

## 4-Methyl-N-[4-[(5-methyl-1,2-oxazol-3-yl)sulfamoyl]phenyl]benzenesulfonamide

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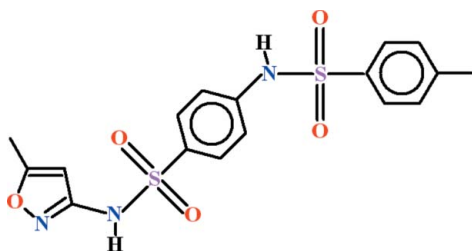
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Key indicators: single-crystal X-ray study;  $T = 296$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.055;  $wR$  factor = 0.141; data-to-parameter ratio = 18.8.

In the title compound,  $\text{C}_{17}\text{H}_{17}\text{N}_3\text{O}_5\text{S}_2$ , the dihedral angle between the two benzene rings is  $81.27(8)^\circ$  and the heterocyclic ring is oriented at  $9.1(2)$  and  $76.01(9)^\circ$  with respect to these rings. Molecules are connected *via*  $\text{N}-\text{H}\cdots\text{N}$  and  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds, generating an  $R_2^2(8)$  motif, into chains running along the  $[001]$  direction. There is also an intramolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bond completing an  $S(6)$  ring motif. The polymeric chains are interlinked through intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds.

### Related literature

For a related crystal structure, see: Ashfaq *et al.* (2010). For graph-set notation, see: Bernstein *et al.* (1995).



### Experimental

#### Crystal data

$\text{C}_{17}\text{H}_{17}\text{N}_3\text{O}_5\text{S}_2$   
 $M_r = 407.46$

Monoclinic,  $P2_1/c$   
 $a = 10.6294(6)$  Å

$b = 12.2394(7)$  Å  
 $c = 14.9673(11)$  Å  
 $\beta = 106.863(2)^\circ$   
 $V = 1863.5(2)$  Å<sup>3</sup>  
 $Z = 4$

Mo  $K\alpha$  radiation  
 $\mu = 0.32$  mm<sup>-1</sup>  
 $T = 296$  K  
 $0.35 \times 0.25 \times 0.22$  mm

#### Data collection

Bruker Kappa APEXII CCD diffractometer  
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)  
 $T_{\min} = 0.915$ ,  $T_{\max} = 0.938$

17943 measured reflections  
4629 independent reflections  
2488 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.056$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.055$   
 $wR(F^2) = 0.141$   
 $S = 1.02$   
4629 reflections

246 parameters  
H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.47$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.37$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{N3}^{\text{i}}$	0.86	2.04	2.859 (3)	159
$\text{N2}-\text{H2A}\cdots\text{O1}^{\text{ii}}$	0.86	2.39	2.979 (3)	127
$\text{C2}-\text{H2}\cdots\text{O4}^{\text{iii}}$	0.93	2.42	3.208 (4)	143
$\text{C13}-\text{H13}\cdots\text{O2}$	0.93	2.49	3.134 (3)	126

Symmetry codes: (i)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (ii)  $x, -y + \frac{1}{2}, z - \frac{1}{2}$ ; (iii)  $-x, y - \frac{1}{2}, -z + \frac{1}{2}$ .

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); 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 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2455).

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## supporting information

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## 4-Methyl-*N*-{4-[(5-methyl-1,2-oxazol-3-yl)sulfamoyl]phenyl}benzenesulfonamide

Muhammad Sohail, Muhammad Nadeem Asghar, M. Nawaz Tahir, Muhammad Shafique and Muhammad Ashfaq

### S1. Comment

The title compound (Fig. 1) has been synthesized as a part of the series of new sulfonamide derivatives. The aim of our research work is to find the potential sulfonamide derivatives possessing anti-microbial activity. The crystal structures of a similar compound, *N*-[4-(*p*-toluenesulfonamido)phenylsulfonyl]acetamide (Ashfaq *et al.*, 2010) has already been reported.

