

4,4,7,7-Tetramethyl-2-tosyl-2,3,3a,4,6,7,8,9-octahydro-1*H*-pyrrolo[3,4-*c*]pyrano[6,5-*b*]cyclohexan-9-one

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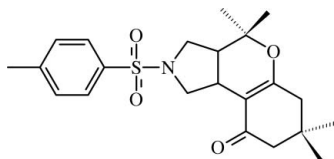
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.046; wR factor = 0.128; data-to-parameter ratio = 26.7.

The title compound, $\text{C}_{22}\text{H}_{29}\text{NO}_4\text{S}$, crystallizes with three independent molecules, *A*, *B* and *C*, in the asymmetric unit. Each of the three independent molecules adopts a folded conformation, with the phenylsulfonyl group lying over the pyranocyclohexanone ring system, and with the pyrrolidine and dihydropyran rings *cis*-fused. In all three molecules, the pyrrolidine ring has a twist conformation and the dihydropyran ring is in a half-chair conformation. The cyclohexenone rings adopt envelope conformations, with the flap atom puckered towards the phenylsulfonyl group in molecule *A*, and away from that group in molecules *B* and *C*, resulting in a different overall conformation for molecule *A* compared to *B* and *C*. In all molecules, the tosyl group is equatorially attached to the pyrrolidine ring. In the crystal structure, the molecules are linked into a three-dimensional framework by $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For related pyrrolo[3,4-*c*]pyran structures, see: Chinnakali *et al.* (2007*a,b*). For biological activities of pyrrolo[3,4-*c*]pyran derivatives, see: Millan *et al.* (2000). For ring puckering parameters, see: Cremer & Pople (1975). For asymmetry parameters, see: Duax *et al.* (1976).



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Experimental

Crystal data

$\text{C}_{22}\text{H}_{29}\text{NO}_4\text{S}$
 $M_r = 403.53$
 Triclinic, $P\bar{1}$
 $a = 14.5021$ (7) Å
 $b = 15.0787$ (7) Å
 $c = 16.9696$ (8) Å
 $\alpha = 64.059$ (2)°
 $\beta = 69.821$ (2)°
 $\gamma = 71.636$ (3)°
 $V = 3072.3$ (3) Å³
 $Z = 6$
 Mo $K\alpha$ radiation
 $\mu = 0.19$ mm⁻¹
 $T = 100.0$ (1) K
 $0.54 \times 0.39 \times 0.20$ mm

Data collection

Bruker SMART APEXII CCD area-detector diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
 $T_{\min} = 0.894$, $T_{\max} = 0.963$
 64276 measured reflections
 20257 independent reflections
 15293 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.042$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.128$
 $S = 1.06$
 20257 reflections
 760 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.51$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| <i>D</i> — <i>H</i> ⋯ <i>A</i> | <i>D</i> — <i>H</i> | <i>H</i> ⋯ <i>A</i> | <i>D</i> ⋯ <i>A</i> | <i>D</i> — <i>H</i> ⋯ <i>A</i> |
|--------------------------------|---------------------|---------------------|---------------------|--------------------------------|
| C2A—H2A⋯O1C | 0.98 | 2.51 | 3.4447 (18) | 160 |
| C2B—H2B⋯O1B ⁱ | 0.98 | 2.52 | 3.4167 (18) | 151 |
| C2C—H2C⋯O1A | 0.98 | 2.51 | 3.4739 (18) | 167 |
| C4A—H4A⋯O4A | 0.97 | 2.48 | 2.9605 (17) | 111 |
| C4B—H4C⋯O4B | 0.97 | 2.48 | 3.0370 (18) | 116 |
| C4C—H4F⋯O4C | 0.97 | 2.43 | 3.0069 (17) | 118 |
| C12A—H12A⋯O2C ⁱⁱ | 0.93 | 2.51 | 3.2130 (16) | 133 |
| C12B—H12B⋯O2A ⁱⁱⁱ | 0.93 | 2.59 | 3.3762 (18) | 143 |
| C14C—H14I⋯O2C ^{iv} | 0.96 | 2.56 | 3.362 (2) | 142 |
| C17A—H17B⋯O4C ⁱⁱ | 0.97 | 2.51 | 3.3480 (18) | 145 |
| C17B—H17C⋯O4A ⁱⁱⁱ | 0.97 | 2.38 | 3.2776 (17) | 154 |
| C21C—H21I⋯O4B ^v | 0.96 | 2.47 | 3.3058 (17) | 146 |
| C22A—H22B⋯O4C ⁱⁱ | 0.96 | 2.55 | 3.4023 (16) | 147 |

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *APEX2*; data reduction: *SAINT* (Bruker, 2005); program(s) used to solve structure: *SHELXTL* (Sheldrick, 1998); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WN2210).

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Acta Cryst. (2007). E63, o4436-o4437 [doi:10.1107/S1600536807052221]

4,4,7,7-Tetramethyl-2-tosyl-2,3,3a,4,6,7,8,9-octahydro-1*H*-pyrrolo[3,4-*c*]pyrano[6,5-*b*]cyclohexan-9-one

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Comment

S33084, a pyrrolo[3,4-*c*]pyran derivative, is a novel, potent, selective, and competitive antagonist at dopamine D3-receptors (Millan *et al.*, 2000). Previously, we have reported the crystal structures of two pyrrolo[3,4-*c*]pyran derivatives (Chinnakali *et al.*, 2007*a,b*). Now we report here the crystal structure of the title compound.

The title compound crystallizes in space group $P\bar{1}$ with three independent molecules, A, B and C, in the asymmetric unit. The conformations of molecules B and C are the same but that of A is different. Views of the three independent molecules, with the atomic numbering schemes, are shown in Fig. 1. The weighted r.m.s. deviations for the superposition of the non-H atoms of any pair of molecules in the title compound using *XP* in *SHELXTL* (Sheldrick, 1998) are 0.675 (Å) (A/B pair), 0.629 (Å) (A/C pair) and 0.095 Å (B/C pair). Views of the superposition of molecular pairs are shown in Fig. 2. The corresponding bond lengths and angles of the three molecules agree with each other and show normal values.

In all three molecules, the pyrrolidine ring (N1/C1—C4) adopts a twist conformation. In molecules B and C, the local twofold rotation axis passes through atom N1 and the mid-point of the opposite bond C2—C3, whereas in molecule A the axis passes through atom C4 and the C1—C2 bond. The puckering parameters (Cremer & Pople, 1975) for the pyrrolidine ring in A/B/C are $q_2 = 0.395$ (1)/0.359 (2)/0.362 (2) Å and $\varphi = 54.9$ (2)/86.0 (2)/267.2 (2)°. The smallest displacement asymmetry parameters (Duax *et al.*, 1976) are $\Delta C_2[C1A—C2A] = 1.0$ (1)°, $\Delta C_2[C2B—C3B] = 4.4$ (2)° and $\Delta C_2[C2C—C3C] = 3.2$ (1)°. In each independent molecule, the N atom exhibits sp^3 hybridization and the tosyl group is equatorially attached to the pyrrolidine ring.

In each of the three independent molecules, the dihydropyran ring (O3/C5/C2/C3/C7/C6) has a half-chair conformation, with a local pseudo-twofold axis running through the midpoints of the C2—C5 and C6—C7 bonds. The puckering parameters Q , θ and φ , and the smallest displacement asymmetry parameters $\Delta C_2[C2—C5]$ for the molecules A/B/C are 0.432 (1)/0.462 (2)/0.459 (1) Å, 51.8 (2)/53.5 (2)/126.5 (2)°, 84.8 (2)/84.4 (2)/267.3 (2)° and 6.9 (2)/7.4 (2)/4.7 (2)°.

The cyclohexenone rings (C6/C7/C20/C19/C18/C17) in molecules A, B and C, adopt envelope conformations with atom C18 at the flap. The asymmetry parameter $\Delta C_5[C18]$ is 5.1 (2), 3.5 (2) and 1.1 (2)° for molecules A, B and C, respectively. The Q , θ and φ values for A/B/C are 0.460 (2)/0.452 (2)/0.438 (2) Å, 55.8 (2)/131.6 (2)/47.0 (2)° and 234.4 (2)/64.8 (2)/241.3 (3)°, respectively. The deviation of atom C18 from the C6/C7/C20/C19/C17 plane in molecule A [−0.641 (2) Å] is in the opposite direction to those in molecules B [0.627 (2) Å] and C [0.609 (2) Å]. This results in a different conformation for molecule A compared to B and C (Fig. 2).

Each of the three independent molecules adopt a folded conformation, with the phenylsulfonyl group lying over the pyranocyclohexanone ring system, and with the pyrrolidine and dihydropyran rings *cis*-fused.

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Each of the independent molecules B and C forms a centrosymmetric dimer linked by paired C—H \cdots O hydrogen bonds *viz.* C2B—H2B \cdots O1Bⁱ and C14C—H14I \cdots O2C^{iv} (symmetry codes are given in Table 1). The B molecules form a dimer centred at (1/2,1/2,0), while the C molecules form a dimer centred at (0,1/2,1/2). The B and C dimers and A molecules are linked into a three-dimensional network by a number of C—H \cdots O hydrogen bonds (Fig. 3).

Experimental

To a solution of 5,5-dimethylcyclohexane-1,3-dione (1 mmol) in dry toluene (20 ml), 2-[*N*-(3-methylbut-2-enyl)-*N*-tosylamino]acetaldehyde (1 mmol) and a catalytic amount of the base ethylenediamine-*N,N'*-diacetate (EDDA) were added and the reaction mixture was refluxed for 12 h. After completion of the reaction, the solvent was evaporated under reduced pressure and the crude product was chromatographed using a hexane-ethyl acetate (8:2 *v/v*) mixture to obtain the title compound. The compound was recrystallized from ethyl acetate solution by slow evaporation.

Refinement

All H atoms were positioned geometrically and allowed to ride on their parent atoms, with C—H = 0.93–0.98 Å. The U_{iso} values were set equal to $1.5U_{\text{eq}}$ of the carrier atom for methyl H atoms and $1.2U_{\text{eq}}$ for the remaining H atoms. A rotating group model was used for the methyl groups attached to aromatic rings.

Figures

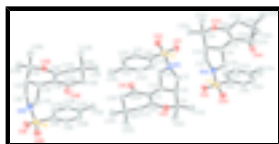


Fig. 1. The three independent molecules of the title compound, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 80% probability level. H atoms have been omitted for clarity.



Fig. 2. Fit of (a) molecule A (dashed lines) on molecule B (solid lines), (b) molecule A (dashed lines) on molecule C (solid lines), and (c) molecule B (dashed lines) on molecule C (solid lines). H atoms have been omitted for clarity.

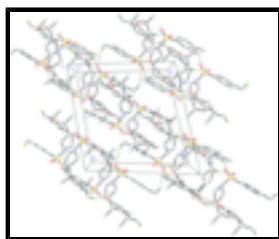


Fig. 3. View of the crystal packing of the title compound. Hydrogen bonds are shown as dashed lines. For the sake of clarity, H atoms not involved in the interactions have been omitted.

