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(3a*R,8b*R**)-3a,8b-Dihydroxy-1-(4-methylphenyl)-2-methylsulfanyl-3-nitro-1,8b-dihydroindeno[1,2-*b*]pyrrol-4(3a*H*)-one**R. A. Nagalakshmi,^a J. Suresh,^a R. Ranjith Kumar,^b
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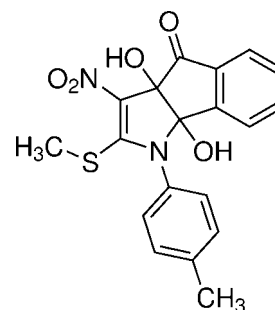
Received 26 January 2014; accepted 5 February 2014

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(C-C) = 0.006$ Å; disorder in main residue; R factor = 0.069; wR factor = 0.188; data-to-parameter ratio = 12.8.

The asymmetric unit of the title compound, $C_{19}H_{16}N_2O_5S$, contains four independent molecules (*A*, *B*, *C* and *D*), with two molecules (*B* and *D*) displaying disorder in their methylsulfanyl groups [occupancy ratios of 0.797 (11):0.203 (11) and 0.85 (2):0.15 (2)]. The nitro groups are twisted slightly out of the planes of the 2-pyrroline rings to which they are bonded with dihedral angles of 10.17 (1), 8.01 (1), 9.44 (1) and 8.87 (1)° in molecules *A*, *B*, *C* and *D*, respectively. The 2-pyrroline rings are almost orthogonal to the attached tolyl rings, forming dihedral angles of 73.44 (1), 81.21 (1), 88.18 (8) and 73.94 (1)° for molecules *A*, *B*, *C* and *D*, respectively. A weak intramolecular O—H...O interaction is observed in molecules *B* and *C*. The two hydroxy groups in each molecule are involved in intermolecular O—H...O hydrogen bonding. In the crystal, molecules are connected *via* O—H...O and C—H...O hydrogen bonds, forming a complex three-dimensional network.

Related literature

For the importance of pyrrolidine and pyrroline derivatives, see: Obniska *et al.* (2002); Stylianakis *et al.* (2003); Coldham & Hufton (2005); Kravchenko *et al.* (2005); Nair & Suja (2007); Pandey *et al.* (2006). For a related structure, see: Nagalakshmi *et al.* (2013). For ring conformation parameters, see: Cremer & Pople (1975).



Experimental

Crystal data

$C_{19}H_{16}N_2O_5S$
 $M_r = 384.40$
 Triclinic, $P\bar{1}$
 $a = 10.0830$ (5) Å
 $b = 10.2775$ (5) Å
 $c = 34.6579$ (17) Å
 $\alpha = 97.540$ (3)°
 $\beta = 96.028$ (3)°

$\gamma = 97.138$ (3)°
 $V = 3505.9$ (3) Å³
 $Z = 8$
 Mo $K\alpha$ radiation
 $\mu = 0.22$ mm⁻¹
 $T = 293$ K
 $0.21 \times 0.19 \times 0.18$ mm

Data collection

Bruker Kappa APEXII
 diffractometer
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.967$, $T_{\max} = 0.974$

53951 measured reflections
 13050 independent reflections
 9856 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.069$
 $wR(F^2) = 0.188$
 $S = 1.06$
 13050 reflections
 1016 parameters

12 restraints
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.68$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.30$ e Å⁻³

Table 1

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> |
|--|-------------|---------------|-----------------------|-------------------------|
| O3 <i>B</i> —H3 <i>B</i> ...O5 <i>B</i> | 0.82 | 2.19 | 2.748 (4) | 125 |
| O3 <i>C</i> —H3 <i>C</i> ...O5 <i>C</i> | 0.82 | 2.17 | 2.728 (4) | 126 |
| O2 <i>A</i> —H2 <i>A</i> ...O3 <i>A</i> ⁱ | 0.82 | 1.97 | 2.785 (3) | 175 |
| O3 <i>A</i> —H3 <i>A</i> ...O3 <i>B</i> ⁱⁱ | 0.82 | 2.04 | 2.854 (3) | 172 |
| O2 <i>B</i> —H2 <i>B</i> ...O1 <i>A</i> ⁱⁱⁱ | 0.82 | 2.01 | 2.787 (3) | 157 |
| O2 <i>C</i> —H2 <i>C</i> ...O5 <i>D</i> ^{iv} | 0.82 | 2.57 | 3.133 (4) | 127 |
| O2 <i>C</i> —H2 <i>C</i> ...O1 <i>D</i> ^{iv} | 0.82 | 2.03 | 2.797 (4) | 155 |
| O3 <i>D</i> —H3 <i>D</i> ...O3 <i>C</i> ^{iv} | 0.82 | 2.04 | 2.857 (3) | 175 |
| C10 <i>A</i> —H10 <i>A</i> ...O5 <i>A</i> ⁱⁱⁱ | 0.93 | 2.54 | 3.347 (5) | 146 |
| C10 <i>C</i> —H10 <i>C</i> ...O5 <i>C</i> ^v | 0.93 | 2.50 | 3.331 (6) | 149 |
| C10 <i>D</i> —H10 <i>D</i> ...O5 <i>D</i> ^v | 0.93 | 2.59 | 3.437 (5) | 151 |
| C10 <i>B</i> —H10 <i>B</i> ...O5 <i>B</i> ⁱⁱ | 0.93 | 2.53 | 3.316 (6) | 142 |
| C2—H2 <i>o</i> ...O1 <i>C</i> ^{vi} | 0.96 | 2.50 | 3.446 (7) | 170 |

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: PLATON (Spek, 2009); software used to prepare material for publication: SHELXL97.

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Supporting information for this paper is available from the IUCr electronic archives (Reference: BH2494).

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supplementary materials

Acta Cryst. (2014). E70, o279–o280 [doi:10.1107/S1600536814002712]

(3aR*,8bR*)-3a,8b-Dihydroxy-1-(4-methylphenyl)-2-methylsulfanyl-3-nitro-1,8b-dihydroindeno[1,2-b]pyrrol-4(3aH)-one

R. A. Nagalakshmi, J. Suresh, R. Ranjith Kumar, V. Jeyachandran and P. L. Nilantha Lakshman

1. Comment

Pyrrolidine-containing compounds form an important class of heterocyclic compounds with wide spread applications to the synthesis of biologically active compounds and natural products (Coldham & Hufton, 2005; Kravchenko *et al.*, 2005; Nair & Suja, 2007; Pandey *et al.*, 2006). Pyrrolidine derivatives possess anti-influenza (Stylianakis *et al.*, 2003) and anti-convulsant (Obniska *et al.*, 2002) activities. The high medicinal values of these compounds, in conjunction with our research interests, prompted us to synthesize and report the X-ray study of the title compound. The same compound with the *p*-tolyl group replaced by a *p*-methoxyphenyl group was previously reported (Nagalakshmi *et al.*, 2013).

The asymmetric unit of the title compound, C₁₉H₁₆N₂O₅S, contains four crystallographically independent molecules *A*, *B*, *C* and *D* (shown in Figs. 1, 2, 3 and 4, respectively), with two molecules (*A* and *C*) exhibiting similar geometries and the other two molecules (*B* and *D*) displaying disorder. The 2-pyrroline rings of molecules *A* and *D* are planar with r.m.s. deviations of 0.0277 (1) and 0.0234 (1) Å; molecules *B* and *C* adopt a slightly twisted conformation for these rings, with puckering parameters $Q = 0.112$ (4) Å, $\varphi_2 = 231$ (2)° and $Q = 0.080$ (4) Å, $\varphi_2 = 226$ (3)°, respectively (Cremer & Pople, 1975). The cyclopentane rings of molecules *A* and *D* are planar. Molecule *B* adopts a twisted cyclopentane conformation, with puckering parameters $Q = 0.098$ (4) Å, $\varphi_2 = 231$ (2)°; molecule *C* adopts an envelope cyclopentane conformation with puckering parameters $Q = 0.078$ (4) Å and $\varphi_2 = 351$ (3)°. The 2-pyrroline rings are almost orthogonal to the attached tolyl rings in all molecules, forming dihedral angles of 73.44 (1), 81.21 (1), 88.18 (8) and 73.94 (1)° for molecules *A*, *B*, *C* and *D*, respectively. The nitro group (N2/O4/O5) is twisted away from the attached 2-pyrroline ring (C1···C4/N1) in all molecules: dihedral angles are 10.17 (1), 8.01 (1), 9.44 (1) and 8.87 (1)° in *A*, *B*, *C* and *D*, respectively. In molecule *B*, the methyl group of the methylsulfanyl functionality is disordered over two positions with occupancies of 0.15 (2):0.85 (2). In molecule *D*, the methylsulfanyl group attached to the 2-pyrroline ring is disordered over two sites, with occupancies of 0.797 (11):0.203 (11).

A weak intra-molecular O—H···O interaction is observed in molecules *B* and *C* (Table 1). The two hydroxy groups in each molecule are involved in inter-molecular O—H···O hydrogen bonding. The C—H···O interactions between molecules form a linear chain along the *b* axis. Intermolecular interactions O2A—H2A···O3A and O2D—H2D···O3D connect the inversion related dimers, forming a ring motif $R_2^2(10)$ (Fig. 5). The C14D—H14D···O4D inter-molecular interactions form a ring motif $R_2^2(16)$. These ring motifs are further connected by C—H···O interactions. The O3D—H3D···O3C interaction forms a linear chain along the *b* axis. The C—H···O interactions involving molecules *A* (Fig. 5) and *B* form a linear chain along the *a* axis. The C—H···O interactions between molecules *C* and *D* form a linear chain along the *b* axis. This combination of O—H···O and C—H···O hydrogen bonds give a complex three-dimensional network for the crystal structure.

2. Experimental

A mixture of (*E*)-4-methyl-*N*-[1-(methylthio)-2-nitrovinyl]aniline (1 mmol) with ninhydrin (2,2-dihydroxyindane-1,3-dione, 1 mmol) in presence of glacial AcOH (3–5 drops) was thoroughly ground in a pestle and mortar at room temperature for 2–10 min. The reaction progress was monitored by thin layer chromatography. After completion of the reaction, the reaction mixture was triturated with crushed ice, the resulting solid filtered off and washed with water to afford the pure product. The compound was further recrystallized from ethanol to obtain suitable crystals for X-ray analysis. Yield 96%; m.p. 471–474 K.

3. Refinement

H atoms were placed at calculated positions and allowed to ride on their carrier atoms with C—H = 0.93 Å (aromatic CH) or 0.96 Å (methyl CH₃) and O—H = 0.82 Å. isotropic displacement parameters for H atoms were defined as $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for aromatic CH and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C}, \text{O})$ for CH₃ and OH groups. Molecules *B* and *C* have disordered parts in their methylsulfanyl fragments. For molecule *B*, the methyl group is disordered over two positions, C2 and C5B, for which occupancies converged to 0.85 (2) and 0.15 (2), respectively. A combination of *SIMU* (similar U_{ij} parameters for C2 and C5B) and *DELU* (rigid bonds for S2—C5B and S2—C2) restraints were applied (Sheldrick, 2008). In the case of molecule *C* the methylsulfanyl group is disordered over two positions, S3—C5C [occupancy: 0.797 (11)] and S3'—C1 [occupancy: 0.203 (11)]. A *SADI* (same distance) restraint was applied for bonds S3—C5C and S3'—C1.

Computing details

Data collection: *APEX2* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); data reduction: *SAINT* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97* (Sheldrick, 2008).

