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3-Benzyl-7-methoxy-9-phenyl-2-tosyl-2,3,3a,4,9,9a-hexahydro-1*H*-pyrrolo-[3,4-*b*]quinoline

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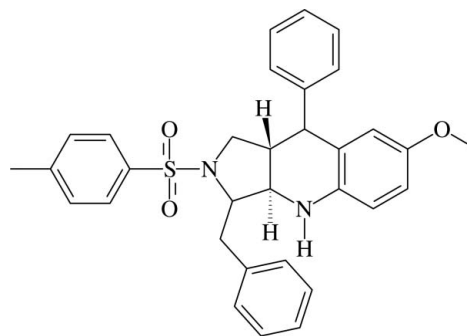
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.054; wR factor = 0.155; data-to-parameter ratio = 17.4.

In the title compound, $\text{C}_{32}\text{H}_{32}\text{N}_2\text{O}_3\text{S}$, the pyrrolidine ring adopts an envelope conformation with the methine C atom nearest to the phenyl ring as the flap atom. The tetrahydropyridine ring has a half-chair conformation. The two rings are *trans*-fused. The phenyl ring bound to the tetrahydropyridine is oriented almost perpendicular [dihedral angle = 86.35 (10°)] to the fused benzene ring. The dihedral angle between the benzylphenyl ring and the sulfonyl-bound phenyl ring is 69.43 (10°). A very weak $\text{N}-\text{H}\cdots\pi$ interaction is observed in the molecular structure. In the crystal, molecules translated one unit along the b axis are linked into $C(10)$ chains by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds; adjacent chains are linked *via* $\text{C}-\text{H}\cdots\pi$ interactions, forming a two-dimensional network parallel to the bc plane.

Related literature

For biological activity of pyrroloquinoline derivatives, see: Ryu *et al.* (2009); Tsuji *et al.* (1995); Ferlin *et al.* (2001). For related structures, see: Sudha *et al.* (2007, 2008*a,b*). For ring puckering parameters, see: Cremer & Pople (1975). For asymmetry parameters, see: Duax *et al.* (1976).



Experimental

Crystal data

 $\text{C}_{32}\text{H}_{32}\text{N}_2\text{O}_3\text{S}$ $M_r = 524.66$ Monoclinic, $P2_1/c$ $a = 21.5063$ (9) Å $b = 11.6188$ (5) Å $c = 10.7616$ (4) Å $\beta = 98.219$ (2°) $V = 2661.46$ (19) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.16$ mm⁻¹ $T = 100$ K $0.32 \times 0.30 \times 0.08$ mm

Data collection

Bruker SMART APEXII CCD

area-detector diffractometer

Absorption correction: multi-scan

(SADABS; Bruker, 2005)

 $T_{\min} = 0.636$, $T_{\max} = 0.987$

28181 measured reflections

6088 independent reflections

4661 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.057$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.054$ $wR(F^2) = 0.155$ $S = 1.02$

6088 reflections

349 parameters

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\text{max}} = 0.46$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.54$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}30-\text{H}30\cdots\text{O}3^{\text{i}}$	0.93	2.53	3.196 (3)	128
$\text{C}24-\text{H}24\cdots\text{C}g1^{\text{ii}}$	0.93	2.56	3.476 (2)	169
$\text{N}2-\text{H}1\text{N}2\cdots\text{C}g2$	0.88 (2)	3.06	3.837 (2)	147

Symmetry codes: (i) $x, y + 1, z$; (ii) $x, -y - \frac{3}{2}, z - \frac{1}{2}$. $\text{C}g1$ and $\text{C}g2$ are the centroids of the $\text{C}4-\text{C}9$ and $\text{C}26-\text{C}31$ rings, respectively.

Data collection: APEX2 (Bruker, 2005); cell refinement: SAINT (Bruker, 2005); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2940).

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supporting information

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3-Benzyl-7-methoxy-9-phenyl-2-tosyl-2,3,3a,4,9,9a-hexahydro-1*H*-pyrrolo[3,4-*b*]quinoline

K. Chinnakali, D. Sudha, M. Jayagobi, R. Raghunathan and Hoong-Kun Fun

S1. Comment

Pyrroloquinoline compounds exhibit antifungal (Ryu *et al.*, 2009), antibacterial (Tsuji *et al.*, 1995) and antiproliferative (Ferlin *et al.*, 2001) activities. We report here the crystal structure of the title compound, a pyrrolo[3,4-*b*]quinoline derivative.