4,4,7,7-Tetramethyl-2-tosyl-2,3,3a,4,6,7,8,9-octahydro-1*H*-pyrrolo[3,4-*c*]pyrano[6,5-*b*]cyclohexan-9-one

Crystal data

C₂₂H₂₉NO₄S

$M_r = 403.53$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$Z = 6$

$F_{000} = 1296$

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

Mo $K\alpha$ radiation

$\lambda = 0.71073 \text{ \AA}$

| | |
|--------------------------------|---|
| $a = 14.5021 (7) \text{ \AA}$ | Cell parameters from 6958 reflections |
| $b = 15.0787 (7) \text{ \AA}$ | $\theta = 2.6\text{--}31.5^\circ$ |
| $c = 16.9696 (8) \text{ \AA}$ | $\mu = 0.19 \text{ mm}^{-1}$ |
| $\alpha = 64.059 (2)^\circ$ | $T = 100.0 (1) \text{ K}$ |
| $\beta = 69.821 (2)^\circ$ | Block, colourless |
| $\gamma = 71.636 (3)^\circ$ | $0.54 \times 0.39 \times 0.20 \text{ mm}$ |
| $V = 3072.3 (3) \text{ \AA}^3$ | |

Data collection

| | |
|--|---|
| Bruker SMART APEX2 CCD area-detector diffractometer | 20257 independent reflections |
| Radiation source: fine-focus sealed tube | 15293 reflections with $I > 2\sigma(I)$ |
| Monochromator: graphite | $R_{\text{int}} = 0.042$ |
| Detector resolution: $8.33 \text{ pixels mm}^{-1}$ | $\theta_{\text{max}} = 31.5^\circ$ |
| $T = 100.0(1) \text{ K}$ | $\theta_{\text{min}} = 1.4^\circ$ |
| ω scans | $h = -21 \rightarrow 21$ |
| Absorption correction: multi-scan (SADABS; Bruker, 2005) | $k = -22 \rightarrow 22$ |
| $T_{\text{min}} = 0.894, T_{\text{max}} = 0.963$ | $l = -24 \rightarrow 24$ |
| 64276 measured reflections | |

Refinement

| | |
|--|--|
| Refinement on F^2 | Secondary atom site location: difference Fourier map |
| Least-squares matrix: full | Hydrogen site location: inferred from neighbouring sites |
| $R[F^2 > 2\sigma(F^2)] = 0.046$ | H-atom parameters constrained |
| $wR(F^2) = 0.128$ | $w = 1/[\sigma^2(F_o^2) + (0.0615P)^2 + 0.432P]$ |
| $S = 1.06$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 20257 reflections | $(\Delta/\sigma)_{\text{max}} = 0.001$ |
| 760 parameters | $\Delta\rho_{\text{max}} = 0.44 \text{ e \AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | $\Delta\rho_{\text{min}} = -0.51 \text{ e \AA}^{-3}$ |
| | Extinction correction: none |

Special details

Experimental. The low-temperature data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

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 $F^2 > 2\sigma(F^2)$ is used only for calculat-

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ing 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)

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|---------------|---------------|--------------|----------------------------------|
| S1A | 0.25730 (2) | 0.00053 (2) | 0.23582 (2) | 0.01537 (7) |
| O1A | 0.28101 (7) | -0.04754 (7) | 0.32238 (7) | 0.0219 (2) |
| O2A | 0.33129 (7) | 0.04251 (7) | 0.15602 (7) | 0.0210 (2) |
| O3A | -0.14661 (7) | 0.12129 (7) | 0.33197 (6) | 0.01629 (18) |
| O4A | 0.01788 (7) | 0.21417 (7) | 0.02177 (6) | 0.01954 (19) |
| N1A | 0.16405 (8) | 0.09352 (8) | 0.24168 (7) | 0.0147 (2) |
| C1A | 0.07741 (9) | 0.07873 (9) | 0.32023 (8) | 0.0156 (2) |
| H1A | 0.0958 | 0.0639 | 0.3752 | 0.019* |
| H1B | 0.0470 | 0.0254 | 0.3286 | 0.019* |
| C2A | 0.00837 (9) | 0.18133 (9) | 0.29219 (8) | 0.0140 (2) |
| H2A | 0.0351 | 0.2296 | 0.2982 | 0.017* |
| C3A | 0.02031 (9) | 0.20865 (9) | 0.19085 (8) | 0.0136 (2) |
| H3A | 0.0098 | 0.2819 | 0.1597 | 0.016* |
| C4A | 0.13049 (9) | 0.16124 (9) | 0.15738 (8) | 0.0161 (2) |
| H4A | 0.1349 | 0.1235 | 0.1219 | 0.019* |
| H4B | 0.1708 | 0.2126 | 0.1210 | 0.019* |
| C5A | -0.10039 (9) | 0.18450 (9) | 0.34712 (8) | 0.0151 (2) |
| C6A | -0.12884 (9) | 0.12720 (9) | 0.24667 (8) | 0.0140 (2) |
| C7A | -0.05380 (9) | 0.16780 (9) | 0.17777 (8) | 0.0139 (2) |
| C8A | 0.21535 (9) | -0.08592 (9) | 0.21879 (8) | 0.0151 (2) |
| C9A | 0.23804 (9) | -0.08690 (10) | 0.13255 (9) | 0.0169 (2) |
| H9A | 0.2754 | -0.0423 | 0.0838 | 0.020* |
| C10A | 0.20415 (10) | -0.15542 (10) | 0.12004 (9) | 0.0182 (2) |
| H10A | 0.2205 | -0.1573 | 0.0627 | 0.022* |
| C11A | 0.14599 (9) | -0.22137 (9) | 0.19239 (9) | 0.0176 (2) |
| C12A | 0.12421 (10) | -0.21869 (10) | 0.27782 (9) | 0.0195 (3) |
| H12A | 0.0856 | -0.2623 | 0.3264 | 0.023* |
| C13A | 0.15886 (10) | -0.15229 (9) | 0.29224 (9) | 0.0180 (2) |
| H13A | 0.1446 | -0.1521 | 0.3500 | 0.022* |
| C14A | 0.10751 (11) | -0.29328 (10) | 0.17736 (10) | 0.0237 (3) |
| H14A | 0.0789 | -0.3401 | 0.2346 | 0.036* |
| H14B | 0.1618 | -0.3292 | 0.1443 | 0.036* |
| H14C | 0.0573 | -0.2563 | 0.1436 | 0.036* |
| C15A | -0.10928 (10) | 0.13859 (10) | 0.44833 (9) | 0.0196 (3) |
| H15A | -0.1787 | 0.1423 | 0.4798 | 0.029* |
| H15B | -0.0793 | 0.1749 | 0.4642 | 0.029* |
| H15C | -0.0754 | 0.0695 | 0.4649 | 0.029* |
| C16A | -0.16078 (10) | 0.29122 (10) | 0.31870 (9) | 0.0203 (3) |
| H16A | -0.2288 | 0.2916 | 0.3539 | 0.030* |
| H16B | -0.1596 | 0.3155 | 0.2558 | 0.030* |
| H16C | -0.1322 | 0.3339 | 0.3284 | 0.030* |
| C17A | -0.20253 (9) | 0.08353 (10) | 0.23786 (8) | 0.0170 (2) |

| | | | | |
|------|---------------|---------------|--------------|--------------|
| H17A | -0.2640 | 0.1332 | 0.2319 | 0.020* |
| H17B | -0.2181 | 0.0260 | 0.2927 | 0.020* |
| C18A | -0.16466 (9) | 0.05037 (10) | 0.15700 (8) | 0.0162 (2) |
| C19A | -0.12275 (10) | 0.13658 (10) | 0.07437 (8) | 0.0174 (2) |
| H19A | -0.0921 | 0.1141 | 0.0239 | 0.021* |
| H19B | -0.1779 | 0.1920 | 0.0585 | 0.021* |
| C20A | -0.04644 (9) | 0.17479 (9) | 0.08724 (8) | 0.0150 (2) |
| C21A | -0.08307 (10) | -0.04498 (10) | 0.17470 (10) | 0.0224 (3) |
| H21A | -0.0600 | -0.0649 | 0.1235 | 0.034* |
| H21B | -0.1101 | -0.0980 | 0.2274 | 0.034* |
| H21C | -0.0281 | -0.0317 | 0.1843 | 0.034* |
| C22A | -0.25169 (10) | 0.02860 (11) | 0.14144 (9) | 0.0219 (3) |
| H22A | -0.3030 | 0.0881 | 0.1303 | 0.033* |
| H22B | -0.2785 | -0.0246 | 0.1940 | 0.033* |
| H22C | -0.2282 | 0.0087 | 0.0902 | 0.033* |
| S1B | 0.58153 (2) | 0.36735 (2) | 0.91676 (2) | 0.01783 (7) |
| O1B | 0.60377 (7) | 0.31500 (7) | 1.00416 (7) | 0.0230 (2) |
| O2B | 0.65469 (7) | 0.41543 (8) | 0.83988 (7) | 0.0246 (2) |
| O3B | 0.20929 (7) | 0.39100 (7) | 1.00821 (6) | 0.01585 (18) |
| O4B | 0.36565 (7) | 0.53637 (7) | 0.69990 (6) | 0.0212 (2) |
| N1B | 0.48350 (8) | 0.45447 (8) | 0.92689 (7) | 0.0166 (2) |
| C1B | 0.39849 (9) | 0.42918 (10) | 1.00667 (8) | 0.0169 (2) |
| H1C | 0.4102 | 0.4313 | 1.0587 | 0.020* |
| H1D | 0.3872 | 0.3630 | 1.0217 | 0.020* |
| C2B | 0.30988 (10) | 0.51128 (9) | 0.97598 (8) | 0.0163 (2) |
| H2B | 0.3087 | 0.5707 | 0.9864 | 0.020* |
| C3B | 0.33392 (9) | 0.53689 (9) | 0.87399 (8) | 0.0152 (2) |
| H3B | 0.3009 | 0.6059 | 0.8442 | 0.018* |
| C4B | 0.44818 (10) | 0.52853 (10) | 0.84526 (9) | 0.0183 (2) |
| H4C | 0.4776 | 0.5050 | 0.7954 | 0.022* |
| H4D | 0.4654 | 0.5930 | 0.8271 | 0.022* |
| C5B | 0.20938 (10) | 0.47821 (9) | 1.02573 (8) | 0.0164 (2) |
| C6B | 0.24455 (9) | 0.39631 (9) | 0.92183 (8) | 0.0133 (2) |
| C7B | 0.30000 (9) | 0.46398 (9) | 0.85579 (8) | 0.0136 (2) |
| C8B | 0.54718 (9) | 0.28174 (9) | 0.89251 (9) | 0.0166 (2) |
| C9B | 0.56429 (10) | 0.29192 (10) | 0.80321 (9) | 0.0193 (3) |
| H9B | 0.5958 | 0.3426 | 0.7562 | 0.023* |
| C10B | 0.53371 (10) | 0.22542 (10) | 0.78513 (9) | 0.0208 (3) |
| H10B | 0.5458 | 0.2315 | 0.7257 | 0.025* |
| C11B | 0.48518 (10) | 0.14986 (10) | 0.85493 (10) | 0.0201 (3) |
| C12B | 0.46883 (10) | 0.14092 (10) | 0.94360 (10) | 0.0203 (3) |
| H12B | 0.4363 | 0.0910 | 0.9906 | 0.024* |
| C13B | 0.50051 (10) | 0.20558 (10) | 0.96285 (9) | 0.0187 (2) |
| H13B | 0.4905 | 0.1980 | 1.0225 | 0.022* |
| C14B | 0.44979 (11) | 0.07989 (11) | 0.83458 (12) | 0.0293 (3) |
| H14D | 0.4512 | 0.0158 | 0.8836 | 0.044* |
| H14E | 0.4930 | 0.0714 | 0.7799 | 0.044* |
| H14F | 0.3826 | 0.1080 | 0.8273 | 0.044* |
| C15B | 0.19143 (11) | 0.44044 (11) | 1.12761 (9) | 0.0218 (3) |

