

[2-[(1*H*-Indol-3-ylmethylidene)amino]-4,5,6,7-tetrahydrobenzo[*b*]thiophen-3-yl](phenyl)methanone

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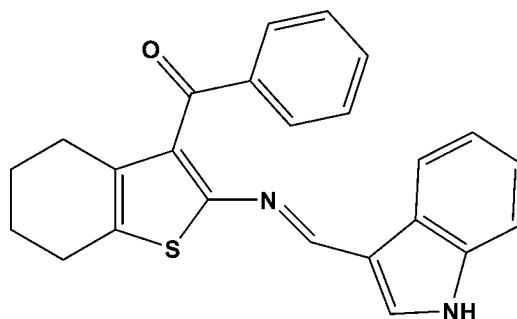
Received 19 March 2014; accepted 21 March 2014

Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.005\text{ \AA}$; R factor = 0.038; wR factor = 0.098; data-to-parameter ratio = 13.6.

The title compound, $C_{24}H_{20}N_2OS$, crystallizes with two independent molecules (*A* and *B*) in the asymmetric unit, in each of which the cyclohexene rings adopt half-chair conformations. The mean plane of the indole ring is twisted from those of the phenyl and thiophene rings by 69.0 (7) and 8.3 (5) $^\circ$, respectively, in molecule *A* and by 65.4 (9) and 6.7 (5) $^\circ$, respectively, in molecule *B*. The dihedral angles between the mean planes of the phenyl and thiophene rings are 63.0 (4) and 58.8 (9) $^\circ$ in molecules *A* and *B*, respectively. In the crystal, N—H \cdots O hydrogen bonds lead to the formation of an infinite chain along [101]. In addition, π — π stacking interactions are observed involving the thiophene and pyrrole rings of the two molecules, with a shortest intercentroid distance of 3.468 (2) \AA .

Related literature

For applications of 2-aminothiophene derivatives, see: Sabnis *et al.* (1999); Puterová *et al.* (2010); Cannito *et al.* (1990); Nikolakopoulos *et al.* (2006); Lütjens *et al.* (2005). For the biological and industrial importance of Schiff bases, see: Desai *et al.* (2001); Karia & Parsania (1999); Samadhiya & Halve (2001); Singh & Dash (1988); Aydogan *et al.* (2001); Taggi *et al.* (2002). For a related structure, see: Kubicki *et al.* (2012). For puckering parameters, see Cremer & Pople (1975). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{24}H_{20}N_2OS$
 $M_r = 384.48$
Monoclinic, $P2_1$
 $a = 8.66858$ (16) \AA
 $b = 21.8200$ (4) \AA
 $c = 10.41956$ (18) \AA
 $\beta = 108.1709$ (19) $^\circ$
 $V = 1872.55$ (6) \AA^3
 $Z = 4$
Cu $K\alpha$ radiation
 $\mu = 1.66\text{ mm}^{-1}$
 $T = 173\text{ K}$
 $0.22 \times 0.18 \times 0.06\text{ mm}$

Data collection

Agilent Eos Gemini diffractometer
Absorption correction: multi-scan
(*CrysAlis PRO* and *CrysAlis RED*; Agilent, 2012)
 $T_{\min} = 0.865$, $T_{\max} = 1.000$
11535 measured reflections
6843 independent reflections
6396 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.098$
 $S = 1.01$
6843 reflections
505 parameters
1 restraint
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.42\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.26\text{ e \AA}^{-3}$
Absolute structure: Flack parameter determined using 2790 quotients $[(I^+)-(I^-)]/[(I^+)+(I^-)]$ (Parsons *et al.*, 2013)
Absolute structure parameter: 0.171 (10)

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|--------------------------------------------------------------------------|--------------|--------------------|-------------|----------------------|
| $\text{N}2\text{A}-\text{H}2\text{A}\cdots\text{O}1\text{A}^{\text{i}}$ | 0.86 | 2.01 | 2.866 (4) | 175 |
| $\text{N}2\text{B}-\text{H}2\text{B}\cdots\text{O}1\text{B}^{\text{ii}}$ | 0.86 | 2.00 | 2.835 (3) | 163 |

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

Data collection: *CrysAlis PRO* (Agilent, 2012); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis RED* (Agilent, 2012); program(s) used to solve structure: *SUPERFLIP* (Palatinus & Chapuis, 2007); program(s) used to refine structure: *SHELXL2012* (Sheldrick, 2008); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *OLEX2*.

MK is grateful to the CPEPA–UGC for the award of a JRF and thanks the University of Mysore for research facilities. JPJ acknowledges the NSF–MRI program (grant No. CHE-1039027) for funds to purchase the X-ray diffractometer.

Supporting information for this paper is available from the IUCr electronic archives (Reference: BT6971).

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supporting information

Acta Cryst. (2014). E70, o501–o502 [doi:10.1107/S1600536814006345]

{2-[(1*H*-Indol-3-ylmethylidene)amino]-4,5,6,7-tetrahydrobenzo[*b*]thiophen-3-yl}(phenyl)methanone

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S1. Comment

2-Aminothiophene derivatives have been used in a number of applications in pesticides, dyes and pharmaceuticals. A review on the synthesis and properties of these compounds was reported by Sabnis *et al.* (1999) and more recently by Puterová *et al.* (2010). Substituted 2-aminothiophenes are active as allosteric enhancers at the human A1 adenosine receptor (Cannito *et al.*, 1990; Nikolakopoulos *et al.*, 2006; Lütjens *et al.*, 2005). Schiff base compounds are an important class of compounds both synthetically and biologically. These compounds show biological activities including antibacterial, antifungal, anticancer and herbicidal activities (Desai *et al.*, 2001; Karia & Parsania, 1999; Samadhiya & Halve, 2001; Singh & Dash, 1988). Furthermore, Schiff bases are utilized as starting materials in the synthesis of compounds of industrial (Aydogan *et al.*, 2001) and biological interest such as β -lactams (Taggi *et al.*, 2002). The crystal structures and molecular structures of two 2-aminothiophenes have been previously reported by our group (Kubicki *et al.*, 2012). In continuation of our work on schiff base derivatives of 2-aminothiophenes, we report here the crystal structure of the title compound, $C_{24}H_{20}N_2OS$.

The title compound crystallizes with two independent molecules in the asymmetric unit (A and B) (Fig. 1). In each of the molecules, the cyclohexene rings adopt half-chair conformations (puckering parameters Q, θ , and $\varphi = 0.508$ (4) \AA , 53.1 (5) $^\circ$ and 149.2 (5) $^\circ$ (A); Q, θ , and $\varphi = 0.492$ (4) \AA , 128.3 (5) $^\circ$ and 327.5 (6) $^\circ$ (B), respectively; Cremer & Pople, 1975). The mean plane of the indole ring is twisted from that of the phenyl and thiophene rings by 69.0 (7) $^\circ$ (A); 65.4 (9) $^\circ$ (B) and 8.3 (5) $^\circ$ (A); 6.7 (5) $^\circ$ (B), respectively. The dihedral angles between the mean plane of the phenyl rings and thiophene rings is 63.0 (4) $^\circ$ (A) and 58.8 (9) $^\circ$ (B), respectively. Bond lengths are in normal ranges (Allen *et al.*, 1987). N—H \cdots O intermolecular hydrogen bonds influence the crystal packing forming an infinite 1D chain along [1 0 1] (Fig. 2). In addition, weak Cg—Cg π — π stacking interactions are observed involving the thiophene rings and pyrrole rings of the two molecules with the shortest intercentroid distance of 3.468 (2) \AA .