The pyrrolidine ring adopts an envelope conformation with C2 as the flap atom. Atom C2 deviates by 0.663 (3) Å from the plane passing through the other four atoms of the ring (r.m.s. deviation 0.020 Å). The asymmetry parameter (Duax *et al.*, 1976) $\Delta C_s[C2] = 5.4 (2)^\circ$ and the puckering parameters (Cremer & Pople, 1975) $q_2 = 0.439 (2) \text{ \AA}$ and $\varphi = 78.2 (3)^\circ$. The tosyl group is attached to the pyrrolidine ring in a biaxial position. The tetrahydropyridine ring adopts a half-chair conformation with an asymmetry parameter $\Delta C_2[C2-C10]$ of $7.6 (2)^\circ$. The phenyl group attached to the tetrahydropyridine ring is also in a biaxial position. The dihedral angle between the C4—C9 and C19—C24 rings is $86.35 (10)^\circ$ and that between the C12—C17 and C26—C31 rings is $69.43 (10)^\circ$. A very weak N—H $\cdots\pi$ interaction (Table 1) is observed in the molecular structure. Bond lengths and angles are comparable with those observed in related structures (Sudha *et al.*, 2007,2008a,b).

In the crystal structure, molecules translated one unit along the *b* axis are linked into C(10) chains by C—H \cdots O hydrogen bonds. Glide-related molecules in adjacent chains are linked *via* C—H $\cdots\pi$ interactions involving the C4—C9 ring, forming a two-dimensional network parallel to the *bc* plane (Fig. 2).

S2. Experimental

InCl₃ (20 mol%) was added to a mixture of 2-(*N*-cinnamyl-*N*-tosylamino)-3-phenyl propanal (1 mmol) and *p*-methoxy aniline (1 mmol) in acetonitrile (20 ml). The reaction mixture was stirred at room temperature for 1 min. On completion of the reaction, as indicated by TLC, the mixture was quenched with water and extracted with ethyl acetate. The organic layer was washed with brine and dried over Na₂SO₄. The solvent was evaporated *in vacuo* and the crude product was chromatographed on silica gel using a hexane-ethyl acetate (8.5:1.5 *v/v*) mixture to obtain the title compound. The compound was recrystallized from ethyl acetate solution by slow evaporation.

S3. Refinement

The N-bound H atom was located in a difference map and refined freely [N—H = 0.88 (2) Å]. The remaining H atoms were positioned geometrically (C—H = 0.93–0.98 Å) and allowed to ride on their parent atoms, with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C_{methyl})$. A rotating group model was used for methyl groups. Reflection 100 was partially obscured by the beam stop and was omitted.

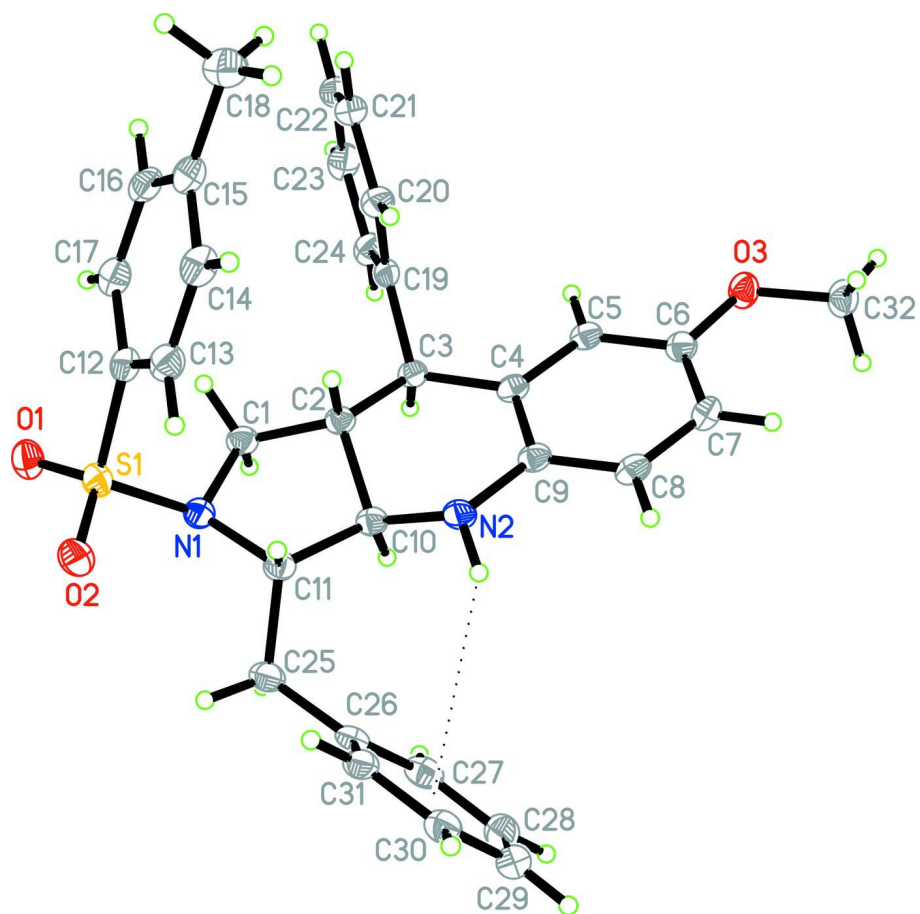


Figure 1

The molecular structure of the title compound. Displacement ellipsoids are drawn at the 50% probability level. The dotted line indicates an N—H \cdots π interaction.