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|------|--------------|--------------|-------------|--------------|
| H15D | 0.1275 | 0.4206 | 1.1557 | 0.033* |
| H15E | 0.1923 | 0.4931 | 1.1446 | 0.033* |
| H15F | 0.2431 | 0.3838 | 1.1470 | 0.033* |
| C16B | 0.12321 (10) | 0.56104 (10) | 0.99339 (9) | 0.0223 (3) |
| H16D | 0.0612 | 0.5379 | 1.0259 | 0.033* |
| H16E | 0.1324 | 0.5783 | 0.9300 | 0.033* |
| H16F | 0.1216 | 0.6192 | 1.0039 | 0.033* |
| C17B | 0.21490 (9) | 0.31860 (9) | 0.90774 (8) | 0.0154 (2) |
| H17C | 0.1542 | 0.3003 | 0.9528 | 0.018* |
| H17D | 0.2672 | 0.2588 | 0.9160 | 0.018* |
| C18B | 0.19713 (9) | 0.35647 (9) | 0.81308 (8) | 0.0145 (2) |
| C19B | 0.28960 (9) | 0.39721 (10) | 0.74443 (8) | 0.0167 (2) |
| H19C | 0.3450 | 0.3409 | 0.7431 | 0.020* |
| H19D | 0.2759 | 0.4297 | 0.6850 | 0.020* |
| C20B | 0.32205 (9) | 0.47148 (9) | 0.76271 (8) | 0.0153 (2) |
| C21B | 0.18315 (10) | 0.26915 (10) | 0.79817 (9) | 0.0200 (3) |
| H21D | 0.1720 | 0.2925 | 0.7391 | 0.030* |
| H21E | 0.1265 | 0.2428 | 0.8430 | 0.030* |
| H21F | 0.2421 | 0.2173 | 0.8031 | 0.030* |
| C22B | 0.10247 (9) | 0.43823 (10) | 0.80570 (9) | 0.0185 (2) |
| H22D | 0.0914 | 0.4618 | 0.7466 | 0.028* |
| H22E | 0.1104 | 0.4931 | 0.8156 | 0.028* |
| H22F | 0.0461 | 0.4111 | 0.8503 | 0.028* |
| S1C | 0.06942 (2) | 0.31075 (2) | 0.43203 (2) | 0.01595 (7) |
| O1C | 0.04312 (7) | 0.36155 (7) | 0.34654 (6) | 0.0209 (2) |
| O2C | -0.00541 (7) | 0.27458 (7) | 0.51339 (6) | 0.0217 (2) |
| O3C | 0.44398 (7) | 0.23966 (7) | 0.32573 (6) | 0.01574 (18) |
| O4C | 0.29178 (7) | 0.12901 (7) | 0.63807 (6) | 0.01970 (19) |
| N1C | 0.15528 (8) | 0.21312 (8) | 0.42480 (7) | 0.0153 (2) |
| C1C | 0.23803 (9) | 0.22488 (10) | 0.34229 (8) | 0.0162 (2) |
| H1E | 0.2597 | 0.2878 | 0.3211 | 0.019* |
| H1F | 0.2179 | 0.2227 | 0.2944 | 0.019* |
| C2C | 0.32110 (9) | 0.13502 (9) | 0.37290 (8) | 0.0144 (2) |
| H2C | 0.3104 | 0.0764 | 0.3686 | 0.017* |
| C3C | 0.30547 (9) | 0.11588 (9) | 0.47298 (8) | 0.0142 (2) |
| H3C | 0.3327 | 0.0455 | 0.5055 | 0.017* |
| C4C | 0.19076 (9) | 0.13798 (9) | 0.50623 (8) | 0.0165 (2) |
| H4E | 0.1641 | 0.0774 | 0.5297 | 0.020* |
| H4F | 0.1706 | 0.1652 | 0.5533 | 0.020* |
| C5C | 0.42593 (9) | 0.15428 (9) | 0.31662 (8) | 0.0155 (2) |
| C6C | 0.41611 (9) | 0.24271 (9) | 0.40928 (8) | 0.0137 (2) |
| C7C | 0.35476 (9) | 0.18503 (9) | 0.48107 (8) | 0.0132 (2) |
| C8C | 0.12120 (9) | 0.39334 (9) | 0.44529 (8) | 0.0156 (2) |
| C9C | 0.10565 (10) | 0.39398 (10) | 0.53056 (9) | 0.0190 (3) |
| H9C | 0.0693 | 0.3500 | 0.5811 | 0.023* |
| C10C | 0.14503 (10) | 0.46115 (10) | 0.53951 (9) | 0.0206 (3) |
| H10C | 0.1336 | 0.4626 | 0.5964 | 0.025* |
| C11C | 0.20147 (10) | 0.52656 (10) | 0.46448 (9) | 0.0188 (3) |
| C12C | 0.21710 (10) | 0.52384 (10) | 0.37981 (9) | 0.0200 (3) |

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|------|--------------|--------------|--------------|------------|
| H12C | 0.2553 | 0.5663 | 0.3295 | 0.024* |
| C13C | 0.17642 (10) | 0.45861 (10) | 0.36939 (9) | 0.0189 (3) |
| H13C | 0.1860 | 0.4586 | 0.3123 | 0.023* |
| C14C | 0.24354 (11) | 0.59926 (11) | 0.47523 (11) | 0.0255 (3) |
| H14G | 0.2928 | 0.6262 | 0.4212 | 0.038* |
| H14H | 0.2741 | 0.5647 | 0.5255 | 0.038* |
| H14I | 0.1905 | 0.6530 | 0.4859 | 0.038* |
| C15C | 0.43654 (10) | 0.18716 (10) | 0.21621 (8) | 0.0192 (3) |
| H15G | 0.5033 | 0.1983 | 0.1838 | 0.029* |
| H15H | 0.3893 | 0.2483 | 0.1965 | 0.029* |
| H15I | 0.4237 | 0.1357 | 0.2048 | 0.029* |
| C16C | 0.50641 (10) | 0.06369 (10) | 0.34852 (9) | 0.0208 (3) |
| H16G | 0.5711 | 0.0781 | 0.3120 | 0.031* |
| H16H | 0.4962 | 0.0068 | 0.3433 | 0.031* |
| H16I | 0.5028 | 0.0489 | 0.4105 | 0.031* |
| C17C | 0.45921 (10) | 0.31889 (9) | 0.41250 (8) | 0.0166 (2) |
| H17E | 0.4135 | 0.3833 | 0.3981 | 0.020* |
| H17F | 0.5219 | 0.3265 | 0.3670 | 0.020* |
| C18C | 0.47790 (9) | 0.28933 (10) | 0.50539 (8) | 0.0159 (2) |
| C19C | 0.38213 (10) | 0.26125 (10) | 0.57732 (8) | 0.0173 (2) |
| H19E | 0.3961 | 0.2337 | 0.6364 | 0.021* |
| H19F | 0.3316 | 0.3221 | 0.5736 | 0.021* |
| C20C | 0.33934 (9) | 0.18652 (9) | 0.57013 (8) | 0.0143 (2) |
| C21C | 0.50086 (11) | 0.37855 (11) | 0.51048 (9) | 0.0239 (3) |
| H21G | 0.5125 | 0.3597 | 0.5686 | 0.036* |
| H21H | 0.4451 | 0.4339 | 0.5018 | 0.036* |
| H21I | 0.5595 | 0.3982 | 0.4642 | 0.036* |
| C22C | 0.56620 (10) | 0.20100 (11) | 0.52004 (10) | 0.0234 (3) |
| H22G | 0.5776 | 0.1828 | 0.5781 | 0.035* |
| H22H | 0.6250 | 0.2200 | 0.4738 | 0.035* |
| H22I | 0.5514 | 0.1447 | 0.5174 | 0.035* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|--------------|--------------|--------------|---------------|---------------|---------------|
| S1A | 0.01382 (13) | 0.01656 (14) | 0.01820 (15) | -0.00132 (11) | -0.00606 (11) | -0.00806 (12) |
| O1A | 0.0239 (5) | 0.0223 (5) | 0.0238 (5) | 0.0022 (4) | -0.0147 (4) | -0.0102 (4) |
| O2A | 0.0147 (4) | 0.0244 (5) | 0.0255 (5) | -0.0060 (4) | -0.0017 (4) | -0.0116 (4) |
| O3A | 0.0179 (4) | 0.0197 (4) | 0.0136 (4) | -0.0086 (3) | -0.0012 (3) | -0.0069 (3) |
| O4A | 0.0192 (4) | 0.0243 (5) | 0.0141 (4) | -0.0104 (4) | -0.0021 (4) | -0.0034 (4) |
| N1A | 0.0151 (5) | 0.0153 (5) | 0.0128 (5) | -0.0018 (4) | -0.0043 (4) | -0.0046 (4) |
| C1A | 0.0155 (5) | 0.0159 (5) | 0.0142 (5) | -0.0023 (4) | -0.0040 (4) | -0.0046 (4) |
| C2A | 0.0148 (5) | 0.0144 (5) | 0.0142 (5) | -0.0038 (4) | -0.0037 (4) | -0.0059 (4) |
| C3A | 0.0138 (5) | 0.0129 (5) | 0.0142 (5) | -0.0041 (4) | -0.0038 (4) | -0.0038 (4) |
| C4A | 0.0149 (5) | 0.0175 (6) | 0.0135 (5) | -0.0043 (4) | -0.0036 (4) | -0.0025 (5) |
| C5A | 0.0161 (5) | 0.0163 (5) | 0.0146 (5) | -0.0047 (4) | -0.0035 (4) | -0.0066 (5) |
| C6A | 0.0151 (5) | 0.0138 (5) | 0.0131 (5) | -0.0041 (4) | -0.0038 (4) | -0.0038 (4) |
| C7A | 0.0145 (5) | 0.0140 (5) | 0.0134 (5) | -0.0047 (4) | -0.0038 (4) | -0.0036 (4) |