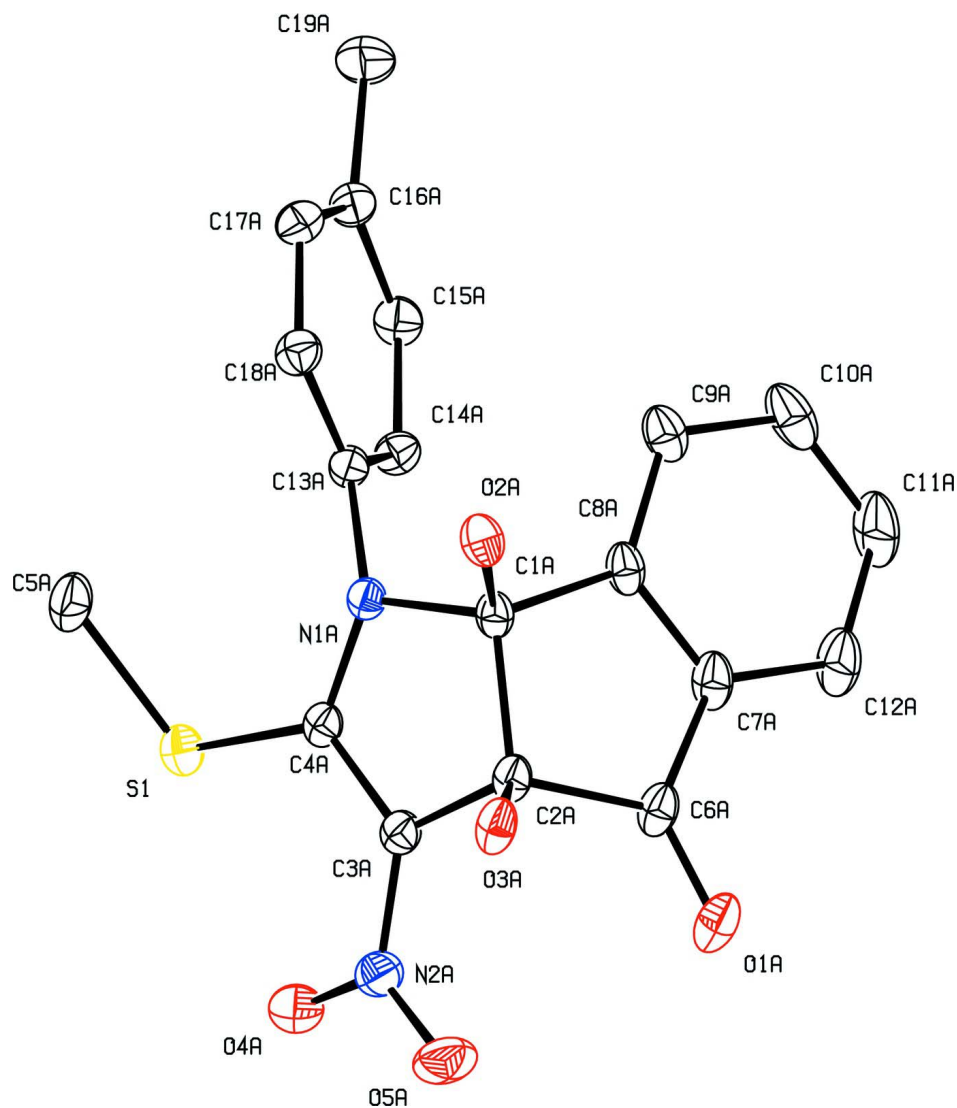


Figure 1

The molecular structure of (I), showing 20% probability displacement ellipsoids and the atom-numbering scheme of molecule *A*. H atoms are omitted for clarity.

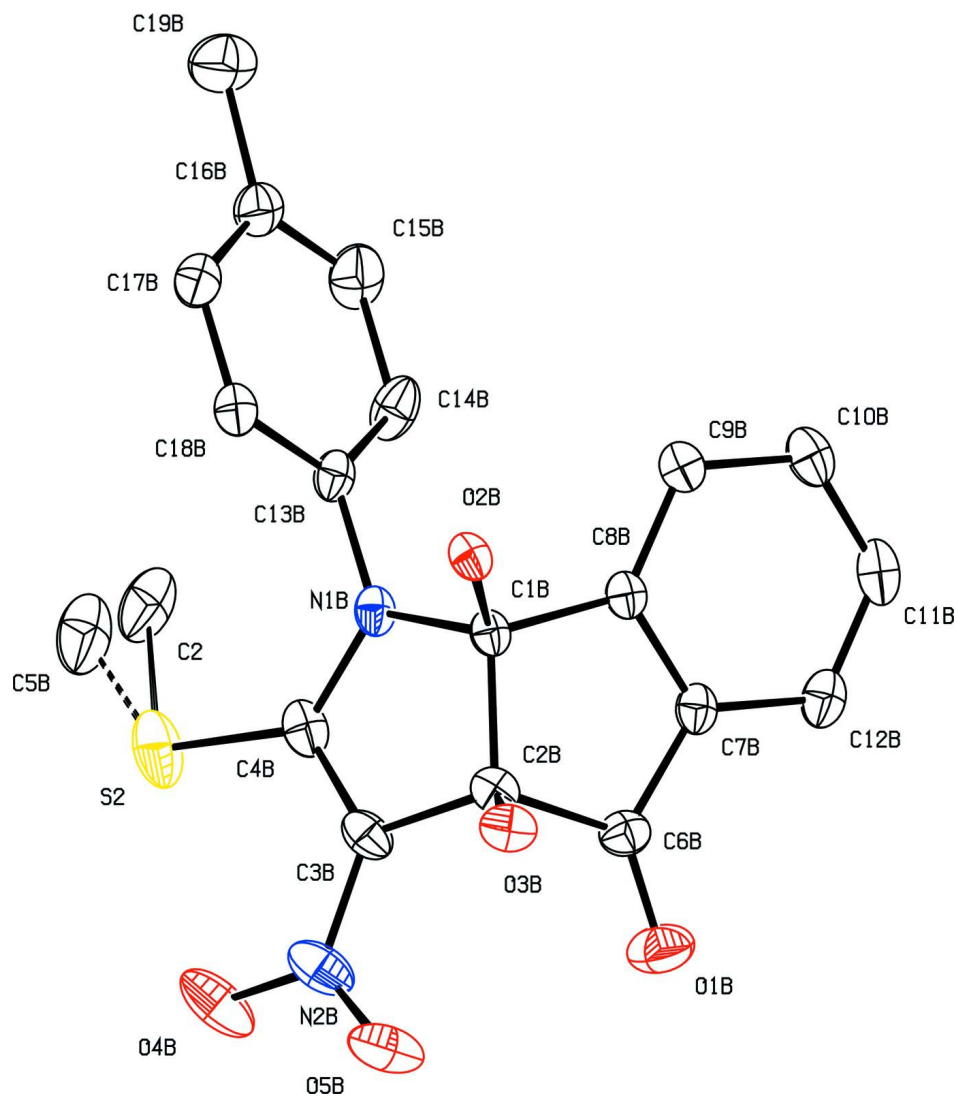


Figure 2

The molecular structure of (I), showing 20% probability displacement ellipsoids and the atom-numbering scheme of molecule *B*. H atoms are omitted for clarity.

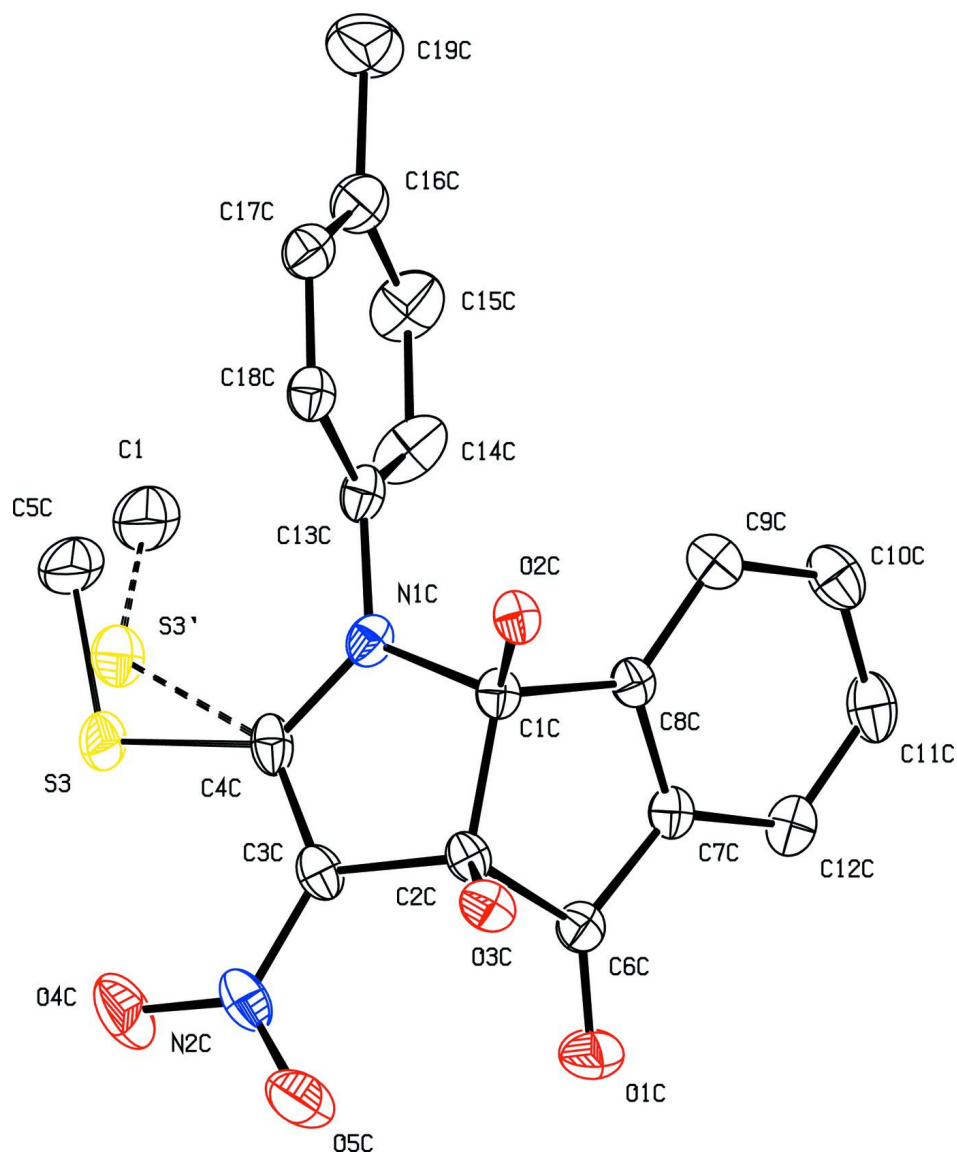


Figure 3

The molecular structure of (I), showing 20% probability displacement ellipsoids and the atom-numbering scheme of molecule C. H atoms are omitted for clarity.

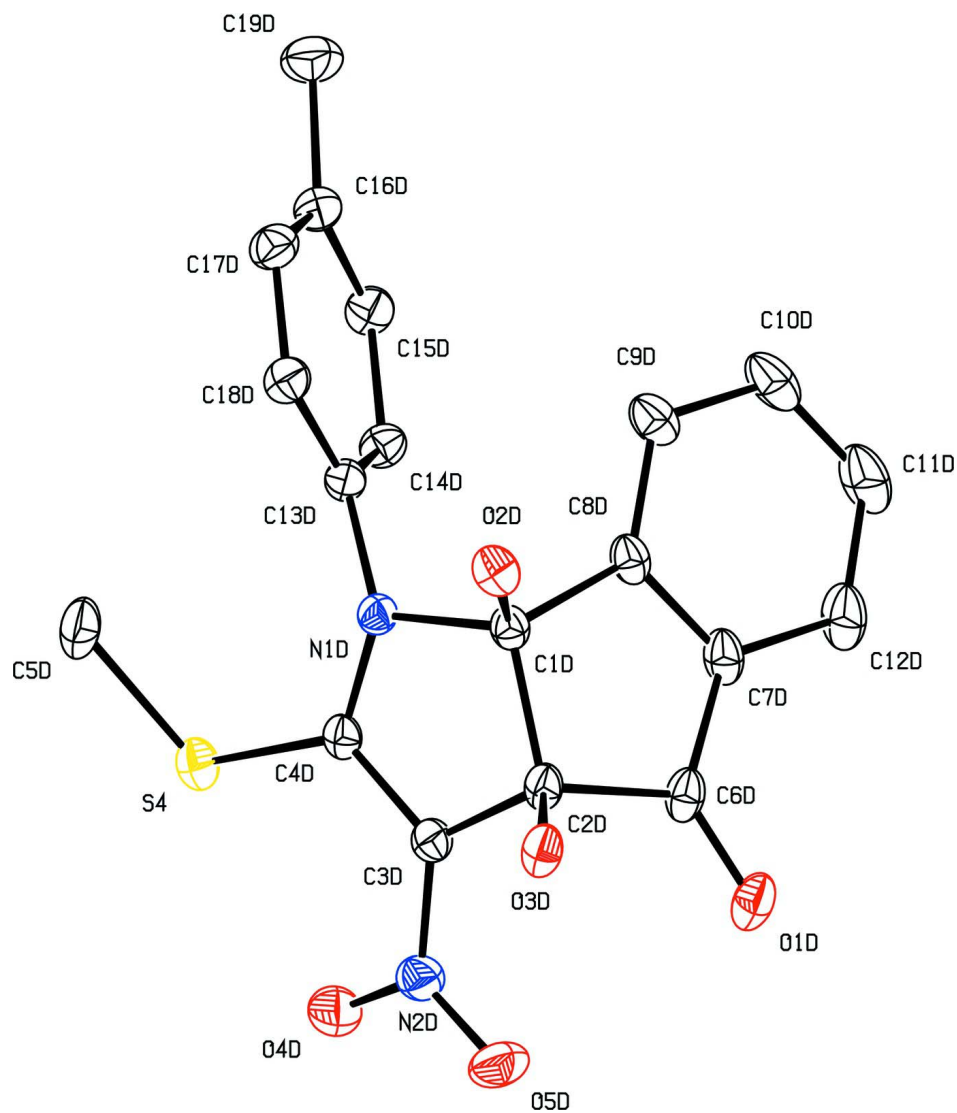
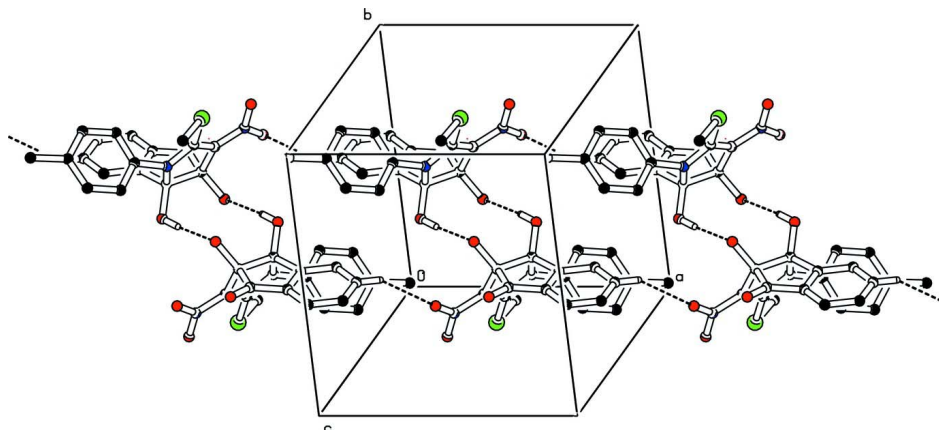


Figure 4

The molecular structure of (I), showing 20% probability displacement ellipsoids and the atom-numbering scheme of molecule *D*. H atoms are omitted for clarity.


