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2-(4,5-Dimethoxy-2-nitrophenyl)-4-methoxy-9-phenylsulfonyl-9H-carbazole-3-carbaldehyde

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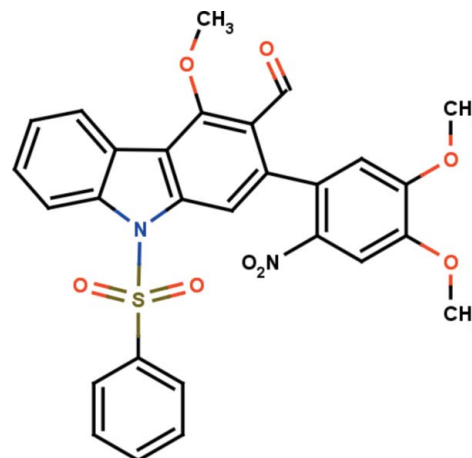
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.043; wR factor = 0.115; data-to-parameter ratio = 15.6.

In the title compound, $\text{C}_{28}\text{H}_{22}\text{N}_2\text{O}_8\text{S}$, the carbazole ring system is roughly planar, with a maximum deviation of 0.084 (3) Å for the C atom connected to the 4,5-dimethoxy-2-nitrophenyl ring. The dihedral angle between the carbazole system and the dimethoxy-substituted nitrophenyl ring is 57.05 (10)°. The aldehyde C atom deviates by 0.164 (5) Å from its attached carbazole ring system. The molecular structure is stabilized by $\text{C}-\text{H}\cdots\text{O}$ interactions which generate two $S(6)$ and one $S(7)$ ring motif. In the crystal, molecules are linked by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming $R_3^2(15)$ ring motifs, which are further crosslinked by $R_3^2(19)$ ring motifs, resulting in (002) layers. The crystal packing also features $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For the biological activities and uses of carbazole derivatives, see: Itoigawa *et al.* (2000); Ramsewak *et al.* (1999). For electronic properties and applications of carbazole derivatives, see: Friend *et al.* (1999); Zhang *et al.* (2004). For related structures, see: Gopinath *et al.* (2013); Narayanan *et al.* (2014). For the Thorpe–Ingold effect, see: Bassindale (1984). For standard bond lengths, see: Allen *et al.* (1987). For graph-set notation: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{28}\text{H}_{22}\text{N}_2\text{O}_8\text{S}$
 $M_r = 546.55$
 Orthorhombic, $Pca2_1$
 $a = 8.3976$ (6) Å
 $b = 13.7584$ (9) Å
 $c = 21.7971$ (12) Å
 $V = 2518.4$ (3) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.19$ mm⁻¹
 $T = 295$ K
 $0.25 \times 0.25 \times 0.20$ mm

Data collection

Bruker Kappa APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2008)
 $T_{\min} = 0.949$, $T_{\max} = 0.971$
 16717 measured reflections
 5538 independent reflections
 4047 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.115$
 $S = 1.01$
 5538 reflections
 355 parameters
 2 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.37$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.35$ e Å⁻³
 Absolute structure: Flack (1983)
 Absolute structure parameter: 0.02 (8)

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}2-\text{H}2\cdots\text{O}1$	0.93	2.35	2.942 (5)	122
$\text{C}11-\text{H}11\cdots\text{O}2$	0.93	2.32	2.927 (3)	122
$\text{C}25-\text{H}25\text{B}\cdots\text{O}8$	0.96	2.60	3.194 (5)	120
$\text{C}17-\text{H}17\cdots\text{O}7^i$	0.93	2.56	3.384 (3)	148
$\text{C}27-\text{H}27\text{B}\cdots\text{O}3^{\text{ii}}$	0.96	2.56	3.239 (5)	128
$\text{C}27-\text{H}27\text{B}\cdots\text{O}8^{\text{iii}}$	0.96	2.58	2.943 (5)	103
$\text{C}25-\text{H}25\text{A}\cdots\text{C}g1^{\text{iv}}$	0.96	2.97	3.675 (4)	131

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

Data collection: APEX2 (Bruker, 2008); cell refinement: APEX2; data reduction: SAINT (Bruker, 2008); 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, 2012) and Mercury (Macrae *et al.*, 2008); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: RK2423).

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supporting information

Acta Cryst. (2014). E70, o424–o425 [doi:10.1107/S1600536814005133]

2-(4,5-Dimethoxy-2-nitrophenyl)-4-methoxy-9-phenylsulfonyl-9H-carbazole-3-carbaldehyde

P. Narayanan, K. Sethusankar, Velu Saravanan and Arasambattu K. Mohanakrishnan

S1. Comment

Carbazole and its derivative have become quite attractive compounds owing to their applications in pharmacy and molecular electronics. It has been reported that carbazole derivatives exhibit various biological activities such as antitumor (Itoigawa *et al.*, 2000), anti-inflammatory and antimutagenic (Ramsewak *et al.*, 1999). Carbazole derivatives also exhibit electroactivity and luminescence and are considered to be potential candidates for electronic applications such as colour displays, organic semiconductors, laser and solar cells (Friend *et al.*, 1999; Zhang *et al.*, 2004).