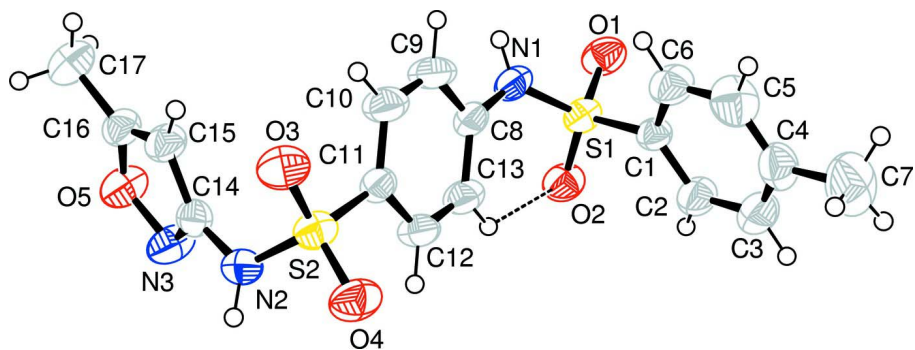
In (I), the phenyl rings A (C1–C6), B (C8–C13) and the heterocyclic five-membered ring C (C14/C15/C16/O5/N3) are planar with r. m. s. deviation of 0.0068 Å, 0.0031 Å and 0.0058 Å, respectively. The dihedral angles between A/B, A/C and B/C are 81.27 (8)°, 9.12 (20)° and 76.01 (9)°, respectively. There exist intramolecular hydrogen bond of C—H···O type (Table 1, Fig. 1) forming an S(6) ring motif (Bernstein *et al.*, 1995). There exist intermolecular hydrogen bonds of N—H···N and N—H···O types (Table 1, Fig. 2) due to which the molecules are connected from the ends to form one-dimensional polymeric network along the [0 0 1] direction and complete  $R_2^2(8)$  ring motifs. The polymeric chains are interlinked through intermolecular hydrogen bond of C—H···O type (Table 1, Fig. 2).

### S2. Experimental

Equimolar amount of 4-amino-*N*-(5-methyl-4,5-dihydro-1,2-oxazol-3-yl)benzenesulfonamide and *p*-toluenesulfonyl chloride was dissolved in 20 ml distilled water. The pH was adjusted to 8–9 using  $\text{Na}_2\text{CO}_3$  (1 *M*) and the solution was stirred at room temperature for 6 h. The progress of reaction was monitored by the consumption of suspended *p*-toluenesulfonyl chloride. On completion, pH was adjusted to 2–3 using HCl (2 *N*). The precipitate formed was filtered, washed with distilled water and recrystallized from methanol to afford colorless prisms (m.p. 403 K).

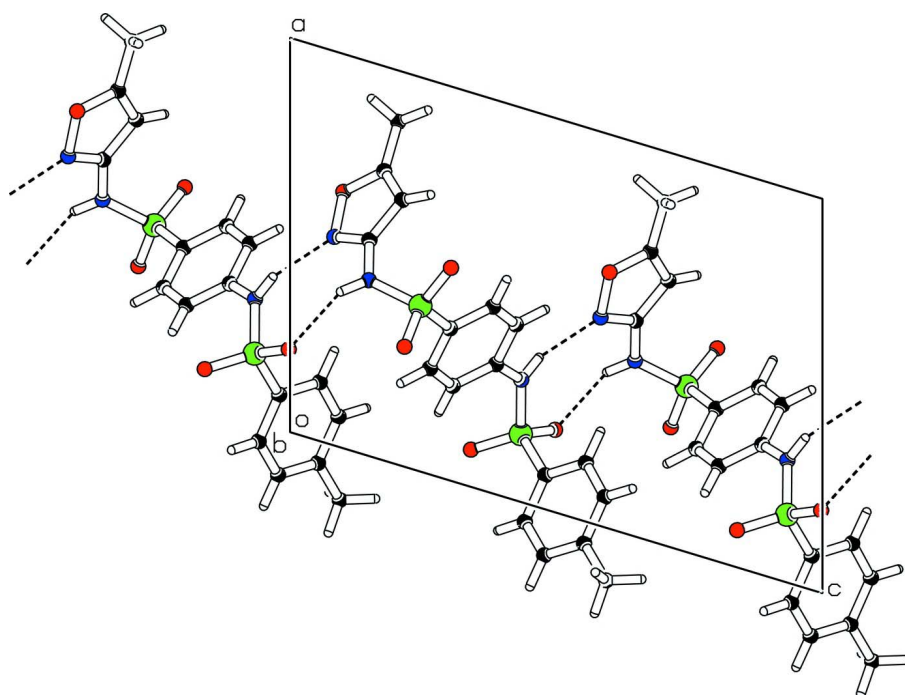
### S3. Refinement

The H-atoms were positioned geometrically (N–H = 0.86 Å, C–H = 0.93–0.96 Å) and refined as riding with  $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C}, \text{N})$ , where  $x = 1.5$  for methyl groups and  $x = 1.2$  for all H-atoms.



