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## Structure Reports

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# 4-[(4-Methoxybenzylidene)amino]-benzenesulfonamide

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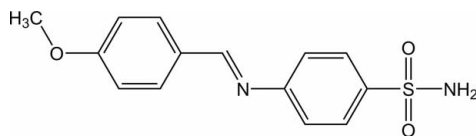
Received 18 April 2012; accepted 26 April 2012

Key indicators: single-crystal X-ray study;  $T = 200$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å;  $R$  factor = 0.035;  $wR$  factor = 0.095; data-to-parameter ratio = 18.6.

The title Schiff base compound,  $\text{C}_{14}\text{H}_{14}\text{N}_2\text{O}_3\text{S}$ , is non-planar, with a dihedral angle of  $24.16(7)^\circ$  between the benzene rings. In the crystal,  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{N}-\text{H}\cdots\text{N}$  hydrogen bonds link the molecules into a layer parallel to (011). Intra- and interlayer  $\text{C}-\text{H}\cdots\text{O}$  interactions and  $\pi-\pi$  interactions [centroid-centroid distances =  $3.8900(9)$  and  $3.9355(8)$  Å] are also present.

## Related literature

For general background to the applications of sulfanilamide Schiff bases, see: Gupta *et al.* (2003); Khalil *et al.* (2009); Nagpal & Singh (2004); Sharaby (2007); Wu *et al.* (2004).



## Experimental

### Crystal data

$\text{C}_{14}\text{H}_{14}\text{N}_2\text{O}_3\text{S}$   
 $M_r = 290.33$

Monoclinic,  $P2_1/c$   
 $a = 16.3315(5)$  Å

$b = 11.1597(3)$  Å  
 $c = 7.6876(3)$  Å  
 $\beta = 100.661(1)^\circ$   
 $V = 1376.92(8)$  Å<sup>3</sup>  
 $Z = 4$

Mo  $K\alpha$  radiation  
 $\mu = 0.24$  mm<sup>-1</sup>  
 $T = 200$  K  
 $0.60 \times 0.33 \times 0.12$  mm

### Data collection

Bruker APEXII CCD diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 2001)  
 $T_{\min} = 0.90$ ,  $T_{\max} = 0.97$

10501 measured reflections  
 3383 independent reflections  
 2821 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.017$

### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.095$   
 $S = 1.08$   
 3383 reflections

182 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\text{max}} = 0.41$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.39$  e Å<sup>-3</sup>

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: PLATON (Spek, 2009) and publCIF (Westrip, 2010).

The authors thank the Department of Chemistry and Govan Mbeki Research and Development Centre (GMRDC), University of Fort Hare, for their support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HY2541).

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## supplementary materials

*Acta Cryst.* (2012). E68, o1599 [doi:10.1107/S1600536812018818]

**4-[(4-Methoxybenzylidene)amino]benzenesulfonamide**

**Omoruyi G. Idemudia, Alexander P. Sadimenko, Anthony J. Afolayan and Eric C. Hosten**

**Comment**

Biological applications of sulfa compounds either alone or as metal complexes are well known (Gupta *et al.*, 2003; Nagpal & Singh, 2004). Their chelating powers towards metal ions tend to increase on forming a Schiff base by way of reaction with a carbonyl (Khalil *et al.*, 2009; Sharaby, 2007; Wu *et al.*, 2004). Herein, we report a new sulfanilamide Schiff base (Fig. 1), as part of our look at developing better chelating ligands from biologically active amine compounds.

The least-squares planes through the phenyl rings of the benzenesulfonamide and methoxybenzaldehyde groups have a dihedral angle of 24.16 (7)°. In the crystal, the molecules are stacked along the *c* axis and linked by N—H···O and N—H···N hydrogen bonds (Table 1 and Fig. 2) into a layer parallel to (0 1 1) (Fig. 3). The least-squares planes through adjacent two methoxybenzaldehyde phenyl rings (C11–C16) are almost parallel with a dihedral angle of 3.97° and a centroid-to-centroid distance of 3.8900 (9) Å. The centroid-to-centroid distance between adjacent two benzenesulfonamide phenyl rings (C21–C26) is 3.9355 (8) Å. C22—H22··· $\pi$  interaction occurs with the adjacent C21–C26 ring (H···Cg distance = 2.81 Å). Intra- and interlayer C—H···O interactions are also observed.