supplementary materials

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|------|--------------|--------------|--------------|---------------|---------------|---------------|
| C8A | 0.0159 (5) | 0.0143 (5) | 0.0158 (6) | -0.0016 (4) | -0.0049 (5) | -0.0064 (5) |
| C9A | 0.0177 (6) | 0.0183 (6) | 0.0149 (6) | -0.0037 (5) | -0.0035 (5) | -0.0065 (5) |
| C10A | 0.0194 (6) | 0.0217 (6) | 0.0162 (6) | -0.0032 (5) | -0.0061 (5) | -0.0085 (5) |
| C11A | 0.0164 (5) | 0.0161 (6) | 0.0228 (6) | -0.0009 (5) | -0.0082 (5) | -0.0084 (5) |
| C12A | 0.0203 (6) | 0.0156 (6) | 0.0191 (6) | -0.0045 (5) | -0.0035 (5) | -0.0038 (5) |
| C13A | 0.0212 (6) | 0.0161 (6) | 0.0155 (6) | -0.0023 (5) | -0.0038 (5) | -0.0061 (5) |
| C14A | 0.0241 (7) | 0.0205 (6) | 0.0322 (8) | -0.0054 (5) | -0.0114 (6) | -0.0105 (6) |
| C15A | 0.0218 (6) | 0.0234 (6) | 0.0156 (6) | -0.0070 (5) | -0.0031 (5) | -0.0084 (5) |
| C16A | 0.0173 (6) | 0.0183 (6) | 0.0236 (6) | -0.0019 (5) | -0.0026 (5) | -0.0093 (5) |
| C17A | 0.0166 (5) | 0.0200 (6) | 0.0155 (6) | -0.0096 (5) | -0.0026 (5) | -0.0044 (5) |
| C18A | 0.0176 (6) | 0.0185 (6) | 0.0152 (6) | -0.0083 (5) | -0.0041 (5) | -0.0052 (5) |
| C19A | 0.0189 (6) | 0.0214 (6) | 0.0141 (5) | -0.0089 (5) | -0.0047 (5) | -0.0043 (5) |
| C20A | 0.0153 (5) | 0.0143 (5) | 0.0146 (5) | -0.0043 (4) | -0.0042 (4) | -0.0031 (4) |
| C21A | 0.0251 (7) | 0.0198 (6) | 0.0249 (7) | -0.0053 (5) | -0.0068 (6) | -0.0092 (5) |
| C22A | 0.0237 (6) | 0.0266 (7) | 0.0189 (6) | -0.0137 (5) | -0.0050 (5) | -0.0057 (5) |
| S1B | 0.01645 (14) | 0.01895 (15) | 0.02195 (16) | -0.00511 (11) | -0.00590 (12) | -0.00887 (12) |
| O1B | 0.0247 (5) | 0.0235 (5) | 0.0275 (5) | -0.0022 (4) | -0.0141 (4) | -0.0110 (4) |
| O2B | 0.0188 (5) | 0.0263 (5) | 0.0306 (5) | -0.0095 (4) | -0.0023 (4) | -0.0114 (4) |
| O3B | 0.0200 (4) | 0.0178 (4) | 0.0106 (4) | -0.0076 (3) | -0.0006 (3) | -0.0056 (3) |
| O4B | 0.0231 (5) | 0.0246 (5) | 0.0133 (4) | -0.0128 (4) | 0.0007 (4) | -0.0031 (4) |
| N1B | 0.0179 (5) | 0.0164 (5) | 0.0174 (5) | -0.0054 (4) | -0.0049 (4) | -0.0061 (4) |
| C1B | 0.0187 (6) | 0.0178 (6) | 0.0160 (6) | -0.0042 (5) | -0.0051 (5) | -0.0066 (5) |
| C2B | 0.0209 (6) | 0.0140 (5) | 0.0165 (6) | -0.0039 (5) | -0.0051 (5) | -0.0072 (5) |
| C3B | 0.0179 (6) | 0.0128 (5) | 0.0153 (5) | -0.0051 (4) | -0.0044 (5) | -0.0039 (4) |
| C4B | 0.0191 (6) | 0.0180 (6) | 0.0184 (6) | -0.0085 (5) | -0.0054 (5) | -0.0031 (5) |
| C5B | 0.0199 (6) | 0.0158 (5) | 0.0142 (5) | -0.0032 (5) | -0.0031 (5) | -0.0072 (5) |
| C6B | 0.0129 (5) | 0.0150 (5) | 0.0120 (5) | -0.0030 (4) | -0.0031 (4) | -0.0046 (4) |
| C7B | 0.0143 (5) | 0.0144 (5) | 0.0130 (5) | -0.0046 (4) | -0.0027 (4) | -0.0049 (4) |
| C8B | 0.0154 (5) | 0.0165 (6) | 0.0196 (6) | -0.0032 (4) | -0.0047 (5) | -0.0077 (5) |
| C9B | 0.0186 (6) | 0.0197 (6) | 0.0184 (6) | -0.0050 (5) | -0.0029 (5) | -0.0065 (5) |
| C10B | 0.0218 (6) | 0.0229 (6) | 0.0215 (6) | -0.0008 (5) | -0.0088 (5) | -0.0114 (5) |
| C11B | 0.0155 (6) | 0.0172 (6) | 0.0315 (7) | 0.0013 (5) | -0.0099 (5) | -0.0123 (5) |
| C12B | 0.0179 (6) | 0.0161 (6) | 0.0261 (7) | -0.0041 (5) | -0.0033 (5) | -0.0081 (5) |
| C13B | 0.0199 (6) | 0.0179 (6) | 0.0177 (6) | -0.0032 (5) | -0.0033 (5) | -0.0075 (5) |
| C14B | 0.0260 (7) | 0.0253 (7) | 0.0487 (10) | 0.0002 (6) | -0.0200 (7) | -0.0197 (7) |
| C15B | 0.0270 (7) | 0.0238 (7) | 0.0144 (6) | -0.0035 (5) | -0.0034 (5) | -0.0091 (5) |
| C16B | 0.0211 (6) | 0.0215 (6) | 0.0196 (6) | 0.0013 (5) | -0.0034 (5) | -0.0082 (5) |
| C17B | 0.0168 (5) | 0.0153 (5) | 0.0149 (5) | -0.0072 (4) | -0.0037 (5) | -0.0036 (4) |
| C18B | 0.0148 (5) | 0.0170 (5) | 0.0136 (5) | -0.0049 (4) | -0.0028 (4) | -0.0067 (5) |
| C19B | 0.0167 (5) | 0.0221 (6) | 0.0135 (5) | -0.0055 (5) | -0.0014 (5) | -0.0091 (5) |
| C20B | 0.0134 (5) | 0.0168 (5) | 0.0142 (5) | -0.0037 (4) | -0.0020 (4) | -0.0049 (5) |
| C21B | 0.0208 (6) | 0.0225 (6) | 0.0221 (6) | -0.0083 (5) | -0.0038 (5) | -0.0111 (5) |
| C22B | 0.0155 (5) | 0.0219 (6) | 0.0195 (6) | -0.0020 (5) | -0.0046 (5) | -0.0097 (5) |
| S1C | 0.01428 (13) | 0.01932 (15) | 0.01696 (14) | -0.00348 (11) | -0.00379 (11) | -0.00892 (12) |
| O1C | 0.0216 (5) | 0.0231 (5) | 0.0218 (5) | 0.0004 (4) | -0.0114 (4) | -0.0104 (4) |
| O2C | 0.0168 (4) | 0.0272 (5) | 0.0227 (5) | -0.0079 (4) | 0.0010 (4) | -0.0127 (4) |
| O3C | 0.0195 (4) | 0.0173 (4) | 0.0121 (4) | -0.0082 (3) | -0.0012 (3) | -0.0057 (3) |
| O4C | 0.0236 (5) | 0.0216 (5) | 0.0130 (4) | -0.0115 (4) | -0.0015 (4) | -0.0029 (4) |
| N1C | 0.0144 (5) | 0.0179 (5) | 0.0139 (5) | -0.0031 (4) | -0.0028 (4) | -0.0066 (4) |

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|------|------------|------------|------------|-------------|-------------|-------------|
| C1C | 0.0168 (5) | 0.0184 (6) | 0.0134 (5) | -0.0032 (5) | -0.0033 (4) | -0.0063 (5) |
| C2C | 0.0160 (5) | 0.0141 (5) | 0.0149 (5) | -0.0043 (4) | -0.0036 (4) | -0.0063 (4) |
| C3C | 0.0163 (5) | 0.0133 (5) | 0.0128 (5) | -0.0053 (4) | -0.0029 (4) | -0.0035 (4) |
| C4C | 0.0167 (5) | 0.0171 (6) | 0.0149 (6) | -0.0071 (5) | -0.0034 (5) | -0.0031 (5) |
| C5C | 0.0169 (5) | 0.0154 (5) | 0.0155 (6) | -0.0043 (4) | -0.0024 (5) | -0.0074 (5) |
| C6C | 0.0147 (5) | 0.0153 (5) | 0.0119 (5) | -0.0043 (4) | -0.0041 (4) | -0.0042 (4) |
| C7C | 0.0139 (5) | 0.0136 (5) | 0.0131 (5) | -0.0045 (4) | -0.0031 (4) | -0.0048 (4) |
| C8C | 0.0157 (5) | 0.0162 (6) | 0.0163 (6) | -0.0018 (4) | -0.0044 (5) | -0.0076 (5) |
| C9C | 0.0206 (6) | 0.0223 (6) | 0.0147 (6) | -0.0069 (5) | -0.0016 (5) | -0.0074 (5) |
| C10C | 0.0241 (6) | 0.0250 (7) | 0.0164 (6) | -0.0048 (5) | -0.0069 (5) | -0.0095 (5) |
| C11C | 0.0185 (6) | 0.0170 (6) | 0.0237 (6) | -0.0004 (5) | -0.0093 (5) | -0.0088 (5) |
| C12C | 0.0200 (6) | 0.0160 (6) | 0.0211 (6) | -0.0045 (5) | -0.0022 (5) | -0.0056 (5) |
| C13C | 0.0242 (6) | 0.0180 (6) | 0.0143 (6) | -0.0045 (5) | -0.0038 (5) | -0.0061 (5) |
| C14C | 0.0247 (7) | 0.0224 (7) | 0.0347 (8) | -0.0055 (5) | -0.0108 (6) | -0.0118 (6) |
| C15C | 0.0224 (6) | 0.0226 (6) | 0.0138 (6) | -0.0064 (5) | -0.0019 (5) | -0.0081 (5) |
| C16C | 0.0176 (6) | 0.0214 (6) | 0.0206 (6) | -0.0008 (5) | -0.0032 (5) | -0.0083 (5) |
| C17C | 0.0200 (6) | 0.0167 (6) | 0.0141 (5) | -0.0089 (5) | -0.0039 (5) | -0.0030 (5) |
| C18C | 0.0175 (5) | 0.0190 (6) | 0.0137 (5) | -0.0087 (5) | -0.0036 (5) | -0.0048 (5) |
| C19C | 0.0192 (6) | 0.0210 (6) | 0.0146 (6) | -0.0086 (5) | -0.0015 (5) | -0.0078 (5) |
| C20C | 0.0134 (5) | 0.0151 (5) | 0.0139 (5) | -0.0041 (4) | -0.0023 (4) | -0.0047 (4) |
| C21C | 0.0305 (7) | 0.0292 (7) | 0.0180 (6) | -0.0184 (6) | -0.0036 (5) | -0.0072 (6) |
| C22C | 0.0188 (6) | 0.0283 (7) | 0.0215 (6) | -0.0066 (5) | -0.0065 (5) | -0.0051 (6) |

Geometric parameters (Å, °)

