

N,N'-Diethyl-N,N'-[1,3-phenylene-bis(methylene)]dibenzenesulfonamide

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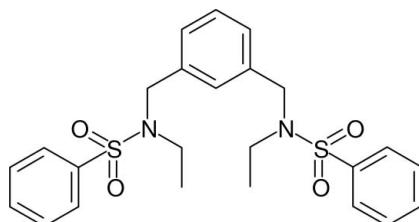
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$; disorder in main residue; R factor = 0.045; wR factor = 0.132; data-to-parameter ratio = 20.7.

In the title compound, $\text{C}_{24}\text{H}_{28}\text{N}_2\text{O}_4\text{S}_2$, the dihedral angles between the central benzene ring and the pendant rings are 77.44 (11) and 79.23 (10) $^\circ$, and the dihedral angle between the pendant rings is 23.31 (12) $^\circ$. Both sulfonamide groups project to the same side of the central benzene ring and the molecule has approximate non-crystallographic mirror symmetry. One of the ethyl side chains is disordered over two sets of sites in a 0.526 (14):0.474 (14) ratio. In the crystal, inversion dimers linked by pairs of weak C–H \cdots O interactions occur, generating $R_2^2(28)$ loops.

Related literature

For a related structure, see: Khan *et al.* (2011).



Experimental

Crystal data

$\text{C}_{24}\text{H}_{28}\text{N}_2\text{O}_4\text{S}_2$	$V = 2421.02\text{ (15) \AA}^3$
$M_r = 472.60$	$Z = 4$
Monoclinic, $P2_1/n$	Mo $K\alpha$ radiation
$a = 9.1865\text{ (3) \AA}$	$\mu = 0.25\text{ mm}^{-1}$
$b = 19.0679\text{ (7) \AA}$	$T = 296\text{ K}$
$c = 14.3870\text{ (5) \AA}$	$0.13 \times 0.10 \times 0.09\text{ mm}$
$\beta = 106.122\text{ (1)}^\circ$	

Data collection

Bruker APEXII CCD diffractometer	6013 independent reflections
23272 measured reflections	4188 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$	290 parameters
$wR(F^2) = 0.132$	H-atom parameters constrained
$S = 1.01$	$\Delta\rho_{\text{max}} = 0.29\text{ e \AA}^{-3}$
6013 reflections	$\Delta\rho_{\text{min}} = -0.38\text{ e \AA}^{-3}$

Table 1

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C17–H17 \cdots O1 ⁱ	0.93	2.57	3.409 (3)	151

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT* (Bruker, 2007); 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: *SHELXL97* (Sheldrick, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2324).

References

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- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
- Khan, I. U., Sheikh, T. A., Ejaz & Harrison, W. T. A. (2011). *Acta Cryst. E67*, o2371.
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supporting information

Acta Cryst. (2011). E67, o3038 [doi:10.1107/S1600536811040700]

N,N'-Diethyl-N,N'-[1,3-phenylenebis(methylene)]dibenzenesulfonamide

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S1. Comment

As part of our ongoing structural studies of sulfonamides (Khan *et al.*, 2011), the synthesis and structure of the title compound, (I), (Fig. 1), are now described.

The dihedral angles between the central benzene ring and the pendant rings are almost equal at 77.44 (11) and 79.23 (10) $^{\circ}$ and the dihedral angle between the pendant rings is 23.31 (12) $^{\circ}$. Both sulfonamide groups project to the same side of the central ring and the molecule has approximate non-crystallographic mirror symmetry. The C8—S1—N1—C7 and C15—S2—N2—C14 torsion angles are -66.87 (15) and 70.98 (15) $^{\circ}$, respectively. The S1—N1—C7—C6 and S2—N2—C14—C2 torsion angles are -148.14 (14) and 144.01 (14) $^{\circ}$, respectively. One of the terminal methyl groups is disordered over two sets of sites in a 0.526 (14):0.474 (14) ratio.