The title compound, C₂₈H₂₂N₂O₈S, comprises a carbazole ring system which is attached to a phenylsulfonyl ring, a dimethoxy substituted nitrophenyl ring, a methoxy group and an aldehyde group. The carbazole ring system is essentially planar with maximum deviation of 0.084 (3) Å for the carbon (C10) atom connected to 4,5-dimethoxy-2-nitrophenyl ring. The aldehyde group carbon (C26) atom and methoxy group oxygen (O5) atom deviate from the adjacent carbazole ring by -0.164 (5) Å and -0.017 (2) Å, respectively. The carbazole ring system is almost orthogonal to phenyl ring (C19–C24) attached to sulfonyl group with dihedral angle of 88.31 (13)°. The dihedral angle between the carbazole ring and the dimethoxy substituted nitrophenyl ring (C13–C18) is 57.05 (10)°.

The atom S1 has a distorted tetrahedral configuration. The widening of angle O2=S1=O1 (119.83 (15)°) and narrowing of angle N1–S1–C19 (104.91 (12)°) from the ideal tetrahedral value are attributed to the Thorpe–Ingold effect (Bassindale, 1984). As a result of electron-withdrawing character of the phenylsulfonyl group, the bond lengths N1–C1 = 1.430 (3) Å and N1–C12 = 1.417 (3) Å in the molecule are longer than the mean value of 1.355 (14) Å (Allen *et al.*, 1987). The sum of the bond angles around N1 (356.4°) indicate the sp² hybridization. The oxygen atoms O6 & O7 are deviated by 0.025 (2) Å and -0.0105 (19) Å, respectively from the phenyl ring (C13–C18). The title compound exhibits the structural similarities with the already reported related structures (Gopinath *et al.*, 2013; Narayanan *et al.*, 2014).

The molecular structure is stabilized by C2–H2⋯O1, C11–H11⋯O2 and C25–H25B⋯O8 intramolecular interactions, which generate two S(6) and one S(7) ring motifs (Fig. 1). In the crystal packing, molecules are linked by C17–H17⋯O7ⁱ and C27–H27B⋯O3ⁱⁱ intermolecular hydrogen bondings form R₃³(15) ring motifs and these ring motifs are further cross linked by C27–H27B⋯O8ⁱⁱⁱ hydrogen bond forms R₃²(19) ring motifs, resulting in two dimensional supramolecular networks (Fig. 2) (Bernstein *et al.*, 1995). The crystal packing is also characterized by C25–H25A⋯Cg1^{iv} interactions. The packing view of the title compound is shown in Fig. 2. Symmetry codes: (i) 1/2+x, 1-y, z; (ii) -1+x, y, z; (iii) -1/2+x, -y, z; (vi) -1/2-x, y, 1/2+z.

S2. Experimental

A solution of 3-(bromomethyl)-2-(4,5-dimethoxy-2-nitrophenyl)-4-methoxy-9-(phenylsulfonyl)-9H-carbazole (1.53 g, 2.5 mmol) and bis(tetrabutylammonium) dichromate (2.63 g, 3.75 mmol) in dry CHCl₃ (50 ml) was refluxed for 10 h. The subsequent removal of solvent in vacuo followed by column chromatographic (silica gel; hexane–ethyl acetate, 4:1)

purification afforded 9-(phenylsulfonyl)-2-(4,5-dimethoxy-2-nitrophenyl)-4-methoxy-9*H*-carbazole-3-carbaldehyde (1.19 g, 86%) as a colourless solid. Single crystal suitable for X-ray diffraction were prepared by slow evaporation of a solution of the title compound in chloroform (CHCl₃) at room temperature. M.p. 471–473 K.

S3. Refinement

The positions of hydrogen atoms were localized from the difference electron density maps and their distances were geometrically constrained. The hydrogen atoms bound to the C atoms are treated as riding atoms, with $d(\text{C}-\text{H})=0.93$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic and aldehyde group, $d(\text{C}-\text{H})=0.96$ and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl groups. The rotation angles for methyl groups were optimized by least squares. In the absence of significant anomalous dispersion effects, an absolute structure was not determined and 1310 Friedel pairs were merged.