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|----------|-------------|-----------|-------------|
| S1A—O1A | 1.4356 (10) | C10B—H10B | 0.93 |
| S1A—O2A | 1.4358 (10) | C11B—C12B | 1.391 (2) |
| S1A—N1A | 1.6245 (11) | C11B—C14B | 1.5094 (19) |
| S1A—C8A | 1.7617 (13) | C12B—C13B | 1.3903 (19) |
| O3A—C6A | 1.3468 (14) | C12B—H12B | 0.93 |
| O3A—C5A | 1.4696 (15) | C13B—H13B | 0.93 |
| O4A—C20A | 1.2305 (14) | C14B—H14D | 0.96 |
| N1A—C1A | 1.4718 (15) | C14B—H14E | 0.96 |
| N1A—C4A | 1.4883 (15) | C14B—H14F | 0.96 |
| C1A—C2A | 1.5267 (17) | C15B—H15D | 0.96 |
| C1A—H1A | 0.97 | C15B—H15E | 0.96 |
| C1A—H1B | 0.97 | C15B—H15F | 0.96 |
| C2A—C5A | 1.5284 (17) | C16B—H16D | 0.96 |
| C2A—C3A | 1.5416 (17) | C16B—H16E | 0.96 |
| C2A—H2A | 0.98 | C16B—H16F | 0.96 |
| C3A—C7A | 1.5115 (17) | C17B—C18B | 1.5371 (17) |
| C3A—C4A | 1.5479 (17) | C17B—H17C | 0.97 |
| C3A—H3A | 0.98 | C17B—H17D | 0.97 |
| C4A—H4A | 0.97 | C18B—C21B | 1.5277 (18) |
| C4A—H4B | 0.97 | C18B—C22B | 1.5289 (17) |
| C5A—C15A | 1.5195 (17) | C18B—C19B | 1.5324 (17) |
| C5A—C16A | 1.5210 (18) | C19B—C20B | 1.5194 (18) |
| C6A—C7A | 1.3571 (16) | C19B—H19C | 0.97 |
| C6A—C17A | 1.4960 (17) | C19B—H19D | 0.97 |

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| C7A—C20A | 1.4611 (17) | C21B—H21D | 0.96 |
| C8A—C9A | 1.3907 (18) | C21B—H21E | 0.96 |
| C8A—C13A | 1.3953 (17) | C21B—H21F | 0.96 |
| C9A—C10A | 1.3920 (18) | C22B—H22D | 0.96 |
| C9A—H9A | 0.93 | C22B—H22E | 0.96 |
| C10A—C11A | 1.3977 (18) | C22B—H22F | 0.96 |
| C10A—H10A | 0.93 | S1C—O2C | 1.4358 (10) |
| C11A—C12A | 1.3895 (19) | S1C—O1C | 1.4360 (10) |
| C11A—C14A | 1.5078 (18) | S1C—N1C | 1.6219 (11) |
| C12A—C13A | 1.3896 (19) | S1C—C8C | 1.7652 (13) |
| C12A—H12A | 0.93 | O3C—C6C | 1.3511 (14) |
| C13A—H13A | 0.93 | O3C—C5C | 1.4712 (15) |
| C14A—H14A | 0.96 | O4C—C20C | 1.2290 (14) |
| C14A—H14B | 0.96 | N1C—C1C | 1.4798 (16) |
| C14A—H14C | 0.96 | N1C—C4C | 1.4871 (16) |
| C15A—H15A | 0.96 | C1C—C2C | 1.5311 (17) |
| C15A—H15B | 0.96 | C1C—H1E | 0.97 |
| C15A—H15C | 0.96 | C1C—H1F | 0.97 |
| C16A—H16A | 0.96 | C2C—C5C | 1.5302 (17) |
| C16A—H16B | 0.96 | C2C—C3C | 1.5382 (17) |
| C16A—H16C | 0.96 | C2C—H2C | 0.98 |
| C17A—C18A | 1.5303 (18) | C3C—C7C | 1.5098 (17) |
| C17A—H17A | 0.97 | C3C—C4C | 1.5377 (17) |
| C17A—H17B | 0.97 | C3C—H3C | 0.98 |
| C18A—C22A | 1.5288 (18) | C4C—H4E | 0.97 |
| C18A—C19A | 1.5299 (17) | C4C—H4F | 0.97 |
| C18A—C21A | 1.5339 (18) | C5C—C15C | 1.5169 (17) |
| C19A—C20A | 1.5108 (17) | C5C—C16C | 1.5185 (18) |
| C19A—H19A | 0.97 | C6C—C7C | 1.3583 (16) |
| C19A—H19B | 0.97 | C6C—C17C | 1.4981 (17) |
| C21A—H21A | 0.96 | C7C—C20C | 1.4587 (17) |
| C21A—H21B | 0.96 | C8C—C9C | 1.3894 (18) |
| C21A—H21C | 0.96 | C8C—C13C | 1.3946 (17) |
| C22A—H22A | 0.96 | C9C—C10C | 1.3906 (19) |
| C22A—H22B | 0.96 | C9C—H9C | 0.93 |
| C22A—H22C | 0.96 | C10C—C11C | 1.3982 (19) |
| S1B—O2B | 1.4337 (10) | C10C—H10C | 0.93 |
| S1B—O1B | 1.4395 (10) | C11C—C12C | 1.3923 (19) |
| S1B—N1B | 1.6212 (11) | C11C—C14C | 1.5102 (19) |
| S1B—C8B | 1.7592 (13) | C12C—C13C | 1.3916 (19) |
| O3B—C6B | 1.3500 (14) | C12C—H12C | 0.93 |
| O3B—C5B | 1.4701 (15) | C13C—H13C | 0.93 |
| O4B—C20B | 1.2271 (15) | C14C—H14G | 0.96 |
| N1B—C1B | 1.4803 (16) | C14C—H14H | 0.96 |
| N1B—C4B | 1.4846 (16) | C14C—H14I | 0.96 |
| C1B—C2B | 1.5331 (17) | C15C—H15G | 0.96 |
| C1B—H1C | 0.97 | C15C—H15H | 0.96 |
| C1B—H1D | 0.97 | C15C—H15I | 0.96 |
| C2B—C5B | 1.5252 (18) | C16C—H16G | 0.96 |

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| C2B—C3B | 1.5329 (17) | C16C—H16H | 0.96 |
| C2B—H2B | 0.98 | C16C—H16I | 0.96 |
| C3B—C7B | 1.5128 (17) | C17C—C18C | 1.5365 (17) |
| C3B—C4B | 1.5376 (18) | C17C—H17E | 0.97 |
| C3B—H3B | 0.98 | C17C—H17F | 0.97 |
| C4B—H4C | 0.97 | C18C—C22C | 1.5267 (19) |
| C4B—H4D | 0.97 | C18C—C21C | 1.5271 (18) |
| C5B—C15B | 1.5175 (17) | C18C—C19C | 1.5320 (17) |
| C5B—C16B | 1.5208 (18) | C19C—C20C | 1.5110 (17) |
| C6B—C7B | 1.3559 (16) | C19C—H19E | 0.97 |
| C6B—C17B | 1.4996 (17) | C19C—H19F | 0.97 |
| C7B—C20B | 1.4588 (17) | C21C—H21G | 0.96 |
| C8B—C13B | 1.3909 (18) | C21C—H21H | 0.96 |
| C8B—C9B | 1.3938 (18) | C21C—H21I | 0.96 |
| C9B—C10B | 1.3926 (19) | C22C—H22G | 0.96 |
| C9B—H9B | 0.93 | C22C—H22H | 0.96 |
| C10B—C11B | 1.3950 (19) | C22C—H22I | 0.96 |
| O1A—S1A—O2A | 120.18 (6) | C13B—C12B—C11B | 120.90 (12) |
| O1A—S1A—N1A | 106.92 (6) | C13B—C12B—H12B | 119.6 |
| O2A—S1A—N1A | 105.63 (6) | C11B—C12B—H12B | 119.6 |
| O1A—S1A—C8A | 107.95 (6) | C12B—C13B—C8B | 119.66 (13) |
| O2A—S1A—C8A | 107.94 (6) | C12B—C13B—H13B | 120.2 |
| N1A—S1A—C8A | 107.64 (6) | C8B—C13B—H13B | 120.2 |
| C6A—O3A—C5A | 117.82 (9) | C11B—C14B—H14D | 109.5 |
| C1A—N1A—C4A | 110.35 (9) | C11B—C14B—H14E | 109.5 |
| C1A—N1A—S1A | 120.18 (8) | H14D—C14B—H14E | 109.5 |
| C4A—N1A—S1A | 117.47 (8) | C11B—C14B—H14F | 109.5 |
| N1A—C1A—C2A | 101.06 (9) | H14D—C14B—H14F | 109.5 |
| N1A—C1A—H1A | 111.6 | H14E—C14B—H14F | 109.5 |
| C2A—C1A—H1A | 111.6 | C5B—C15B—H15D | 109.5 |
| N1A—C1A—H1B | 111.6 | C5B—C15B—H15E | 109.5 |
| C2A—C1A—H1B | 111.6 | H15D—C15B—H15E | 109.5 |
| H1A—C1A—H1B | 109.4 | C5B—C15B—H15F | 109.5 |
| C1A—C2A—C5A | 114.93 (10) | H15D—C15B—H15F | 109.5 |
| C1A—C2A—C3A | 103.59 (10) | H15E—C15B—H15F | 109.5 |
| C5A—C2A—C3A | 113.38 (10) | C5B—C16B—H16D | 109.5 |
| C1A—C2A—H2A | 108.2 | C5B—C16B—H16E | 109.5 |
| C5A—C2A—H2A | 108.2 | H16D—C16B—H16E | 109.5 |
| C3A—C2A—H2A | 108.2 | C5B—C16B—H16F | 109.5 |
| C7A—C3A—C2A | 110.51 (10) | H16D—C16B—H16F | 109.5 |
| C7A—C3A—C4A | 112.84 (10) | H16E—C16B—H16F | 109.5 |
| C2A—C3A—C4A | 104.08 (9) | C6B—C17B—C18B | 112.09 (10) |
| C7A—C3A—H3A | 109.8 | C6B—C17B—H17C | 109.2 |
| C2A—C3A—H3A | 109.8 | C18B—C17B—H17C | 109.2 |
| C4A—C3A—H3A | 109.8 | C6B—C17B—H17D | 109.2 |
| N1A—C4A—C3A | 104.35 (9) | C18B—C17B—H17D | 109.2 |
| N1A—C4A—H4A | 110.9 | H17C—C17B—H17D | 107.9 |
| C3A—C4A—H4A | 110.9 | C21B—C18B—C22B | 108.94 (10) |
| N1A—C4A—H4B | 110.9 | C21B—C18B—C19B | 110.35 (10) |