**Experimental**

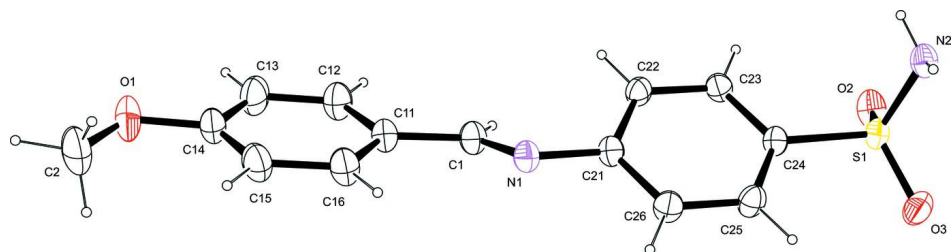
A mixture of (4-aminobenzenesulfonamide)sulfadiazine and 4-methoxybenzaldehyde (anisaldehyde) (molar ratio 1:1) in methanol was refluxed for 15 h. The resultant pale yellow precipitate was isolated by filtration and recrystallized from methanol. Yield 68% and melting point 199–201°C. Single crystals suitable for X-ray analysis were obtained from methanol by slow evaporation at room temperature.

**Refinement**

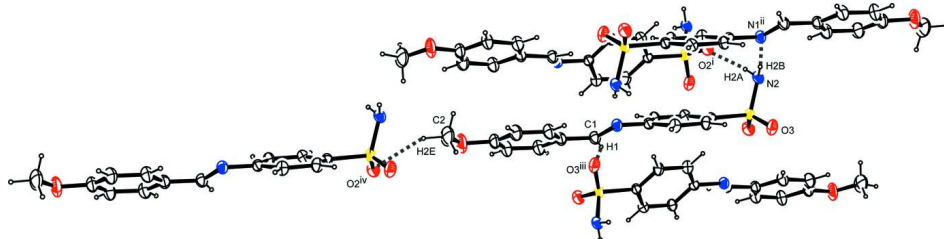
C-bound H atoms were placed in calculated positions and refined as riding atoms, with C—H = 0.95 (CH), 0.98 (CH<sub>3</sub>) Å and with  $U_{\text{iso}}(\text{H}) = 1.2(1.5 \text{ for methyl})U_{\text{eq}}(\text{C})$ . N-bound H atoms were located on a difference Fourier map and refined as riding with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$ .

**Computing details**

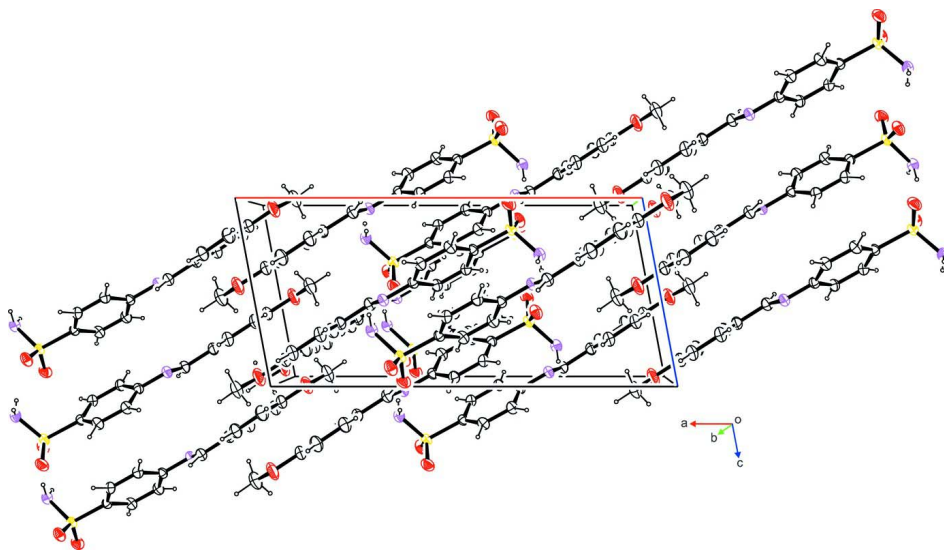
Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINTE* (Bruker, 2007); data reduction: *SAINTE* (Bruker, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *PLATON* (Spek, 2009) and *pubCIF* (Westrip, 2010).


**Figure 1**

The molecular structure of the title compound, with displacement ellipsoids drawn at the 50% probability level.


**Figure 2**

Hydrogen bonds in the title compound. [Symmetry codes: (i)  $x, -y+3/2, z-1/2$ ; (ii)  $-x+1, -y+2, -z+1$ ; (iii)  $-x+1, y-1/2, -z+3/2$ ; (iv)  $x-1, y, z-1$ .]