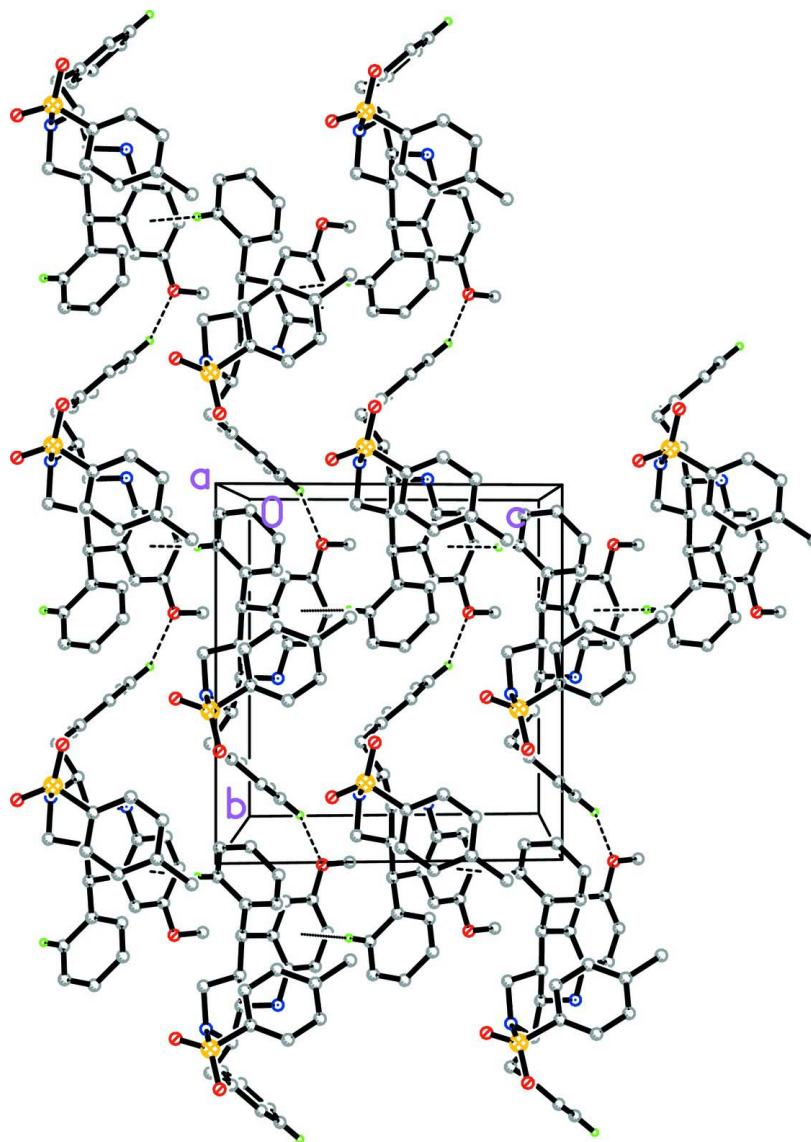


Figure 2

Crystal packing of the title compound. C—H...O hydrogen bonds and C—H... π interactions are shown as dashed lines. For the sake of clarity, H atoms not involved in the interactions have been omitted.

3-Benzyl-7-methoxy-9-phenyl-2-tosyl-2,3,3a,4,9,9a-hexahydro-1H-pyrrolo[3,4-b]quinoline

Crystal data

$C_{32}H_{32}N_2O_3S$

$M_r = 524.66$

Monoclinic, $P2_1/c$

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

$a = 21.5063\ (9)\ \text{\AA}$

$b = 11.6188\ (5)\ \text{\AA}$

$c = 10.7616\ (4)\ \text{\AA}$

$\beta = 98.219\ (2)^\circ$

$V = 2661.46\ (19)\ \text{\AA}^3$

$Z = 4$

$F(000) = 1112$

$D_x = 1.309\ \text{Mg m}^{-3}$

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

Cell parameters from 6305 reflections

$\theta = 2.6\text{--}28.5^\circ$

$\mu = 0.16\ \text{mm}^{-1}$

$T = 100\ \text{K}$

Plate, colourless

$0.32 \times 0.30 \times 0.08\ \text{mm}$

Data collection

Bruker SMART APEXII CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2005)
 $T_{\min} = 0.636$, $T_{\max} = 0.987$