S2. Experimental

To a solution of (2-Amino-4,5,6,7-tetrahydrobenzo[*b*]thiophen-3-yl)phenyl methanone (200 mg, 0.79 mmol) in 10 ml of methanol an equimolar amount of 1*H*-Indole-3-carbaldehyde (115 mg, 0.79 mmol) was added with constant stirring. The mixture was refluxed for 6 hours. An orange colored precipitate was obtained. The reaction completion was confirmed by thin layer chromatography. The precipitate was filtered and dried at room temperature overnight. The solid was recrystallized using methanol and the crystals were used as such for X-ray diffraction studies.

S3. Refinement

All of the H atoms were placed in their calculated positions and then refined using the riding model with atom—H lengths of 0.93 \AA (CH); 0.97 \AA (CH₂) or 0.86 \AA (NH). Isotropic displacement parameters for these atoms were set to 1.2

(CH, CH₂, NH) times U_{eq} of the parent atom.

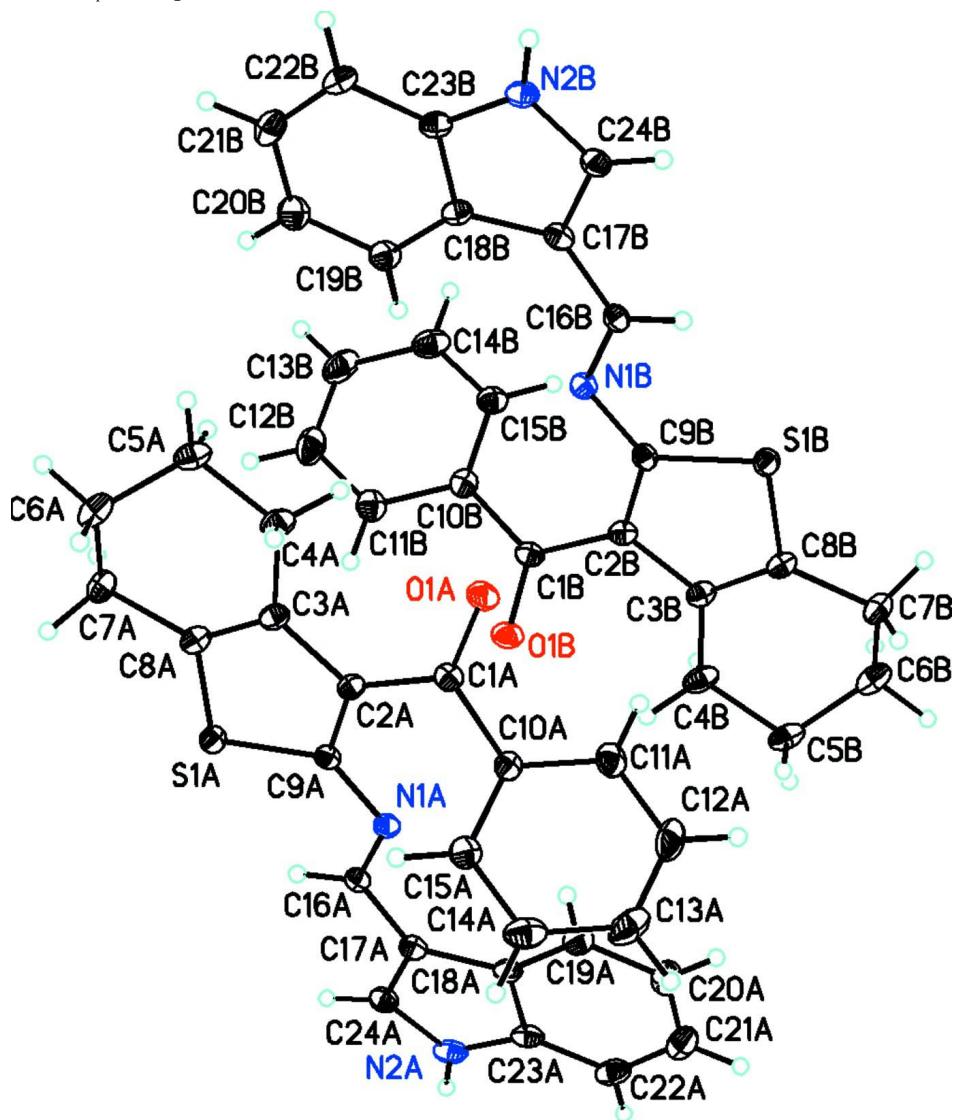
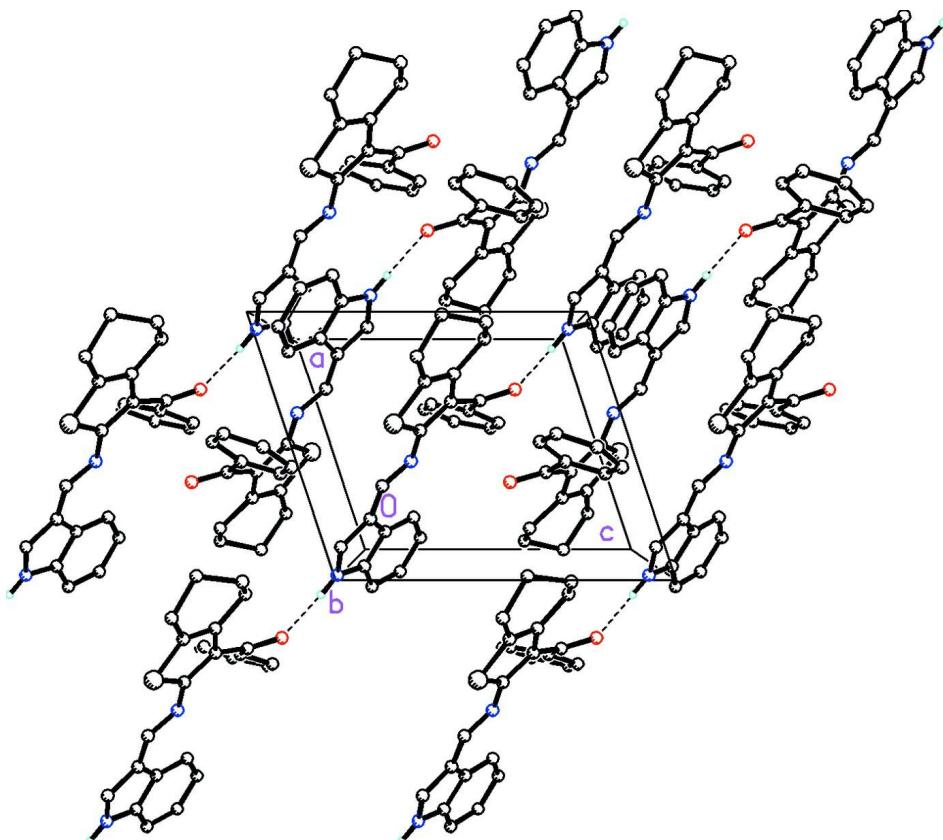


Figure 1

ORTEP drawing of the title compound showing the labeling scheme of the two molecules (A and B) within the asymmetric unit with 30% probability displacement ellipsoids.