**Figure 3**

Crystal packing of the title compound viewed along  $[0\ 1\ 0]$ .

#### 4-[(4-Methoxybenzylidene)amino]benzenesulfonamide

##### Crystal data

$C_{14}H_{14}N_2O_3S$

$M_r = 290.33$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P\ 2ybc$

$a = 16.3315\ (5)\ \text{\AA}$

$b = 11.1597\ (3)\ \text{\AA}$

$c = 7.6876\ (3)\ \text{\AA}$

$\beta = 100.661\ (1)^\circ$

$V = 1376.92\ (8)\ \text{\AA}^3$

$Z = 4$

$F(000) = 608$

$D_x = 1.401\ \text{Mg m}^{-3}$

Melting point: 473.15 K  
 Mo  $K\alpha$  radiation,  $\lambda = 0.71073 \text{ \AA}$   
 Cell parameters from 118 reflections  
 $\theta = 3.1\text{--}29.3^\circ$

$\mu = 0.24 \text{ mm}^{-1}$   
 $T = 200 \text{ K}$   
 Platelet, yellow  
 $0.60 \times 0.33 \times 0.12 \text{ mm}$

*Data collection*

Bruker APEXII CCD  
 diffractometer  
 Radiation source: sealed tube  
 Graphite monochromator  
 Detector resolution:  $8.3333 \text{ pixels mm}^{-1}$   
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan  
 (SADABS; Bruker, 2001)  
 $T_{\min} = 0.90$ ,  $T_{\max} = 0.97$

10501 measured reflections  
 3383 independent reflections  
 2821 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.017$   
 $\theta_{\max} = 28.3^\circ$ ,  $\theta_{\min} = 2.2^\circ$   
 $h = -21 \rightarrow 21$   
 $k = -14 \rightarrow 14$   
 $l = -10 \rightarrow 8$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.035$   
 $wR(F^2) = 0.095$   
 $S = 1.08$   
 3383 reflections  
 182 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  
 $w = 1/[\sigma^2(F_o^2) + (0.0391P)^2 + 0.7392P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.41 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.39 \text{ e \AA}^{-3}$

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.663198 (19)	0.87003 (3)	0.83688 (5)	0.02211 (10)
O1	-0.04751 (7)	0.67929 (13)	0.0325 (2)	0.0479 (4)
O2	0.68589 (6)	0.74895 (9)	0.88958 (16)	0.0306 (3)
O3	0.66935 (7)	0.96150 (10)	0.96956 (16)	0.0333 (3)
N1	0.31032 (7)	0.85378 (10)	0.45630 (17)	0.0231 (2)
N2	0.72148 (7)	0.90797 (11)	0.69963 (17)	0.0248 (3)
H2A	0.719	0.8559	0.6172	0.03*
H2B	0.7141	0.9792	0.6628	0.03*
C1	0.27140 (8)	0.75396 (13)	0.4333 (2)	0.0250 (3)
H1	0.2986	0.6843	0.4867	0.03*
C2	-0.10221 (11)	0.7766 (2)	-0.0293 (3)	0.0553 (6)
H2C	-0.0756	0.8292	-0.1044	0.083*