28181 measured reflections
6088 independent reflections
4661 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.057$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.0^\circ$
 $h = -23 \rightarrow 27$
 $k = -15 \rightarrow 14$
 $l = -13 \rightarrow 13$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.054$
 $wR(F^2) = 0.155$
 $S = 1.02$
6088 reflections
349 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 atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0891P)^2 + 0.9992P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.46 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.54 \text{ e } \text{\AA}^{-3}$

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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.36811 (2)	0.64135 (4)	-0.09053 (5)	0.02080 (15)
O1	0.39510 (7)	0.60291 (13)	-0.19725 (13)	0.0264 (3)
O2	0.36734 (7)	0.76149 (12)	-0.06064 (14)	0.0266 (3)
O3	0.07988 (7)	0.14332 (12)	0.25445 (14)	0.0235 (3)
N1	0.29441 (8)	0.59952 (14)	-0.11262 (15)	0.0197 (4)
N2	0.18416 (8)	0.55576 (14)	0.11176 (16)	0.0202 (4)
H1N2	0.1626 (11)	0.619 (2)	0.122 (2)	0.024 (6)*
C1	0.28306 (10)	0.47333 (16)	-0.12239 (18)	0.0195 (4)
H1A	0.3222	0.4312	-0.1194	0.023*
H1B	0.2557	0.4539	-0.1991	0.023*
C2	0.25163 (9)	0.44865 (15)	-0.00752 (17)	0.0172 (4)
H2	0.2834	0.4530	0.0673	0.021*
C3	0.21612 (9)	0.33569 (15)	-0.00428 (17)	0.0169 (4)
H3	0.1899	0.3268	-0.0860	0.020*
C4	0.17183 (9)	0.34490 (16)	0.09480 (17)	0.0176 (4)

C5	0.14457 (9)	0.24516 (16)	0.13452 (18)	0.0182 (4)
H5	0.1546	0.1747	0.1016	0.022*
C6	0.10267 (9)	0.24795 (16)	0.22227 (18)	0.0194 (4)
C7	0.08772 (10)	0.35345 (17)	0.27171 (19)	0.0213 (4)
H7	0.0602	0.3569	0.3308	0.026*
C8	0.11424 (10)	0.45329 (17)	0.23202 (19)	0.0213 (4)
H8	0.1037	0.5236	0.2648	0.026*
C9	0.15627 (9)	0.45154 (16)	0.14423 (18)	0.0181 (4)
C10	0.20811 (9)	0.55138 (16)	-0.00791 (18)	0.0179 (4)
H10	0.1731	0.5435	-0.0764	0.022*
C11	0.24967 (10)	0.65334 (16)	-0.03459 (18)	0.0192 (4)
H11	0.2727	0.6828	0.0441	0.023*
C12	0.40517 (9)	0.56541 (17)	0.04217 (18)	0.0206 (4)
C13	0.40440 (10)	0.60984 (18)	0.16170 (19)	0.0231 (4)
H13	0.3878	0.6825	0.1723	0.028*
C14	0.42880 (10)	0.54439 (18)	0.2653 (2)	0.0249 (5)
H14	0.4288	0.5744	0.3454	0.030*
C15	0.45327 (10)	0.43488 (19)	0.2518 (2)	0.0251 (5)
C16	0.45382 (10)	0.39268 (18)	0.1305 (2)	0.0238 (4)
H16	0.4702	0.3199	0.1197	0.029*
C17	0.43044 (10)	0.45702 (17)	0.0261 (2)	0.0233 (4)
H17	0.4316	0.4281	-0.0541	0.028*
C18	0.47876 (12)	0.3636 (2)	0.3642 (2)	0.0334 (5)
H18A	0.4656	0.3963	0.4382	0.050*
H18B	0.5238	0.3627	0.3732	0.050*
H18C	0.4631	0.2864	0.3531	0.050*
C19	0.26098 (9)	0.23397 (16)	0.01226 (18)	0.0177 (4)
C20	0.30745 (10)	0.22449 (17)	0.11657 (19)	0.0220 (4)
H20	0.3105	0.2803	0.1791	0.026*
C21	0.34910 (10)	0.13259 (18)	0.1278 (2)	0.0254 (5)
H21	0.3800	0.1277	0.1974	0.030*
C22	0.34495 (11)	0.04784 (17)	0.0358 (2)	0.0268 (5)
H22	0.3725	-0.0143	0.0441	0.032*
C23	0.29953 (11)	0.05698 (17)	-0.0680 (2)	0.0269 (5)
H23	0.2968	0.0013	-0.1306	0.032*
C24	0.25761 (10)	0.14916 (17)	-0.0797 (2)	0.0223 (4)
H24	0.2270	0.1540	-0.1499	0.027*
C25	0.21481 (10)	0.75164 (17)	-0.10939 (19)	0.0227 (4)
H25A	0.2447	0.8112	-0.1230	0.027*
H25B	0.1962	0.7229	-0.1909	0.027*
C26	0.16400 (10)	0.80340 (16)	-0.04408 (18)	0.0202 (4)
C27	0.10099 (10)	0.77711 (17)	-0.0814 (2)	0.0249 (5)
H27	0.0900	0.7246	-0.1461	0.030*
C28	0.05413 (11)	0.82820 (19)	-0.0235 (2)	0.0276 (5)
H28	0.0122	0.8108	-0.0503	0.033*
C29	0.07005 (11)	0.90497 (18)	0.0741 (2)	0.0275 (5)
H29	0.0388	0.9400	0.1124	0.033*
C30	0.13279 (10)	0.92955 (17)	0.1147 (2)	0.0237 (4)