Figure 5

The partial packing diagram showing O—H...O and C—H...O interactions in molecule *A*.

(3a*R,8b*R**)-3a,8b-Dihydroxy-1-(4-methylphenyl)-2-methylsulfanyl-3-nitro-1,8b-dihydroindeno[1,2-*b*]pyrrol-4(3a*H*)-one**

Crystal data

C₁₉H₁₆N₂O₅S

M_r = 384.40

Triclinic, *P* $\bar{1}$

Hall symbol: -P 1

a = 10.0830 (5) Å

b = 10.2775 (5) Å

c = 34.6579 (17) Å

α = 97.540 (3)°

β = 96.028 (3)°

γ = 97.138 (3)°

V = 3505.9 (3) Å³

Z = 8

F(000) = 1600

D_x = 1.457 Mg m⁻³

Melting point: 471 K

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 2000 reflections

θ = 2–31°

μ = 0.22 mm⁻¹

T = 293 K

Block, yellow

0.21 × 0.19 × 0.18 mm

Data collection

Bruker Kappa APEXII
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 0 pixels mm⁻¹

ω and φ scans

Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)

T_{min} = 0.967, *T_{max}* = 0.974

53951 measured reflections

13050 independent reflections

9856 reflections with *I* > 2σ(*I*)

R_{int} = 0.041

θ_{\max} = 25.5°, θ_{\min} = 0.6°

h = -12→12

k = -12→12

l = -41→41

Refinement

Refinement on *F*²

Least-squares matrix: full

R [*F*² > 2σ(*F*²)] = 0.069

wR (*F*²) = 0.188

S = 1.06

13050 reflections

1016 parameters

12 restraints

0 constraints

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/[σ²(*F_o*²) + (0.0785*P*)² + 4.0405*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.68 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|------|--------------|-------------|--------------|----------------------------------|------------|
| C5C | -0.0868 (7) | 0.8339 (10) | 0.7819 (3) | 0.101 (3) | 0.797 (11) |
| H7 | -0.1013 | 0.7976 | 0.8056 | 0.152* | 0.797 (11) |
| H8 | -0.0665 | 0.7659 | 0.7626 | 0.152* | 0.797 (11) |
| H9 | -0.1666 | 0.8673 | 0.7722 | 0.152* | 0.797 (11) |
| S3 | 0.0486 (3) | 0.9635 (4) | 0.79199 (5) | 0.0627 (8) | 0.797 (11) |
| C1 | -0.074 (3) | 0.738 (2) | 0.7612 (9) | 0.108 (6) | 0.203 (11) |
| H16 | -0.1206 | 0.6849 | 0.7777 | 0.162* | 0.203 (11) |
| H17 | -0.0078 | 0.6911 | 0.7499 | 0.162* | 0.203 (11) |
| H18 | -0.1374 | 0.7570 | 0.7407 | 0.162* | 0.203 (11) |
| S3' | 0.0055 (12) | 0.8874 (18) | 0.7892 (2) | 0.072 (3) | 0.203 (11) |
| C1A | 0.3058 (3) | 0.6338 (3) | 0.48764 (9) | 0.0393 (7) | |
| C1B | -0.4152 (3) | 0.5374 (3) | 0.34840 (9) | 0.0408 (7) | |
| C2A | 0.4370 (3) | 0.6700 (3) | 0.46797 (9) | 0.0408 (7) | |
| C2B | -0.5618 (4) | 0.4670 (4) | 0.34575 (10) | 0.0482 (8) | |
| C3A | 0.5131 (3) | 0.7805 (3) | 0.49689 (9) | 0.0403 (7) | |
| C3B | -0.6424 (4) | 0.5603 (4) | 0.32766 (11) | 0.0558 (9) | |
| C4A | 0.4468 (3) | 0.8042 (3) | 0.52945 (9) | 0.0369 (7) | |
| C4B | -0.5613 (4) | 0.6593 (4) | 0.31486 (10) | 0.0542 (9) | |
| C5A | 0.4410 (4) | 0.8588 (4) | 0.60993 (11) | 0.0636 (11) | |
| H1 | 0.4460 | 0.7655 | 0.6075 | 0.095* | |
| H2 | 0.3488 | 0.8733 | 0.6099 | 0.095* | |
| H3 | 0.4923 | 0.9027 | 0.6341 | 0.095* | |
| C5B | -0.540 (7) | 0.893 (5) | 0.282 (2) | 0.106 (6) | 0.15 (2) |
| H4 | -0.4535 | 0.8672 | 0.2771 | 0.159* | 0.15 (2) |
| H5 | -0.5284 | 0.9640 | 0.3030 | 0.159* | 0.15 (2) |
| H6 | -0.5805 | 0.9207 | 0.2584 | 0.159* | 0.15 (2) |
| C2 | -0.4941 (10) | 0.8501 (10) | 0.2651 (3) | 0.094 (3) | 0.85 (2) |
| H19 | -0.4300 | 0.9119 | 0.2831 | 0.142* | 0.85 (2) |
| H20 | -0.5329 | 0.8950 | 0.2451 | 0.142* | 0.85 (2) |
| H21 | -0.4497 | 0.7804 | 0.2533 | 0.142* | 0.85 (2) |
| C6A | 0.3863 (4) | 0.7211 (4) | 0.43000 (10) | 0.0467 (8) | |
| C6B | -0.5640 (4) | 0.3353 (4) | 0.31823 (11) | 0.0527 (9) | |
| C7A | 0.2434 (4) | 0.7296 (4) | 0.43012 (10) | 0.0520 (9) | |
| C7B | -0.4258 (4) | 0.3248 (3) | 0.31042 (10) | 0.0469 (8) | |
| C8A | 0.1947 (3) | 0.6753 (3) | 0.46124 (9) | 0.0444 (8) | |
| C8B | -0.3394 (3) | 0.4343 (3) | 0.32883 (9) | 0.0409 (7) | |
| C9A | 0.0578 (4) | 0.6616 (4) | 0.46453 (12) | 0.0557 (9) | |
| H9A | 0.0234 | 0.6243 | 0.4851 | 0.067* | |
| C9B | -0.2021 (4) | 0.4392 (4) | 0.32877 (10) | 0.0510 (9) | |
| H9B | -0.1431 | 0.5130 | 0.3413 | 0.061* | |
| C10A | -0.0254 (4) | 0.7053 (5) | 0.43626 (14) | 0.0724 (13) | |
| H10A | -0.1175 | 0.6969 | 0.4378 | 0.087* | |
| C10B | -0.1548 (5) | 0.3312 (4) | 0.30955 (12) | 0.0639 (11) | |
| H10B | -0.0625 | 0.3312 | 0.3097 | 0.077* | |

| | | | | |
|------|-------------|------------|--------------|-------------|
| C11A | 0.0245 (5) | 0.7610 (5) | 0.40576 (14) | 0.0820 (15) |
| H11A | -0.0344 | 0.7901 | 0.3873 | 0.098* |
| C11B | -0.2432 (5) | 0.2229 (4) | 0.29000 (13) | 0.0687 (12) |
| H11B | -0.2092 | 0.1523 | 0.2766 | 0.082* |
| C12A | 0.1586 (5) | 0.7740 (5) | 0.40212 (12) | 0.0701 (12) |
| H12A | 0.1922 | 0.8116 | 0.3815 | 0.084* |
| C12B | -0.3785 (5) | 0.2177 (4) | 0.29002 (11) | 0.0597 (10) |
| H12B | -0.4376 | 0.1450 | 0.2768 | 0.072* |
| C13A | 0.2158 (3) | 0.7433 (3) | 0.54657 (9) | 0.0383 (7) |
| C13B | -0.3150 (4) | 0.7404 (3) | 0.32166 (10) | 0.0472 (8) |
| C14A | 0.1567 (3) | 0.8551 (3) | 0.54164 (10) | 0.0438 (8) |
| H14A | 0.1977 | 0.9204 | 0.5289 | 0.053* |
| C14B | -0.2340 (5) | 0.7157 (4) | 0.29249 (12) | 0.0696 (12) |
| H14B | -0.2553 | 0.6394 | 0.2741 | 0.084* |
| C15B | -0.1219 (5) | 0.8049 (5) | 0.29082 (15) | 0.0800 (14) |
| H15B | -0.0675 | 0.7865 | 0.2713 | 0.096* |
| C16A | -0.0278 (3) | 0.7719 (4) | 0.57473 (10) | 0.0498 (8) |
| C16B | -0.0875 (4) | 0.9188 (4) | 0.31660 (13) | 0.0625 (11) |
| C15A | 0.0347 (4) | 0.8687 (4) | 0.55607 (11) | 0.0498 (8) |
| H15A | -0.0053 | 0.9442 | 0.5531 | 0.060* |
| C17A | 0.0366 (4) | 0.6634 (4) | 0.58047 (11) | 0.0524 (9) |
| H17A | -0.0022 | 0.5996 | 0.5941 | 0.063* |
| C18A | 0.1573 (3) | 0.6482 (4) | 0.56627 (10) | 0.0468 (8) |
| H18A | 0.1988 | 0.5742 | 0.5700 | 0.056* |
| C17B | -0.1718 (4) | 0.9422 (4) | 0.34562 (11) | 0.0576 (10) |
| H17B | -0.1522 | 1.0200 | 0.3634 | 0.069* |
| C18B | -0.2816 (4) | 0.8545 (4) | 0.34848 (10) | 0.0496 (8) |
| H18B | -0.3342 | 0.8714 | 0.3686 | 0.060* |
| C19A | -0.1661 (4) | 0.7826 (5) | 0.58720 (14) | 0.0724 (12) |
| H19A | -0.2325 | 0.7221 | 0.5695 | 0.109* |
| H19B | -0.1857 | 0.8715 | 0.5868 | 0.109* |
| H19C | -0.1679 | 0.7608 | 0.6133 | 0.109* |
| C19B | 0.0391 (5) | 1.0120 (6) | 0.31561 (17) | 0.0941 (17) |
| H19D | 0.0805 | 0.9835 | 0.2928 | 0.141* |
| H19E | 0.1004 | 1.0120 | 0.3388 | 0.141* |
| H19F | 0.0173 | 1.0999 | 0.3146 | 0.141* |
| N1A | 0.3318 (2) | 0.7197 (3) | 0.52688 (7) | 0.0366 (6) |
| N1B | -0.4299 (3) | 0.6477 (3) | 0.32421 (8) | 0.0446 (7) |
| N2A | 0.6351 (3) | 0.8495 (3) | 0.49127 (9) | 0.0523 (7) |
| N2B | -0.7813 (4) | 0.5448 (5) | 0.32615 (11) | 0.0795 (12) |
| O1A | 0.4557 (3) | 0.7450 (3) | 0.40438 (7) | 0.0596 (7) |
| O1B | -0.6620 (3) | 0.2552 (3) | 0.30708 (10) | 0.0868 (10) |
| O2A | 0.2734 (2) | 0.5022 (2) | 0.49147 (7) | 0.0484 (6) |
| H2A | 0.3352 | 0.4785 | 0.5049 | 0.073* |
| O3B | -0.5922 (3) | 0.4394 (3) | 0.38327 (7) | 0.0563 (7) |
| H3B | -0.6712 | 0.4060 | 0.3816 | 0.084* |
| O3A | 0.5110 (3) | 0.5622 (3) | 0.46113 (7) | 0.0504 (6) |
| H3A | 0.4885 | 0.5238 | 0.4387 | 0.076* |
| O2B | -0.3484 (2) | 0.5870 (2) | 0.38529 (6) | 0.0448 (5) |