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| C3A—C4A—H4B | 110.9 | C22B—C18B—C19B | 111.29 (10) |
| H4A—C4A—H4B | 108.9 | C21B—C18B—C17B | 109.25 (10) |
| O3A—C5A—C15A | 103.84 (9) | C22B—C18B—C17B | 109.54 (10) |
| O3A—C5A—C16A | 107.99 (10) | C19B—C18B—C17B | 107.44 (10) |
| C15A—C5A—C16A | 111.47 (11) | C20B—C19B—C18B | 115.13 (10) |
| O3A—C5A—C2A | 109.93 (10) | C20B—C19B—H19C | 108.5 |
| C15A—C5A—C2A | 112.56 (10) | C18B—C19B—H19C | 108.5 |
| C16A—C5A—C2A | 110.73 (10) | C20B—C19B—H19D | 108.5 |
| O3A—C6A—C7A | 124.19 (11) | C18B—C19B—H19D | 108.5 |
| O3A—C6A—C17A | 111.25 (10) | H19C—C19B—H19D | 107.5 |
| C7A—C6A—C17A | 124.56 (11) | O4B—C20B—C7B | 121.17 (12) |
| C6A—C7A—C20A | 118.57 (11) | O4B—C20B—C19B | 120.14 (11) |
| C6A—C7A—C3A | 122.42 (11) | C7B—C20B—C19B | 118.69 (10) |
| C20A—C7A—C3A | 118.98 (10) | C18B—C21B—H21D | 109.5 |
| C9A—C8A—C13A | 120.71 (12) | C18B—C21B—H21E | 109.5 |
| C9A—C8A—S1A | 119.88 (10) | H21D—C21B—H21E | 109.5 |
| C13A—C8A—S1A | 119.40 (10) | C18B—C21B—H21F | 109.5 |
| C8A—C9A—C10A | 119.26 (12) | H21D—C21B—H21F | 109.5 |
| C8A—C9A—H9A | 120.4 | H21E—C21B—H21F | 109.5 |
| C10A—C9A—H9A | 120.4 | C18B—C22B—H22D | 109.5 |
| C9A—C10A—C11A | 120.96 (12) | C18B—C22B—H22E | 109.5 |
| C9A—C10A—H10A | 119.5 | H22D—C22B—H22E | 109.5 |
| C11A—C10A—H10A | 119.5 | C18B—C22B—H22F | 109.5 |
| C12A—C11A—C10A | 118.61 (12) | H22D—C22B—H22F | 109.5 |
| C12A—C11A—C14A | 121.03 (12) | H22E—C22B—H22F | 109.5 |
| C10A—C11A—C14A | 120.36 (12) | O2C—S1C—O1C | 120.06 (6) |
| C11A—C12A—C13A | 121.44 (12) | O2C—S1C—N1C | 106.19 (6) |
| C11A—C12A—H12A | 119.3 | O1C—S1C—N1C | 106.49 (6) |
| C13A—C12A—H12A | 119.3 | O2C—S1C—C8C | 107.60 (6) |
| C12A—C13A—C8A | 119.00 (12) | O1C—S1C—C8C | 107.15 (6) |
| C12A—C13A—H13A | 120.5 | N1C—S1C—C8C | 109.01 (6) |
| C8A—C13A—H13A | 120.5 | C6C—O3C—C5C | 117.58 (9) |
| C11A—C14A—H14A | 109.5 | C1C—N1C—C4C | 111.13 (10) |
| C11A—C14A—H14B | 109.5 | C1C—N1C—S1C | 118.85 (8) |
| H14A—C14A—H14B | 109.5 | C4C—N1C—S1C | 119.80 (8) |
| C11A—C14A—H14C | 109.5 | N1C—C1C—C2C | 103.70 (10) |
| H14A—C14A—H14C | 109.5 | N1C—C1C—H1E | 111.0 |
| H14B—C14A—H14C | 109.5 | C2C—C1C—H1E | 111.0 |
| C5A—C15A—H15A | 109.5 | N1C—C1C—H1F | 111.0 |
| C5A—C15A—H15B | 109.5 | C2C—C1C—H1F | 111.0 |
| H15A—C15A—H15B | 109.5 | H1E—C1C—H1F | 109.0 |
| C5A—C15A—H15C | 109.5 | C5C—C2C—C1C | 112.87 (10) |
| H15A—C15A—H15C | 109.5 | C5C—C2C—C3C | 112.68 (10) |
| H15B—C15A—H15C | 109.5 | C1C—C2C—C3C | 104.27 (10) |
| C5A—C16A—H16A | 109.5 | C5C—C2C—H2C | 109.0 |
| C5A—C16A—H16B | 109.5 | C1C—C2C—H2C | 109.0 |
| H16A—C16A—H16B | 109.5 | C3C—C2C—H2C | 109.0 |
| C5A—C16A—H16C | 109.5 | C7C—C3C—C4C | 112.76 (10) |
| H16A—C16A—H16C | 109.5 | C7C—C3C—C2C | 110.05 (10) |

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| H16B—C16A—H16C | 109.5 | C4C—C3C—C2C | 102.80 (10) |
| C6A—C17A—C18A | 113.35 (10) | C7C—C3C—H3C | 110.3 |
| C6A—C17A—H17A | 108.9 | C4C—C3C—H3C | 110.3 |
| C18A—C17A—H17A | 108.9 | C2C—C3C—H3C | 110.3 |
| C6A—C17A—H17B | 108.9 | N1C—C4C—C3C | 104.62 (9) |
| C18A—C17A—H17B | 108.9 | N1C—C4C—H4E | 110.8 |
| H17A—C17A—H17B | 107.7 | C3C—C4C—H4E | 110.8 |
| C22A—C18A—C19A | 109.92 (10) | N1C—C4C—H4F | 110.8 |
| C22A—C18A—C17A | 109.41 (10) | C3C—C4C—H4F | 110.8 |
| C19A—C18A—C17A | 107.65 (10) | H4E—C4C—H4F | 108.9 |
| C22A—C18A—C21A | 109.09 (11) | O3C—C5C—C15C | 104.35 (10) |
| C19A—C18A—C21A | 110.41 (11) | O3C—C5C—C16C | 107.86 (10) |
| C17A—C18A—C21A | 110.33 (10) | C15C—C5C—C16C | 111.01 (11) |
| C20A—C19A—C18A | 114.55 (10) | O3C—C5C—C2C | 109.23 (10) |
| C20A—C19A—H19A | 108.6 | C15C—C5C—C2C | 112.52 (10) |
| C18A—C19A—H19A | 108.6 | C16C—C5C—C2C | 111.51 (10) |
| C20A—C19A—H19B | 108.6 | O3C—C6C—C7C | 123.96 (11) |
| C18A—C19A—H19B | 108.6 | O3C—C6C—C17C | 111.81 (10) |
| H19A—C19A—H19B | 107.6 | C7C—C6C—C17C | 124.22 (11) |
| O4A—C20A—C7A | 121.31 (11) | C6C—C7C—C20C | 118.82 (11) |
| O4A—C20A—C19A | 120.05 (11) | C6C—C7C—C3C | 122.50 (11) |
| C7A—C20A—C19A | 118.58 (10) | C20C—C7C—C3C | 118.63 (10) |
| C18A—C21A—H21A | 109.5 | C9C—C8C—C13C | 120.54 (12) |
| C18A—C21A—H21B | 109.5 | C9C—C8C—S1C | 120.07 (10) |
| H21A—C21A—H21B | 109.5 | C13C—C8C—S1C | 119.39 (10) |
| C18A—C21A—H21C | 109.5 | C8C—C9C—C10C | 119.30 (12) |
| H21A—C21A—H21C | 109.5 | C8C—C9C—H9C | 120.3 |
| H21B—C21A—H21C | 109.5 | C10C—C9C—H9C | 120.3 |
| C18A—C22A—H22A | 109.5 | C9C—C10C—C11C | 121.13 (12) |
| C18A—C22A—H22B | 109.5 | C9C—C10C—H10C | 119.4 |
| H22A—C22A—H22B | 109.5 | C11C—C10C—H10C | 119.4 |
| C18A—C22A—H22C | 109.5 | C12C—C11C—C10C | 118.60 (12) |
| H22A—C22A—H22C | 109.5 | C12C—C11C—C14C | 120.87 (12) |
| H22B—C22A—H22C | 109.5 | C10C—C11C—C14C | 120.52 (13) |
| O2B—S1B—O1B | 120.15 (6) | C13C—C12C—C11C | 121.00 (12) |
| O2B—S1B—N1B | 106.45 (6) | C13C—C12C—H12C | 119.5 |
| O1B—S1B—N1B | 106.65 (6) | C11C—C12C—H12C | 119.5 |
| O2B—S1B—C8B | 108.33 (6) | C12C—C13C—C8C | 119.40 (12) |
| O1B—S1B—C8B | 107.35 (6) | C12C—C13C—H13C | 120.3 |
| N1B—S1B—C8B | 107.29 (6) | C8C—C13C—H13C | 120.3 |
| C6B—O3B—C5B | 116.93 (9) | C11C—C14C—H14G | 109.5 |
| C1B—N1B—C4B | 111.34 (10) | C11C—C14C—H14H | 109.5 |
| C1B—N1B—S1B | 118.83 (8) | H14G—C14C—H14H | 109.5 |
| C4B—N1B—S1B | 119.41 (9) | C11C—C14C—H14I | 109.5 |
| N1B—C1B—C2B | 103.30 (10) | H14G—C14C—H14I | 109.5 |
| N1B—C1B—H1C | 111.1 | H14H—C14C—H14I | 109.5 |
| C2B—C1B—H1C | 111.1 | C5C—C15C—H15G | 109.5 |
| N1B—C1B—H1D | 111.1 | C5C—C15C—H15H | 109.5 |
| C2B—C1B—H1D | 111.1 | H15G—C15C—H15H | 109.5 |