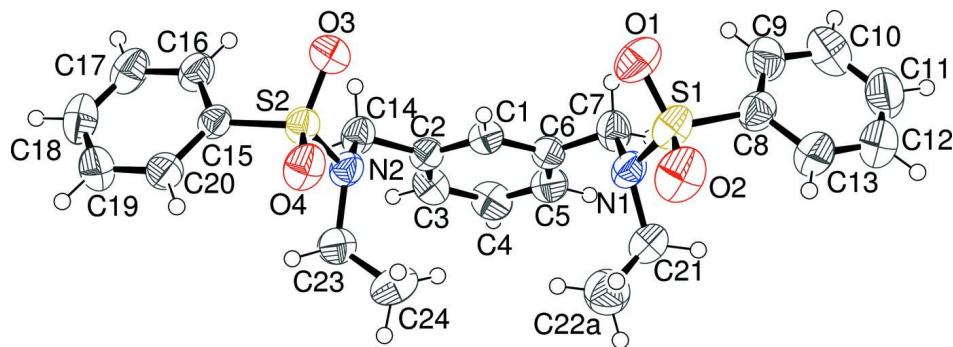
In the crystal, inversion dimers linked by pairs of weak C—H \cdots O interactions occur (Fig. 2, Table 1).

S2. Experimental

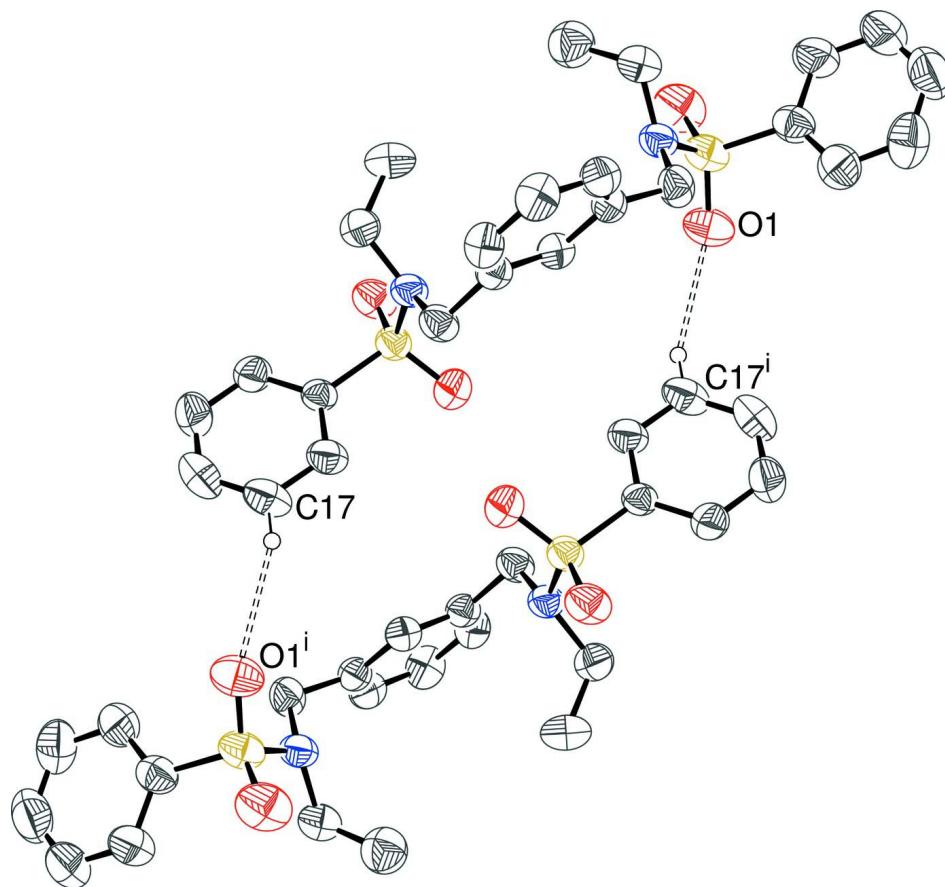
A mixture of *N,N'*-diethyl-(benzene-1,3-diylmethanediyl)dibenzenesulfonamide (0.2 g, 0.43 mmol), sodium hydride (0.21 g; 0.88 mmol) and *N,N*-dimethylformamide (10.0 ml) was stirred in a 100-ml RB flask at room temperature for half an hour followed by the addition of ethyl iodide (0.134 g; 0.86 mmol). The reaction mixture was further stirred for five hours, and its completion was monitored by TLC. After completion, the contents were poured over crushed ice. The precipitated product was isolated, washed and crystallized from methanol to yield colourless blocks of (I).

S3. Refinement

The N-bound H atom was located in a difference Fourier map and its position was freely refined with the constraint $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$. The C-bound hydrogen atoms were placed in calculated positions (C—H = 0.93–0.97 Å) and refined as riding atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. One of the ethyl side chains is disordered over two sets of sites in a 0.526 (14):0.474 (14) ratio for atoms C22A:C22B, which were refined isotropically.