**Figure 2**

Molecular packing of the title compound viewed along the b axis. Dashed lines indicate $\text{N}—\text{H}\cdots\text{O}$ intermolecular hydrogen bonds forming an infinite 1D-chain along $[1\ 0\ 1]$. H atoms not involved in hydrogen bonding have been removed for clarity.

{2-[$(1\text{H-Indol-3-ylmethylidene)amino$]-4,5,6,7-tetrahydrobenzo[*b*]thiophen-3-yl}(phenyl)methanone

Crystal data

$\text{C}_{24}\text{H}_{20}\text{N}_2\text{OS}$

$M_r = 384.48$

Monoclinic, $P2_1$

$a = 8.66858$ (16) Å

$b = 21.8200$ (4) Å

$c = 10.41956$ (18) Å

$\beta = 108.1709$ (19)°

$V = 1872.55$ (6) Å³

$Z = 4$

$F(000) = 808$

$D_x = 1.364$ Mg m⁻³

$\text{Cu } K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 6047 reflections

$\theta = 4.1\text{--}71.4^\circ$

$\mu = 1.66$ mm⁻¹

$T = 173$ K

Block, orange

$0.22 \times 0.18 \times 0.06$ mm

Data collection

Agilent Eos Gemini
diffractometer

Radiation source: Enhance (Cu) X-ray Source

Detector resolution: 16.0416 pixels mm⁻¹
 ω scans

Absorption correction: multi-scan

(*CrysAlis PRO* and *CrysAlis RED*; Agilent,
2012)

$T_{\min} = 0.865$, $T_{\max} = 1.000$

11535 measured reflections

6843 independent reflections

6396 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.039$
 $\theta_{\text{max}} = 71.4^\circ$, $\theta_{\text{min}} = 4.1^\circ$
 $h = -10 \rightarrow 10$

$k = -26 \rightarrow 24$
 $l = -10 \rightarrow 12$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.098$
 $S = 1.01$
6843 reflections
505 parameters
1 restraint
Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0599P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.42 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26 \text{ e } \text{\AA}^{-3}$
Absolute structure: Flack parameter determined using 2790 quotients $[(I^+)-(I^-)]/[(I^+)+(I^-)]$
(Parsons *et al.*, 2013)
Absolute structure parameter: 0.171 (10)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds involving l.s. planes.

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|-------------|--------------|-------------|----------------------------------|
| S1A | 0.47092 (9) | 0.61450 (3) | 1.00227 (7) | 0.02402 (17) |
| O1A | 0.3541 (3) | 0.75120 (12) | 0.6132 (2) | 0.0308 (5) |
| N1A | 0.6290 (3) | 0.72540 (12) | 1.0124 (3) | 0.0214 (5) |
| N2A | 1.0929 (3) | 0.76763 (15) | 1.3669 (3) | 0.0294 (6) |
| H2A | 1.1732 | 0.7651 | 1.4402 | 0.035* |
| C1A | 0.3945 (4) | 0.75388 (16) | 0.7365 (3) | 0.0220 (6) |
| C2A | 0.3944 (4) | 0.69809 (14) | 0.8176 (3) | 0.0209 (6) |
| C3A | 0.2860 (4) | 0.64697 (14) | 0.7671 (3) | 0.0232 (6) |
| C4A | 0.1599 (4) | 0.64209 (16) | 0.6297 (3) | 0.0304 (7) |
| H4AA | 0.2062 | 0.6558 | 0.5611 | 0.037* |
| H4AB | 0.0686 | 0.6685 | 0.6263 | 0.037* |
| C5A | 0.1007 (4) | 0.57594 (17) | 0.6006 (4) | 0.0326 (8) |
| H5AA | 0.0047 | 0.5752 | 0.5218 | 0.039* |
| H5AB | 0.1841 | 0.5518 | 0.5803 | 0.039* |
| C6A | 0.0609 (5) | 0.54746 (18) | 0.7194 (4) | 0.0360 (8) |
| H6AA | 0.0138 | 0.5072 | 0.6944 | 0.043* |
| H6AB | -0.0183 | 0.5727 | 0.7432 | 0.043* |
| C7A | 0.2142 (4) | 0.54194 (16) | 0.8409 (4) | 0.0302 (7) |
| H7AA | 0.1847 | 0.5341 | 0.9219 | 0.036* |
| H7AB | 0.2794 | 0.5079 | 0.8277 | 0.036* |
| C8A | 0.3107 (4) | 0.60052 (15) | 0.8572 (3) | 0.0249 (6) |
| C9A | 0.5045 (4) | 0.68713 (14) | 0.9440 (3) | 0.0207 (6) |
| C10A | 0.4374 (4) | 0.81499 (15) | 0.8025 (3) | 0.0225 (6) |
| C11A | 0.5143 (4) | 0.85813 (17) | 0.7429 (4) | 0.0319 (7) |

| | | | | |
|------|-------------|--------------|-------------|--------------|
| H11A | 0.5438 | 0.8475 | 0.6673 | 0.038* |
| C12A | 0.5459 (5) | 0.91625 (18) | 0.7967 (4) | 0.0409 (9) |
| H12A | 0.6015 | 0.9441 | 0.7599 | 0.049* |
| C13A | 0.4957 (5) | 0.93340 (17) | 0.9049 (4) | 0.0442 (10) |
| H13A | 0.5137 | 0.9732 | 0.9384 | 0.053* |
| C14A | 0.4184 (5) | 0.89116 (19) | 0.9637 (4) | 0.0393 (9) |
| H14A | 0.3846 | 0.9026 | 1.0366 | 0.047* |
| C15A | 0.3916 (4) | 0.83197 (16) | 0.9137 (3) | 0.0291 (7) |
| H15A | 0.3427 | 0.8035 | 0.9549 | 0.035* |
| C16A | 0.7273 (4) | 0.70843 (15) | 1.1267 (3) | 0.0213 (6) |
| H16A | 0.7091 | 0.6710 | 1.1621 | 0.026* |
| C17A | 0.8636 (4) | 0.74433 (15) | 1.2023 (3) | 0.0227 (6) |
| C18A | 0.9206 (4) | 0.80258 (15) | 1.1706 (3) | 0.0226 (6) |
| C19A | 0.8656 (4) | 0.84400 (16) | 1.0639 (3) | 0.0289 (7) |
| H19A | 0.7702 | 0.8366 | 0.9940 | 0.035* |
| C20A | 0.9562 (5) | 0.89634 (18) | 1.0644 (4) | 0.0373 (8) |
| H20A | 0.9212 | 0.9241 | 0.9935 | 0.045* |
| C21A | 1.0994 (5) | 0.90821 (18) | 1.1692 (4) | 0.0378 (8) |
| H21A | 1.1577 | 0.9438 | 1.1670 | 0.045* |
| C22A | 1.1558 (4) | 0.86806 (19) | 1.2761 (4) | 0.0344 (8) |
| H22A | 1.2510 | 0.8759 | 1.3459 | 0.041* |
| C23A | 1.0650 (4) | 0.81545 (17) | 1.2756 (3) | 0.0268 (7) |
| C24A | 0.9739 (4) | 0.72543 (17) | 1.3226 (3) | 0.0258 (7) |
| H24A | 0.9673 | 0.6890 | 1.3668 | 0.031* |
| S1B | 0.58516 (9) | 0.82206 (3) | 0.31301 (7) | 0.02433 (17) |
| O1B | 0.7245 (3) | 0.70323 (12) | 0.7261 (2) | 0.0310 (5) |
| N1B | 0.4361 (3) | 0.71109 (13) | 0.3214 (3) | 0.0221 (5) |
| N2B | -0.0273 (3) | 0.66696 (14) | -0.0369 (3) | 0.0255 (6) |
| H2B | -0.1076 | 0.6698 | -0.1102 | 0.031* |
| C1B | 0.6804 (4) | 0.69303 (15) | 0.6038 (3) | 0.0224 (6) |
| C2B | 0.6739 (4) | 0.74464 (15) | 0.5103 (3) | 0.0222 (6) |
| C3B | 0.7811 (4) | 0.79705 (15) | 0.5483 (3) | 0.0244 (6) |
| C4B | 0.9186 (4) | 0.80540 (16) | 0.6792 (4) | 0.0328 (8) |
| H4BA | 1.0085 | 0.7790 | 0.6788 | 0.039* |
| H4BB | 0.8821 | 0.7934 | 0.7545 | 0.039* |
| C5B | 0.9759 (5) | 0.87144 (17) | 0.6973 (4) | 0.0330 (8) |
| H5BA | 1.0771 | 0.8738 | 0.7709 | 0.040* |
| H5BB | 0.8962 | 0.8961 | 0.7216 | 0.040* |
| C6B | 1.0013 (5) | 0.89755 (18) | 0.5699 (4) | 0.0365 (8) |
| H6BA | 1.0791 | 0.8724 | 0.5441 | 0.044* |
| H6BB | 1.0454 | 0.9386 | 0.5880 | 0.044* |
| C7B | 0.8411 (5) | 0.89950 (17) | 0.4535 (4) | 0.0315 (8) |
| H7BA | 0.7765 | 0.9341 | 0.4651 | 0.038* |
| H7BB | 0.8632 | 0.9045 | 0.3683 | 0.038* |
| C8B | 0.7491 (4) | 0.84082 (16) | 0.4515 (3) | 0.0254 (7) |
| C9B | 0.5592 (4) | 0.75132 (15) | 0.3840 (3) | 0.0211 (6) |
| C10B | 0.6424 (4) | 0.62857 (15) | 0.5582 (3) | 0.0250 (6) |
| C11B | 0.5927 (4) | 0.58857 (17) | 0.6427 (4) | 0.0314 (7) |