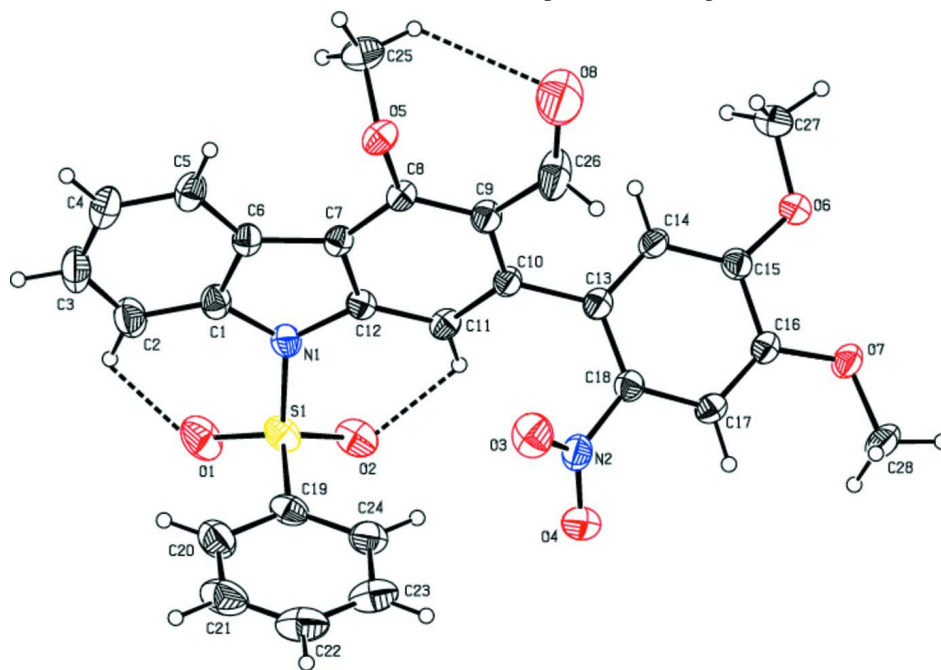


Figure 1

The molecular structure of the title compound with the atom numbering scheme. Displacement ellipsoids are drawn at 30% probability level. H atoms are present as small spheres of arbitrary radius. The intramolecular non-classical C—H...O hydrogen bonds, which are generate S(6) and S(7) ring motifs, shown as a dashed lines.

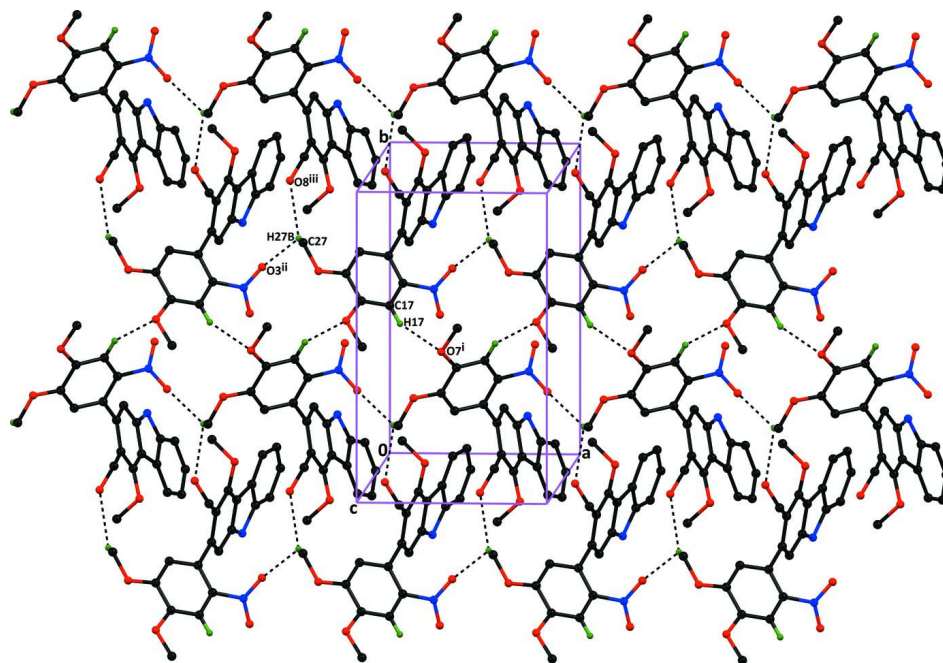


Figure 2

The packing arrangement of the title compound viewed down *b* axis. The dashed lines indicate C—H \cdots O intermolecular interactions, which are results in $R_3^3(15)$ and $R_3^2(19)$ ring motifs. The hydrogen atoms not involved in the hydrogen bonding and phenylsulfonyl group have been excluded for clarity. Symmetry codes: (i) $1/2+x, 1-y, z$; (ii) $-1+x, y, z$; (iii) $-1/2+x, -y, z$.