**Figure 1**

The molecular structure of the title molecule, showing the numbering scheme and 50% displacement ellipsoids. Only one orientation of the disordered atom C22 and its attached H atoms is shown.

**Figure 2**

A view of the inversion dimer in the crystal structure of the title compound. All H atoms except H17 have been omitted for clarity, and the C—H···O interactions are shown as dashed lines [Symmetry code: (i) 1-x, -y, 1-z].

N,N'-Diethyl-N,N'-[1,3- phenylenebis(methylene)]dibenzenesulfonamide*Crystal data*

C₂₄H₂₈N₂O₄S₂
 $M_r = 472.60$
 Monoclinic, P2₁/n
 Hall symbol: -P 2yn
 $a = 9.1865 (3)$ Å
 $b = 19.0679 (7)$ Å
 $c = 14.3870 (5)$ Å
 $\beta = 106.122 (1)^\circ$
 $V = 2421.02 (15)$ Å³
 $Z = 4$

$F(000) = 1000$
 $D_x = 1.297 \text{ Mg m}^{-3}$
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 6013 reflections
 $\theta = 2.6\text{--}28.3^\circ$
 $\mu = 0.25 \text{ mm}^{-1}$
 $T = 296 \text{ K}$
 Block, colourless
 $0.13 \times 0.10 \times 0.09 \text{ mm}$

Data collection

Bruker APEXII CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 ω scans
 23272 measured reflections
 6013 independent reflections

4188 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$
 $\theta_{\text{max}} = 28.3^\circ, \theta_{\text{min}} = 2.6^\circ$
 $h = -12 \rightarrow 8$
 $k = -24 \rightarrow 25$
 $l = -17 \rightarrow 19$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.045$
 $wR(F^2) = 0.132$
 $S = 1.01$
 6013 reflections
 290 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
 $w = 1/[\sigma^2(F_o^2) + (0.060P)^2 + 0.7693P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.29 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.38 \text{ e } \text{\AA}^{-3}$

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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.78689 (19)	0.13374 (9)	0.68904 (13)	0.0448 (4)	
H1	0.7268	0.1480	0.6288	0.054*	
C2	0.73697 (19)	0.08014 (9)	0.73745 (14)	0.0449 (4)	
C3	0.8266 (2)	0.05935 (11)	0.82682 (15)	0.0554 (5)	
H3	0.7942	0.0233	0.8598	0.066*	