| | | | | |
|------|-------------|--------------|------------|-------------|
| H11B | 0.5740 | 0.6035 | 0.7201 | 0.038* |
| C12B | 0.5714 (5) | 0.52710 (19) | 0.6113 (4) | 0.0433 (9) |
| H12B | 0.5373 | 0.5008 | 0.6673 | 0.052* |
| C13B | 0.6003 (6) | 0.50458 (18) | 0.4975 (4) | 0.0448 (10) |
| H13B | 0.5871 | 0.4630 | 0.4775 | 0.054* |
| C14B | 0.6494 (5) | 0.54392 (18) | 0.4123 (4) | 0.0394 (9) |
| H14B | 0.6695 | 0.5286 | 0.3358 | 0.047* |
| C15B | 0.6681 (4) | 0.60575 (17) | 0.4415 (3) | 0.0291 (7) |
| H15B | 0.6979 | 0.6322 | 0.3833 | 0.035* |
| C16B | 0.3368 (4) | 0.72624 (15) | 0.2058 (3) | 0.0219 (6) |
| H16B | 0.3536 | 0.7636 | 0.1692 | 0.026* |
| C17B | 0.2019 (4) | 0.68968 (15) | 0.1295 (3) | 0.0225 (6) |
| C18B | 0.1459 (4) | 0.63051 (14) | 0.1581 (3) | 0.0221 (6) |
| C19B | 0.2011 (4) | 0.58836 (17) | 0.2628 (3) | 0.0282 (7) |
| H19B | 0.2962 | 0.5954 | 0.3333 | 0.034* |
| C20B | 0.1103 (5) | 0.53570 (17) | 0.2591 (4) | 0.0346 (8) |
| H20B | 0.1458 | 0.5070 | 0.3281 | 0.042* |
| C21B | -0.0332 (5) | 0.52460 (18) | 0.1541 (4) | 0.0361 (8) |
| H21B | -0.0916 | 0.4890 | 0.1553 | 0.043* |
| C22B | -0.0899 (4) | 0.56541 (17) | 0.0489 (4) | 0.0295 (7) |
| H22B | -0.1851 | 0.5580 | -0.0212 | 0.035* |
| C23B | 0.0021 (4) | 0.61853 (16) | 0.0524 (3) | 0.0237 (6) |
| C24B | 0.0910 (4) | 0.70917 (16) | 0.0096 (3) | 0.0248 (6) |
| H24B | 0.0971 | 0.7461 | -0.0330 | 0.030* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| S1A | 0.0249 (4) | 0.0210 (4) | 0.0232 (3) | 0.0012 (3) | 0.0033 (3) | 0.0041 (3) |
| O1A | 0.0297 (12) | 0.0381 (14) | 0.0193 (11) | -0.0017 (10) | -0.0002 (9) | 0.0041 (10) |
| N1A | 0.0206 (13) | 0.0236 (13) | 0.0180 (12) | 0.0000 (10) | 0.0032 (10) | -0.0002 (10) |
| N2A | 0.0220 (13) | 0.0421 (17) | 0.0186 (12) | 0.0033 (12) | -0.0016 (10) | -0.0068 (11) |
| C1A | 0.0155 (13) | 0.0277 (16) | 0.0197 (14) | 0.0028 (11) | 0.0009 (11) | 0.0023 (12) |
| C2A | 0.0176 (14) | 0.0209 (15) | 0.0225 (14) | 0.0003 (11) | 0.0037 (11) | -0.0015 (11) |
| C3A | 0.0213 (15) | 0.0211 (16) | 0.0259 (15) | 0.0023 (12) | 0.0053 (12) | -0.0040 (11) |
| C4A | 0.0289 (17) | 0.0267 (17) | 0.0282 (16) | 0.0002 (13) | -0.0020 (13) | -0.0019 (13) |
| C5A | 0.0288 (17) | 0.0314 (19) | 0.0304 (17) | 0.0001 (14) | -0.0009 (13) | -0.0074 (14) |
| C6A | 0.0280 (17) | 0.0291 (18) | 0.047 (2) | -0.0054 (14) | 0.0068 (15) | -0.0081 (15) |
| C7A | 0.0305 (17) | 0.0210 (16) | 0.0379 (18) | -0.0019 (13) | 0.0091 (14) | -0.0004 (13) |
| C8A | 0.0213 (14) | 0.0217 (16) | 0.0290 (15) | 0.0018 (12) | 0.0040 (12) | -0.0024 (12) |
| C9A | 0.0224 (15) | 0.0210 (15) | 0.0191 (14) | 0.0022 (11) | 0.0070 (12) | 0.0003 (11) |
| C10A | 0.0214 (13) | 0.0208 (15) | 0.0208 (13) | 0.0034 (11) | 0.0000 (11) | 0.0060 (11) |
| C11A | 0.0312 (18) | 0.0325 (19) | 0.0280 (16) | 0.0019 (14) | 0.0034 (14) | 0.0096 (14) |
| C12A | 0.040 (2) | 0.0287 (19) | 0.046 (2) | -0.0044 (15) | 0.0013 (17) | 0.0151 (16) |
| C13A | 0.049 (2) | 0.0215 (18) | 0.046 (2) | 0.0053 (15) | -0.0089 (18) | 0.0000 (15) |
| C14A | 0.044 (2) | 0.036 (2) | 0.0315 (18) | 0.0114 (16) | 0.0022 (16) | -0.0061 (15) |
| C15A | 0.0310 (17) | 0.0261 (17) | 0.0271 (15) | 0.0058 (13) | 0.0044 (13) | 0.0052 (13) |
| C16A | 0.0220 (15) | 0.0228 (15) | 0.0189 (13) | 0.0017 (12) | 0.0062 (11) | 0.0007 (11) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C17A | 0.0206 (15) | 0.0260 (16) | 0.0201 (14) | 0.0036 (12) | 0.0045 (12) | 0.0007 (11) |
| C18A | 0.0207 (14) | 0.0249 (16) | 0.0218 (14) | 0.0034 (12) | 0.0060 (12) | -0.0033 (12) |
| C19A | 0.0279 (16) | 0.0279 (18) | 0.0277 (16) | 0.0026 (14) | 0.0040 (13) | -0.0002 (13) |
| C20A | 0.040 (2) | 0.0300 (19) | 0.043 (2) | 0.0006 (15) | 0.0150 (17) | 0.0023 (15) |
| C21A | 0.0339 (19) | 0.0302 (19) | 0.053 (2) | -0.0053 (15) | 0.0188 (17) | -0.0076 (16) |
| C22A | 0.0235 (17) | 0.041 (2) | 0.0380 (19) | -0.0042 (14) | 0.0086 (14) | -0.0160 (16) |
| C23A | 0.0223 (15) | 0.0343 (18) | 0.0229 (15) | 0.0036 (13) | 0.0059 (12) | -0.0088 (13) |
| C24A | 0.0225 (14) | 0.0328 (18) | 0.0207 (15) | 0.0023 (13) | 0.0045 (12) | -0.0005 (12) |
| S1B | 0.0254 (3) | 0.0217 (4) | 0.0228 (3) | -0.0004 (3) | 0.0030 (3) | 0.0029 (3) |
| O1B | 0.0328 (13) | 0.0339 (13) | 0.0211 (11) | 0.0015 (10) | 0.0012 (9) | 0.0000 (10) |
| N1B | 0.0209 (14) | 0.0230 (13) | 0.0214 (12) | 0.0019 (11) | 0.0054 (10) | 0.0016 (10) |
| N2B | 0.0211 (13) | 0.0312 (15) | 0.0196 (12) | 0.0012 (11) | -0.0005 (10) | -0.0007 (11) |
| C1B | 0.0182 (14) | 0.0271 (17) | 0.0189 (15) | 0.0042 (11) | 0.0015 (11) | 0.0010 (12) |
| C2B | 0.0202 (14) | 0.0233 (16) | 0.0212 (14) | 0.0035 (12) | 0.0039 (11) | -0.0013 (12) |
| C3B | 0.0224 (15) | 0.0212 (15) | 0.0263 (15) | 0.0027 (12) | 0.0028 (12) | -0.0026 (12) |
| C4B | 0.0294 (17) | 0.0255 (18) | 0.0329 (17) | 0.0036 (13) | -0.0057 (14) | -0.0063 (13) |
| C5B | 0.0284 (18) | 0.0277 (18) | 0.0350 (18) | 0.0012 (13) | -0.0015 (14) | -0.0104 (14) |
| C6B | 0.0287 (18) | 0.0307 (19) | 0.049 (2) | -0.0063 (14) | 0.0111 (16) | -0.0138 (16) |
| C7B | 0.0325 (19) | 0.0227 (17) | 0.0376 (19) | -0.0051 (14) | 0.0085 (15) | -0.0031 (13) |
| C8B | 0.0238 (15) | 0.0218 (16) | 0.0279 (16) | -0.0006 (12) | 0.0045 (13) | -0.0050 (12) |
| C9B | 0.0218 (14) | 0.0205 (14) | 0.0216 (14) | 0.0030 (12) | 0.0078 (12) | 0.0029 (11) |
| C10B | 0.0194 (14) | 0.0252 (17) | 0.0274 (15) | 0.0037 (11) | 0.0029 (12) | 0.0032 (12) |
| C11B | 0.0297 (17) | 0.0314 (18) | 0.0314 (17) | 0.0037 (14) | 0.0068 (13) | 0.0058 (13) |
| C12B | 0.041 (2) | 0.033 (2) | 0.049 (2) | -0.0053 (16) | 0.0049 (17) | 0.0124 (17) |
| C13B | 0.050 (2) | 0.0216 (17) | 0.049 (2) | 0.0006 (15) | -0.0041 (18) | 0.0004 (15) |
| C14B | 0.042 (2) | 0.0323 (19) | 0.0360 (19) | 0.0076 (16) | 0.0002 (16) | -0.0066 (15) |
| C15B | 0.0277 (16) | 0.0301 (18) | 0.0262 (15) | 0.0032 (13) | 0.0037 (12) | 0.0006 (13) |
| C16B | 0.0219 (15) | 0.0235 (15) | 0.0207 (14) | 0.0029 (12) | 0.0071 (12) | 0.0032 (11) |
| C17B | 0.0218 (15) | 0.0262 (16) | 0.0191 (14) | 0.0052 (12) | 0.0060 (11) | 0.0010 (11) |
| C18B | 0.0195 (14) | 0.0231 (16) | 0.0231 (14) | 0.0029 (12) | 0.0057 (12) | -0.0049 (12) |
| C19B | 0.0286 (17) | 0.0259 (17) | 0.0268 (16) | 0.0043 (13) | 0.0039 (13) | 0.0018 (13) |
| C20B | 0.043 (2) | 0.0249 (18) | 0.0335 (18) | 0.0040 (15) | 0.0083 (15) | 0.0054 (14) |
| C21B | 0.039 (2) | 0.0236 (17) | 0.047 (2) | -0.0053 (15) | 0.0145 (16) | -0.0045 (15) |
| C22B | 0.0255 (16) | 0.0292 (17) | 0.0306 (16) | -0.0038 (13) | 0.0044 (13) | -0.0082 (13) |
| C23B | 0.0231 (15) | 0.0243 (16) | 0.0226 (14) | 0.0045 (13) | 0.0054 (12) | -0.0041 (12) |
| C24B | 0.0243 (15) | 0.0279 (16) | 0.0204 (14) | 0.0017 (13) | 0.0045 (12) | 0.0024 (12) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|----------|-----------|----------|-----------|
| S1A—C8A | 1.732 (3) | S1B—C8B | 1.729 (3) |
| S1A—C9A | 1.754 (3) | S1B—C9B | 1.756 (3) |
| O1A—C1A | 1.223 (4) | O1B—C1B | 1.232 (4) |
| N1A—C9A | 1.375 (4) | N1B—C9B | 1.379 (4) |
| N1A—C16A | 1.284 (4) | N1B—C16B | 1.286 (4) |
| N2A—H2A | 0.8600 | N2B—H2B | 0.8600 |
| N2A—C23A | 1.381 (5) | N2B—C23B | 1.378 (4) |
| N2A—C24A | 1.352 (5) | N2B—C24B | 1.351 (4) |
| C1A—C2A | 1.482 (4) | C1B—C2B | 1.479 (4) |