2-(4,5-Dimethoxy-2-nitrophenyl)-4-methoxy-9-phenylsulfonyl-9H-carbazole-3-carbaldehyde

Crystal data

$C_{28}H_{22}N_2O_8S$

$M_r = 546.55$

Orthorhombic, $Pca2_1$

Hall symbol: $P\ 2c\ -2ac$

$a = 8.3976$ (6) Å

$b = 13.7584$ (9) Å

$c = 21.7971$ (12) Å

$V = 2518.4$ (3) Å³

$Z = 4$

$F(000) = 1136$

$D_x = 1.441$ Mg m⁻³

Melting point = 471–473 K

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

Cell parameters from 4047 reflections

$\theta = 2.4$ – 29.9°

$\mu = 0.19$ mm⁻¹

$T = 295$ K

Block, colourless

$0.25 \times 0.25 \times 0.20$ mm

Data collection

Bruker Kappa APEX-II CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω - & φ -scans

Absorption correction: multi-scan
(*SADABS*; Bruker, 2008)

$T_{\min} = 0.949$, $T_{\max} = 0.971$

16717 measured reflections

5538 independent reflections

4047 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 29.9^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -11 \rightarrow 11$

$k = -19 \rightarrow 12$

$l = -16 \rightarrow 30$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.115$
 $S = 1.01$
 5538 reflections
 355 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-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.059P)^2 + 0.2808P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.37 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.35 \text{ e } \text{\AA}^{-3}$
 Absolute structure: Flack (1983)
 Absolute structure parameter: 0.02 (8)

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 $F^2 > \sigma(F^2)$ 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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.8650 (3)	0.1054 (2)	0.40810 (13)	0.0448 (6)
C2	0.9302 (4)	0.0894 (3)	0.35094 (14)	0.0615 (8)
H2	0.9377	0.1391	0.3221	0.074*
C3	0.9838 (4)	-0.0031 (3)	0.33812 (16)	0.0708 (10)
H3	1.0298	-0.0156	0.3001	0.085*
C4	0.9712 (4)	-0.0772 (3)	0.37992 (15)	0.0598 (8)
H4	1.0067	-0.1391	0.3694	0.072*
C5	0.9070 (3)	-0.0617 (2)	0.43698 (14)	0.0511 (7)
H5	0.8994	-0.1121	0.4652	0.061*
C6	0.8536 (3)	0.0312 (2)	0.45165 (12)	0.0396 (6)
C7	0.7830 (3)	0.07281 (18)	0.50642 (12)	0.0366 (5)
C8	0.7490 (3)	0.03425 (17)	0.56383 (12)	0.0371 (5)
C9	0.6929 (3)	0.09573 (19)	0.61001 (12)	0.0434 (6)
C10	0.6676 (3)	0.19546 (18)	0.59770 (12)	0.0382 (5)
C11	0.6948 (3)	0.23323 (19)	0.53996 (12)	0.0391 (6)
H11	0.6732	0.2981	0.5313	0.047*
C12	0.7553 (3)	0.17175 (18)	0.49529 (11)	0.0365 (5)
C13	0.6053 (3)	0.26215 (18)	0.64630 (12)	0.0387 (5)
C14	0.4585 (3)	0.24500 (19)	0.67360 (12)	0.0419 (6)
H14	0.4016	0.1895	0.6630	0.050*
C15	0.3949 (3)	0.30877 (18)	0.71640 (13)	0.0417 (6)
C16	0.4779 (3)	0.39378 (19)	0.73222 (11)	0.0393 (5)
C17	0.6221 (3)	0.41252 (19)	0.70510 (12)	0.0408 (6)