C4	0.9639 (2)	0.09163 (13)	0.86768 (15)	0.0614 (5)
H4	1.0237	0.0775	0.9281	0.074*
C5	1.0123 (2)	0.14489 (11)	0.81882 (15)	0.0545 (5)
H5	1.1049	0.1666	0.8466	0.065*
C6	0.92480 (19)	0.16639 (9)	0.72889 (14)	0.0452 (4)
C7	0.9798 (2)	0.22228 (10)	0.67292 (17)	0.0557 (5)
H7A	1.0849	0.2329	0.7054	0.067*
H7B	0.9751	0.2045	0.6090	0.067*
C8	1.0539 (3)	0.36406 (11)	0.56934 (15)	0.0582 (5)
C9	1.1599 (3)	0.32027 (13)	0.54866 (18)	0.0702 (6)
H9	1.1345	0.2745	0.5280	0.084*
C10	1.3050 (3)	0.34546 (17)	0.5591 (2)	0.0881 (8)
H10	1.3785	0.3162	0.5467	0.106*
C11	1.3404 (4)	0.41342 (19)	0.5877 (2)	0.0957 (10)
H11	1.4381	0.4299	0.5947	0.115*
C12	1.2346 (4)	0.45677 (18)	0.6058 (2)	0.1002 (10)
H12	1.2595	0.5030	0.6238	0.120*
C13	1.0910 (3)	0.43270 (13)	0.5975 (2)	0.0796 (7)
H13	1.0188	0.4623	0.6107	0.096*
C14	0.5875 (2)	0.04462 (10)	0.69110 (16)	0.0517 (5)
H14A	0.5908	0.0235	0.6304	0.062*
H14B	0.5708	0.0076	0.7332	0.062*
C15	0.22235 (19)	0.00993 (9)	0.59280 (13)	0.0437 (4)
C16	0.2729 (2)	-0.05564 (10)	0.57526 (16)	0.0553 (5)
H16	0.3584	-0.0602	0.5531	0.066*
C17	0.1955 (3)	-0.11417 (11)	0.59097 (18)	0.0676 (6)
H17	0.2282	-0.1586	0.5790	0.081*
C18	0.0694 (3)	-0.10715 (13)	0.62447 (19)	0.0728 (7)
H18	0.0180	-0.1469	0.6356	0.087*
C19	0.0198 (3)	-0.04250 (14)	0.64141 (18)	0.0697 (6)
H19	-0.0661	-0.0383	0.6633	0.084*
C20	0.0957 (2)	0.01662 (11)	0.62643 (16)	0.0558 (5)
H20	0.0623	0.0608	0.6388	0.067*
C21	0.9119 (3)	0.32830 (13)	0.75365 (18)	0.0700 (6)
H21A	0.9693	0.3704	0.7501	0.084*
H21B	0.9692	0.3008	0.8083	0.084*
C22A	0.7607 (7)	0.3480 (4)	0.7682 (5)	0.079 (2)*
H22A	0.7765	0.3746	0.8269	0.119*
H22B	0.7046	0.3063	0.7725	0.119*
H22C	0.7048	0.3759	0.7145	0.119*
C22B	0.7963 (9)	0.3175 (6)	0.8056 (7)	0.095 (3)*
H22D	0.8201	0.3458	0.8630	0.143*
H22E	0.7947	0.2690	0.8231	0.143*
H22F	0.6987	0.3306	0.7645	0.143*
C23	0.4252 (2)	0.12225 (12)	0.76072 (16)	0.0597 (5)
H23A	0.3194	0.1129	0.7553	0.072*
H23B	0.4857	0.0974	0.8171	0.072*
C24	0.4538 (4)	0.19853 (14)	0.7750 (2)	0.0879 (8)