| | | | |
|--------------|------------|--------------|------------|
| C1A—C10A | 1.493 (5) | C1B—C10B | 1.488 (5) |
| C2A—C3A | 1.447 (4) | C2B—C3B | 1.448 (5) |
| C2A—C9A | 1.386 (4) | C2B—C9B | 1.388 (4) |
| C3A—C4A | 1.510 (4) | C3B—C4B | 1.517 (4) |
| C3A—C8A | 1.352 (5) | C3B—C8B | 1.353 (5) |
| C4A—H4AA | 0.9700 | C4B—H4BA | 0.9700 |
| C4A—H4AB | 0.9700 | C4B—H4BB | 0.9700 |
| C4A—C5A | 1.530 (5) | C4B—C5B | 1.517 (5) |
| C5A—H5AA | 0.9700 | C5B—H5BA | 0.9700 |
| C5A—H5AB | 0.9700 | C5B—H5BB | 0.9700 |
| C5A—C6A | 1.519 (6) | C5B—C6B | 1.522 (6) |
| C6A—H6AA | 0.9700 | C6B—H6BA | 0.9700 |
| C6A—H6AB | 0.9700 | C6B—H6BB | 0.9700 |
| C6A—C7A | 1.528 (5) | C6B—C7B | 1.533 (5) |
| C7A—H7AA | 0.9700 | C7B—H7BA | 0.9700 |
| C7A—H7AB | 0.9700 | C7B—H7BB | 0.9700 |
| C7A—C8A | 1.508 (5) | C7B—C8B | 1.506 (5) |
| C10A—C11A | 1.405 (5) | C10B—C11B | 1.399 (5) |
| C10A—C15A | 1.387 (5) | C10B—C15B | 1.394 (5) |
| C11A—H11A | 0.9300 | C11B—H11B | 0.9300 |
| C11A—C12A | 1.379 (6) | C11B—C12B | 1.379 (6) |
| C12A—H12A | 0.9300 | C12B—H12B | 0.9300 |
| C12A—C13A | 1.380 (7) | C12B—C13B | 1.377 (7) |
| C13A—H13A | 0.9300 | C13B—H13B | 0.9300 |
| C13A—C14A | 1.389 (7) | C13B—C14B | 1.393 (6) |
| C14A—H14A | 0.9300 | C14B—H14B | 0.9300 |
| C14A—C15A | 1.385 (5) | C14B—C15B | 1.382 (5) |
| C15A—H15A | 0.9300 | C15B—H15B | 0.9300 |
| C16A—H16A | 0.9300 | C16B—H16B | 0.9300 |
| C16A—C17A | 1.432 (4) | C16B—C17B | 1.434 (5) |
| C17A—C18A | 1.439 (5) | C17B—C18B | 1.443 (5) |
| C17A—C24A | 1.381 (4) | C17B—C24B | 1.384 (4) |
| C18A—C19A | 1.396 (5) | C18B—C19B | 1.393 (5) |
| C18A—C23A | 1.410 (4) | C18B—C23B | 1.407 (4) |
| C19A—H19A | 0.9300 | C19B—H19B | 0.9300 |
| C19A—C20A | 1.385 (5) | C19B—C20B | 1.386 (5) |
| C20A—H20A | 0.9300 | C20B—H20B | 0.9300 |
| C20A—C21A | 1.399 (6) | C20B—C21B | 1.398 (6) |
| C21A—H21A | 0.9300 | C21B—H21B | 0.9300 |
| C21A—C22A | 1.381 (6) | C21B—C22B | 1.378 (5) |
| C22A—H22A | 0.9300 | C22B—H22B | 0.9300 |
| C22A—C23A | 1.391 (5) | C22B—C23B | 1.401 (5) |
| C24A—H24A | 0.9300 | C24B—H24B | 0.9300 |
| | | | |
| C8A—S1A—C9A | 91.80 (15) | C8B—S1B—C9B | 92.09 (16) |
| C16A—N1A—C9A | 119.5 (3) | C16B—N1B—C9B | 118.2 (3) |
| C23A—N2A—H2A | 125.4 | C23B—N2B—H2B | 125.6 |
| C24A—N2A—H2A | 125.4 | C24B—N2B—H2B | 125.6 |