H30	0.1437	0.9797	0.1817	0.028*
C31	0.17939 (10)	0.87961 (16)	0.05583 (19)	0.0213 (4)
H31	0.2213	0.8971	0.0832	0.026*
C32	0.03712 (10)	0.14275 (18)	0.3447 (2)	0.0255 (5)
H32A	0.0236	0.0653	0.3568	0.038*
H32B	0.0014	0.1897	0.3150	0.038*
H32C	0.0577	0.1728	0.4230	0.038*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0206 (3)	0.0180 (3)	0.0242 (3)	-0.00236 (19)	0.0046 (2)	0.00132 (18)
O1	0.0265 (9)	0.0282 (8)	0.0259 (8)	-0.0026 (6)	0.0091 (6)	0.0015 (6)
O2	0.0252 (9)	0.0192 (7)	0.0353 (8)	-0.0044 (6)	0.0042 (7)	0.0022 (6)
O3	0.0211 (8)	0.0197 (7)	0.0313 (8)	-0.0037 (6)	0.0089 (6)	-0.0016 (6)
N1	0.0190 (9)	0.0161 (8)	0.0245 (9)	0.0001 (7)	0.0044 (7)	0.0006 (6)
N2	0.0250 (10)	0.0124 (8)	0.0243 (9)	0.0029 (7)	0.0075 (7)	-0.0008 (6)
C1	0.0199 (11)	0.0162 (9)	0.0222 (10)	0.0009 (7)	0.0029 (8)	-0.0006 (7)
C2	0.0164 (10)	0.0145 (9)	0.0204 (9)	0.0006 (7)	0.0014 (8)	-0.0007 (7)
C3	0.0156 (10)	0.0159 (9)	0.0189 (9)	0.0004 (7)	0.0015 (7)	-0.0002 (7)
C4	0.0150 (10)	0.0182 (9)	0.0188 (9)	0.0020 (7)	-0.0007 (7)	-0.0015 (7)
C5	0.0161 (10)	0.0155 (9)	0.0223 (10)	0.0012 (7)	0.0008 (8)	-0.0015 (7)
C6	0.0154 (10)	0.0179 (9)	0.0242 (10)	-0.0018 (7)	0.0007 (8)	0.0005 (7)
C7	0.0158 (10)	0.0246 (10)	0.0245 (10)	0.0012 (8)	0.0056 (8)	-0.0012 (8)
C8	0.0189 (11)	0.0180 (10)	0.0270 (10)	0.0039 (8)	0.0030 (8)	-0.0035 (8)
C9	0.0144 (10)	0.0169 (9)	0.0222 (10)	0.0002 (7)	-0.0009 (8)	-0.0010 (7)
C10	0.0182 (10)	0.0163 (9)	0.0190 (9)	0.0006 (7)	0.0017 (8)	-0.0003 (7)
C11	0.0196 (11)	0.0159 (9)	0.0224 (10)	0.0010 (7)	0.0038 (8)	-0.0009 (7)
C12	0.0152 (10)	0.0221 (10)	0.0248 (10)	-0.0020 (8)	0.0038 (8)	0.0008 (8)
C13	0.0208 (11)	0.0205 (10)	0.0282 (11)	-0.0011 (8)	0.0045 (8)	-0.0022 (8)
C14	0.0212 (11)	0.0296 (11)	0.0239 (10)	-0.0007 (9)	0.0035 (8)	-0.0036 (8)
C15	0.0151 (11)	0.0296 (11)	0.0301 (11)	-0.0011 (8)	0.0022 (8)	0.0031 (9)
C16	0.0151 (10)	0.0226 (10)	0.0336 (11)	0.0003 (8)	0.0035 (8)	-0.0014 (8)
C17	0.0205 (11)	0.0236 (10)	0.0266 (11)	-0.0016 (8)	0.0058 (8)	-0.0043 (8)
C18	0.0320 (14)	0.0355 (13)	0.0325 (12)	0.0057 (10)	0.0036 (10)	0.0063 (10)
C19	0.0160 (10)	0.0147 (9)	0.0235 (10)	-0.0002 (7)	0.0068 (8)	0.0028 (7)
C20	0.0225 (11)	0.0212 (10)	0.0228 (10)	0.0027 (8)	0.0046 (8)	0.0006 (8)
C21	0.0212 (11)	0.0280 (11)	0.0277 (11)	0.0056 (9)	0.0058 (9)	0.0086 (8)
C22	0.0267 (12)	0.0169 (10)	0.0399 (12)	0.0061 (8)	0.0152 (10)	0.0061 (8)
C23	0.0274 (12)	0.0169 (10)	0.0386 (12)	-0.0010 (8)	0.0127 (10)	-0.0058 (9)
C24	0.0199 (11)	0.0197 (10)	0.0277 (11)	-0.0021 (8)	0.0047 (8)	-0.0026 (8)
C25	0.0269 (12)	0.0178 (10)	0.0237 (10)	0.0034 (8)	0.0044 (9)	0.0038 (8)
C26	0.0241 (11)	0.0126 (9)	0.0238 (10)	0.0023 (7)	0.0032 (8)	0.0059 (7)
C27	0.0287 (12)	0.0176 (9)	0.0271 (11)	-0.0024 (8)	-0.0006 (9)	0.0027 (8)
C28	0.0189 (11)	0.0301 (11)	0.0331 (12)	-0.0022 (9)	0.0011 (9)	0.0079 (9)
C29	0.0248 (12)	0.0253 (11)	0.0342 (12)	0.0049 (9)	0.0099 (9)	0.0075 (9)
C30	0.0280 (12)	0.0173 (9)	0.0264 (10)	0.0005 (8)	0.0057 (9)	0.0021 (8)
C31	0.0193 (11)	0.0158 (9)	0.0282 (11)	0.0005 (8)	0.0015 (8)	0.0038 (8)