H2D	-0.1142	0.8221	0.0721	0.083*
H2E	-0.1543	0.7451	-0.0979	0.083*
C11	0.18743 (8)	0.74102 (13)	0.3295 (2)	0.0257 (3)
C12	0.15750 (9)	0.62521 (14)	0.2867 (2)	0.0319 (3)
H12	0.1914	0.558	0.3275	0.038*
C13	0.07929 (10)	0.60763 (15)	0.1859 (2)	0.0365 (4)
H13	0.0598	0.5287	0.1558	0.044*
C14	0.02902 (9)	0.70602 (16)	0.1285 (2)	0.0335 (4)
C15	0.05726 (9)	0.82138 (15)	0.1709 (2)	0.0344 (4)
H15	0.0228	0.8884	0.1321	0.041*
C16	0.13644 (9)	0.83808 (14)	0.2707 (2)	0.0310 (3)
H16	0.1561	0.9171	0.2993	0.037*
C21	0.39382 (8)	0.85383 (12)	0.55004 (19)	0.0206 (3)
C22	0.44881 (8)	0.75989 (12)	0.5357 (2)	0.0225 (3)
H22	0.43	0.6923	0.4643	0.027*
C23	0.53057 (8)	0.76496 (12)	0.6253 (2)	0.0223 (3)
H23	0.5675	0.7002	0.6176	0.027*
C24	0.55836 (8)	0.86504 (11)	0.72618 (19)	0.0199 (3)
C25	0.50485 (9)	0.96019 (12)	0.7391 (2)	0.0244 (3)
H25	0.5244	1.0287	0.8077	0.029*
C26	0.42248 (8)	0.95441 (12)	0.6508 (2)	0.0246 (3)
H26	0.3857	1.0192	0.6592	0.03*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.01897 (16)	0.02111 (17)	0.02420 (19)	-0.00156 (11)	-0.00138 (12)	0.00123 (13)
O1	0.0217 (5)	0.0634 (9)	0.0533 (9)	-0.0056 (5)	-0.0072 (5)	-0.0087 (7)
O2	0.0250 (5)	0.0263 (5)	0.0377 (7)	0.0009 (4)	-0.0015 (4)	0.0113 (5)
O3	0.0301 (5)	0.0363 (6)	0.0304 (6)	-0.0027 (4)	-0.0023 (4)	-0.0097 (5)
N1	0.0194 (5)	0.0234 (6)	0.0252 (6)	0.0002 (4)	0.0007 (4)	0.0015 (5)
N2	0.0224 (5)	0.0197 (5)	0.0314 (7)	-0.0027 (4)	0.0029 (5)	0.0025 (5)
C1	0.0216 (6)	0.0235 (6)	0.0284 (8)	-0.0005 (5)	0.0010 (5)	0.0029 (5)
C2	0.0239 (8)	0.0838 (16)	0.0532 (13)	0.0076 (9)	-0.0063 (8)	-0.0003 (12)
C11	0.0198 (6)	0.0279 (7)	0.0282 (8)	-0.0023 (5)	0.0013 (5)	0.0010 (6)
C12	0.0263 (7)	0.0282 (7)	0.0391 (9)	-0.0030 (6)	0.0005 (6)	0.0016 (7)
C13	0.0282 (7)	0.0346 (8)	0.0440 (10)	-0.0092 (6)	-0.0002 (7)	-0.0053 (7)
C14	0.0188 (6)	0.0477 (9)	0.0323 (9)	-0.0032 (6)	0.0005 (6)	-0.0036 (7)
C15	0.0226 (7)	0.0388 (9)	0.0398 (10)	0.0041 (6)	0.0008 (6)	0.0013 (7)
C16	0.0234 (7)	0.0292 (7)	0.0388 (9)	-0.0004 (6)	0.0014 (6)	0.0009 (7)
C21	0.0191 (6)	0.0199 (6)	0.0217 (7)	-0.0003 (5)	0.0013 (5)	0.0036 (5)
C22	0.0218 (6)	0.0191 (6)	0.0259 (7)	-0.0024 (5)	0.0026 (5)	-0.0029 (5)
C23	0.0202 (6)	0.0193 (6)	0.0274 (7)	0.0002 (5)	0.0045 (5)	-0.0008 (5)
C24	0.0183 (5)	0.0192 (6)	0.0213 (7)	-0.0016 (4)	0.0014 (5)	0.0022 (5)
C25	0.0254 (6)	0.0176 (6)	0.0281 (7)	-0.0009 (5)	0.0000 (5)	-0.0023 (5)
C26	0.0238 (6)	0.0190 (6)	0.0297 (8)	0.0026 (5)	0.0015 (5)	-0.0011 (5)

Geometric parameters (Å, °)