**Figure 1**

View of the title molecule with displacement ellipsoids drawn at the 50% probability level. H-atoms are shown as small circles of arbitrary radii. The dotted line represents an intramolecular C-H...O hydrogen bond



**Figure 2**

The partial packing (*PLATON*; Spek, 2009) which shows that molecules form polymeric chains extending along [0 0 1] direction.

#### 4-Methyl-N-{4-[(5-methyl-1,2-oxazol-3-yl)sulfamoyl]phenyl}benzenesulfonamide

##### Crystal data

$C_{17}H_{17}N_3O_5S_2$

$M_r = 407.46$

Monoclinic,  $P2_1/c$

Hall symbol: -P 2ybc

$a = 10.6294$  (6) Å

$b = 12.2394$  (7) Å

$c = 14.9673$  (11) Å

$\beta = 106.863$  (2)°

$V = 1863.5$  (2) Å<sup>3</sup>

$Z = 4$

$F(000) = 848$

$D_x = 1.452$  Mg m<sup>-3</sup>

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

Cell parameters from 2488 reflections

$\theta = 2.2$ – $28.3$ °

$\mu = 0.32$  mm<sup>-1</sup>

$T = 296$  K  $0.35 \times 0.25 \times 0.22$  mm  
 Prism, colorless

*Data collection*

Bruker Kappa APEXII CCD diffractometer Radiation source: fine-focus sealed tube Graphite monochromator Detector resolution: $7.50$ pixels $\text{mm}^{-1}$ $\omega$ scans Absorption correction: multi-scan (SADABS; Bruker, 2005) $T_{\min} = 0.915$ , $T_{\max} = 0.938$	17943 measured reflections 4629 independent reflections 2488 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.056$ $\theta_{\max} = 28.3^\circ$ , $\theta_{\min} = 2.2^\circ$ $h = -14 \rightarrow 13$ $k = -14 \rightarrow 16$ $l = -19 \rightarrow 19$
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*Refinement*

Refinement on $F^2$ Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.055$ $wR(F^2) = 0.141$ $S = 1.02$ 4629 reflections 246 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 $w = 1/[\sigma^2(F_o^2) + (0.0577P)^2 + 0.1435P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.47 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.37 \text{ e } \text{\AA}^{-3}$
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*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 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	0.17178 (7)	0.12476 (6)	0.43330 (5)	0.0475 (3)
S2	0.42023 (7)	0.58034 (6)	0.24478 (5)	0.0485 (3)
O1	0.20899 (18)	0.03512 (15)	0.49755 (15)	0.0596 (8)
O2	0.09435 (18)	0.10311 (16)	0.33965 (14)	0.0575 (7)
O3	0.54004 (19)	0.62547 (15)	0.30218 (14)	0.0594 (7)
O4	0.30436 (19)	0.64578 (16)	0.21489 (15)	0.0642 (8)
O5	0.65024 (19)	0.33654 (16)	0.10010 (15)	0.0618 (8)
N1	0.3101 (2)	0.18092 (19)	0.43406 (16)	0.0519 (8)
N2	0.4456 (2)	0.53685 (18)	0.14776 (16)	0.0486 (8)
N3	0.5291 (2)	0.3918 (2)	0.08307 (18)	0.0605 (9)
C1	0.0863 (2)	0.2224 (2)	0.47933 (18)	0.0398 (8)
C2	-0.0473 (3)	0.2308 (3)	0.4409 (2)	0.0573 (11)
C3	-0.1174 (3)	0.3048 (3)	0.4774 (2)	0.0684 (14)
C4	-0.0567 (3)	0.3690 (3)	0.5522 (2)	0.0615 (12)