C32	0.0210 (11)	0.0261 (11)	0.0307 (11)	-0.0041 (8)	0.0080 (9)	0.0006 (9)
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Geometric parameters (Å, °)

S1—O1	1.4303 (15)	C14—H14	0.93
S1—O2	1.4332 (15)	C15—C16	1.396 (3)
S1—N1	1.6424 (18)	C15—C18	1.504 (3)
S1—C12	1.768 (2)	C16—C17	1.383 (3)
O3—C6	1.374 (2)	C16—H16	0.93
O3—C32	1.430 (2)	C17—H17	0.93
N1—C1	1.488 (2)	C18—H18A	0.96
N1—C11	1.501 (2)	C18—H18B	0.96
N2—C9	1.417 (2)	C18—H18C	0.96
N2—C10	1.455 (2)	C19—C24	1.391 (3)
N2—H1N2	0.88 (2)	C19—C20	1.396 (3)
C1—C2	1.519 (3)	C20—C21	1.388 (3)
C1—H1A	0.97	C20—H20	0.93
C1—H1B	0.97	C21—C22	1.390 (3)
C2—C10	1.516 (3)	C21—H21	0.93
C2—C3	1.521 (3)	C22—C23	1.379 (3)
C2—H2	0.98	C22—H22	0.93
C3—C19	1.520 (3)	C23—C24	1.394 (3)
C3—C4	1.532 (3)	C23—H23	0.93
C3—H3	0.98	C24—H24	0.93
C4—C5	1.393 (3)	C25—C26	1.506 (3)
C4—C9	1.407 (3)	C25—H25A	0.97
C5—C6	1.396 (3)	C25—H25B	0.97
C5—H5	0.93	C26—C27	1.391 (3)
C6—C7	1.392 (3)	C26—C31	1.396 (3)
C7—C8	1.387 (3)	C27—C28	1.392 (3)
C7—H7	0.93	C27—H27	0.93
C8—C9	1.398 (3)	C28—C29	1.383 (3)
C8—H8	0.93	C28—H28	0.93
C10—C11	1.536 (3)	C29—C30	1.387 (3)
C10—H10	0.98	C29—H29	0.93
C11—C25	1.530 (3)	C30—C31	1.387 (3)
C11—H11	0.98	C30—H30	0.93
C12—C13	1.388 (3)	C31—H31	0.93
C12—C17	1.392 (3)	C32—H32A	0.96
C13—C14	1.389 (3)	C32—H32B	0.96
C13—H13	0.93	C32—H32C	0.96
C14—C15	1.392 (3)		
O1—S1—O2	120.22 (9)	C13—C14—C15	121.46 (19)
O1—S1—N1	106.58 (9)	C13—C14—H14	119.3
O2—S1—N1	106.18 (9)	C15—C14—H14	119.3
O1—S1—C12	107.76 (9)	C14—C15—C16	118.19 (19)
O2—S1—C12	108.85 (9)	C14—C15—C18	121.25 (19)