S1—O3	1.4333 (11)	C12—H12	0.95
S1—O2	1.4393 (10)	C13—C14	1.393 (2)
S1—N2	1.6032 (13)	C13—H13	0.95
S1—C24	1.7663 (13)	C14—C15	1.386 (2)
O1—C14	1.3617 (17)	C15—C16	1.389 (2)
O1—C2	1.430 (2)	C15—H15	0.95
N1—C1	1.2786 (18)	C16—H16	0.95
N1—C21	1.4196 (16)	C21—C26	1.3945 (19)
N2—H2A	0.8551	C21—C22	1.3978 (18)
N2—H2B	0.8452	C22—C23	1.3856 (18)
C1—C11	1.4607 (18)	C22—H22	0.95
C1—H1	0.95	C23—C24	1.3875 (19)
C2—H2C	0.98	C23—H23	0.95
C2—H2D	0.98	C24—C25	1.3904 (18)
C2—H2E	0.98	C25—C26	1.3916 (19)
C11—C16	1.390 (2)	C25—H25	0.95
C11—C12	1.399 (2)	C26—H26	0.95
C12—C13	1.380 (2)		
O3—S1—O2	119.20 (7)	C14—C13—H13	120.1
O3—S1—N2	107.96 (7)	O1—C14—C15	124.27 (15)
O2—S1—N2	106.25 (7)	O1—C14—C13	115.28 (15)
O3—S1—C24	107.42 (6)	C15—C14—C13	120.44 (14)
O2—S1—C24	106.39 (6)	C14—C15—C16	119.35 (15)
N2—S1—C24	109.38 (6)	C14—C15—H15	120.3
C14—O1—C2	117.90 (15)	C16—C15—H15	120.3
C1—N1—C21	118.48 (12)	C15—C16—C11	121.04 (15)
S1—N2—H2A	110.9	C15—C16—H16	119.5
S1—N2—H2B	113.8	C11—C16—H16	119.5
H2A—N2—H2B	114.0	C26—C21—C22	119.53 (12)
N1—C1—C11	123.66 (13)	C26—C21—N1	118.32 (12)
N1—C1—H1	118.2	C22—C21—N1	122.04 (12)
C11—C1—H1	118.2	C23—C22—C21	120.29 (12)
O1—C2—H2C	109.5	C23—C22—H22	119.9
O1—C2—H2D	109.5	C21—C22—H22	119.9
H2C—C2—H2D	109.5	C22—C23—C24	119.74 (12)
O1—C2—H2E	109.5	C22—C23—H23	120.1
H2C—C2—H2E	109.5	C24—C23—H23	120.1
H2D—C2—H2E	109.5	C23—C24—C25	120.64 (12)
C16—C11—C12	118.77 (13)	C23—C24—S1	118.85 (10)
C16—C11—C1	123.09 (13)	C25—C24—S1	120.51 (10)
C12—C11—C1	118.13 (13)	C24—C25—C26	119.57 (13)
C13—C12—C11	120.67 (14)	C24—C25—H25	120.2
C13—C12—H12	119.7	C26—C25—H25	120.2
C11—C12—H12	119.7	C25—C26—C21	120.19 (12)
C12—C13—C14	119.73 (15)	C25—C26—H26	119.9
C12—C13—H13	120.1	C21—C26—H26	119.9

C21—N1—C1—C11	-175.87 (13)	C26—C21—C22—C23	1.9 (2)
N1—C1—C11—C16	-11.2 (2)	N1—C21—C22—C23	178.05 (13)
N1—C1—C11—C12	168.42 (16)	C21—C22—C23—C24	-1.5 (2)
C16—C11—C12—C13	0.9 (3)	C22—C23—C24—C25	0.3 (2)
C1—C11—C12—C13	-178.67 (16)	C22—C23—C24—S1	-179.27 (11)
C11—C12—C13—C14	-1.1 (3)	O3—S1—C24—C23	-163.03 (12)
C2—O1—C14—C15	0.3 (3)	O2—S1—C24—C23	-34.34 (13)
C2—O1—C14—C13	179.64 (17)	N2—S1—C24—C23	80.04 (12)
C12—C13—C14—O1	-178.90 (16)	O3—S1—C24—C25	17.41 (14)
C12—C13—C14—C15	0.4 (3)	O2—S1—C24—C25	146.11 (12)
O1—C14—C15—C16	179.57 (16)	N2—S1—C24—C25	-99.52 (13)
C13—C14—C15—C16	0.3 (3)	C23—C24—C25—C26	0.5 (2)
C14—C15—C16—C11	-0.4 (3)	S1—C24—C25—C26	180.00 (11)
C12—C11—C16—C15	-0.2 (2)	C24—C25—C26—C21	0.0 (2)
C1—C11—C16—C15	179.39 (15)	C22—C21—C26—C25	-1.2 (2)
C1—N1—C21—C26	-148.41 (14)	N1—C21—C26—C25	-177.44 (13)
C1—N1—C21—C22	35.4 (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>
N2—H2 <i>A</i> ...O2 <sup>i</sup>	0.86	2.09	2.9280 (17)	166
N2—H2 <i>B</i> ...N1 <sup>ii</sup>	0.85	2.08	2.9233 (17)	173
C1—H1...O3 <sup>iii</sup>	0.95	2.55	3.4466 (18)	157
C2—H2 <i>E</i> ...O2 <sup>iv</sup>	0.98	2.59	3.415 (2)	141

Symmetry codes: (i)  $x, -y+3/2, z-1/2$ ; (ii)  $-x+1, -y+2, -z+1$ ; (iii)  $-x+1, y-1/2, -z+3/2$ ; (iv)  $x-1, y, z-1$ .