| | | | |
|----------------|-----------|----------------|-----------|
| C24A—N2A—C23A | 109.2 (3) | C24B—N2B—C23B | 108.7 (3) |
| O1A—C1A—C2A | 120.6 (3) | O1B—C1B—C2B | 118.7 (3) |
| O1A—C1A—C10A | 118.4 (3) | O1B—C1B—C10B | 117.9 (3) |
| C2A—C1A—C10A | 121.0 (3) | C2B—C1B—C10B | 123.4 (3) |
| C3A—C2A—C1A | 122.8 (3) | C3B—C2B—C1B | 122.1 (3) |
| C9A—C2A—C1A | 124.1 (3) | C9B—C2B—C1B | 125.0 (3) |
| C9A—C2A—C3A | 112.9 (3) | C9B—C2B—C3B | 112.7 (3) |
| C2A—C3A—C4A | 126.5 (3) | C2B—C3B—C4B | 126.7 (3) |
| C8A—C3A—C2A | 112.4 (3) | C8B—C3B—C2B | 112.9 (3) |
| C8A—C3A—C4A | 121.1 (3) | C8B—C3B—C4B | 120.4 (3) |
| C3A—C4A—H4AA | 109.5 | C3B—C4B—H4BA | 109.4 |
| C3A—C4A—H4AB | 109.5 | C3B—C4B—H4BB | 109.4 |
| C3A—C4A—C5A | 110.7 (3) | C3B—C4B—C5B | 111.2 (3) |
| H4AA—C4A—H4AB | 108.1 | H4BA—C4B—H4BB | 108.0 |
| C5A—C4A—H4AA | 109.5 | C5B—C4B—H4BA | 109.4 |
| C5A—C4A—H4AB | 109.5 | C5B—C4B—H4BB | 109.4 |
| C4A—C5A—H5AA | 109.3 | C4B—C5B—H5BA | 109.2 |
| C4A—C5A—H5AB | 109.3 | C4B—C5B—H5BB | 109.2 |
| H5AA—C5A—H5AB | 107.9 | C4B—C5B—C6B | 112.2 (3) |
| C6A—C5A—C4A | 111.8 (3) | H5BA—C5B—H5BB | 107.9 |
| C6A—C5A—H5AA | 109.3 | C6B—C5B—H5BA | 109.2 |
| C6A—C5A—H5AB | 109.3 | C6B—C5B—H5BB | 109.2 |
| C5A—C6A—H6AA | 109.6 | C5B—C6B—H6BA | 109.4 |
| C5A—C6A—H6AB | 109.6 | C5B—C6B—H6BB | 109.4 |
| C5A—C6A—C7A | 110.3 (3) | C5B—C6B—C7B | 111.0 (3) |
| H6AA—C6A—H6AB | 108.1 | H6BA—C6B—H6BB | 108.0 |
| C7A—C6A—H6AA | 109.6 | C7B—C6B—H6BA | 109.4 |
| C7A—C6A—H6AB | 109.6 | C7B—C6B—H6BB | 109.4 |
| C6A—C7A—H7AA | 109.8 | C6B—C7B—H7BA | 109.8 |
| C6A—C7A—H7AB | 109.8 | C6B—C7B—H7BB | 109.8 |
| H7AA—C7A—H7AB | 108.2 | H7BA—C7B—H7BB | 108.2 |
| C8A—C7A—C6A | 109.6 (3) | C8B—C7B—C6B | 109.4 (3) |
| C8A—C7A—H7AA | 109.8 | C8B—C7B—H7BA | 109.8 |
| C8A—C7A—H7AB | 109.8 | C8B—C7B—H7BB | 109.8 |
| C3A—C8A—S1A | 112.6 (2) | C3B—C8B—S1B | 112.3 (3) |
| C3A—C8A—C7A | 126.5 (3) | C3B—C8B—C7B | 127.3 (3) |
| C7A—C8A—S1A | 120.9 (2) | C7B—C8B—S1B | 120.4 (3) |
| N1A—C9A—S1A | 123.9 (2) | N1B—C9B—S1B | 122.8 (2) |
| N1A—C9A—C2A | 125.8 (3) | N1B—C9B—C2B | 127.0 (3) |
| C2A—C9A—S1A | 110.2 (2) | C2B—C9B—S1B | 110.1 (2) |
| C11A—C10A—C1A | 118.6 (3) | C11B—C10B—C1B | 118.0 (3) |
| C15A—C10A—C1A | 121.9 (3) | C15B—C10B—C1B | 122.4 (3) |
| C15A—C10A—C11A | 119.3 (3) | C15B—C10B—C11B | 119.3 (3) |
| C10A—C11A—H11A | 120.1 | C10B—C11B—H11B | 119.9 |
| C12A—C11A—C10A | 119.8 (4) | C12B—C11B—C10B | 120.2 (4) |
| C12A—C11A—H11A | 120.1 | C12B—C11B—H11B | 119.9 |
| C11A—C12A—H12A | 119.7 | C11B—C12B—H12B | 119.9 |
| C11A—C12A—C13A | 120.6 (4) | C13B—C12B—C11B | 120.3 (4) |