N1—S1—C12	106.47 (9)	C16—C15—C18	120.6 (2)
C6—O3—C32	117.56 (15)	C17—C16—C15	121.27 (19)
C1—N1—C11	109.76 (15)	C17—C16—H16	119.4
C1—N1—S1	116.56 (13)	C15—C16—H16	119.4
C11—N1—S1	118.58 (13)	C16—C17—C12	119.42 (19)
C9—N2—C10	113.85 (15)	C16—C17—H17	120.3
C9—N2—H1N2	115.6 (15)	C12—C17—H17	120.3
C10—N2—H1N2	113.5 (15)	C15—C18—H18A	109.5
N1—C1—C2	102.50 (15)	C15—C18—H18B	109.5
N1—C1—H1A	111.3	H18A—C18—H18B	109.5
C2—C1—H1A	111.3	C15—C18—H18C	109.5
N1—C1—H1B	111.3	H18A—C18—H18C	109.5
C2—C1—H1B	111.3	H18B—C18—H18C	109.5
H1A—C1—H1B	109.2	C24—C19—C20	118.32 (18)
C10—C2—C1	101.27 (15)	C24—C19—C3	120.00 (18)
C10—C2—C3	111.56 (16)	C20—C19—C3	121.65 (17)
C1—C2—C3	117.70 (15)	C21—C20—C19	120.63 (19)
C10—C2—H2	108.6	C21—C20—H20	119.7
C1—C2—H2	108.6	C19—C20—H20	119.7
C3—C2—H2	108.6	C20—C21—C22	120.5 (2)
C19—C3—C2	111.18 (16)	C20—C21—H21	119.7
C19—C3—C4	114.85 (15)	C22—C21—H21	119.7
C2—C3—C4	108.65 (15)	C23—C22—C21	119.24 (19)
C19—C3—H3	107.3	C23—C22—H22	120.4
C2—C3—H3	107.3	C21—C22—H22	120.4
C4—C3—H3	107.3	C22—C23—C24	120.41 (19)
C5—C4—C9	118.88 (18)	C22—C23—H23	119.8
C5—C4—C3	119.10 (16)	C24—C23—H23	119.8
C9—C4—C3	121.99 (17)	C19—C24—C23	120.9 (2)
C4—C5—C6	121.94 (17)	C19—C24—H24	119.6
C4—C5—H5	119.0	C23—C24—H24	119.6
C6—C5—H5	119.0	C26—C25—C11	112.71 (16)
O3—C6—C7	124.94 (18)	C26—C25—H25A	109.1
O3—C6—C5	115.99 (17)	C11—C25—H25A	109.1
C7—C6—C5	119.07 (18)	C26—C25—H25B	109.1
C8—C7—C6	119.42 (18)	C11—C25—H25B	109.1
C8—C7—H7	120.3	H25A—C25—H25B	107.8
C6—C7—H7	120.3	C27—C26—C31	118.40 (19)
C7—C8—C9	122.02 (18)	C27—C26—C25	121.28 (19)
C7—C8—H8	119.0	C31—C26—C25	120.32 (19)
C9—C8—H8	119.0	C26—C27—C28	121.0 (2)
C8—C9—C4	118.68 (17)	C26—C27—H27	119.5
C8—C9—N2	119.44 (17)	C28—C27—H27	119.5
C4—C9—N2	121.81 (17)	C29—C28—C27	119.9 (2)
N2—C10—C2	108.76 (15)	C29—C28—H28	120.0
N2—C10—C11	115.32 (16)	C27—C28—H28	120.0
C2—C10—C11	103.43 (16)	C28—C29—C30	119.7 (2)
N2—C10—H10	109.7	C28—C29—H29	120.1