C5	0.0766 (4)	0.3572 (3)	0.5909 (3)	0.0703 (14)
C6	0.1495 (3)	0.2850 (3)	0.5556 (2)	0.0592 (11)
C7	-0.1339 (4)	0.4484 (3)	0.5932 (3)	0.1003 (19)
C8	0.3303 (2)	0.2762 (2)	0.38676 (19)	0.0440 (9)
C9	0.4465 (3)	0.3330 (3)	0.4242 (2)	0.0607 (11)
C10	0.4738 (3)	0.4247 (3)	0.3811 (2)	0.0578 (11)
C11	0.3859 (2)	0.4623 (2)	0.30027 (18)	0.0427 (9)
C12	0.2695 (3)	0.4061 (2)	0.2627 (2)	0.0509 (10)
C13	0.2425 (3)	0.3132 (2)	0.3054 (2)	0.0513 (10)
C14	0.5480 (3)	0.4644 (2)	0.14888 (19)	0.0430 (9)
C15	0.6744 (3)	0.4591 (2)	0.2104 (2)	0.0498 (10)
C16	0.7342 (3)	0.3799 (2)	0.1768 (2)	0.0482 (10)
C17	0.8701 (3)	0.3364 (3)	0.2031 (2)	0.0661 (11)
H1	0.37992	0.14872	0.46739	0.0623*
H2	-0.09040	0.18700	0.39052	0.0687*
H2A	0.39496	0.55907	0.09507	0.0583*
H3	-0.20773	0.31110	0.45045	0.0821*
H5	0.11898	0.39923	0.64258	0.0842*
H6	0.23972	0.27863	0.58283	0.0707*
H7A	-0.22555	0.42985	0.57205	0.1503*
H7B	-0.12163	0.52121	0.57344	0.1503*
H7C	-0.10358	0.44443	0.66009	0.1503*
H9	0.50638	0.30849	0.47898	0.0729*
H10	0.55227	0.46190	0.40663	0.0696*
H12	0.20926	0.43125	0.20833	0.0610*
H13	0.16469	0.27516	0.27928	0.0614*
H15	0.70953	0.50155	0.26337	0.0597*
H17A	0.90803	0.34876	0.15310	0.0992*
H17B	0.86837	0.25935	0.21492	0.0992*
H17C	0.92184	0.37272	0.25840	0.0992*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0381 (4)	0.0495 (4)	0.0522 (5)	0.0024 (3)	0.0088 (3)	-0.0024 (4)
S2	0.0440 (4)	0.0468 (4)	0.0504 (5)	0.0060 (3)	0.0071 (3)	-0.0020 (3)
O1	0.0506 (12)	0.0490 (12)	0.0743 (15)	0.0042 (10)	0.0104 (10)	0.0135 (11)
O2	0.0497 (11)	0.0649 (13)	0.0521 (13)	0.0017 (10)	0.0054 (10)	-0.0180 (10)
O3	0.0567 (12)	0.0535 (12)	0.0586 (14)	-0.0064 (10)	0.0019 (11)	-0.0103 (10)
O4	0.0588 (13)	0.0586 (13)	0.0694 (15)	0.0221 (11)	0.0095 (11)	0.0017 (11)
O5	0.0484 (12)	0.0645 (13)	0.0666 (15)	0.0060 (11)	0.0075 (11)	-0.0149 (11)
N1	0.0332 (12)	0.0610 (15)	0.0591 (16)	0.0086 (11)	0.0097 (11)	0.0108 (13)
N2	0.0456 (13)	0.0547 (14)	0.0425 (14)	0.0071 (12)	0.0083 (11)	0.0061 (11)
N3	0.0435 (14)	0.0665 (17)	0.0629 (18)	0.0073 (13)	0.0018 (13)	-0.0123 (14)
C1	0.0350 (14)	0.0425 (15)	0.0391 (15)	-0.0024 (12)	0.0063 (12)	-0.0003 (12)
C2	0.0443 (16)	0.074 (2)	0.0498 (19)	0.0014 (16)	0.0076 (14)	-0.0161 (16)
C3	0.0527 (19)	0.087 (3)	0.065 (2)	0.0168 (18)	0.0165 (17)	-0.005 (2)
C4	0.073 (2)	0.058 (2)	0.062 (2)	0.0090 (18)	0.0330 (19)	0.0019 (17)