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|------|---------------|--------------|--------------|-------------|
| H2B | -0.3891 | 0.6430 | 0.3960 | 0.067* |
| O4A | 0.6824 (3) | 0.9511 (3) | 0.51445 (9) | 0.0707 (8) |
| O4B | -0.8460 (3) | 0.6210 (5) | 0.30993 (11) | 0.1101 (14) |
| O5A | 0.6916 (3) | 0.8092 (3) | 0.46307 (9) | 0.0767 (9) |
| O5B | -0.8371 (3) | 0.4541 (5) | 0.34225 (12) | 0.0987 (12) |
| S1 | 0.50842 (9) | 0.92446 (9) | 0.56943 (3) | 0.0504 (2) |
| S2 | -0.62570 (16) | 0.78016 (15) | 0.29113 (4) | 0.0902 (4) |
| C1D | 0.6383 (3) | 0.8124 (3) | 1.01223 (9) | 0.0405 (7) |
| C2D | 0.6902 (3) | 0.9583 (3) | 1.03256 (9) | 0.0422 (7) |
| C3D | 0.7771 (3) | 1.0098 (3) | 1.00404 (9) | 0.0417 (7) |
| C4D | 0.7760 (3) | 0.9169 (3) | 0.97153 (9) | 0.0379 (7) |
| C5D | 0.7641 (5) | 0.8362 (5) | 0.89118 (11) | 0.0664 (11) |
| H10 | 0.6699 | 0.8350 | 0.8935 | 0.100* |
| H11 | 0.7871 | 0.7483 | 0.8912 | 0.100* |
| H12 | 0.7833 | 0.8668 | 0.8671 | 0.100* |
| C6D | 0.7756 (3) | 0.9422 (4) | 1.07046 (10) | 0.0475 (8) |
| C7D | 0.7847 (4) | 0.8024 (4) | 1.07031 (10) | 0.0513 (9) |
| C8D | 0.7035 (3) | 0.7262 (4) | 1.03883 (10) | 0.0451 (8) |
| C9D | 0.6886 (4) | 0.5892 (4) | 1.03508 (12) | 0.0578 (10) |
| H9D | 0.6333 | 0.5371 | 1.0141 | 0.069* |
| C10D | 0.7581 (5) | 0.5323 (5) | 1.06343 (15) | 0.0710 (13) |
| H10D | 0.7495 | 0.4405 | 1.0614 | 0.085* |
| C11D | 0.8397 (5) | 0.6084 (5) | 1.09461 (15) | 0.0810 (15) |
| H11D | 0.8860 | 0.5671 | 1.1131 | 0.097* |
| C12D | 0.8538 (5) | 0.7432 (5) | 1.09896 (12) | 0.0701 (12) |
| H12D | 0.9078 | 0.7944 | 1.1203 | 0.084* |
| C13D | 0.7059 (3) | 0.6746 (3) | 0.95415 (9) | 0.0404 (7) |
| C14D | 0.8240 (3) | 0.6224 (3) | 0.95949 (10) | 0.0461 (8) |
| H14D | 0.9012 | 0.6754 | 0.9726 | 0.055* |
| C15D | 0.8284 (4) | 0.4909 (4) | 0.94538 (11) | 0.0512 (9) |
| H15D | 0.9088 | 0.4560 | 0.9493 | 0.061* |
| C16D | 0.7156 (4) | 0.4109 (4) | 0.92571 (11) | 0.0533 (9) |
| C17D | 0.5980 (4) | 0.4670 (4) | 0.91898 (11) | 0.0539 (9) |
| H17D | 0.5220 | 0.4152 | 0.9047 | 0.065* |
| C18D | 0.5921 (4) | 0.5971 (4) | 0.93292 (10) | 0.0477 (8) |
| H18D | 0.5128 | 0.6331 | 0.9282 | 0.057* |
| C19D | 0.7187 (5) | 0.2650 (4) | 0.91262 (15) | 0.0778 (13) |
| H19J | 0.6730 | 0.2137 | 0.9298 | 0.117* |
| H19K | 0.8104 | 0.2483 | 0.9135 | 0.117* |
| H19L | 0.6744 | 0.2404 | 0.8863 | 0.117* |
| N1D | 0.6938 (3) | 0.8045 (3) | 0.97352 (7) | 0.0379 (6) |
| N2D | 0.8481 (3) | 1.1362 (3) | 1.00990 (9) | 0.0531 (7) |
| O1D | 0.8237 (3) | 1.0340 (3) | 1.09637 (7) | 0.0595 (7) |
| O2D | 0.5001 (2) | 0.7761 (3) | 1.00695 (8) | 0.0506 (6) |
| H2D | 0.4646 | 0.8271 | 0.9942 | 0.076* |
| O3D | 0.5852 (2) | 1.0344 (3) | 1.03922 (7) | 0.0498 (6) |
| H3D | 0.5644 | 1.0295 | 1.0613 | 0.075* |
| O4D | 0.9295 (3) | 1.1659 (3) | 0.98715 (9) | 0.0699 (8) |
| O5D | 0.8301 (3) | 1.2162 (3) | 1.03853 (9) | 0.0744 (8) |