H17	0.6778	0.4690	0.7146	0.049*
C18	0.6841 (3)	0.34688 (19)	0.66353 (12)	0.0401 (6)
C19	1.0216 (3)	0.3359 (2)	0.43564 (14)	0.0496 (7)
C20	1.1564 (4)	0.3137 (2)	0.4026 (2)	0.0661 (9)
H20	1.1486	0.2835	0.3646	0.079*
C21	1.3031 (4)	0.3368 (3)	0.4268 (2)	0.0798 (12)
H21	1.3949	0.3226	0.4047	0.096*
C22	1.3149 (4)	0.3804 (3)	0.4830 (2)	0.0792 (12)
H22	1.4146	0.3946	0.4993	0.095*
C23	1.1808 (5)	0.4032 (3)	0.5151 (2)	0.0798 (11)
H23	1.1896	0.4341	0.5529	0.096*
C24	1.0324 (4)	0.3809 (2)	0.49226 (17)	0.0599 (8)
H24	0.9410	0.3957	0.5145	0.072*
C25	0.6625 (4)	-0.1291 (2)	0.55932 (18)	0.0603 (8)
H25A	0.6311	-0.1181	0.5176	0.090*
H25B	0.5721	-0.1210	0.5858	0.090*
H25C	0.7031	-0.1940	0.5634	0.090*
C26	0.6670 (6)	0.0594 (3)	0.67271 (17)	0.0833 (12)
H26	0.7082	0.0952	0.7052	0.100*
C27	0.1723 (4)	0.2068 (2)	0.73733 (17)	0.0636 (9)
H27A	0.2407	0.1538	0.7485	0.095*
H27B	0.1418	0.2005	0.6951	0.095*
H27C	0.0788	0.2058	0.7627	0.095*
C28	0.4937 (4)	0.5310 (2)	0.79770 (17)	0.0663 (9)
H28A	0.5144	0.5765	0.7653	0.099*
H28B	0.5928	0.5081	0.8142	0.099*
H28C	0.4335	0.5623	0.8295	0.099*
N1	0.8002 (3)	0.19235 (16)	0.43400 (11)	0.0443 (5)
N2	0.8427 (3)	0.3687 (2)	0.63994 (11)	0.0508 (6)
O1	0.8401 (3)	0.2956 (2)	0.34239 (11)	0.0744 (7)
O2	0.7169 (2)	0.36484 (15)	0.43482 (11)	0.0597 (5)
O3	0.9392 (3)	0.30255 (18)	0.63549 (13)	0.0732 (7)
O4	0.8724 (3)	0.45314 (18)	0.62628 (12)	0.0744 (7)
O5	0.7846 (2)	-0.06046 (13)	0.57617 (9)	0.0473 (5)
O6	0.2548 (3)	0.29660 (14)	0.74591 (10)	0.0550 (5)
O7	0.4055 (2)	0.45109 (14)	0.77418 (9)	0.0497 (5)
O8	0.5939 (8)	-0.0159 (2)	0.68447 (18)	0.168 (2)
S1	0.83364 (8)	0.30432 (5)	0.40724 (4)	0.05068 (19)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0412 (13)	0.0577 (17)	0.0356 (13)	-0.0009 (12)	-0.0036 (11)	-0.0048 (13)
C2	0.0640 (19)	0.079 (2)	0.0416 (17)	-0.0063 (17)	0.0054 (14)	-0.0041 (15)
C3	0.063 (2)	0.102 (3)	0.0476 (19)	0.0011 (19)	0.0092 (15)	-0.0243 (19)
C4	0.0575 (18)	0.069 (2)	0.0532 (19)	0.0106 (15)	-0.0038 (14)	-0.0212 (16)
C5	0.0435 (14)	0.0604 (18)	0.0495 (17)	0.0090 (13)	-0.0059 (12)	-0.0139 (14)
C6	0.0319 (12)	0.0496 (16)	0.0372 (14)	0.0014 (11)	-0.0048 (9)	-0.0059 (11)