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|--------------------|------------|--------------------|------------|
| C13A—C12A—H12A | 119.7 | C13B—C12B—H12B | 119.9 |
| C12A—C13A—H13A | 120.1 | C12B—C13B—H13B | 119.9 |
| C12A—C13A—C14A | 119.9 (4) | C12B—C13B—C14B | 120.2 (4) |
| C14A—C13A—H13A | 120.1 | C14B—C13B—H13B | 119.9 |
| C13A—C14A—H14A | 120.0 | C13B—C14B—H14B | 120.0 |
| C15A—C14A—C13A | 120.0 (4) | C15B—C14B—C13B | 120.0 (4) |
| C15A—C14A—H14A | 120.0 | C15B—C14B—H14B | 120.0 |
| C10A—C15A—H15A | 119.8 | C10B—C15B—H15B | 120.0 |
| C14A—C15A—C10A | 120.4 (3) | C14B—C15B—C10B | 120.1 (4) |
| C14A—C15A—H15A | 119.8 | C14B—C15B—H15B | 120.0 |
| N1A—C16A—H16A | 118.4 | N1B—C16B—H16B | 117.7 |
| N1A—C16A—C17A | 123.2 (3) | N1B—C16B—C17B | 124.6 (3) |
| C17A—C16A—H16A | 118.4 | C17B—C16B—H16B | 117.7 |
| C16A—C17A—C18A | 129.8 (3) | C16B—C17B—C18B | 130.7 (3) |
| C24A—C17A—C16A | 123.7 (3) | C24B—C17B—C16B | 123.1 (3) |
| C24A—C17A—C18A | 106.4 (3) | C24B—C17B—C18B | 106.1 (3) |
| C19A—C18A—C17A | 134.3 (3) | C19B—C18B—C17B | 134.2 (3) |
| C19A—C18A—C23A | 119.3 (3) | C19B—C18B—C23B | 119.7 (3) |
| C23A—C18A—C17A | 106.4 (3) | C23B—C18B—C17B | 106.1 (3) |
| C18A—C19A—H19A | 120.7 | C18B—C19B—H19B | 120.9 |
| C20A—C19A—C18A | 118.6 (3) | C20B—C19B—C18B | 118.1 (3) |
| C20A—C19A—H19A | 120.7 | C20B—C19B—H19B | 120.9 |
| C19A—C20A—H20A | 119.4 | C19B—C20B—H20B | 119.2 |
| C19A—C20A—C21A | 121.3 (4) | C19B—C20B—C21B | 121.6 (4) |
| C21A—C20A—H20A | 119.4 | C21B—C20B—H20B | 119.2 |
| C20A—C21A—H21A | 119.4 | C20B—C21B—H21B | 119.3 |
| C22A—C21A—C20A | 121.1 (4) | C22B—C21B—C20B | 121.3 (4) |
| C22A—C21A—H21A | 119.4 | C22B—C21B—H21B | 119.3 |
| C21A—C22A—H22A | 121.2 | C21B—C22B—H22B | 121.5 |
| C21A—C22A—C23A | 117.6 (3) | C21B—C22B—C23B | 117.1 (3) |
| C23A—C22A—H22A | 121.2 | C23B—C22B—H22B | 121.5 |
| N2A—C23A—C18A | 107.8 (3) | N2B—C23B—C18B | 108.6 (3) |
| N2A—C23A—C22A | 130.0 (3) | N2B—C23B—C22B | 129.3 (3) |
| C22A—C23A—C18A | 122.1 (3) | C22B—C23B—C18B | 122.1 (3) |
| N2A—C24A—C17A | 110.2 (3) | N2B—C24B—C17B | 110.5 (3) |
| N2A—C24A—H24A | 124.9 | N2B—C24B—H24B | 124.8 |
| C17A—C24A—H24A | 124.9 | C17B—C24B—H24B | 124.8 |
| | | | |
| O1A—C1A—C2A—C3A | -28.7 (5) | O1B—C1B—C2B—C3B | 29.5 (5) |
| O1A—C1A—C2A—C9A | 146.2 (3) | O1B—C1B—C2B—C9B | -144.7 (3) |
| O1A—C1A—C10A—C11A | -34.1 (4) | O1B—C1B—C10B—C11B | 26.2 (4) |
| O1A—C1A—C10A—C15A | 140.5 (3) | O1B—C1B—C10B—C15B | -148.1 (3) |
| N1A—C16A—C17A—C18A | -1.1 (5) | N1B—C16B—C17B—C18B | -0.6 (6) |
| N1A—C16A—C17A—C24A | -177.3 (3) | N1B—C16B—C17B—C24B | 175.6 (3) |
| C1A—C2A—C3A—C4A | -0.2 (5) | C1B—C2B—C3B—C4B | 4.6 (5) |
| C1A—C2A—C3A—C8A | 178.3 (3) | C1B—C2B—C3B—C8B | -176.2 (3) |
| C1A—C2A—C9A—S1A | -176.7 (2) | C1B—C2B—C9B—S1B | 174.9 (3) |
| C1A—C2A—C9A—N1A | -0.3 (5) | C1B—C2B—C9B—N1B | -2.4 (5) |