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| H1C—C1B—H1D | 109.1 | C5C—C15C—H15I | 109.5 |
| C5B—C2B—C3B | 112.70 (10) | H15G—C15C—H15I | 109.5 |
| C5B—C2B—C1B | 112.84 (10) | H15H—C15C—H15I | 109.5 |
| C3B—C2B—C1B | 104.55 (10) | C5C—C16C—H16G | 109.5 |
| C5B—C2B—H2B | 108.9 | C5C—C16C—H16H | 109.5 |
| C3B—C2B—H2B | 108.9 | H16G—C16C—H16H | 109.5 |
| C1B—C2B—H2B | 108.9 | C5C—C16C—H16I | 109.5 |
| C7B—C3B—C2B | 110.04 (10) | H16G—C16C—H16I | 109.5 |
| C7B—C3B—C4B | 113.36 (11) | H16H—C16C—H16I | 109.5 |
| C2B—C3B—C4B | 102.89 (10) | C6C—C17C—C18C | 112.49 (10) |
| C7B—C3B—H3B | 110.1 | C6C—C17C—H17E | 109.1 |
| C2B—C3B—H3B | 110.1 | C18C—C17C—H17E | 109.1 |
| C4B—C3B—H3B | 110.1 | C6C—C17C—H17F | 109.1 |
| N1B—C4B—C3B | 104.67 (10) | C18C—C17C—H17F | 109.1 |
| N1B—C4B—H4C | 110.8 | H17E—C17C—H17F | 107.8 |
| C3B—C4B—H4C | 110.8 | C22C—C18C—C21C | 109.50 (11) |
| N1B—C4B—H4D | 110.8 | C22C—C18C—C19C | 110.95 (10) |
| C3B—C4B—H4D | 110.8 | C21C—C18C—C19C | 108.92 (11) |
| H4C—C4B—H4D | 108.9 | C22C—C18C—C17C | 109.72 (11) |
| O3B—C5B—C15B | 104.31 (10) | C21C—C18C—C17C | 109.80 (10) |
| O3B—C5B—C16B | 107.68 (10) | C19C—C18C—C17C | 107.92 (10) |
| C15B—C5B—C16B | 111.02 (11) | C20C—C19C—C18C | 115.16 (10) |
| O3B—C5B—C2B | 109.24 (10) | C20C—C19C—H19E | 108.5 |
| C15B—C5B—C2B | 112.69 (11) | C18C—C19C—H19E | 108.5 |
| C16B—C5B—C2B | 111.52 (10) | C20C—C19C—H19F | 108.5 |
| O3B—C6B—C7B | 123.96 (11) | C18C—C19C—H19F | 108.5 |
| O3B—C6B—C17B | 111.91 (10) | H19E—C19C—H19F | 107.5 |
| C7B—C6B—C17B | 124.13 (11) | O4C—C20C—C7C | 120.99 (11) |
| C6B—C7B—C20B | 118.76 (11) | O4C—C20C—C19C | 120.38 (11) |
| C6B—C7B—C3B | 122.51 (11) | C7C—C20C—C19C | 118.63 (10) |
| C20B—C7B—C3B | 118.60 (10) | C18C—C21C—H21G | 109.5 |
| C13B—C8B—C9B | 120.30 (12) | C18C—C21C—H21H | 109.5 |
| C13B—C8B—S1B | 119.57 (10) | H21G—C21C—H21H | 109.5 |
| C9B—C8B—S1B | 120.11 (10) | C18C—C21C—H21I | 109.5 |
| C10B—C9B—C8B | 119.34 (12) | H21G—C21C—H21I | 109.5 |
| C10B—C9B—H9B | 120.3 | H21H—C21C—H21I | 109.5 |
| C8B—C9B—H9B | 120.3 | C18C—C22C—H22G | 109.5 |
| C9B—C10B—C11B | 120.95 (13) | C18C—C22C—H22H | 109.5 |
| C9B—C10B—H10B | 119.5 | H22G—C22C—H22H | 109.5 |
| C11B—C10B—H10B | 119.5 | C18C—C22C—H22I | 109.5 |
| C12B—C11B—C10B | 118.83 (12) | H22G—C22C—H22I | 109.5 |
| C12B—C11B—C14B | 120.55 (13) | H22H—C22C—H22I | 109.5 |
| C10B—C11B—C14B | 120.61 (13) | | |
| O1A—S1A—N1A—C1A | -45.96 (11) | C4B—C3B—C7B—C6B | 122.83 (13) |
| O2A—S1A—N1A—C1A | -175.06 (9) | C2B—C3B—C7B—C20B | -176.04 (10) |
| C8A—S1A—N1A—C1A | 69.81 (10) | C4B—C3B—C7B—C20B | -61.45 (14) |
| O1A—S1A—N1A—C4A | 174.96 (9) | O2B—S1B—C8B—C13B | 161.51 (10) |
| O2A—S1A—N1A—C4A | 45.86 (10) | O1B—S1B—C8B—C13B | 30.38 (12) |
| C8A—S1A—N1A—C4A | -69.27 (10) | N1B—S1B—C8B—C13B | -83.93 (11) |

| | | | |
|---------------------|--------------|---------------------|--------------|
| C4A—N1A—C1A—C2A | -34.00 (12) | O2B—S1B—C8B—C9B | -20.19 (12) |
| S1A—N1A—C1A—C2A | -175.70 (8) | O1B—S1B—C8B—C9B | -151.33 (11) |
| N1A—C1A—C2A—C5A | 164.73 (10) | N1B—S1B—C8B—C9B | 94.36 (11) |
| N1A—C1A—C2A—C3A | 40.49 (11) | C13B—C8B—C9B—C10B | 0.33 (19) |
| C1A—C2A—C3A—C7A | 88.38 (11) | S1B—C8B—C9B—C10B | -177.95 (10) |
| C5A—C2A—C3A—C7A | -36.86 (14) | C8B—C9B—C10B—C11B | 0.9 (2) |
| C1A—C2A—C3A—C4A | -33.03 (12) | C9B—C10B—C11B—C12B | -0.87 (19) |
| C5A—C2A—C3A—C4A | -158.27 (10) | C9B—C10B—C11B—C14B | 178.33 (12) |
| C1A—N1A—C4A—C3A | 13.58 (13) | C10B—C11B—C12B—C13B | -0.30 (19) |
| S1A—N1A—C4A—C3A | 156.44 (8) | C14B—C11B—C12B—C13B | -179.50 (12) |
| C7A—C3A—C4A—N1A | -107.31 (11) | C11B—C12B—C13B—C8B | 1.5 (2) |
| C2A—C3A—C4A—N1A | 12.53 (12) | C9B—C8B—C13B—C12B | -1.48 (19) |
| C6A—O3A—C5A—C15A | -164.07 (10) | S1B—C8B—C13B—C12B | 176.82 (10) |
| C6A—O3A—C5A—C16A | 77.48 (13) | O3B—C6B—C17B—C18B | 146.48 (10) |
| C6A—O3A—C5A—C2A | -43.43 (14) | C7B—C6B—C17B—C18B | -33.42 (16) |
| C1A—C2A—C5A—O3A | -65.07 (13) | C6B—C17B—C18B—C21B | 171.38 (10) |
| C3A—C2A—C5A—O3A | 53.83 (13) | C6B—C17B—C18B—C22B | -69.36 (13) |
| C1A—C2A—C5A—C15A | 50.16 (14) | C6B—C17B—C18B—C19B | 51.67 (13) |
| C3A—C2A—C5A—C15A | 169.05 (10) | C21B—C18B—C19B—C20B | -169.58 (11) |
| C1A—C2A—C5A—C16A | 175.69 (11) | C22B—C18B—C19B—C20B | 69.35 (14) |
| C3A—C2A—C5A—C16A | -65.41 (13) | C17B—C18B—C19B—C20B | -50.57 (14) |
| C5A—O3A—C6A—C7A | 16.51 (17) | C6B—C7B—C20B—O4B | 174.67 (12) |
| C5A—O3A—C6A—C17A | -163.36 (10) | C3B—C7B—C20B—O4B | -1.21 (18) |
| O3A—C6A—C7A—C20A | -176.47 (11) | C6B—C7B—C20B—C19B | -5.13 (17) |
| C17A—C6A—C7A—C20A | 3.39 (19) | C3B—C7B—C20B—C19B | 178.99 (11) |
| O3A—C6A—C7A—C3A | 1.69 (19) | C18B—C19B—C20B—O4B | -151.47 (12) |
| C17A—C6A—C7A—C3A | -178.45 (11) | C18B—C19B—C20B—C7B | 28.34 (16) |
| C2A—C3A—C7A—C6A | 9.35 (16) | O2C—S1C—N1C—C1C | 173.66 (9) |
| C4A—C3A—C7A—C6A | 125.42 (12) | O1C—S1C—N1C—C1C | 44.61 (11) |
| C2A—C3A—C7A—C20A | -172.49 (10) | C8C—S1C—N1C—C1C | -70.68 (10) |
| C4A—C3A—C7A—C20A | -56.42 (14) | O2C—S1C—N1C—C4C | -44.19 (10) |
| O1A—S1A—C8A—C9A | -144.95 (10) | O1C—S1C—N1C—C4C | -173.23 (9) |
| O2A—S1A—C8A—C9A | -13.62 (12) | C8C—S1C—N1C—C4C | 71.48 (10) |
| N1A—S1A—C8A—C9A | 99.96 (11) | C4C—N1C—C1C—C2C | 13.39 (13) |
| O1A—S1A—C8A—C13A | 35.54 (12) | S1C—N1C—C1C—C2C | 158.58 (8) |
| O2A—S1A—C8A—C13A | 166.87 (10) | N1C—C1C—C2C—C5C | -153.75 (10) |
| N1A—S1A—C8A—C13A | -79.54 (11) | N1C—C1C—C2C—C3C | -31.13 (12) |
| C13A—C8A—C9A—C10A | -0.31 (19) | C5C—C2C—C3C—C7C | 39.41 (13) |
| S1A—C8A—C9A—C10A | -179.81 (9) | C1C—C2C—C3C—C7C | -83.33 (11) |
| C8A—C9A—C10A—C11A | 1.44 (19) | C5C—C2C—C3C—C4C | 159.76 (10) |
| C9A—C10A—C11A—C12A | -1.30 (19) | C1C—C2C—C3C—C4C | 37.02 (12) |
| C9A—C10A—C11A—C14A | 178.36 (12) | C1C—N1C—C4C—C3C | 9.68 (13) |
| C10A—C11A—C12A—C13A | 0.03 (19) | S1C—N1C—C4C—C3C | -135.15 (9) |
| C14A—C11A—C12A—C13A | -179.63 (12) | C7C—C3C—C4C—N1C | 89.95 (12) |
| C11A—C12A—C13A—C8A | 1.07 (19) | C2C—C3C—C4C—N1C | -28.52 (12) |
| C9A—C8A—C13A—C12A | -0.92 (19) | C6C—O3C—C5C—C15C | 164.37 (10) |
| S1A—C8A—C13A—C12A | 178.58 (10) | C6C—O3C—C5C—C16C | -77.51 (13) |
| O3A—C6A—C17A—C18A | -157.46 (10) | C6C—O3C—C5C—C2C | 43.85 (13) |
| C7A—C6A—C17A—C18A | 22.67 (18) | C1C—C2C—C5C—O3C | 61.06 (13) |