C7	0.0321 (11)	0.0412 (13)	0.0367 (13)	0.0015 (10)	-0.0052 (9)	-0.0046 (10)
C8	0.0418 (13)	0.0336 (13)	0.0358 (13)	0.0038 (11)	-0.0059 (10)	-0.0001 (10)
C9	0.0535 (15)	0.0396 (14)	0.0369 (14)	0.0042 (12)	-0.0008 (11)	0.0023 (11)
C10	0.0444 (13)	0.0344 (13)	0.0358 (14)	0.0015 (11)	0.0004 (10)	-0.0023 (10)
C11	0.0442 (13)	0.0333 (13)	0.0397 (14)	0.0042 (11)	-0.0053 (11)	0.0003 (11)
C12	0.0371 (11)	0.0414 (14)	0.0311 (13)	-0.0015 (11)	-0.0049 (9)	0.0038 (10)
C13	0.0473 (14)	0.0354 (13)	0.0334 (13)	0.0037 (11)	-0.0032 (10)	0.0016 (10)
C14	0.0547 (15)	0.0324 (13)	0.0387 (14)	-0.0034 (11)	-0.0003 (11)	-0.0033 (10)
C15	0.0460 (13)	0.0398 (14)	0.0395 (14)	0.0009 (11)	0.0019 (11)	0.0026 (11)
C16	0.0452 (13)	0.0356 (13)	0.0370 (14)	0.0048 (11)	-0.0026 (10)	-0.0033 (10)
C17	0.0459 (13)	0.0328 (12)	0.0437 (15)	0.0005 (11)	-0.0039 (11)	-0.0010 (11)
C18	0.0422 (13)	0.0408 (15)	0.0372 (14)	0.0006 (11)	-0.0010 (10)	-0.0009 (11)
C19	0.0480 (14)	0.0421 (15)	0.0588 (19)	0.0000 (12)	0.0029 (13)	0.0186 (14)
C20	0.0587 (18)	0.068 (2)	0.072 (2)	0.0017 (16)	0.0180 (16)	0.0060 (18)
C21	0.0464 (18)	0.070 (2)	0.123 (4)	-0.0021 (17)	0.020 (2)	0.011 (2)
C22	0.0528 (19)	0.056 (2)	0.129 (4)	-0.0050 (16)	-0.009 (2)	0.013 (2)
C23	0.073 (2)	0.062 (2)	0.104 (3)	-0.0222 (19)	-0.012 (2)	-0.004 (2)
C24	0.0561 (17)	0.0543 (18)	0.069 (2)	-0.0086 (14)	0.0060 (15)	0.0004 (16)
C25	0.0645 (19)	0.0419 (16)	0.075 (2)	-0.0013 (14)	-0.0089 (16)	0.0048 (15)
C26	0.157 (4)	0.0522 (18)	0.0410 (18)	0.0338 (19)	0.011 (2)	0.0073 (16)
C27	0.0669 (19)	0.058 (2)	0.065 (2)	-0.0191 (16)	0.0083 (15)	0.0023 (15)
C28	0.0626 (19)	0.0521 (19)	0.084 (3)	-0.0076 (15)	0.0101 (16)	-0.0333 (17)
N1	0.0502 (12)	0.0490 (13)	0.0336 (12)	-0.0035 (10)	0.0008 (9)	0.0015 (10)
N2	0.0500 (14)	0.0579 (16)	0.0446 (14)	-0.0007 (12)	0.0018 (10)	-0.0125 (11)
O1	0.0811 (17)	0.0983 (19)	0.0439 (14)	-0.0043 (13)	-0.0030 (11)	0.0244 (12)
O2	0.0505 (11)	0.0575 (12)	0.0710 (14)	0.0094 (9)	-0.0002 (10)	0.0204 (11)
O3	0.0489 (12)	0.0819 (17)	0.0890 (19)	0.0147 (12)	0.0014 (12)	-0.0089 (13)
O4	0.0733 (15)	0.0623 (15)	0.0878 (19)	-0.0204 (12)	0.0230 (13)	-0.0121 (13)
O5	0.0556 (11)	0.0359 (10)	0.0504 (12)	0.0089 (8)	-0.0117 (8)	-0.0006 (8)
O6	0.0559 (11)	0.0523 (12)	0.0570 (13)	-0.0130 (10)	0.0151 (9)	-0.0105 (9)
O7	0.0508 (10)	0.0424 (11)	0.0559 (12)	0.0031 (9)	0.0052 (9)	-0.0143 (9)
O8	0.346 (7)	0.068 (2)	0.090 (3)	-0.015 (3)	0.072 (3)	0.0203 (17)
S1	0.0495 (4)	0.0595 (5)	0.0431 (4)	0.0009 (3)	-0.0010 (3)	0.0174 (3)

Geometric parameters (Å, °)

C1—C2	1.379 (4)	C18—N2	1.459 (4)
C1—C6	1.398 (4)	C19—C20	1.376 (4)
C1—N1	1.430 (3)	C19—C24	1.383 (5)
C2—C3	1.379 (5)	C19—S1	1.751 (3)
C2—H2	0.9300	C20—C21	1.377 (5)
C3—C4	1.371 (5)	C20—H20	0.9300
C3—H3	0.9300	C21—C22	1.368 (7)
C4—C5	1.372 (4)	C21—H21	0.9300
C4—H4	0.9300	C22—C23	1.363 (6)
C5—C6	1.391 (4)	C22—H22	0.9300
C5—H5	0.9300	C23—C24	1.377 (5)
C6—C7	1.451 (3)	C23—H23	0.9300