H24A	0.5597	0.2078	0.7849	0.132*
H24B	0.4240	0.2138	0.8307	0.132*
H24C	0.3961	0.2234	0.7189	0.132*
S1	0.87268 (6)	0.33183 (3)	0.56484 (4)	0.05927 (16)
S2	0.32295 (5)	0.08483 (2)	0.57589 (4)	0.04724 (14)
N1	0.88989 (17)	0.28716 (8)	0.66325 (12)	0.0512 (4)
N2	0.46183 (16)	0.09500 (8)	0.67317 (12)	0.0484 (4)
O1	0.82899 (19)	0.28392 (9)	0.48611 (12)	0.0765 (5)
O2	0.77734 (19)	0.39036 (9)	0.56817 (15)	0.0850 (5)
O3	0.38878 (17)	0.07049 (8)	0.49896 (11)	0.0615 (4)
O4	0.22630 (16)	0.14401 (7)	0.57007 (12)	0.0634 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0399 (8)	0.0470 (10)	0.0441 (10)	0.0051 (7)	0.0063 (7)	-0.0007 (8)
C2	0.0399 (8)	0.0434 (10)	0.0506 (10)	0.0043 (7)	0.0113 (8)	-0.0030 (8)
C3	0.0532 (11)	0.0572 (12)	0.0570 (12)	0.0080 (9)	0.0173 (9)	0.0120 (9)
C4	0.0490 (10)	0.0816 (15)	0.0479 (11)	0.0100 (10)	0.0041 (9)	0.0077 (10)
C5	0.0358 (8)	0.0674 (13)	0.0560 (12)	0.0034 (8)	0.0057 (8)	-0.0092 (10)
C6	0.0385 (8)	0.0441 (10)	0.0540 (10)	0.0041 (7)	0.0145 (8)	-0.0031 (8)
C7	0.0427 (9)	0.0502 (11)	0.0763 (14)	0.0030 (8)	0.0200 (9)	0.0027 (10)
C8	0.0683 (12)	0.0491 (11)	0.0539 (12)	-0.0051 (9)	0.0117 (10)	0.0015 (9)
C9	0.0817 (16)	0.0574 (13)	0.0753 (15)	-0.0027 (11)	0.0281 (13)	0.0091 (11)
C10	0.0849 (18)	0.096 (2)	0.094 (2)	0.0008 (16)	0.0428 (16)	0.0184 (16)
C11	0.091 (2)	0.110 (3)	0.089 (2)	-0.0403 (19)	0.0303 (17)	0.0000 (18)
C12	0.119 (2)	0.087 (2)	0.106 (2)	-0.0455 (19)	0.049 (2)	-0.0221 (17)
C13	0.0994 (19)	0.0578 (14)	0.0856 (18)	-0.0164 (13)	0.0322 (15)	-0.0105 (13)
C14	0.0459 (9)	0.0407 (10)	0.0667 (12)	0.0010 (8)	0.0123 (9)	-0.0021 (9)
C15	0.0380 (8)	0.0434 (9)	0.0468 (10)	-0.0011 (7)	0.0072 (7)	-0.0072 (8)
C16	0.0461 (10)	0.0473 (11)	0.0698 (13)	0.0003 (8)	0.0116 (9)	-0.0116 (9)
C17	0.0636 (13)	0.0445 (11)	0.0858 (17)	-0.0032 (10)	0.0061 (12)	-0.0072 (11)
C18	0.0654 (14)	0.0633 (14)	0.0829 (17)	-0.0207 (11)	0.0091 (12)	0.0073 (12)
C19	0.0542 (12)	0.0799 (17)	0.0783 (16)	-0.0141 (11)	0.0240 (11)	-0.0028 (13)
C20	0.0475 (10)	0.0573 (12)	0.0639 (13)	-0.0006 (9)	0.0174 (9)	-0.0110 (10)
C21	0.0774 (15)	0.0623 (14)	0.0712 (15)	-0.0154 (11)	0.0223 (12)	-0.0171 (11)
C23	0.0568 (11)	0.0632 (13)	0.0605 (13)	-0.0029 (10)	0.0187 (10)	-0.0101 (10)
C24	0.107 (2)	0.0684 (16)	0.0941 (19)	-0.0111 (15)	0.0384 (17)	-0.0337 (15)
S1	0.0548 (3)	0.0474 (3)	0.0663 (3)	0.0028 (2)	0.0013 (2)	-0.0028 (2)
S2	0.0431 (2)	0.0417 (2)	0.0565 (3)	0.00114 (18)	0.0131 (2)	-0.0028 (2)
N1	0.0474 (8)	0.0412 (8)	0.0643 (10)	-0.0011 (6)	0.0140 (7)	-0.0090 (7)
N2	0.0399 (7)	0.0448 (8)	0.0598 (10)	-0.0005 (6)	0.0127 (7)	-0.0110 (7)
O1	0.0781 (10)	0.0737 (11)	0.0641 (10)	-0.0086 (8)	-0.0031 (8)	-0.0132 (8)
O2	0.0706 (10)	0.0608 (10)	0.1129 (15)	0.0211 (8)	0.0075 (10)	0.0060 (10)
O3	0.0645 (9)	0.0673 (9)	0.0570 (8)	-0.0022 (7)	0.0239 (7)	-0.0007 (7)
O4	0.0549 (8)	0.0441 (8)	0.0862 (11)	0.0089 (6)	0.0111 (7)	-0.0004 (7)

Geometric parameters (\AA , $\text{^{\circ}}$)