C2—C10—H10	109.7	C30—C29—H29	120.1
C11—C10—H10	109.7	C31—C30—C29	120.2 (2)
N1—C11—C25	108.76 (15)	C31—C30—H30	119.9
N1—C11—C10	102.71 (14)	C29—C30—H30	119.9
C25—C11—C10	114.82 (17)	C30—C31—C26	120.7 (2)
N1—C11—H11	110.1	C30—C31—H31	119.7
C25—C11—H11	110.1	C26—C31—H31	119.7
C10—C11—H11	110.1	O3—C32—H32A	109.5
C13—C12—C17	120.53 (19)	O3—C32—H32B	109.5
C13—C12—S1	119.93 (16)	H32A—C32—H32B	109.5
C17—C12—S1	119.33 (15)	O3—C32—H32C	109.5
C12—C13—C14	119.12 (19)	H32A—C32—H32C	109.5
C12—C13—H13	120.4	H32B—C32—H32C	109.5
C14—C13—H13	120.4		
O1—S1—N1—C1	61.38 (15)	N2—C10—C11—N1	148.55 (16)
O2—S1—N1—C1	-169.33 (13)	C2—C10—C11—N1	29.94 (19)
C12—S1—N1—C1	-53.44 (16)	N2—C10—C11—C25	-93.6 (2)
O1—S1—N1—C11	-163.96 (13)	C2—C10—C11—C25	147.81 (16)
O2—S1—N1—C11	-34.67 (16)	O1—S1—C12—C13	158.49 (16)
C12—S1—N1—C11	81.22 (15)	O2—S1—C12—C13	26.59 (19)
C11—N1—C1—C2	-23.1 (2)	N1—S1—C12—C13	-87.49 (18)
S1—N1—C1—C2	115.35 (15)	O1—S1—C12—C17	-26.68 (19)
N1—C1—C2—C10	40.98 (18)	O2—S1—C12—C17	-158.58 (16)
N1—C1—C2—C3	162.85 (16)	N1—S1—C12—C17	87.34 (18)
C10—C2—C3—C19	-173.02 (15)	C17—C12—C13—C14	-0.4 (3)
C1—C2—C3—C19	70.6 (2)	S1—C12—C13—C14	174.34 (16)
C10—C2—C3—C4	-45.7 (2)	C12—C13—C14—C15	-0.8 (3)
C1—C2—C3—C4	-162.10 (16)	C13—C14—C15—C16	1.2 (3)
C19—C3—C4—C5	-40.9 (2)	C13—C14—C15—C18	-179.1 (2)
C2—C3—C4—C5	-166.12 (17)	C14—C15—C16—C17	-0.4 (3)
C19—C3—C4—C9	141.09 (18)	C18—C15—C16—C17	179.9 (2)
C2—C3—C4—C9	15.9 (2)	C15—C16—C17—C12	-0.8 (3)
C9—C4—C5—C6	-0.4 (3)	C13—C12—C17—C16	1.2 (3)
C3—C4—C5—C6	-178.46 (18)	S1—C12—C17—C16	-173.61 (16)
C32—O3—C6—C7	0.6 (3)	C2—C3—C19—C24	-118.83 (19)
C32—O3—C6—C5	179.61 (18)	C4—C3—C19—C24	117.29 (19)
C4—C5—C6—O3	-179.17 (17)	C2—C3—C19—C20	59.2 (2)
C4—C5—C6—C7	-0.1 (3)	C4—C3—C19—C20	-64.7 (2)
O3—C6—C7—C8	179.58 (19)	C24—C19—C20—C21	0.0 (3)
C5—C6—C7—C8	0.6 (3)	C3—C19—C20—C21	-178.04 (18)
C6—C7—C8—C9	-0.6 (3)	C19—C20—C21—C22	-0.5 (3)
C7—C8—C9—C4	0.1 (3)	C20—C21—C22—C23	1.0 (3)
C7—C8—C9—N2	-177.01 (19)	C21—C22—C23—C24	-1.0 (3)
C5—C4—C9—C8	0.4 (3)	C20—C19—C24—C23	0.0 (3)
C3—C4—C9—C8	178.37 (17)	C3—C19—C24—C23	178.09 (18)
C5—C4—C9—N2	177.43 (18)	C22—C23—C24—C19	0.5 (3)
C3—C4—C9—N2	-4.6 (3)	N1—C11—C25—C26	174.16 (16)

C10—N2—C9—C8	-159.30 (18)	C10—C11—C25—C26	59.8 (2)
C10—N2—C9—C4	23.6 (3)	C11—C25—C26—C27	-102.6 (2)
C9—N2—C10—C2	-53.1 (2)	C11—C25—C26—C31	77.6 (2)
C9—N2—C10—C11	-168.66 (16)	C31—C26—C27—C28	2.0 (3)
C1—C2—C10—N2	-167.42 (15)	C25—C26—C27—C28	-177.81 (18)
C3—C2—C10—N2	66.5 (2)	C26—C27—C28—C29	-1.1 (3)
C1—C2—C10—C11	-44.36 (18)	C27—C28—C29—C30	-0.8 (3)
C3—C2—C10—C11	-170.41 (15)	C28—C29—C30—C31	1.6 (3)
C1—N1—C11—C25	-126.22 (17)	C29—C30—C31—C26	-0.5 (3)
S1—N1—C11—C25	96.32 (18)	C27—C26—C31—C30	-1.2 (3)
C1—N1—C11—C10	-4.1 (2)	C25—C26—C31—C30	178.62 (18)
S1—N1—C11—C10	-141.61 (14)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
C30—H30 \cdots O3 ⁱ	0.93	2.53	3.196 (3)	128
C24—H24 \cdots Cg1 ⁱⁱ	0.93	2.56	3.476 (2)	169
N2—H1N2 \cdots Cg2	0.88 (2)	3.06	3.837 (2)	147

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