C7—C8	1.389 (4)	C24—H24	0.9300
C7—C12	1.402 (3)	C25—O5	1.442 (4)
C8—O5	1.364 (3)	C25—H25A	0.9600
C8—C9	1.397 (4)	C25—H25B	0.9600
C9—C10	1.414 (3)	C25—H25C	0.9600
C9—C26	1.471 (4)	C26—O8	1.231 (5)
C10—C11	1.381 (4)	C26—H26	0.9300
C10—C13	1.496 (4)	C27—O6	1.429 (3)
C11—C12	1.386 (4)	C27—H27A	0.9600
C11—H11	0.9300	C27—H27B	0.9600
C12—N1	1.417 (3)	C27—H27C	0.9600
C13—C14	1.388 (4)	C28—O7	1.422 (3)
C13—C18	1.392 (4)	C28—H28A	0.9600
C14—C15	1.388 (4)	C28—H28B	0.9600
C14—H14	0.9300	C28—H28C	0.9600
C15—O6	1.351 (3)	N1—S1	1.671 (2)
C15—C16	1.404 (4)	N2—O3	1.223 (3)
C16—O7	1.352 (3)	N2—O4	1.225 (3)
C16—C17	1.372 (4)	O1—S1	1.420 (2)
C17—C18	1.381 (4)	O2—S1	1.420 (2)
C17—H17	0.9300		
C2—C1—C6	121.6 (3)	C20—C19—C24	120.8 (3)
C2—C1—N1	129.9 (3)	C20—C19—S1	120.1 (3)
C6—C1—N1	108.5 (2)	C24—C19—S1	119.0 (2)
C1—C2—C3	117.4 (3)	C19—C20—C21	119.0 (4)
C1—C2—H2	121.3	C19—C20—H20	120.5
C3—C2—H2	121.3	C21—C20—H20	120.5
C4—C3—C2	121.8 (3)	C22—C21—C20	120.6 (4)
C4—C3—H3	119.1	C22—C21—H21	119.7
C2—C3—H3	119.1	C20—C21—H21	119.7
C3—C4—C5	121.1 (3)	C23—C22—C21	120.1 (4)
C3—C4—H4	119.4	C23—C22—H22	120.0
C5—C4—H4	119.4	C21—C22—H22	120.0
C4—C5—C6	118.6 (3)	C22—C23—C24	120.7 (4)
C4—C5—H5	120.7	C22—C23—H23	119.6
C6—C5—H5	120.7	C24—C23—H23	119.6
C5—C6—C1	119.5 (2)	C23—C24—C19	118.8 (3)
C5—C6—C7	133.1 (3)	C23—C24—H24	120.6
C1—C6—C7	107.4 (2)	C19—C24—H24	120.6
C8—C7—C12	119.5 (2)	O5—C25—H25A	109.5
C8—C7—C6	132.4 (2)	O5—C25—H25B	109.5
C12—C7—C6	108.0 (2)	H25A—C25—H25B	109.5
O5—C8—C7	119.9 (2)	O5—C25—H25C	109.5
O5—C8—C9	120.7 (2)	H25A—C25—H25C	109.5
C7—C8—C9	119.2 (2)	H25B—C25—H25C	109.5
C8—C9—C10	120.1 (2)	O8—C26—C9	123.6 (4)
C8—C9—C26	120.9 (2)	O8—C26—H26	118.2

C10—C9—C26	118.9 (2)	C9—C26—H26	118.2
C11—C10—C9	120.9 (2)	O6—C27—H27A	109.5
C11—C10—C13	118.2 (2)	O6—C27—H27B	109.5
C9—C10—C13	120.9 (2)	H27A—C27—H27B	109.5
C10—C11—C12	118.1 (2)	O6—C27—H27C	109.5
C10—C11—H11	120.9	H27A—C27—H27C	109.5
C12—C11—H11	120.9	H27B—C27—H27C	109.5
C11—C12—C7	122.1 (2)	O7—C28—H28A	109.5
C11—C12—N1	129.6 (2)	O7—C28—H28B	109.5
C7—C12—N1	108.3 (2)	H28A—C28—H28B	109.5
C14—C13—C18	116.7 (2)	O7—C28—H28C	109.5
C14—C13—C10	120.6 (2)	H28A—C28—H28C	109.5
C18—C13—C10	122.6 (2)	H28B—C28—H28C	109.5
C15—C14—C13	121.5 (2)	C12—N1—C1	107.8 (2)
C15—C14—H14	119.3	C12—N1—S1	123.94 (18)
C13—C14—H14	119.3	C1—N1—S1	124.68 (19)
O6—C15—C14	125.2 (2)	O3—N2—O4	123.5 (3)
O6—C15—C16	114.7 (2)	O3—N2—C18	118.7 (3)
C14—C15—C16	120.0 (2)	O4—N2—C18	117.9 (2)
O7—C16—C17	125.4 (2)	C8—O5—C25	114.8 (2)
O7—C16—C15	115.4 (2)	C15—O6—C27	117.9 (2)
C17—C16—C15	119.2 (2)	C16—O7—C28	117.4 (2)
C16—C17—C18	119.5 (2)	O1—S1—O2	119.83 (15)
C16—C17—H17	120.3	O1—S1—N1	106.05 (14)
C18—C17—H17	120.3	O2—S1—N1	106.06 (12)
C17—C18—C13	123.1 (2)	O1—S1—C19	109.79 (15)
C17—C18—N2	116.1 (2)	O2—S1—C19	109.09 (15)
C13—C18—N2	120.8 (2)	N1—S1—C19	104.91 (12)
C6—C1—C2—C3	0.1 (4)	C15—C16—C17—C18	-1.0 (4)
N1—C1—C2—C3	-179.8 (3)	C16—C17—C18—C13	1.4 (4)
C1—C2—C3—C4	1.0 (5)	C16—C17—C18—N2	-175.9 (2)
C2—C3—C4—C5	-1.3 (5)	C14—C13—C18—C17	-0.6 (4)
C3—C4—C5—C6	0.4 (4)	C10—C13—C18—C17	175.2 (2)
C4—C5—C6—C1	0.6 (4)	C14—C13—C18—N2	176.7 (2)
C4—C5—C6—C7	-179.2 (3)	C10—C13—C18—N2	-7.6 (4)
C2—C1—C6—C5	-0.9 (4)	C24—C19—C20—C21	0.0 (5)
N1—C1—C6—C5	179.0 (2)	S1—C19—C20—C21	-178.1 (3)
C2—C1—C6—C7	178.9 (2)	C19—C20—C21—C22	0.6 (5)
N1—C1—C6—C7	-1.1 (3)	C20—C21—C22—C23	-1.3 (6)
C5—C6—C7—C8	2.3 (4)	C21—C22—C23—C24	1.4 (6)
C1—C6—C7—C8	-177.5 (2)	C22—C23—C24—C19	-0.8 (6)
C5—C6—C7—C12	178.9 (3)	C20—C19—C24—C23	0.1 (5)
C1—C6—C7—C12	-0.9 (3)	S1—C19—C24—C23	178.2 (3)
C12—C7—C8—O5	-176.0 (2)	C8—C9—C26—O8	-48.4 (6)
C6—C7—C8—O5	0.3 (4)	C10—C9—C26—O8	134.7 (4)
C12—C7—C8—C9	-2.1 (3)	C11—C12—N1—C1	176.3 (2)
C6—C7—C8—C9	174.2 (2)	C7—C12—N1—C1	-3.2 (3)