C1—C2	1.385 (3)	C16—C17	1.376 (3)
C1—C6	1.385 (2)	C16—H16	0.9300
C1—H1	0.9300	C17—C18	1.379 (3)
C2—C3	1.378 (3)	C17—H17	0.9300
C2—C14	1.510 (3)	C18—C19	1.360 (4)
C3—C4	1.380 (3)	C18—H18	0.9300
C3—H3	0.9300	C19—C20	1.374 (3)
C4—C5	1.377 (3)	C19—H19	0.9300
C4—H4	0.9300	C20—H20	0.9300
C5—C6	1.382 (3)	C21—C22B	1.473 (7)
C5—H5	0.9300	C21—N1	1.484 (3)
C6—C7	1.505 (3)	C21—C22A	1.509 (6)
C7—N1	1.473 (2)	C21—H21A	0.9700
C7—H7A	0.9700	C21—H21B	0.9700
C7—H7B	0.9700	C22A—H22A	0.9600
C8—C9	1.377 (3)	C22A—H22B	0.9600
C8—C13	1.384 (3)	C22A—H22C	0.9600
C8—S1	1.759 (2)	C22B—H22D	0.9600
C9—C10	1.385 (4)	C22B—H22E	0.9600
C9—H9	0.9300	C22B—H22F	0.9600
C10—C11	1.371 (4)	C23—C24	1.482 (3)
C10—H10	0.9300	C23—N2	1.485 (3)
C11—C12	1.355 (4)	C23—H23A	0.9700
C11—H11	0.9300	C23—H23B	0.9700
C12—C13	1.370 (4)	C24—H24A	0.9600
C12—H12	0.9300	C24—H24B	0.9600
C13—H13	0.9300	C24—H24C	0.9600
C14—N2	1.469 (2)	S1—O1	1.4235 (16)
C14—H14A	0.9700	S1—O2	1.4276 (16)
C14—H14B	0.9700	S1—N1	1.6220 (18)
C15—C16	1.381 (3)	S2—O4	1.4239 (14)
C15—C20	1.385 (3)	S2—O3	1.4274 (15)
C15—S2	1.7543 (18)	S2—N2	1.6224 (16)
C2—C1—C6	121.08 (17)	C17—C18—H18	119.8
C2—C1—H1	119.5	C18—C19—C20	120.4 (2)
C6—C1—H1	119.5	C18—C19—H19	119.8
C3—C2—C1	119.08 (17)	C20—C19—H19	119.8
C3—C2—C14	121.14 (18)	C19—C20—C15	119.5 (2)
C1—C2—C14	119.76 (17)	C19—C20—H20	120.3
C2—C3—C4	120.50 (19)	C15—C20—H20	120.3
C2—C3—H3	119.7	C22B—C21—N1	115.4 (3)
C4—C3—H3	119.7	C22B—C21—C22A	31.0 (3)
C5—C4—C3	119.87 (19)	N1—C21—C22A	110.3 (3)
C5—C4—H4	120.1	C22B—C21—H21A	128.4
C3—C4—H4	120.1	N1—C21—H21A	109.6

C4—C5—C6	120.67 (18)	C22A—C21—H21A	109.6
C4—C5—H5	119.7	C22B—C21—H21B	79.5
C6—C5—H5	119.7	N1—C21—H21B	109.6
C5—C6—C1	118.79 (18)	C22A—C21—H21B	109.6
C5—C6—C7	121.08 (17)	H21A—C21—H21B	108.1
C1—C6—C7	120.09 (17)	C21—C22A—H22A	109.5
N1—C7—C6	112.47 (15)	C21—C22A—H22B	109.5
N1—C7—H7A	109.1	H22A—C22A—H22B	109.5
C6—C7—H7A	109.1	C21—C22A—H22C	109.5
N1—C7—H7B	109.1	H22A—C22A—H22C	109.5
C6—C7—H7B	109.1	H22B—C22A—H22C	109.5
H7A—C7—H7B	107.8	C21—C22B—H22D	109.5
C9—C8—C13	120.4 (2)	C21—C22B—H22E	109.5
C9—C8—S1	119.95 (17)	H22D—C22B—H22E	109.5
C13—C8—S1	119.6 (2)	C21—C22B—H22F	109.5
C8—C9—C10	118.9 (2)	H22D—C22B—H22F	109.5
C8—C9—H9	120.5	H22E—C22B—H22F	109.5
C10—C9—H9	120.5	C24—C23—N2	112.85 (19)
C11—C10—C9	120.1 (3)	C24—C23—H23A	109.0
C11—C10—H10	120.0	N2—C23—H23A	109.0
C9—C10—H10	120.0	C24—C23—H23B	109.0
C12—C11—C10	120.7 (3)	N2—C23—H23B	109.0
C12—C11—H11	119.6	H23A—C23—H23B	107.8
C10—C11—H11	119.6	C23—C24—H24A	109.5
C11—C12—C13	120.2 (3)	C23—C24—H24B	109.5
C11—C12—H12	119.9	H24A—C24—H24B	109.5
C13—C12—H12	119.9	C23—C24—H24C	109.5
C12—C13—C8	119.7 (3)	H24A—C24—H24C	109.5
C12—C13—H13	120.2	H24B—C24—H24C	109.5
C8—C13—H13	120.2	O1—S1—O2	119.23 (11)
N2—C14—C2	111.00 (15)	O1—S1—N1	107.17 (10)
N2—C14—H14A	109.4	O2—S1—N1	107.05 (10)
C2—C14—H14A	109.4	O1—S1—C8	107.96 (11)
N2—C14—H14B	109.4	O2—S1—C8	107.99 (11)
C2—C14—H14B	109.4	N1—S1—C8	106.83 (9)
H14A—C14—H14B	108.0	O4—S2—O3	119.85 (10)
C16—C15—C20	120.34 (18)	O4—S2—N2	106.51 (9)
C16—C15—S2	119.65 (14)	O3—S2—N2	106.90 (9)
C20—C15—S2	119.98 (14)	O4—S2—C15	108.02 (8)
C17—C16—C15	119.26 (19)	O3—S2—C15	107.73 (9)
C17—C16—H16	120.4	N2—S2—C15	107.22 (8)
C15—C16—H16	120.4	C7—N1—C21	115.15 (17)
C16—C17—C18	120.1 (2)	C7—N1—S1	116.09 (13)
C16—C17—H17	119.9	C21—N1—S1	116.36 (14)
C18—C17—H17	119.9	C14—N2—C23	115.60 (17)
C19—C18—C17	120.4 (2)	C14—N2—S2	117.71 (13)
C19—C18—H18	119.8	C23—N2—S2	117.68 (13)