| | | | | |
|------|--------------|--------------|--------------|-------------|
| S4 | 0.86152 (10) | 0.94568 (10) | 0.93187 (3) | 0.0532 (2) |
| C1C | 0.3302 (3) | 0.7827 (3) | 0.84914 (9) | 0.0427 (7) |
| C2C | 0.4071 (4) | 0.9207 (4) | 0.84560 (10) | 0.0465 (8) |
| C3C | 0.3014 (4) | 0.9892 (4) | 0.82676 (10) | 0.0531 (9) |
| C4C | 0.1834 (4) | 0.9043 (4) | 0.81611 (10) | 0.0535 (9) |
| C6C | 0.5135 (4) | 0.8886 (4) | 0.81767 (10) | 0.0505 (9) |
| C7C | 0.5025 (3) | 0.7446 (4) | 0.80862 (9) | 0.0457 (8) |
| C8C | 0.4066 (3) | 0.6835 (3) | 0.82799 (9) | 0.0436 (8) |
| C9C | 0.3899 (4) | 0.5475 (4) | 0.82721 (11) | 0.0593 (10) |
| H9C | 0.3254 | 0.5054 | 0.8404 | 0.071* |
| C10C | 0.4731 (5) | 0.4761 (5) | 0.80591 (13) | 0.0739 (12) |
| H10C | 0.4655 | 0.3847 | 0.8054 | 0.089* |
| C11C | 0.5665 (5) | 0.5370 (5) | 0.78551 (13) | 0.0716 (12) |
| H11C | 0.6191 | 0.4863 | 0.7709 | 0.086* |
| C12C | 0.5829 (4) | 0.6709 (5) | 0.78655 (11) | 0.0595 (10) |
| H12C | 0.6462 | 0.7124 | 0.7728 | 0.071* |
| C13C | 0.0970 (3) | 0.6701 (4) | 0.82203 (10) | 0.0499 (9) |
| C14C | 0.0850 (5) | 0.5777 (6) | 0.78867 (13) | 0.0871 (16) |
| H14C | 0.1385 | 0.5917 | 0.7689 | 0.105* |
| C15C | -0.0070 (6) | 0.4646 (6) | 0.78502 (16) | 0.0958 (18) |
| H15C | -0.0155 | 0.4032 | 0.7624 | 0.115* |
| C16C | -0.0872 (5) | 0.4396 (5) | 0.81398 (15) | 0.0755 (13) |
| C17C | -0.0760 (4) | 0.5352 (4) | 0.84590 (12) | 0.0609 (10) |
| H17C | -0.1322 | 0.5235 | 0.8651 | 0.073* |
| C18C | 0.0158 (4) | 0.6483 (4) | 0.85049 (11) | 0.0529 (9) |
| H18C | 0.0228 | 0.7102 | 0.8730 | 0.063* |
| C19C | -0.1815 (6) | 0.3122 (6) | 0.8111 (2) | 0.116 (2) |
| H19G | -0.1368 | 0.2500 | 0.8242 | 0.174* |
| H19H | -0.2078 | 0.2761 | 0.7841 | 0.174* |
| H19I | -0.2599 | 0.3293 | 0.8233 | 0.174* |
| N1C | 0.1946 (3) | 0.7845 (3) | 0.82644 (8) | 0.0491 (7) |
| N2C | 0.3258 (4) | 1.1234 (4) | 0.82451 (11) | 0.0724 (10) |
| O1C | 0.5932 (3) | 0.9711 (3) | 0.80765 (10) | 0.0791 (9) |
| O2C | 0.3173 (2) | 0.7522 (2) | 0.88630 (6) | 0.0471 (6) |
| H2C | 0.2759 | 0.8057 | 0.8979 | 0.071* |
| O3C | 0.4712 (3) | 0.9862 (3) | 0.88256 (7) | 0.0550 (6) |
| H3C | 0.5082 | 1.0599 | 0.8801 | 0.083* |
| O4C | 0.2372 (4) | 1.1761 (4) | 0.80671 (12) | 0.1060 (13) |
| O5C | 0.4335 (4) | 1.1883 (3) | 0.84059 (12) | 0.0951 (11) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|--------------|-------------|
| C5C | 0.057 (4) | 0.118 (7) | 0.130 (7) | -0.005 (5) | -0.020 (4) | 0.070 (6) |
| S3 | 0.0530 (11) | 0.075 (2) | 0.0635 (9) | 0.0199 (12) | -0.0072 (7) | 0.0244 (9) |
| C1 | 0.070 (9) | 0.125 (12) | 0.131 (12) | 0.003 (10) | -0.026 (9) | 0.063 (10) |
| S3' | 0.061 (5) | 0.084 (9) | 0.071 (4) | 0.014 (6) | -0.021 (3) | 0.029 (4) |
| C1A | 0.0416 (17) | 0.0419 (19) | 0.0355 (16) | 0.0149 (14) | -0.0010 (13) | 0.0056 (13) |
| C1B | 0.0479 (18) | 0.0448 (19) | 0.0313 (16) | 0.0139 (14) | 0.0000 (13) | 0.0085 (13) |
| C2A | 0.0441 (18) | 0.049 (2) | 0.0339 (16) | 0.0247 (15) | 0.0046 (13) | 0.0079 (14) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C2B | 0.0484 (19) | 0.061 (2) | 0.0351 (17) | 0.0101 (16) | 0.0030 (14) | 0.0077 (15) |
| C3A | 0.0396 (17) | 0.0465 (19) | 0.0388 (17) | 0.0160 (14) | 0.0046 (13) | 0.0115 (14) |
| C3B | 0.047 (2) | 0.076 (3) | 0.046 (2) | 0.0205 (19) | -0.0019 (16) | 0.0096 (18) |
| C4A | 0.0402 (17) | 0.0377 (17) | 0.0368 (16) | 0.0163 (13) | 0.0035 (13) | 0.0107 (13) |
| C4B | 0.069 (2) | 0.064 (2) | 0.0310 (17) | 0.026 (2) | -0.0041 (16) | 0.0033 (16) |
| C5A | 0.076 (3) | 0.076 (3) | 0.0365 (19) | 0.015 (2) | 0.0023 (18) | 0.0005 (18) |
| C5B | 0.156 (11) | 0.073 (10) | 0.080 (11) | 0.032 (8) | -0.044 (9) | 0.009 (9) |
| C2 | 0.147 (5) | 0.059 (5) | 0.070 (5) | 0.007 (4) | -0.034 (3) | 0.028 (4) |
| C6A | 0.060 (2) | 0.050 (2) | 0.0351 (17) | 0.0277 (17) | 0.0061 (16) | 0.0086 (15) |
| C6B | 0.052 (2) | 0.059 (2) | 0.0442 (19) | 0.0011 (18) | -0.0030 (16) | 0.0087 (16) |
| C7A | 0.061 (2) | 0.058 (2) | 0.0409 (19) | 0.0327 (18) | -0.0027 (16) | 0.0042 (16) |
| C7B | 0.060 (2) | 0.045 (2) | 0.0383 (17) | 0.0141 (16) | 0.0007 (15) | 0.0113 (15) |
| C8A | 0.0475 (19) | 0.046 (2) | 0.0382 (17) | 0.0197 (15) | -0.0056 (14) | -0.0029 (14) |
| C8B | 0.0508 (19) | 0.0456 (19) | 0.0291 (15) | 0.0136 (15) | 0.0019 (13) | 0.0114 (13) |
| C9A | 0.050 (2) | 0.059 (2) | 0.055 (2) | 0.0192 (17) | -0.0079 (17) | -0.0032 (17) |
| C9B | 0.056 (2) | 0.057 (2) | 0.0424 (19) | 0.0153 (17) | 0.0079 (16) | 0.0082 (16) |
| C10A | 0.053 (2) | 0.076 (3) | 0.082 (3) | 0.031 (2) | -0.020 (2) | -0.012 (2) |
| C10B | 0.071 (3) | 0.073 (3) | 0.054 (2) | 0.031 (2) | 0.013 (2) | 0.008 (2) |
| C11A | 0.087 (3) | 0.100 (4) | 0.064 (3) | 0.057 (3) | -0.018 (2) | 0.012 (3) |
| C11B | 0.093 (3) | 0.061 (3) | 0.060 (3) | 0.038 (2) | 0.015 (2) | 0.010 (2) |
| C12A | 0.085 (3) | 0.083 (3) | 0.051 (2) | 0.046 (2) | -0.005 (2) | 0.018 (2) |
| C12B | 0.083 (3) | 0.047 (2) | 0.048 (2) | 0.014 (2) | 0.0016 (19) | 0.0047 (17) |
| C13A | 0.0371 (16) | 0.0443 (18) | 0.0353 (16) | 0.0112 (14) | 0.0052 (13) | 0.0061 (13) |
| C13B | 0.064 (2) | 0.0405 (19) | 0.0389 (18) | 0.0115 (16) | 0.0007 (16) | 0.0121 (14) |
| C14A | 0.0467 (19) | 0.0424 (19) | 0.0449 (18) | 0.0088 (15) | 0.0101 (15) | 0.0099 (14) |
| C14B | 0.109 (4) | 0.048 (2) | 0.054 (2) | 0.007 (2) | 0.029 (2) | 0.0027 (18) |
| C15B | 0.101 (4) | 0.071 (3) | 0.082 (3) | 0.018 (3) | 0.044 (3) | 0.031 (3) |
| C16A | 0.0438 (19) | 0.063 (2) | 0.0443 (19) | 0.0110 (17) | 0.0106 (15) | 0.0050 (16) |
| C16B | 0.069 (3) | 0.058 (3) | 0.064 (3) | 0.010 (2) | 0.001 (2) | 0.028 (2) |
| C15A | 0.048 (2) | 0.051 (2) | 0.054 (2) | 0.0198 (16) | 0.0075 (16) | 0.0055 (16) |
| C17A | 0.052 (2) | 0.058 (2) | 0.050 (2) | 0.0025 (17) | 0.0115 (16) | 0.0175 (17) |
| C18A | 0.0486 (19) | 0.047 (2) | 0.0482 (19) | 0.0121 (15) | 0.0079 (16) | 0.0146 (15) |
| C17B | 0.070 (3) | 0.050 (2) | 0.048 (2) | 0.0104 (19) | -0.0110 (19) | 0.0051 (17) |
| C18B | 0.062 (2) | 0.051 (2) | 0.0365 (18) | 0.0173 (18) | -0.0005 (16) | 0.0034 (15) |
| C19A | 0.050 (2) | 0.095 (3) | 0.075 (3) | 0.015 (2) | 0.020 (2) | 0.009 (2) |
| C19B | 0.085 (3) | 0.103 (4) | 0.098 (4) | 0.000 (3) | 0.003 (3) | 0.048 (3) |
| N1A | 0.0367 (14) | 0.0413 (15) | 0.0333 (13) | 0.0112 (11) | 0.0042 (11) | 0.0051 (11) |
| N1B | 0.0565 (17) | 0.0438 (16) | 0.0341 (14) | 0.0151 (13) | -0.0032 (12) | 0.0069 (12) |
| N2A | 0.0452 (17) | 0.064 (2) | 0.0539 (18) | 0.0153 (15) | 0.0097 (15) | 0.0206 (16) |
| N2B | 0.052 (2) | 0.126 (4) | 0.062 (2) | 0.028 (2) | -0.0060 (18) | 0.015 (2) |
| O1A | 0.0809 (18) | 0.0702 (18) | 0.0390 (13) | 0.0384 (14) | 0.0152 (13) | 0.0165 (12) |
| O1B | 0.069 (2) | 0.088 (2) | 0.089 (2) | -0.0166 (18) | 0.0012 (17) | -0.0097 (18) |
| O2A | 0.0506 (14) | 0.0431 (14) | 0.0499 (14) | 0.0124 (11) | -0.0046 (11) | 0.0044 (10) |
| O3B | 0.0530 (14) | 0.0763 (18) | 0.0403 (13) | 0.0035 (13) | 0.0076 (11) | 0.0151 (12) |
| O3A | 0.0618 (15) | 0.0618 (16) | 0.0353 (12) | 0.0368 (13) | 0.0058 (11) | 0.0083 (11) |
| O2B | 0.0478 (13) | 0.0540 (14) | 0.0326 (11) | 0.0169 (11) | -0.0012 (9) | 0.0018 (10) |
| O4A | 0.0630 (18) | 0.078 (2) | 0.0673 (18) | -0.0103 (15) | 0.0089 (14) | 0.0168 (16) |
| O4B | 0.071 (2) | 0.181 (4) | 0.091 (3) | 0.062 (2) | -0.0104 (19) | 0.040 (3) |
| O5A | 0.0607 (17) | 0.100 (2) | 0.079 (2) | 0.0227 (16) | 0.0366 (16) | 0.0176 (17) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| O5B | 0.0479 (18) | 0.145 (4) | 0.106 (3) | 0.007 (2) | 0.0063 (18) | 0.036 (3) |
| S1 | 0.0530 (5) | 0.0503 (5) | 0.0445 (5) | 0.0062 (4) | -0.0008 (4) | 0.0011 (4) |
| S2 | 0.1209 (11) | 0.0944 (10) | 0.0656 (7) | 0.0605 (8) | -0.0119 (7) | 0.0250 (7) |
| C1D | 0.0443 (18) | 0.0436 (19) | 0.0372 (17) | 0.0137 (14) | 0.0080 (14) | 0.0093 (13) |
| C2D | 0.0489 (19) | 0.047 (2) | 0.0342 (16) | 0.0232 (15) | 0.0019 (14) | 0.0062 (14) |
| C3D | 0.0431 (18) | 0.0421 (19) | 0.0405 (17) | 0.0116 (14) | -0.0010 (14) | 0.0083 (14) |
| C4D | 0.0376 (16) | 0.0435 (18) | 0.0362 (16) | 0.0175 (14) | 0.0035 (13) | 0.0093 (13) |
| C5D | 0.084 (3) | 0.085 (3) | 0.0343 (19) | 0.026 (2) | 0.0094 (19) | 0.0071 (19) |
| C6D | 0.0464 (19) | 0.066 (2) | 0.0342 (17) | 0.0251 (17) | 0.0038 (14) | 0.0061 (16) |
| C7D | 0.055 (2) | 0.067 (2) | 0.0415 (19) | 0.0314 (18) | 0.0075 (16) | 0.0180 (17) |
| C8D | 0.0483 (19) | 0.053 (2) | 0.0415 (18) | 0.0206 (16) | 0.0113 (15) | 0.0167 (15) |
| C9D | 0.065 (2) | 0.054 (2) | 0.065 (2) | 0.0217 (18) | 0.0222 (19) | 0.0232 (19) |
| C10D | 0.082 (3) | 0.064 (3) | 0.086 (3) | 0.038 (2) | 0.030 (3) | 0.040 (2) |
| C11D | 0.089 (3) | 0.094 (4) | 0.080 (3) | 0.053 (3) | 0.013 (3) | 0.046 (3) |
| C12D | 0.074 (3) | 0.095 (4) | 0.052 (2) | 0.045 (2) | 0.000 (2) | 0.022 (2) |
| C13D | 0.0440 (18) | 0.0398 (18) | 0.0384 (17) | 0.0104 (14) | 0.0043 (14) | 0.0052 (13) |
| C14D | 0.0437 (18) | 0.045 (2) | 0.0478 (19) | 0.0084 (15) | 0.0036 (15) | 0.0016 (15) |
| C15D | 0.051 (2) | 0.052 (2) | 0.053 (2) | 0.0197 (17) | 0.0053 (16) | 0.0043 (16) |
| C16D | 0.066 (2) | 0.044 (2) | 0.050 (2) | 0.0116 (18) | 0.0083 (18) | 0.0019 (16) |
| C17D | 0.054 (2) | 0.050 (2) | 0.052 (2) | 0.0018 (17) | -0.0026 (17) | 0.0001 (17) |
| C18D | 0.0448 (19) | 0.049 (2) | 0.049 (2) | 0.0120 (15) | -0.0014 (15) | 0.0043 (16) |
| C19D | 0.096 (3) | 0.048 (2) | 0.085 (3) | 0.014 (2) | 0.000 (3) | -0.002 (2) |
| N1D | 0.0417 (14) | 0.0375 (15) | 0.0352 (14) | 0.0085 (11) | 0.0055 (11) | 0.0041 (11) |
| N2D | 0.0568 (19) | 0.0457 (18) | 0.0555 (19) | 0.0094 (14) | -0.0030 (15) | 0.0089 (15) |
| O1D | 0.0647 (16) | 0.0750 (19) | 0.0386 (13) | 0.0288 (14) | -0.0048 (12) | -0.0017 (13) |
| O2D | 0.0445 (13) | 0.0559 (16) | 0.0567 (15) | 0.0154 (11) | 0.0105 (11) | 0.0164 (12) |
| O3D | 0.0569 (14) | 0.0604 (16) | 0.0372 (12) | 0.0334 (12) | 0.0022 (11) | 0.0046 (11) |
| O4D | 0.0740 (19) | 0.0618 (18) | 0.0677 (18) | -0.0120 (15) | 0.0015 (15) | 0.0127 (14) |
| O5D | 0.096 (2) | 0.0473 (16) | 0.074 (2) | 0.0147 (15) | 0.0043 (17) | -0.0111 (14) |
| S4 | 0.0569 (5) | 0.0618 (6) | 0.0452 (5) | 0.0105 (4) | 0.0132 (4) | 0.0156 (4) |
| C1C | 0.0469 (18) | 0.053 (2) | 0.0286 (15) | 0.0117 (15) | -0.0013 (13) | 0.0077 (14) |
| C2C | 0.055 (2) | 0.048 (2) | 0.0377 (17) | 0.0136 (16) | -0.0001 (15) | 0.0109 (14) |
| C3C | 0.065 (2) | 0.054 (2) | 0.0433 (19) | 0.0186 (19) | 0.0000 (17) | 0.0135 (16) |
| C4C | 0.062 (2) | 0.071 (3) | 0.0363 (18) | 0.033 (2) | 0.0064 (16) | 0.0172 (17) |
| C6C | 0.053 (2) | 0.060 (2) | 0.0411 (19) | 0.0109 (18) | 0.0034 (16) | 0.0167 (16) |
| C7C | 0.0479 (19) | 0.056 (2) | 0.0331 (16) | 0.0120 (16) | -0.0023 (14) | 0.0062 (14) |
| C8C | 0.0475 (19) | 0.047 (2) | 0.0354 (17) | 0.0100 (15) | -0.0030 (14) | 0.0053 (14) |
| C9C | 0.075 (3) | 0.056 (2) | 0.046 (2) | 0.010 (2) | 0.0071 (19) | 0.0058 (17) |
| C10C | 0.102 (4) | 0.061 (3) | 0.061 (3) | 0.025 (2) | 0.011 (3) | 0.004 (2) |
| C11C | 0.086 (3) | 0.075 (3) | 0.057 (2) | 0.039 (2) | 0.011 (2) | -0.003 (2) |
| C12C | 0.057 (2) | 0.078 (3) | 0.045 (2) | 0.019 (2) | 0.0059 (17) | 0.0065 (19) |
| C13C | 0.0452 (19) | 0.070 (2) | 0.0357 (18) | 0.0179 (17) | -0.0005 (15) | 0.0069 (16) |
| C14C | 0.067 (3) | 0.130 (5) | 0.051 (3) | -0.007 (3) | 0.009 (2) | -0.017 (3) |
| C15C | 0.086 (4) | 0.106 (4) | 0.075 (3) | -0.006 (3) | -0.002 (3) | -0.033 (3) |
| C16C | 0.060 (3) | 0.077 (3) | 0.081 (3) | 0.002 (2) | -0.018 (2) | 0.009 (3) |
| C17C | 0.054 (2) | 0.080 (3) | 0.052 (2) | 0.010 (2) | 0.0001 (18) | 0.026 (2) |
| C18C | 0.055 (2) | 0.063 (2) | 0.0428 (19) | 0.0152 (18) | 0.0033 (16) | 0.0103 (17) |
| C19C | 0.105 (4) | 0.089 (4) | 0.136 (6) | -0.017 (3) | -0.033 (4) | 0.019 (4) |
| N1C | 0.0429 (16) | 0.067 (2) | 0.0382 (15) | 0.0101 (14) | 0.0003 (12) | 0.0143 (14) |