O5—C8—C9—C10	175.3 (2)	C11—C12—N1—S1	16.8 (4)
C7—C8—C9—C10	1.5 (4)	C7—C12—N1—S1	-162.79 (18)
O5—C8—C9—C26	-1.5 (4)	C2—C1—N1—C12	-177.3 (3)
C7—C8—C9—C26	-175.3 (3)	C6—C1—N1—C12	2.7 (3)
C8—C9—C10—C11	1.2 (4)	C2—C1—N1—S1	-18.0 (4)
C26—C9—C10—C11	178.1 (3)	C6—C1—N1—S1	162.05 (18)
C8—C9—C10—C13	179.1 (2)	C17—C18—N2—O3	137.1 (3)
C26—C9—C10—C13	-4.0 (4)	C13—C18—N2—O3	-40.3 (4)
C9—C10—C11—C12	-3.2 (4)	C17—C18—N2—O4	-42.0 (3)
C13—C10—C11—C12	178.8 (2)	C13—C18—N2—O4	140.5 (3)
C10—C11—C12—C7	2.6 (3)	C7—C8—O5—C25	-86.9 (3)
C10—C11—C12—N1	-176.9 (2)	C9—C8—O5—C25	99.3 (3)
C8—C7—C12—C11	0.1 (4)	C14—C15—O6—C27	6.6 (4)
C6—C7—C12—C11	-177.0 (2)	C16—C15—O6—C27	-173.2 (3)
C8—C7—C12—N1	179.7 (2)	C17—C16—O7—C28	-8.0 (4)
C6—C7—C12—N1	2.6 (3)	C15—C16—O7—C28	171.9 (3)
C11—C10—C13—C14	118.4 (3)	C12—N1—S1—O1	-167.4 (2)
C9—C10—C13—C14	-59.5 (3)	C1—N1—S1—O1	36.5 (3)
C11—C10—C13—C18	-57.2 (3)	C12—N1—S1—O2	-38.9 (2)
C9—C10—C13—C18	124.9 (3)	C1—N1—S1—O2	164.9 (2)
C18—C13—C14—C15	-0.7 (4)	C12—N1—S1—C19	76.5 (2)
C10—C13—C14—C15	-176.5 (2)	C1—N1—S1—C19	-79.7 (2)
C13—C14—C15—O6	-178.7 (2)	C20—C19—S1—O1	-24.5 (3)
C13—C14—C15—C16	1.1 (4)	C24—C19—S1—O1	157.3 (2)
O6—C15—C16—O7	-0.4 (3)	C20—C19—S1—O2	-157.6 (2)
C14—C15—C16—O7	179.8 (2)	C24—C19—S1—O2	24.2 (3)
O6—C15—C16—C17	179.5 (2)	C20—C19—S1—N1	89.1 (3)
C14—C15—C16—C17	-0.3 (4)	C24—C19—S1—N1	-89.1 (3)
O7—C16—C17—C18	178.9 (2)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C2—H2 \cdots O1	0.93	2.35	2.942 (5)	122
C11—H11 \cdots O2	0.93	2.32	2.927 (3)	122
C25—H25B \cdots O8	0.96	2.60	3.194 (5)	120
C17—H17 \cdots O7 ⁱ	0.93	2.56	3.384 (3)	148
C27—H27B \cdots O3 ⁱⁱ	0.96	2.56	3.239 (5)	128
C27—H27B \cdots O8 ⁱⁱⁱ	0.96	2.58	2.943 (5)	103
C25—H25A \cdots Cg1 ^{iv}	0.96	2.97	3.675 (4)	131

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