| | | | |
|---------------------|------------|---------------------|------------|
| C1A—C10A—C11A—C12A | 175.9 (3) | C1B—C10B—C11B—C12B | −173.7 (3) |
| C1A—C10A—C15A—C14A | −173.2 (3) | C1B—C10B—C15B—C14B | 172.2 (3) |
| C2A—C1A—C10A—C11A | 148.3 (3) | C2B—C1B—C10B—C11B | −156.3 (3) |
| C2A—C1A—C10A—C15A | −37.1 (4) | C2B—C1B—C10B—C15B | 29.4 (5) |
| C2A—C3A—C4A—C5A | 165.5 (3) | C2B—C3B—C4B—C5B | −166.0 (3) |
| C2A—C3A—C8A—S1A | −3.0 (4) | C2B—C3B—C8B—S1B | 1.8 (4) |
| C2A—C3A—C8A—C7A | 176.6 (3) | C2B—C3B—C8B—C7B | −176.5 (3) |
| C3A—C2A—C9A—S1A | −1.4 (3) | C3B—C2B—C9B—S1B | 0.2 (3) |
| C3A—C2A—C9A—N1A | 175.0 (3) | C3B—C2B—C9B—N1B | −177.1 (3) |
| C3A—C4A—C5A—C6A | 46.8 (4) | C3B—C4B—C5B—C6B | −47.0 (4) |
| C4A—C3A—C8A—S1A | 175.6 (3) | C4B—C3B—C8B—S1B | −178.9 (3) |
| C4A—C3A—C8A—C7A | −4.8 (5) | C4B—C3B—C8B—C7B | 2.8 (6) |
| C4A—C5A—C6A—C7A | −64.8 (4) | C4B—C5B—C6B—C7B | 63.3 (4) |
| C5A—C6A—C7A—C8A | 44.4 (4) | C5B—C6B—C7B—C8B | −42.8 (4) |
| C6A—C7A—C8A—S1A | 168.1 (2) | C6B—C7B—C8B—S1B | −166.6 (3) |
| C6A—C7A—C8A—C3A | −11.4 (5) | C6B—C7B—C8B—C3B | 11.6 (5) |
| C8A—S1A—C9A—N1A | −176.7 (3) | C8B—S1B—C9B—N1B | 178.1 (3) |
| C8A—S1A—C9A—C2A | −0.2 (3) | C8B—S1B—C9B—C2B | 0.7 (3) |
| C8A—C3A—C4A—C5A | −12.9 (5) | C8B—C3B—C4B—C5B | 14.8 (5) |
| C9A—S1A—C8A—C3A | 1.8 (3) | C9B—S1B—C8B—C3B | −1.5 (3) |
| C9A—S1A—C8A—C7A | −177.8 (3) | C9B—S1B—C8B—C7B | 177.0 (3) |
| C9A—N1A—C16A—C17A | 177.6 (3) | C9B—N1B—C16B—C17B | −179.1 (3) |
| C9A—C2A—C3A—C4A | −175.6 (3) | C9B—C2B—C3B—C4B | 179.4 (3) |
| C9A—C2A—C3A—C8A | 2.9 (4) | C9B—C2B—C3B—C8B | −1.4 (4) |
| C10A—C1A—C2A—C3A | 148.8 (3) | C10B—C1B—C2B—C3B | −148.0 (3) |
| C10A—C1A—C2A—C9A | −36.3 (4) | C10B—C1B—C2B—C9B | 37.8 (5) |
| C10A—C11A—C12A—C13A | −3.1 (5) | C10B—C11B—C12B—C13B | 0.6 (6) |
| C11A—C10A—C15A—C14A | 1.3 (5) | C11B—C10B—C15B—C14B | −2.0 (5) |
| C11A—C12A—C13A—C14A | 2.6 (6) | C11B—C12B—C13B—C14B | −0.8 (6) |
| C12A—C13A—C14A—C15A | −0.1 (6) | C12B—C13B—C14B—C15B | −0.4 (6) |
| C13A—C14A—C15A—C10A | −1.9 (5) | C13B—C14B—C15B—C10B | 1.8 (5) |
| C15A—C10A—C11A—C12A | 1.1 (5) | C15B—C10B—C11B—C12B | 0.8 (5) |
| C16A—N1A—C9A—S1A | −0.8 (4) | C16B—N1B—C9B—S1B | 0.2 (4) |
| C16A—N1A—C9A—C2A | −176.8 (3) | C16B—N1B—C9B—C2B | 177.2 (3) |
| C16A—C17A—C18A—C19A | 2.3 (6) | C16B—C17B—C18B—C19B | −1.9 (6) |
| C16A—C17A—C18A—C23A | −176.5 (3) | C16B—C17B—C18B—C23B | 176.2 (3) |
| C16A—C17A—C24A—N2A | 177.2 (3) | C16B—C17B—C24B—N2B | −177.0 (3) |
| C17A—C18A—C19A—C20A | −178.0 (4) | C17B—C18B—C19B—C20B | 177.6 (4) |
| C17A—C18A—C23A—N2A | −0.6 (3) | C17B—C18B—C23B—N2B | 0.9 (3) |
| C17A—C18A—C23A—C22A | 178.5 (3) | C17B—C18B—C23B—C22B | −177.8 (3) |
| C18A—C17A—C24A—N2A | 0.3 (4) | C18B—C17B—C24B—N2B | 0.0 (4) |
| C18A—C19A—C20A—C21A | −0.4 (6) | C18B—C19B—C20B—C21B | −0.2 (6) |
| C19A—C18A—C23A—N2A | −179.5 (3) | C19B—C18B—C23B—N2B | 179.3 (3) |
| C19A—C18A—C23A—C22A | −0.5 (5) | C19B—C18B—C23B—C22B | 0.7 (5) |
| C19A—C20A—C21A—C22A | 0.1 (6) | C19B—C20B—C21B—C22B | 0.6 (6) |
| C20A—C21A—C22A—C23A | 0.0 (6) | C20B—C21B—C22B—C23B | −0.3 (6) |
| C21A—C22A—C23A—N2A | 179.0 (3) | C21B—C22B—C23B—N2B | −178.7 (3) |
| C21A—C22A—C23A—C18A | 0.2 (5) | C21B—C22B—C23B—C18B | −0.3 (5) |

| | | | |
|---------------------|------------|---------------------|------------|
| C23A—N2A—C24A—C17A | −0.6 (4) | C23B—N2B—C24B—C17B | 0.6 (4) |
| C23A—C18A—C19A—C20A | 0.6 (5) | C23B—C18B—C19B—C20B | −0.4 (5) |
| C24A—N2A—C23A—C18A | 0.7 (4) | C24B—N2B—C23B—C18B | −0.9 (4) |
| C24A—N2A—C23A—C22A | −178.2 (3) | C24B—N2B—C23B—C22B | 177.6 (3) |
| C24A—C17A—C18A—C19A | 178.9 (4) | C24B—C17B—C18B—C19B | −178.6 (4) |
| C24A—C17A—C18A—C23A | 0.2 (3) | C24B—C17B—C18B—C23B | −0.5 (3) |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|-----------------------------|------|-------|-----------|---------|
| N2A—H2A···O1A ⁱ | 0.86 | 2.01 | 2.866 (4) | 175 |
| N2B—H2B···O1B ⁱⁱ | 0.86 | 2.00 | 2.835 (3) | 163 |

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