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|-----|-------------|-------------|-------------|--------------|--------------|-------------|
| N2C | 0.093 (3) | 0.061 (2) | 0.071 (2) | 0.029 (2) | 0.004 (2) | 0.0287 (19) |
| O1C | 0.083 (2) | 0.073 (2) | 0.088 (2) | 0.0013 (17) | 0.0293 (18) | 0.0309 (17) |
| O2C | 0.0581 (14) | 0.0550 (15) | 0.0312 (11) | 0.0168 (11) | 0.0055 (10) | 0.0096 (10) |
| O3C | 0.0694 (16) | 0.0526 (15) | 0.0407 (13) | 0.0047 (12) | -0.0028 (12) | 0.0089 (11) |
| O4C | 0.131 (3) | 0.083 (2) | 0.111 (3) | 0.041 (2) | -0.018 (2) | 0.042 (2) |
| O5C | 0.108 (3) | 0.061 (2) | 0.111 (3) | -0.0012 (19) | -0.017 (2) | 0.0289 (19) |

Geometric parameters (Å, °)

| | | | |
|----------|------------|-----------|-----------|
| C5C—S3 | 1.757 (8) | N2A—O4A | 1.241 (4) |
| S3—C4C | 1.737 (4) | N2B—O4B | 1.233 (5) |
| C1—S3' | 1.755 (10) | N2B—O5B | 1.253 (5) |
| S3'—C4C | 1.909 (10) | C1D—O2D | 1.384 (4) |
| C1A—O2A | 1.379 (4) | C1D—N1D | 1.503 (4) |
| C1A—N1A | 1.500 (4) | C1D—C8D | 1.516 (4) |
| C1A—C8A | 1.509 (4) | C1D—C2D | 1.571 (5) |
| C1A—C2A | 1.578 (5) | C2D—O3D | 1.412 (4) |
| C1B—O2B | 1.382 (4) | C2D—C3D | 1.494 (5) |
| C1B—N1B | 1.508 (4) | C2D—C6D | 1.533 (4) |
| C1B—C8B | 1.513 (5) | C3D—C4D | 1.376 (5) |
| C1B—C2B | 1.551 (5) | C3D—N2D | 1.382 (5) |
| C2A—O3A | 1.419 (4) | C4D—N1D | 1.349 (4) |
| C2A—C3A | 1.485 (5) | C4D—S4 | 1.736 (3) |
| C2A—C6A | 1.541 (4) | C5D—S4 | 1.800 (4) |
| C2B—O3B | 1.423 (4) | C6D—O1D | 1.226 (4) |
| C2B—C3B | 1.489 (5) | C6D—C7D | 1.451 (5) |
| C2B—C6B | 1.547 (5) | C7D—C8D | 1.379 (5) |
| C3A—C4A | 1.381 (4) | C7D—C12D | 1.400 (5) |
| C3A—N2A | 1.387 (4) | C8D—C9D | 1.384 (5) |
| C3B—C4B | 1.375 (6) | C9D—C10D | 1.381 (6) |
| C3B—N2B | 1.384 (5) | C10D—C11D | 1.374 (7) |
| C4A—N1A | 1.348 (4) | C11D—C12D | 1.361 (7) |
| C4A—S1 | 1.735 (3) | C13D—C14D | 1.371 (5) |
| C4B—N1B | 1.354 (5) | C13D—C18D | 1.391 (5) |
| C4B—S2 | 1.731 (4) | C13D—N1D | 1.439 (4) |
| C5A—S1 | 1.795 (4) | C14D—C15D | 1.383 (5) |
| C5B—S2 | 1.45 (6) | C15D—C16D | 1.376 (5) |
| C2—S2 | 1.809 (10) | C16D—C17D | 1.392 (5) |
| C6A—O1A | 1.218 (4) | C16D—C19D | 1.514 (6) |
| C6A—C7A | 1.455 (5) | C17D—C18D | 1.372 (5) |
| C6B—O1B | 1.199 (4) | N2D—O4D | 1.235 (4) |
| C6B—C7B | 1.462 (5) | N2D—O5D | 1.247 (4) |
| C7A—C8A | 1.385 (5) | C1C—O2C | 1.380 (4) |
| C7A—C12A | 1.385 (5) | C1C—N1C | 1.508 (4) |
| C7B—C8B | 1.372 (5) | C1C—C8C | 1.512 (5) |
| C7B—C12B | 1.391 (5) | C1C—C2C | 1.556 (5) |
| C8A—C9A | 1.389 (5) | C2C—O3C | 1.415 (4) |
| C8B—C9B | 1.379 (5) | C2C—C3C | 1.489 (5) |
| C9A—C10A | 1.379 (6) | C2C—C6C | 1.556 (5) |
| C9B—C10B | 1.381 (5) | C3C—C4C | 1.371 (6) |

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|-------------|------------|--------------|-----------|
| C10A—C11A | 1.378 (7) | C3C—N2C | 1.384 (5) |
| C10B—C11B | 1.386 (6) | C4C—N1C | 1.341 (5) |
| C11A—C12A | 1.364 (7) | C6C—O1C | 1.204 (4) |
| C11B—C12B | 1.359 (6) | C6C—C7C | 1.460 (5) |
| C13A—C18A | 1.373 (5) | C7C—C8C | 1.370 (5) |
| C13A—C14A | 1.380 (5) | C7C—C12C | 1.395 (5) |
| C13A—N1A | 1.445 (4) | C8C—C9C | 1.384 (5) |
| C13B—C18B | 1.380 (5) | C9C—C10C | 1.389 (6) |
| C13B—C14B | 1.382 (5) | C10C—C11C | 1.374 (7) |
| C13B—N1B | 1.424 (5) | C11C—C12C | 1.361 (6) |
| C14A—C15A | 1.391 (5) | C13C—C18C | 1.370 (5) |
| C14B—C15B | 1.375 (6) | C13C—C14C | 1.381 (6) |
| C15B—C16B | 1.360 (7) | C13C—N1C | 1.417 (5) |
| C16A—C15A | 1.380 (5) | C14C—C15C | 1.377 (7) |
| C16A—C17A | 1.385 (5) | C15C—C16C | 1.384 (7) |
| C16A—C19A | 1.515 (5) | C16C—C17C | 1.365 (7) |
| C16B—C17B | 1.400 (6) | C16C—C19C | 1.504 (7) |
| C16B—C19B | 1.504 (6) | C17C—C18C | 1.372 (6) |
| C17A—C18A | 1.378 (5) | N2C—O5C | 1.237 (5) |
| C17B—C18B | 1.358 (5) | N2C—O4C | 1.251 (5) |
| N2A—O5A | 1.232 (4) | | |
| | | | |
| C4C—S3—C5C | 107.6 (3) | C4B—S2—C2 | 106.8 (3) |
| C1—S3'—C4C | 122.1 (12) | O2D—C1D—N1D | 111.0 (3) |
| O2A—C1A—N1A | 111.3 (2) | O2D—C1D—C8D | 109.7 (3) |
| O2A—C1A—C8A | 109.3 (3) | N1D—C1D—C8D | 111.5 (3) |
| N1A—C1A—C8A | 111.8 (3) | O2D—C1D—C2D | 115.9 (3) |
| O2A—C1A—C2A | 116.1 (3) | N1D—C1D—C2D | 104.0 (2) |
| N1A—C1A—C2A | 103.9 (2) | C8D—C1D—C2D | 104.6 (3) |
| C8A—C1A—C2A | 104.3 (3) | O3D—C2D—C3D | 112.4 (3) |
| O2B—C1B—N1B | 110.6 (3) | O3D—C2D—C6D | 113.1 (3) |
| O2B—C1B—C8B | 109.1 (3) | C3D—C2D—C6D | 110.8 (3) |
| N1B—C1B—C8B | 111.4 (3) | O3D—C2D—C1D | 113.3 (3) |
| O2B—C1B—C2B | 117.7 (3) | C3D—C2D—C1D | 102.1 (3) |
| N1B—C1B—C2B | 103.0 (3) | C6D—C2D—C1D | 104.4 (3) |
| C8B—C1B—C2B | 104.9 (3) | C4D—C3D—N2D | 124.9 (3) |
| O3A—C2A—C3A | 112.3 (3) | C4D—C3D—C2D | 111.7 (3) |
| O3A—C2A—C6A | 113.0 (3) | N2D—C3D—C2D | 123.4 (3) |
| C3A—C2A—C6A | 110.6 (3) | N1D—C4D—C3D | 111.3 (3) |
| O3A—C2A—C1A | 113.2 (3) | N1D—C4D—S4 | 124.7 (2) |
| C3A—C2A—C1A | 102.3 (2) | C3D—C4D—S4 | 123.8 (3) |
| C6A—C2A—C1A | 104.5 (3) | O1D—C6D—C7D | 127.3 (3) |
| O3B—C2B—C3B | 115.9 (3) | O1D—C6D—C2D | 124.3 (3) |
| O3B—C2B—C6B | 109.5 (3) | C7D—C6D—C2D | 108.4 (3) |
| C3B—C2B—C6B | 112.7 (3) | C8D—C7D—C12D | 120.9 (4) |
| O3B—C2B—C1B | 110.7 (3) | C8D—C7D—C6D | 111.2 (3) |
| C3B—C2B—C1B | 102.9 (3) | C12D—C7D—C6D | 127.7 (4) |
| C6B—C2B—C1B | 104.3 (3) | C7D—C8D—C9D | 120.4 (3) |
| C4A—C3A—N2A | 125.3 (3) | C7D—C8D—C1D | 111.1 (3) |

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|----------------|-----------|----------------|-------------|
| C4A—C3A—C2A | 111.3 (3) | C9D—C8D—C1D | 128.5 (3) |
| N2A—C3A—C2A | 123.4 (3) | C10D—C9D—C8D | 118.0 (4) |
| C4B—C3B—N2B | 127.3 (4) | C11D—C10D—C9D | 121.5 (4) |
| C4B—C3B—C2B | 111.5 (3) | C12D—C11D—C10D | 121.1 (4) |
| N2B—C3B—C2B | 121.2 (4) | C11D—C12D—C7D | 118.1 (4) |
| N1A—C4A—C3A | 111.7 (3) | C14D—C13D—C18D | 120.0 (3) |
| N1A—C4A—S1 | 124.5 (2) | C14D—C13D—N1D | 120.8 (3) |
| C3A—C4A—S1 | 123.7 (3) | C18D—C13D—N1D | 119.0 (3) |
| N1B—C4B—C3B | 110.3 (3) | C13D—C14D—C15D | 120.0 (3) |
| N1B—C4B—S2 | 127.2 (3) | C16D—C15D—C14D | 120.9 (3) |
| C3B—C4B—S2 | 122.4 (3) | C15D—C16D—C17D | 118.3 (3) |
| O1A—C6A—C7A | 127.5 (3) | C15D—C16D—C19D | 120.7 (4) |
| O1A—C6A—C2A | 124.4 (3) | C17D—C16D—C19D | 121.0 (4) |
| C7A—C6A—C2A | 108.0 (3) | C18D—C17D—C16D | 121.3 (3) |
| O1B—C6B—C7B | 127.3 (4) | C17D—C18D—C13D | 119.3 (3) |
| O1B—C6B—C2B | 125.0 (4) | C4D—N1D—C13D | 125.5 (3) |
| C7B—C6B—C2B | 107.6 (3) | C4D—N1D—C1D | 110.6 (3) |
| C8A—C7A—C12A | 121.5 (4) | C13D—N1D—C1D | 117.1 (2) |
| C8A—C7A—C6A | 110.9 (3) | O4D—N2D—O5D | 121.9 (3) |
| C12A—C7A—C6A | 127.4 (4) | O4D—N2D—C3D | 119.2 (3) |
| C8B—C7B—C12B | 121.4 (4) | O5D—N2D—C3D | 118.9 (3) |
| C8B—C7B—C6B | 110.9 (3) | C4D—S4—C5D | 103.93 (19) |
| C12B—C7B—C6B | 127.5 (4) | O2C—C1C—N1C | 111.3 (3) |
| C7A—C8A—C9A | 120.1 (3) | O2C—C1C—C8C | 109.1 (3) |
| C7A—C8A—C1A | 111.8 (3) | N1C—C1C—C8C | 110.9 (3) |
| C9A—C8A—C1A | 128.1 (3) | O2C—C1C—C2C | 117.8 (3) |
| C7B—C8B—C9B | 120.6 (3) | N1C—C1C—C2C | 102.4 (3) |
| C7B—C8B—C1B | 111.3 (3) | C8C—C1C—C2C | 105.0 (3) |
| C9B—C8B—C1B | 128.1 (3) | O3C—C2C—C3C | 115.0 (3) |
| C10A—C9A—C8A | 117.7 (4) | O3C—C2C—C6C | 110.4 (3) |
| C8B—C9B—C10B | 118.0 (4) | C3C—C2C—C6C | 111.7 (3) |
| C11A—C10A—C9A | 121.7 (4) | O3C—C2C—C1C | 111.2 (3) |
| C9B—C10B—C11B | 120.8 (4) | C3C—C2C—C1C | 103.6 (3) |
| C12A—C11A—C10A | 121.1 (4) | C6C—C2C—C1C | 104.2 (3) |
| C12B—C11B—C10B | 121.3 (4) | C4C—C3C—N2C | 128.4 (4) |
| C11A—C12A—C7A | 117.9 (4) | C4C—C3C—C2C | 110.6 (3) |
| C11B—C12B—C7B | 117.8 (4) | N2C—C3C—C2C | 120.7 (4) |
| C18A—C13A—C14A | 120.8 (3) | N1C—C4C—C3C | 111.3 (3) |
| C18A—C13A—N1A | 119.8 (3) | N1C—C4C—S3 | 130.8 (4) |
| C14A—C13A—N1A | 119.1 (3) | C3C—C4C—S3 | 117.9 (3) |
| C18B—C13B—C14B | 119.1 (4) | N1C—C4C—S3' | 104.9 (6) |
| C18B—C13B—N1B | 120.7 (3) | C3C—C4C—S3' | 143.7 (6) |
| C14B—C13B—N1B | 120.1 (3) | O1C—C6C—C7C | 128.1 (4) |
| C13A—C14A—C15A | 119.0 (3) | O1C—C6C—C2C | 124.1 (4) |
| C15B—C14B—C13B | 119.5 (4) | C7C—C6C—C2C | 107.7 (3) |
| C16B—C15B—C14B | 122.6 (4) | C8C—C7C—C12C | 120.9 (4) |
| C15A—C16A—C17A | 118.5 (3) | C8C—C7C—C6C | 111.1 (3) |
| C15A—C16A—C19A | 120.3 (4) | C12C—C7C—C6C | 127.8 (4) |
| C17A—C16A—C19A | 121.1 (4) | C7C—C8C—C9C | 120.8 (3) |

