1'—H1'	120.00
N1—C7—C8	123.93 (19)	C6'—C1'—H1'	120.00

N1—C7—C12	116.10 (16)	C1'—C2'—H2'	120.00
C7—C8—C9	119.5 (2)	C3'—C2'—H2'	120.00
C8—C9—C10	120.6 (3)	C2'—C3'—H3'	120.00
C9—C10—C11	120.7 (3)	C4'—C3'—H3'	120.00
C10—C11—C12	119.5 (2)	C3'—C4'—H4'	120.00
C7—C12—C11	119.7 (2)	C5'—C4'—H4'	120.00
O3—C12—C7	115.04 (16)	C4'—C5'—H5'	120.00
O3—C12—C11	125.23 (19)	C6'—C5'—H5'	120.00
C6—C1—H1	120.00	C7'—C8'—H8'	120.00
C2—C1—H1	120.00	C9'—C8'—H8'	120.00
C3—C2—H2	120.00	C8'—C9'—H9'	120.00
C1—C2—H2	120.00	C10'—C9'—H9'	120.00
C4—C3—H3	119.00	C9'—C10'—H10'	119.00
C2—C3—H3	119.00	C11'—C10'—H10'	120.00
C5—C4—H4	120.00	C10'—C11'—H11'	120.00
C3—C4—H4	120.00	C12'—C11'—H11'	120.00
C4—C5—H5	120.00	O3'—C13'—H13D	109.00
C6—C5—H5	120.00	O3'—C13'—H13E	109.00
C7—C8—H8	120.00	O3'—C13'—H13F	109.00
C9—C8—H8	120.00	H13D—C13'—H13E	109.00
C10—C9—H9	120.00	H13D—C13'—H13F	109.00
C8—C9—H9	120.00	H13E—C13'—H13F	110.00
O1—S1—N1—C7	-49.05 (18)	C4—C5—C6—S1	177.86 (17)
O2—S1—N1—C7	-177.75 (15)	C4—C5—C6—C1	0.0 (3)
C6—S1—N1—C7	67.25 (18)	C12—C7—C8—C9	1.0 (3)
O1—S1—C6—C1	-14.8 (2)	N1—C7—C12—O3	-0.7 (2)
O1—S1—C6—C5	167.32 (15)	N1—C7—C8—C9	-178.4 (2)
O2—S1—C6—C1	115.07 (18)	C8—C7—C12—O3	179.93 (17)
O2—S1—C6—C5	-62.83 (17)	C8—C7—C12—C11	-1.2 (3)
N1—S1—C6—C1	-132.41 (18)	N1—C7—C12—C11	178.21 (18)
N1—S1—C6—C5	49.70 (18)	C7—C8—C9—C10	-0.5 (4)
N1'—S1'—C6'—C5'	95.33 (18)	C8—C9—C10—C11	0.1 (4)
O1'—S1'—N1'—C7'	163.92 (14)	C9—C10—C11—C12	-0.3 (4)
O2'—S1'—N1'—C7'	35.80 (17)	C10—C11—C12—C7	0.8 (3)
C6'—S1'—N1'—C7'	-81.17 (16)	C10—C11—C12—O3	179.6 (2)
O1'—S1'—C6'—C1'	30.5 (2)	C6'—C1'—C2'—C3'	-0.7 (4)
O1'—S1'—C6'—C5'	-150.54 (17)	C2'—C1'—C6'—S1'	179.66 (19)
O2'—S1'—C6'—C1'	160.64 (17)	C2'—C1'—C6'—C5'	0.7 (3)
O2'—S1'—C6'—C5'	-20.4 (2)	C1'—C2'—C3'—C4'	0.0 (4)
N1'—S1'—C6'—C1'	-83.61 (19)	C2'—C3'—C4'—C5'	0.7 (4)
C13—O3—C12—C7	174.60 (17)	C3'—C4'—C5'—C6'	-0.6 (4)
C13—O3—C12—C11	-4.2 (3)	C4'—C5'—C6'—S1'	-179.01 (18)
C13'—O3'—C12'—C7'	-172.1 (2)	C4'—C5'—C6'—C1'	-0.1 (3)
C13'—O3'—C12'—C11'	8.2 (3)	N1'—C7'—C8'—C9'	-178.5 (2)
S1—N1—C7—C8	-8.1 (3)	C12'—C7'—C8'—C9'	-0.3 (3)
S1—N1—C7—C12	172.47 (14)	N1'—C7'—C12'—O3'	-1.5 (3)
S1'—N1'—C7'—C8'	-88.1 (2)	N1'—C7'—C12'—C11'	178.22 (18)

S1'—N1'—C7'—C12'	93.8 (2)	C8'—C7'—C12'—O3'	-179.62 (18)
C2—C1—C6—C5	-0.7 (4)	C8'—C7'—C12'—C11'	0.1 (3)
C6—C1—C2—C3	1.6 (4)	C7'—C8'—C9'—C10'	0.3 (4)
C2—C1—C6—S1	-178.6 (2)	C8'—C9'—C10'—C11'	0.1 (4)
C1—C2—C3—C4	-1.9 (5)	C9'—C10'—C11'—C12'	-0.3 (4)
C2—C3—C4—C5	1.2 (4)	C10'—C11'—C12'—O3'	179.9 (2)
C3—C4—C5—C6	-0.2 (4)	C10'—C11'—C12'—C7'	0.2 (3)

Hydrogen-bond geometry (Å, °)

Cg4 is the centroid of the C7'—C12' phenyl ring.

<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'—H1N'...O2	0.828 (18)	2.310 (17)	3.074 (2)	153.6 (17)
N1—H1N...O1'	0.843 (17)	2.129 (17)	2.961 (2)	168.7 (17)
N1—H1N...O3	0.843 (17)	2.258 (18)	2.592 (2)	103.8 (14)
C4—H4...O2 ⁱ	0.93	2.47	3.377 (3)	167
C4'—H4'...Cg4 ⁱⁱ	0.93	2.85	3.601 (2)	138

Symmetry codes: (i) $-x+1, -y+2, -z+1$; (ii) $-x, -y+2, -z$.