C6—C1—C2—C3	0.1 (3)	C9—C8—S1—O2	−167.52 (19)
C6—C1—C2—C14	−178.75 (16)	C13—C8—S1—O2	15.1 (2)
C1—C2—C3—C4	0.2 (3)	C9—C8—S1—N1	77.6 (2)
C14—C2—C3—C4	179.06 (18)	C13—C8—S1—N1	−99.8 (2)
C2—C3—C4—C5	−0.2 (3)	C16—C15—S2—O4	162.80 (16)
C3—C4—C5—C6	−0.1 (3)	C20—C15—S2—O4	−19.20 (19)
C4—C5—C6—C1	0.5 (3)	C16—C15—S2—O3	31.99 (18)
C4—C5—C6—C7	−177.20 (18)	C20—C15—S2—O3	−150.01 (16)
C2—C1—C6—C5	−0.5 (3)	C16—C15—S2—N2	−82.76 (17)
C2—C1—C6—C7	177.22 (16)	C20—C15—S2—N2	95.24 (17)
C5—C6—C7—N1	−112.7 (2)	C6—C7—N1—C21	71.0 (2)
C1—C6—C7—N1	69.7 (2)	C6—C7—N1—S1	−148.14 (14)
C13—C8—C9—C10	1.8 (4)	C22B—C21—N1—C7	−98.2 (6)
S1—C8—C9—C10	−175.61 (19)	C22A—C21—N1—C7	−131.5 (4)
C8—C9—C10—C11	−1.3 (4)	C22B—C21—N1—S1	121.0 (6)
C9—C10—C11—C12	−0.2 (5)	C22A—C21—N1—S1	87.7 (4)
C10—C11—C12—C13	1.3 (5)	O1—S1—N1—C7	48.65 (16)
C11—C12—C13—C8	−0.8 (5)	O2—S1—N1—C7	177.65 (14)
C9—C8—C13—C12	−0.7 (4)	C8—S1—N1—C7	−66.87 (15)
S1—C8—C13—C12	176.7 (2)	O1—S1—N1—C21	−170.94 (15)
C3—C2—C14—N2	120.3 (2)	O2—S1—N1—C21	−41.94 (17)
C1—C2—C14—N2	−60.8 (2)	C8—S1—N1—C21	73.54 (16)
C20—C15—C16—C17	0.4 (3)	C2—C14—N2—C23	−69.5 (2)
S2—C15—C16—C17	178.43 (17)	C2—C14—N2—S2	144.01 (14)
C15—C16—C17—C18	−0.4 (3)	C24—C23—N2—C14	114.9 (2)
C16—C17—C18—C19	0.6 (4)	C24—C23—N2—S2	−98.7 (2)
C17—C18—C19—C20	−0.8 (4)	O4—S2—N2—C14	−173.57 (14)
C18—C19—C20—C15	0.8 (3)	O3—S2—N2—C14	−44.33 (16)
C16—C15—C20—C19	−0.6 (3)	C15—S2—N2—C14	70.98 (15)
S2—C15—C20—C19	−178.58 (17)	O4—S2—N2—C23	40.68 (17)
C9—C8—S1—O1	−37.4 (2)	O3—S2—N2—C23	169.93 (15)
C13—C8—S1—O1	145.2 (2)	C15—S2—N2—C23	−74.77 (16)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
C17—H17···O1 ⁱ	0.93	2.57	3.409 (3)	151

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