C5	0.082 (3)	0.063 (2)	0.065 (2)	-0.014 (2)	0.020 (2)	-0.0241 (18)
C6	0.0465 (16)	0.066 (2)	0.059 (2)	-0.0070 (16)	0.0059 (15)	-0.0122 (17)
C7	0.128 (4)	0.084 (3)	0.105 (3)	0.030 (3)	0.059 (3)	-0.007 (2)
C8	0.0335 (14)	0.0518 (17)	0.0465 (17)	0.0084 (14)	0.0113 (13)	-0.0003 (14)
C9	0.0451 (17)	0.071 (2)	0.053 (2)	-0.0002 (16)	-0.0063 (15)	0.0160 (17)
C10	0.0416 (16)	0.068 (2)	0.054 (2)	-0.0052 (16)	-0.0018 (14)	0.0076 (17)
C11	0.0353 (14)	0.0511 (16)	0.0409 (16)	0.0056 (13)	0.0098 (12)	-0.0021 (13)
C12	0.0399 (16)	0.0602 (19)	0.0451 (17)	0.0064 (15)	0.0007 (14)	0.0041 (15)
C13	0.0353 (14)	0.061 (2)	0.0507 (18)	-0.0007 (14)	0.0017 (13)	-0.0019 (16)
C14	0.0414 (16)	0.0438 (16)	0.0438 (16)	-0.0055 (14)	0.0126 (13)	0.0034 (14)
C15	0.0392 (15)	0.0574 (19)	0.0480 (17)	-0.0066 (14)	0.0052 (14)	-0.0071 (15)
C16	0.0384 (15)	0.0499 (17)	0.0537 (19)	-0.0048 (14)	0.0094 (14)	0.0042 (15)
C17	0.0455 (17)	0.067 (2)	0.083 (2)	0.0075 (16)	0.0141 (17)	-0.0025 (18)

*Geometric parameters (Å, °)*

S1—O1	1.437 (2)	C8—C9	1.386 (4)
S1—O2	1.429 (2)	C9—C10	1.367 (5)
S1—N1	1.620 (2)	C10—C11	1.376 (4)
S1—C1	1.758 (3)	C11—C12	1.384 (4)
S2—O3	1.425 (2)	C12—C13	1.375 (4)
S2—O4	1.428 (2)	C14—C15	1.393 (4)
S2—N2	1.641 (2)	C15—C16	1.335 (4)
S2—C11	1.757 (3)	C16—C17	1.482 (5)
O5—N3	1.411 (3)	C2—H2	0.9300
O5—C16	1.342 (4)	C3—H3	0.9300
N1—C8	1.413 (3)	C5—H5	0.9300
N2—C14	1.400 (4)	C6—H6	0.9300
N3—C14	1.298 (4)	C7—H7A	0.9600
N1—H1	0.8600	C7—H7B	0.9600
N2—H2A	0.8600	C7—H7C	0.9600
C1—C2	1.373 (4)	C9—H9	0.9300
C1—C6	1.377 (4)	C10—H10	0.9300
C2—C3	1.382 (5)	C12—H12	0.9300
C3—C4	1.367 (4)	C13—H13	0.9300
C4—C7	1.512 (5)	C15—H15	0.9300
C4—C5	1.374 (5)	C17—H17A	0.9600
C5—C6	1.378 (5)	C17—H17B	0.9600
C8—C13	1.378 (4)	C17—H17C	0.9600
O1—S1—O2	118.90 (12)	C8—C13—C12	120.2 (3)
O1—S1—N1	104.26 (12)	N2—C14—N3	118.4 (3)
O1—S1—C1	109.27 (12)	N2—C14—C15	129.4 (2)
O2—S1—N1	110.41 (12)	N3—C14—C15	112.2 (3)
O2—S1—C1	107.11 (12)	C14—C15—C16	105.0 (3)
N1—S1—C1	106.26 (12)	O5—C16—C17	116.1 (2)
O3—S2—O4	120.68 (12)	C15—C16—C17	134.0 (3)
O3—S2—N2	108.17 (12)	O5—C16—C15	109.8 (3)