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| C15B—C16B—C17B | 116.8 (4) | C7C—C8C—C1C | 111.5 (3) |
| C15B—C16B—C19B | 121.9 (5) | C9C—C8C—C1C | 127.7 (3) |
| C17B—C16B—C19B | 121.2 (5) | C8C—C9C—C10C | 117.5 (4) |
| C16A—C15A—C14A | 121.0 (3) | C11C—C10C—C9C | 121.7 (4) |
| C18A—C17A—C16A | 121.2 (3) | C12C—C11C—C10C | 120.6 (4) |
| C13A—C18A—C17A | 119.5 (3) | C11C—C12C—C7C | 118.6 (4) |
| C18B—C17B—C16B | 121.9 (4) | C18C—C13C—C14C | 119.3 (4) |
| C17B—C18B—C13B | 120.1 (4) | C18C—C13C—N1C | 121.3 (3) |
| C4A—N1A—C13A | 126.1 (3) | C14C—C13C—N1C | 119.4 (4) |
| C4A—N1A—C1A | 110.4 (2) | C15C—C14C—C13C | 119.2 (4) |
| C13A—N1A—C1A | 116.9 (2) | C14C—C15C—C16C | 122.1 (5) |
| C4B—N1B—C13B | 127.9 (3) | C17C—C16C—C15C | 117.0 (4) |
| C4B—N1B—C1B | 111.0 (3) | C17C—C16C—C19C | 120.9 (5) |
| C13B—N1B—C1B | 119.4 (3) | C15C—C16C—C19C | 122.1 (5) |
| O5A—N2A—O4A | 122.3 (3) | C16C—C17C—C18C | 122.1 (4) |
| O5A—N2A—C3A | 119.0 (3) | C13C—C18C—C17C | 120.2 (4) |
| O4A—N2A—C3A | 118.7 (3) | C4C—N1C—C13C | 129.8 (3) |
| O4B—N2B—O5B | 122.0 (4) | C4C—N1C—C1C | 111.4 (3) |
| O4B—N2B—C3B | 119.8 (5) | C13C—N1C—C1C | 118.6 (3) |
| O5B—N2B—C3B | 118.1 (4) | O5C—N2C—O4C | 121.8 (4) |
| C4A—S1—C5A | 103.90 (18) | O5C—N2C—C3C | 119.3 (4) |
| C5B—S2—C4B | 122 (2) | O4C—N2C—C3C | 119.0 (4) |
| O2A—C1A—C2A—O3A | -6.8 (4) | O2D—C1D—C2D—C3D | -126.5 (3) |
| N1A—C1A—C2A—O3A | 115.7 (3) | N1D—C1D—C2D—C3D | -4.4 (3) |
| C8A—C1A—C2A—O3A | -127.1 (3) | C8D—C1D—C2D—C3D | 112.7 (3) |
| O2A—C1A—C2A—C3A | -127.9 (3) | O2D—C1D—C2D—C6D | 118.1 (3) |
| N1A—C1A—C2A—C3A | -5.5 (3) | N1D—C1D—C2D—C6D | -119.9 (3) |
| C8A—C1A—C2A—C3A | 111.7 (3) | C8D—C1D—C2D—C6D | -2.8 (3) |
| O2A—C1A—C2A—C6A | 116.7 (3) | O3D—C2D—C3D—C4D | -119.6 (3) |
| N1A—C1A—C2A—C6A | -120.9 (3) | C6D—C2D—C3D—C4D | 112.8 (3) |
| C8A—C1A—C2A—C6A | -3.7 (3) | C1D—C2D—C3D—C4D | 2.1 (3) |
| O2B—C1B—C2B—O3B | -13.3 (4) | O3D—C2D—C3D—N2D | 59.7 (4) |
| N1B—C1B—C2B—O3B | -135.2 (3) | C6D—C2D—C3D—N2D | -67.9 (4) |
| C8B—C1B—C2B—O3B | 108.1 (3) | C1D—C2D—C3D—N2D | -178.6 (3) |
| O2B—C1B—C2B—C3B | 111.2 (3) | N2D—C3D—C4D—N1D | -177.8 (3) |
| N1B—C1B—C2B—C3B | -10.8 (3) | C2D—C3D—C4D—N1D | 1.5 (4) |
| C8B—C1B—C2B—C3B | -127.4 (3) | N2D—C3D—C4D—S4 | -1.5 (5) |
| O2B—C1B—C2B—C6B | -131.0 (3) | C2D—C3D—C4D—S4 | 177.8 (2) |
| N1B—C1B—C2B—C6B | 107.1 (3) | O3D—C2D—C6D—O1D | -49.3 (5) |
| C8B—C1B—C2B—C6B | -9.6 (3) | C3D—C2D—C6D—O1D | 77.9 (4) |
| O3A—C2A—C3A—C4A | -118.8 (3) | C1D—C2D—C6D—O1D | -172.9 (3) |
| C6A—C2A—C3A—C4A | 113.9 (3) | O3D—C2D—C6D—C7D | 129.0 (3) |
| C1A—C2A—C3A—C4A | 3.0 (3) | C3D—C2D—C6D—C7D | -103.8 (3) |
| O3A—C2A—C3A—N2A | 61.1 (4) | C1D—C2D—C6D—C7D | 5.4 (4) |
| C6A—C2A—C3A—N2A | -66.2 (4) | O1D—C6D—C7D—C8D | 171.8 (4) |
| C1A—C2A—C3A—N2A | -177.1 (3) | C2D—C6D—C7D—C8D | -6.4 (4) |
| O3B—C2B—C3B—C4B | 130.0 (3) | O1D—C6D—C7D—C12D | -2.4 (6) |
| C6B—C2B—C3B—C4B | -102.7 (4) | C2D—C6D—C7D—C12D | 179.4 (4) |

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| C1B—C2B—C3B—C4B | 9.1 (4) | C12D—C7D—C8D—C9D | 0.2 (5) |
| O3B—C2B—C3B—N2B | -48.1 (5) | C6D—C7D—C8D—C9D | -174.4 (3) |
| C6B—C2B—C3B—N2B | 79.1 (4) | C12D—C7D—C8D—C1D | 179.2 (3) |
| C1B—C2B—C3B—N2B | -169.1 (3) | C6D—C7D—C8D—C1D | 4.6 (4) |
| N2A—C3A—C4A—N1A | -178.8 (3) | O2D—C1D—C8D—C7D | -125.8 (3) |
| C2A—C3A—C4A—N1A | 1.1 (4) | N1D—C1D—C8D—C7D | 110.8 (3) |
| N2A—C3A—C4A—S1 | -1.1 (5) | C2D—C1D—C8D—C7D | -0.9 (4) |
| C2A—C3A—C4A—S1 | 178.8 (2) | O2D—C1D—C8D—C9D | 53.1 (5) |
| N2B—C3B—C4B—N1B | 174.8 (4) | N1D—C1D—C8D—C9D | -70.2 (4) |
| C2B—C3B—C4B—N1B | -3.1 (4) | C2D—C1D—C8D—C9D | 178.0 (3) |
| N2B—C3B—C4B—S2 | -4.1 (6) | C7D—C8D—C9D—C10D | -0.6 (5) |
| C2B—C3B—C4B—S2 | 177.9 (3) | C1D—C8D—C9D—C10D | -179.5 (3) |
| O3A—C2A—C6A—O1A | -47.8 (5) | C8D—C9D—C10D—C11D | 0.2 (6) |
| C3A—C2A—C6A—O1A | 79.2 (4) | C9D—C10D—C11D—C12D | 0.7 (7) |
| C1A—C2A—C6A—O1A | -171.3 (3) | C10D—C11D—C12D—C7D | -1.2 (7) |
| O3A—C2A—C6A—C7A | 130.0 (3) | C8D—C7D—C12D—C11D | 0.7 (6) |
| C3A—C2A—C6A—C7A | -103.0 (3) | C6D—C7D—C12D—C11D | 174.4 (4) |
| C1A—C2A—C6A—C7A | 6.5 (4) | C18D—C13D—C14D—C15D | -3.1 (5) |
| O3B—C2B—C6B—O1B | 65.6 (5) | N1D—C13D—C14D—C15D | 170.6 (3) |
| C3B—C2B—C6B—O1B | -65.0 (5) | C13D—C14D—C15D—C16D | 0.4 (6) |
| C1B—C2B—C6B—O1B | -175.9 (4) | C14D—C15D—C16D—C17D | 2.5 (6) |
| O3B—C2B—C6B—C7B | -111.5 (3) | C14D—C15D—C16D—C19D | -176.1 (4) |
| C3B—C2B—C6B—C7B | 117.9 (3) | C15D—C16D—C17D—C18D | -2.8 (6) |
| C1B—C2B—C6B—C7B | 7.0 (4) | C19D—C16D—C17D—C18D | 175.8 (4) |
| O1A—C6A—C7A—C8A | 170.6 (4) | C16D—C17D—C18D—C13D | 0.1 (6) |
| C2A—C6A—C7A—C8A | -7.1 (4) | C14D—C13D—C18D—C17D | 2.8 (5) |
| O1A—C6A—C7A—C12A | -4.1 (7) | N1D—C13D—C18D—C17D | -171.0 (3) |
| C2A—C6A—C7A—C12A | 178.2 (4) | C3D—C4D—N1D—C13D | -154.6 (3) |
| O1B—C6B—C7B—C8B | -178.4 (4) | S4—C4D—N1D—C13D | 29.2 (4) |
| C2B—C6B—C7B—C8B | -1.4 (4) | C3D—C4D—N1D—C1D | -4.7 (4) |
| O1B—C6B—C7B—C12B | -3.2 (6) | S4—C4D—N1D—C1D | 179.1 (2) |
| C2B—C6B—C7B—C12B | 173.9 (3) | C14D—C13D—N1D—C4D | 54.7 (4) |
| C12A—C7A—C8A—C9A | 1.3 (6) | C18D—C13D—N1D—C4D | -131.5 (3) |
| C6A—C7A—C8A—C9A | -173.7 (3) | C14D—C13D—N1D—C1D | -93.5 (4) |
| C12A—C7A—C8A—C1A | 179.8 (4) | C18D—C13D—N1D—C1D | 80.3 (4) |
| C6A—C7A—C8A—C1A | 4.7 (4) | O2D—C1D—N1D—C4D | 130.9 (3) |
| O2A—C1A—C8A—C7A | -125.2 (3) | C8D—C1D—N1D—C4D | -106.5 (3) |
| N1A—C1A—C8A—C7A | 111.2 (3) | C2D—C1D—N1D—C4D | 5.7 (3) |
| C2A—C1A—C8A—C7A | -0.4 (4) | O2D—C1D—N1D—C13D | -76.4 (3) |
| O2A—C1A—C8A—C9A | 53.1 (4) | C8D—C1D—N1D—C13D | 46.3 (4) |
| N1A—C1A—C8A—C9A | -70.6 (4) | C2D—C1D—N1D—C13D | 158.4 (3) |
| C2A—C1A—C8A—C9A | 177.9 (3) | C4D—C3D—N2D—O4D | -9.0 (5) |
| C12B—C7B—C8B—C9B | -2.3 (5) | C2D—C3D—N2D—O4D | 171.8 (3) |
| C6B—C7B—C8B—C9B | 173.3 (3) | C4D—C3D—N2D—O5D | 172.5 (3) |
| C12B—C7B—C8B—C1B | 179.2 (3) | C2D—C3D—N2D—O5D | -6.7 (5) |
| C6B—C7B—C8B—C1B | -5.2 (4) | N1D—C4D—S4—C5D | 23.7 (3) |
| O2B—C1B—C8B—C7B | 136.3 (3) | C3D—C4D—S4—C5D | -152.1 (3) |
| N1B—C1B—C8B—C7B | -101.3 (3) | O2C—C1C—C2C—O3C | 9.4 (4) |
| C2B—C1B—C8B—C7B | 9.4 (3) | N1C—C1C—C2C—O3C | 131.9 (3) |