supplementary materials

| | | | |
|---------------------|--------------|---------------------|--------------|
| C6A—C17A—C18A—C22A | -167.78 (11) | C3C—C2C—C5C—O3C | -56.72 (13) |
| C6A—C17A—C18A—C19A | -48.36 (14) | C1C—C2C—C5C—C15C | -54.34 (14) |
| C6A—C17A—C18A—C21A | 72.19 (13) | C3C—C2C—C5C—C15C | -172.12 (10) |
| C22A—C18A—C19A—C20A | 171.53 (11) | C1C—C2C—C5C—C16C | -179.82 (11) |
| C17A—C18A—C19A—C20A | 52.44 (14) | C3C—C2C—C5C—C16C | 62.40 (14) |
| C21A—C18A—C19A—C20A | -68.06 (14) | C5C—O3C—C6C—C7C | -14.29 (17) |
| C6A—C7A—C20A—O4A | 177.36 (12) | C5C—O3C—C6C—C17C | 166.96 (10) |
| C3A—C7A—C20A—O4A | -0.87 (18) | O3C—C6C—C7C—C20C | 172.98 (11) |
| C6A—C7A—C20A—C19A | 0.04 (17) | C17C—C6C—C7C—C20C | -8.43 (18) |
| C3A—C7A—C20A—C19A | -178.19 (11) | O3C—C6C—C7C—C3C | -4.26 (19) |
| C18A—C19A—C20A—O4A | 153.01 (12) | C17C—C6C—C7C—C3C | 174.34 (11) |
| C18A—C19A—C20A—C7A | -29.63 (16) | C4C—C3C—C7C—C6C | -123.50 (13) |
| O2B—S1B—N1B—C1B | -173.78 (9) | C2C—C3C—C7C—C6C | -9.36 (16) |
| O1B—S1B—N1B—C1B | -44.36 (11) | C4C—C3C—C7C—C20C | 59.26 (14) |
| C8B—S1B—N1B—C1B | 70.41 (10) | C2C—C3C—C7C—C20C | 173.40 (10) |
| O2B—S1B—N1B—C4B | 44.31 (11) | O2C—S1C—C8C—C9C | 15.46 (12) |
| O1B—S1B—N1B—C4B | 173.73 (9) | O1C—S1C—C8C—C9C | 145.84 (11) |
| C8B—S1B—N1B—C4B | -71.49 (10) | N1C—S1C—C8C—C9C | -99.30 (11) |
| C4B—N1B—C1B—C2B | -13.96 (13) | O2C—S1C—C8C—C13C | -163.64 (10) |
| S1B—N1B—C1B—C2B | -158.72 (9) | O1C—S1C—C8C—C13C | -33.26 (12) |
| N1B—C1B—C2B—C5B | 154.09 (10) | N1C—S1C—C8C—C13C | 81.61 (11) |
| N1B—C1B—C2B—C3B | 31.27 (12) | C13C—C8C—C9C—C10C | 0.6 (2) |
| C5B—C2B—C3B—C7B | -38.52 (14) | S1C—C8C—C9C—C10C | -178.52 (10) |
| C1B—C2B—C3B—C7B | 84.39 (12) | C8C—C9C—C10C—C11C | -1.2 (2) |
| C5B—C2B—C3B—C4B | -159.61 (10) | C9C—C10C—C11C—C12C | 0.5 (2) |
| C1B—C2B—C3B—C4B | -36.70 (12) | C9C—C10C—C11C—C14C | 179.85 (12) |
| C1B—N1B—C4B—C3B | -8.77 (14) | C10C—C11C—C12C—C13C | 0.9 (2) |
| S1B—N1B—C4B—C3B | 135.76 (9) | C14C—C11C—C12C—C13C | -178.45 (12) |
| C7B—C3B—C4B—N1B | -91.02 (12) | C11C—C12C—C13C—C8C | -1.6 (2) |
| C2B—C3B—C4B—N1B | 27.78 (12) | C9C—C8C—C13C—C12C | 0.80 (19) |
| C6B—O3B—C5B—C15B | -166.34 (10) | S1C—C8C—C13C—C12C | 179.89 (10) |
| C6B—O3B—C5B—C16B | 75.64 (13) | O3C—C6C—C17C—C18C | -149.76 (10) |
| C6B—O3B—C5B—C2B | -45.63 (13) | C7C—C6C—C17C—C18C | 31.49 (17) |
| C3B—C2B—C5B—O3B | 57.36 (13) | C6C—C17C—C18C—C22C | 71.25 (13) |
| C1B—C2B—C5B—O3B | -60.78 (13) | C6C—C17C—C18C—C21C | -168.35 (11) |
| C3B—C2B—C5B—C15B | 172.81 (10) | C6C—C17C—C18C—C19C | -49.75 (14) |
| C1B—C2B—C5B—C15B | 54.66 (14) | C22C—C18C—C19C—C20C | -69.78 (14) |
| C3B—C2B—C5B—C16B | -61.54 (14) | C21C—C18C—C19C—C20C | 169.61 (11) |
| C1B—C2B—C5B—C16B | -179.68 (11) | C17C—C18C—C19C—C20C | 50.45 (14) |
| C5B—O3B—C6B—C7B | 15.91 (17) | C6C—C7C—C20C—O4C | -173.46 (12) |
| C5B—O3B—C6B—C17B | -164.00 (10) | C3C—C7C—C20C—O4C | 3.88 (18) |
| O3B—C6B—C7B—C20B | -171.63 (11) | C6C—C7C—C20C—C19C | 7.21 (17) |
| C17B—C6B—C7B—C20B | 8.27 (18) | C3C—C7C—C20C—C19C | -175.45 (10) |
| O3B—C6B—C7B—C3B | 4.08 (19) | C18C—C19C—C20C—O4C | 150.50 (12) |
| C17B—C6B—C7B—C3B | -176.02 (11) | C18C—C19C—C20C—C7C | -30.17 (16) |
| C2B—C3B—C7B—C6B | 8.24 (16) | | |

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> |
|--------------------------------|-------------|---------------|-----------------------|-------------------------|
| C2A—H2A···O1C | 0.98 | 2.51 | 3.4447 (18) | 160 |
| C2B—H2B···O1B ⁱ | 0.98 | 2.52 | 3.4167 (18) | 151 |
| C2C—H2C···O1A | 0.98 | 2.51 | 3.4739 (18) | 167 |
| C4A—H4A···O4A | 0.97 | 2.48 | 2.9605 (17) | 111 |
| C4B—H4C···O4B | 0.97 | 2.48 | 3.0370 (18) | 116 |
| C4C—H4F···O4C | 0.97 | 2.43 | 3.0069 (17) | 118 |
| C12A—H12A···O2C ⁱⁱ | 0.93 | 2.51 | 3.2130 (16) | 133 |
| C12B—H12B···O2A ⁱⁱⁱ | 0.93 | 2.59 | 3.3762 (18) | 143 |
| C14C—H14I···O2C ^{iv} | 0.96 | 2.56 | 3.362 (2) | 142 |
| C17A—H17B···O4C ⁱⁱ | 0.97 | 2.51 | 3.3480 (18) | 145 |
| C17B—H17C···O4A ⁱⁱⁱ | 0.97 | 2.38 | 3.2776 (17) | 154 |
| C21C—H21I···O4B ^v | 0.96 | 2.47 | 3.3058 (17) | 146 |
| C22A—H22B···O4C ⁱⁱ | 0.96 | 2.55 | 3.4023 (16) | 147 |

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

Fig. 1

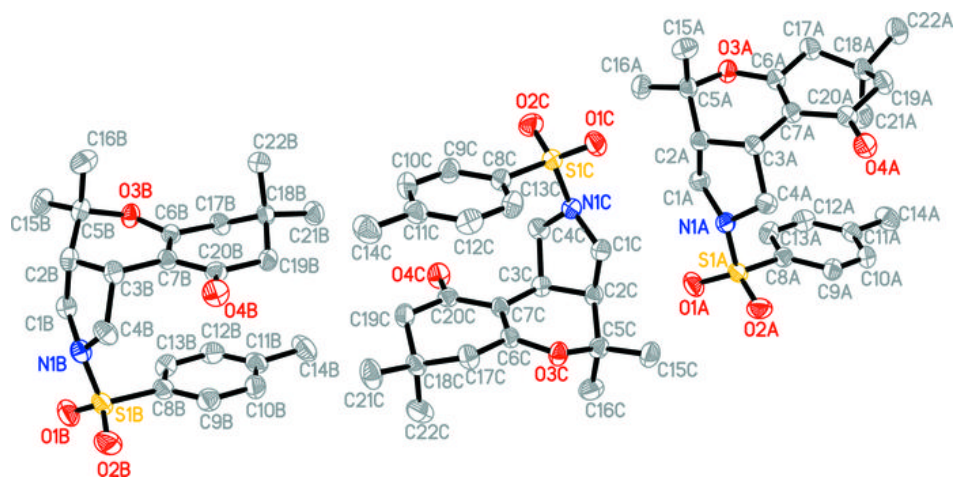


Fig. 2

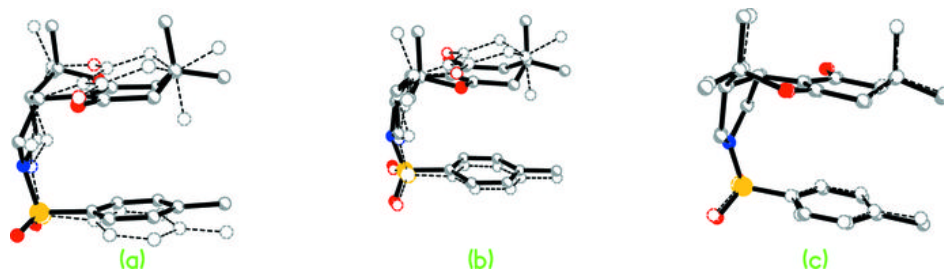
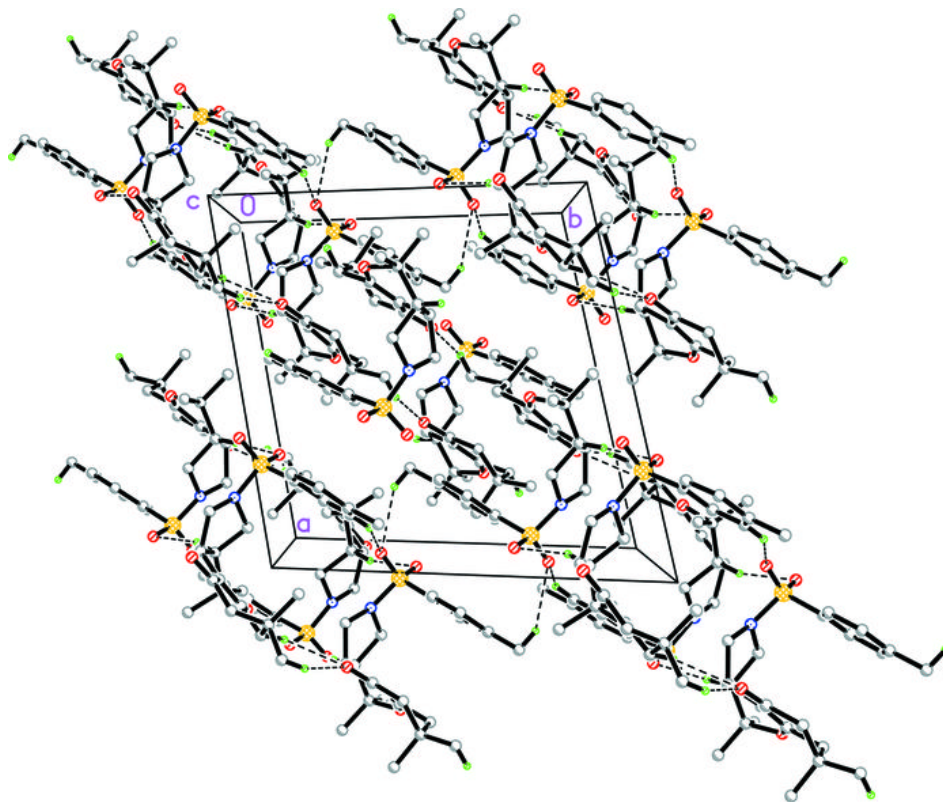


Fig. 3



Seven papers on fused-ring heterocyclic ketones containing an *N*-tosyl-pyrrolo[3,4-*c*]pyrano moiety. Corrigenda

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Corrections are made to the name of an author in seven papers by Chinnakali *et al.* [*Acta Cryst.* (2007), E**63**, o4363, o4364, o4434–o4435, o4436–o4437, o4438, o4489–o4490 and o4491–o4492].

In the papers by Chinnakali, Jayagopi *et al.* (2007*a,b*) and Chinnakali, Sudha *et al.* (2007*a,b,c,d,e*), the name of the author M. Jayagopi is given incorrectly. The correct name should be M. Jayagobi, as given above.

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