O3—S2—C11	107.80 (12)	C1—C2—H2	120.00
O4—S2—N2	104.65 (12)	C3—C2—H2	120.00
O4—S2—C11	109.32 (12)	C2—C3—H3	119.00
N2—S2—C11	105.17 (12)	C4—C3—H3	119.00
N3—O5—C16	108.1 (2)	C4—C5—H5	119.00
S1—N1—C8	128.05 (18)	C6—C5—H5	119.00
S2—N2—C14	121.44 (19)	C1—C6—H6	121.00
O5—N3—C14	105.0 (2)	C5—C6—H6	121.00
C8—N1—H1	116.00	C4—C7—H7A	109.00
S1—N1—H1	116.00	C4—C7—H7B	109.00
S2—N2—H2A	119.00	C4—C7—H7C	109.00
C14—N2—H2A	119.00	H7A—C7—H7B	109.00
S1—C1—C6	121.3 (2)	H7A—C7—H7C	109.00
C2—C1—C6	120.2 (3)	H7B—C7—H7C	110.00
S1—C1—C2	118.4 (2)	C8—C9—H9	120.00
C1—C2—C3	119.6 (3)	C10—C9—H9	120.00
C2—C3—C4	121.3 (3)	C9—C10—H10	120.00
C3—C4—C5	118.0 (3)	C11—C10—H10	120.00
C3—C4—C7	121.3 (3)	C11—C12—H12	120.00
C5—C4—C7	120.7 (3)	C13—C12—H12	120.00
C4—C5—C6	122.1 (4)	C8—C13—H13	120.00
C1—C6—C5	118.8 (3)	C12—C13—H13	120.00
N1—C8—C9	117.3 (2)	C14—C15—H15	128.00
N1—C8—C13	123.5 (2)	C16—C15—H15	128.00
C9—C8—C13	119.2 (3)	C16—C17—H17A	110.00
C8—C9—C10	120.5 (3)	C16—C17—H17B	109.00
C9—C10—C11	120.4 (3)	C16—C17—H17C	109.00
S2—C11—C12	120.0 (2)	H17A—C17—H17B	109.00
C10—C11—C12	119.3 (3)	H17A—C17—H17C	110.00
S2—C11—C10	120.6 (2)	H17B—C17—H17C	109.00
C11—C12—C13	120.3 (3)		
O1—S1—N1—C8	-176.4 (2)	O5—N3—C14—C15	-1.4 (3)
O2—S1—N1—C8	54.8 (3)	S1—C1—C2—C3	178.3 (2)
C1—S1—N1—C8	-61.0 (3)	C6—C1—C2—C3	2.0 (5)
O1—S1—C1—C2	-106.2 (2)	S1—C1—C6—C5	-177.5 (3)
O1—S1—C1—C6	70.1 (3)	C2—C1—C6—C5	-1.3 (5)
O2—S1—C1—C2	23.9 (3)	C1—C2—C3—C4	-1.0 (5)
O2—S1—C1—C6	-159.9 (2)	C2—C3—C4—C5	-0.6 (5)
N1—S1—C1—C2	141.9 (2)	C2—C3—C4—C7	-178.8 (3)
N1—S1—C1—C6	-41.9 (3)	C3—C4—C5—C6	1.2 (6)
O3—S2—N2—C14	53.7 (2)	C7—C4—C5—C6	179.5 (3)
O4—S2—N2—C14	-176.5 (2)	C4—C5—C6—C1	-0.3 (5)
C11—S2—N2—C14	-61.3 (2)	N1—C8—C9—C10	178.7 (3)
O3—S2—C11—C10	-4.6 (3)	C13—C8—C9—C10	0.1 (5)
O3—S2—C11—C12	175.4 (2)	N1—C8—C13—C12	-179.3 (3)
O4—S2—C11—C10	-137.5 (2)	C9—C8—C13—C12	-0.8 (4)
O4—S2—C11—C12	42.5 (3)	C8—C9—C10—C11	0.4 (5)

N2—S2—C11—C10	110.6 (2)	C9—C10—C11—S2	179.8 (2)
N2—S2—C11—C12	-69.4 (2)	C9—C10—C11—C12	-0.2 (4)
C16—O5—N3—C14	0.7 (3)	S2—C11—C12—C13	179.5 (2)
N3—O5—C16—C15	0.4 (3)	C10—C11—C12—C13	-0.5 (4)
N3—O5—C16—C17	-176.7 (2)	C11—C12—C13—C8	1.0 (4)
S1—N1—C8—C9	154.9 (2)	N2—C14—C15—C16	-176.3 (3)
S1—N1—C8—C13	-26.6 (4)	N3—C14—C15—C16	1.6 (3)
S2—N2—C14—N3	145.2 (2)	C14—C15—C16—O5	-1.1 (3)
S2—N2—C14—C15	-37.0 (4)	C14—C15—C16—C17	175.2 (3)
O5—N3—C14—N2	176.8 (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—H1...N3 <sup>i</sup>	0.86	2.04	2.859 (3)	159
N2—H2 <i>A</i> ...O1 <sup>ii</sup>	0.86	2.39	2.979 (3)	127
C2—H2...O4 <sup>iii</sup>	0.93	2.42	3.208 (4)	143
C13—H13...O2	0.93	2.49	3.134 (3)	126

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