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| O2B—C1B—C8B—C9B | -42.0 (4) | C8C—C1C—C2C—O3C | -112.2 (3) |
| N1B—C1B—C8B—C9B | 80.4 (4) | O2C—C1C—C2C—C3C | -114.7 (3) |
| C2B—C1B—C8B—C9B | -168.9 (3) | N1C—C1C—C2C—C3C | 7.8 (3) |
| C7A—C8A—C9A—C10A | -0.8 (5) | C8C—C1C—C2C—C3C | 123.7 (3) |
| C1A—C8A—C9A—C10A | -178.9 (4) | O2C—C1C—C2C—C6C | 128.4 (3) |
| C7B—C8B—C9B—C10B | 0.3 (5) | N1C—C1C—C2C—C6C | -109.2 (3) |
| C1B—C8B—C9B—C10B | 178.5 (3) | C8C—C1C—C2C—C6C | 6.8 (3) |
| C8A—C9A—C10A—C11A | -0.2 (6) | O3C—C2C—C3C—C4C | -127.7 (3) |
| C8B—C9B—C10B—C11B | 1.8 (6) | C6C—C2C—C3C—C4C | 105.4 (4) |
| C9A—C10A—C11A—C12A | 0.6 (7) | C1C—C2C—C3C—C4C | -6.1 (4) |
| C9B—C10B—C11B—C12B | -1.9 (6) | O3C—C2C—C3C—N2C | 47.7 (5) |
| C10A—C11A—C12A—C7A | -0.1 (7) | C6C—C2C—C3C—N2C | -79.2 (4) |
| C8A—C7A—C12A—C11A | -0.9 (6) | C1C—C2C—C3C—N2C | 169.3 (3) |
| C6A—C7A—C12A—C11A | 173.3 (4) | N2C—C3C—C4C—N1C | -173.4 (4) |
| C10B—C11B—C12B—C7B | -0.1 (6) | C2C—C3C—C4C—N1C | 1.6 (4) |
| C8B—C7B—C12B—C11B | 2.2 (5) | N2C—C3C—C4C—S3 | 7.3 (6) |
| C6B—C7B—C12B—C11B | -172.6 (4) | C2C—C3C—C4C—S3 | -177.8 (3) |
| C18A—C13A—C14A—C15A | -1.7 (5) | N2C—C3C—C4C—S3' | 10.8 (9) |
| N1A—C13A—C14A—C15A | 171.2 (3) | C2C—C3C—C4C—S3' | -174.2 (5) |
| C18B—C13B—C14B—C15B | 0.2 (6) | C5C—S3—C4C—N1C | 0.5 (5) |
| N1B—C13B—C14B—C15B | -178.8 (4) | C5C—S3—C4C—C3C | 179.7 (4) |
| C13B—C14B—C15B—C16B | -1.0 (7) | C5C—S3—C4C—S3' | 4.6 (7) |
| C14B—C15B—C16B—C17B | 0.3 (7) | C1—S3'—C4C—N1C | -32.4 (16) |
| C14B—C15B—C16B—C19B | 176.9 (5) | C1—S3'—C4C—C3C | 143.5 (17) |
| C17A—C16A—C15A—C14A | 3.0 (6) | C1—S3'—C4C—S3 | 150.8 (18) |
| C19A—C16A—C15A—C14A | -174.8 (3) | O3C—C2C—C6C—O1C | -60.7 (5) |
| C13A—C14A—C15A—C16A | -0.6 (5) | C3C—C2C—C6C—O1C | 68.7 (5) |
| C15A—C16A—C17A—C18A | -3.2 (6) | C1C—C2C—C6C—O1C | 179.9 (3) |
| C19A—C16A—C17A—C18A | 174.6 (4) | O3C—C2C—C6C—C7C | 116.0 (3) |
| C14A—C13A—C18A—C17A | 1.5 (5) | C3C—C2C—C6C—C7C | -114.6 (3) |
| N1A—C13A—C18A—C17A | -171.4 (3) | C1C—C2C—C6C—C7C | -3.4 (3) |
| C16A—C17A—C18A—C13A | 0.9 (6) | O1C—C6C—C7C—C8C | 174.8 (4) |
| C15B—C16B—C17B—C18B | 1.2 (6) | C2C—C6C—C7C—C8C | -1.7 (4) |
| C19B—C16B—C17B—C18B | -175.5 (4) | O1C—C6C—C7C—C12C | 0.5 (6) |
| C16B—C17B—C18B—C13B | -2.0 (6) | C2C—C6C—C7C—C12C | -176.0 (3) |
| C14B—C13B—C18B—C17B | 1.2 (5) | C12C—C7C—C8C—C9C | 2.1 (5) |
| N1B—C13B—C18B—C17B | -179.8 (3) | C6C—C7C—C8C—C9C | -172.6 (3) |
| C3A—C4A—N1A—C13A | -155.3 (3) | C12C—C7C—C8C—C1C | -178.8 (3) |
| S1—C4A—N1A—C13A | 27.0 (4) | C6C—C7C—C8C—C1C | 6.5 (4) |
| C3A—C4A—N1A—C1A | -5.1 (3) | O2C—C1C—C8C—C7C | -135.5 (3) |
| S1—C4A—N1A—C1A | 177.2 (2) | N1C—C1C—C8C—C7C | 101.5 (3) |
| C18A—C13A—N1A—C4A | -132.1 (3) | C2C—C1C—C8C—C7C | -8.4 (3) |
| C14A—C13A—N1A—C4A | 54.9 (4) | O2C—C1C—C8C—C9C | 43.5 (5) |
| C18A—C13A—N1A—C1A | 79.4 (4) | N1C—C1C—C8C—C9C | -79.5 (4) |
| C14A—C13A—N1A—C1A | -93.6 (4) | C2C—C1C—C8C—C9C | 170.6 (3) |
| O2A—C1A—N1A—C4A | 132.2 (3) | C7C—C8C—C9C—C10C | -0.4 (6) |
| C8A—C1A—N1A—C4A | -105.3 (3) | C1C—C8C—C9C—C10C | -179.4 (4) |
| C2A—C1A—N1A—C4A | 6.6 (3) | C8C—C9C—C10C—C11C | -1.6 (7) |
| O2A—C1A—N1A—C13A | -74.6 (3) | C9C—C10C—C11C—C12C | 1.9 (7) |

| | | | |
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| C8A—C1A—N1A—C13A | 48.0 (4) | C10C—C11C—C12C—C7C | -0.2 (6) |
| C2A—C1A—N1A—C13A | 159.8 (2) | C8C—C7C—C12C—C11C | -1.8 (5) |
| C3B—C4B—N1B—C13B | -169.6 (3) | C6C—C7C—C12C—C11C | 172.0 (4) |
| S2—C4B—N1B—C13B | 9.3 (5) | C18C—C13C—C14C—C15C | -1.0 (7) |
| C3B—C4B—N1B—C1B | -4.7 (4) | N1C—C13C—C14C—C15C | 178.0 (5) |
| S2—C4B—N1B—C1B | 174.2 (3) | C13C—C14C—C15C—C16C | -0.7 (9) |
| C18B—C13B—N1B—C4B | 74.3 (4) | C14C—C15C—C16C—C17C | 3.0 (8) |
| C14B—C13B—N1B—C4B | -106.8 (5) | C14C—C15C—C16C—C19C | -176.1 (6) |
| C18B—C13B—N1B—C1B | -89.5 (4) | C15C—C16C—C17C—C18C | -3.6 (7) |
| C14B—C13B—N1B—C1B | 89.4 (4) | C19C—C16C—C17C—C18C | 175.4 (4) |
| O2B—C1B—N1B—C4B | -116.6 (3) | C14C—C13C—C18C—C17C | 0.4 (6) |
| C8B—C1B—N1B—C4B | 121.9 (3) | N1C—C13C—C18C—C17C | -178.6 (3) |
| C2B—C1B—N1B—C4B | 10.0 (3) | C16C—C17C—C18C—C13C | 2.0 (6) |
| O2B—C1B—N1B—C13B | 49.7 (4) | C3C—C4C—N1C—C13C | 178.1 (3) |
| C8B—C1B—N1B—C13B | -71.8 (4) | S3—C4C—N1C—C13C | -2.7 (6) |
| C2B—C1B—N1B—C13B | 176.3 (3) | S3'—C4C—N1C—C13C | -4.5 (5) |
| C4A—C3A—N2A—O5A | 171.8 (3) | C3C—C4C—N1C—C1C | 4.1 (4) |
| C2A—C3A—N2A—O5A | -8.1 (5) | S3—C4C—N1C—C1C | -176.7 (3) |
| C4A—C3A—N2A—O4A | -9.6 (5) | S3'—C4C—N1C—C1C | -178.5 (3) |
| C2A—C3A—N2A—O4A | 170.5 (3) | C18C—C13C—N1C—C4C | -89.6 (5) |
| C4B—C3B—N2B—O4B | 4.4 (7) | C14C—C13C—N1C—C4C | 91.4 (5) |
| C2B—C3B—N2B—O4B | -177.8 (4) | C18C—C13C—N1C—C1C | 84.0 (4) |
| C4B—C3B—N2B—O5B | -174.2 (4) | C14C—C13C—N1C—C1C | -94.9 (4) |
| C2B—C3B—N2B—O5B | 3.6 (6) | O2C—C1C—N1C—C4C | 119.2 (3) |
| N1A—C4A—S1—C5A | 26.9 (3) | C8C—C1C—N1C—C4C | -119.1 (3) |
| C3A—C4A—S1—C5A | -150.5 (3) | C2C—C1C—N1C—C4C | -7.5 (3) |
| N1B—C4B—S2—C5B | -6 (4) | O2C—C1C—N1C—C13C | -55.5 (4) |
| C3B—C4B—S2—C5B | 173 (4) | C8C—C1C—N1C—C13C | 66.2 (4) |
| N1B—C4B—S2—C2 | 21.0 (5) | C2C—C1C—N1C—C13C | 177.7 (3) |
| C3B—C4B—S2—C2 | -160.3 (5) | C4C—C3C—N2C—O5C | 169.8 (4) |
| O2D—C1D—C2D—O3D | -5.4 (4) | C2C—C3C—N2C—O5C | -4.7 (6) |
| N1D—C1D—C2D—O3D | 116.7 (3) | C4C—C3C—N2C—O4C | -9.1 (7) |
| C8D—C1D—C2D—O3D | -126.2 (3) | C2C—C3C—N2C—O4C | 176.3 (4) |

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> |
|--------------------------------|-------------|---------------|-----------------------|-------------------------|
| O3B—H3B...O5B | 0.82 | 2.19 | 2.748 (4) | 125 |
| O3C—H3C...O5C | 0.82 | 2.17 | 2.728 (4) | 126 |
| O2A—H2A...O3A ⁱ | 0.82 | 1.97 | 2.785 (3) | 175 |
| O3A—H3A...O3B ⁱⁱ | 0.82 | 2.04 | 2.854 (3) | 172 |
| O2B—H2B...O1A ⁱⁱⁱ | 0.82 | 2.01 | 2.787 (3) | 157 |
| O2C—H2C...O5D ^{iv} | 0.82 | 2.57 | 3.133 (4) | 127 |
| O2C—H2C...O1D ^{iv} | 0.82 | 2.03 | 2.797 (4) | 155 |
| O3D—H3D...O3C ^v | 0.82 | 2.04 | 2.857 (3) | 175 |
| C10A—H10A...O5A ⁱⁱⁱ | 0.93 | 2.54 | 3.347 (5) | 146 |
| C10C—H10C...O5C ^v | 0.93 | 2.50 | 3.331 (6) | 149 |
| C10D—H10D...O5D ^v | 0.93 | 2.59 | 3.437 (5) | 151 |

| | | | | |
|------------------------------------|------|------|-----------|-----|
| $C10B \cdots H10B \cdots O5B^{ii}$ | 0.93 | 2.53 | 3.316 (6) | 142 |
| $C2 \cdots H20 \cdots O1C^{vi}$ | 0.96 | 2.50 | 3.446 (7) | 170 |

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