

(1*R*,6*R*)-1-Methyl-8-azaspiro[5.6]-dodecan-7-one

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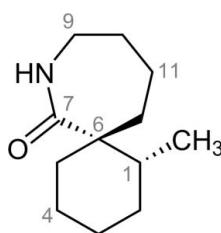
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Key indicators: single-crystal X-ray study; $T = 90$ K; mean $\sigma(C-C) = 0.003$ Å; R factor = 0.040; wR factor = 0.104; data-to-parameter ratio = 10.1.

The crystal structure of the title compound, C₁₂H₂₁NO, has been investigated to establish the absolute stereochemistry at position 1. The absolute stereochemistry at the quaternary centre at position 6 is established to be *R* using an asymmetric Birch reductive alkylation reaction for which the stereochemical outcome is known. The crystal structure indicates the presence of two conformers of the bicyclic (1*R*,6*R*)-spiro-lactam ring system that differ in the conformation adopted by the six-membered ring. In one conformer, the methyl group adopts an axial position whereas in the other conformer, the same methyl group adopts an equatorial position. In both conformers, the seven-membered ring adopts a chair conformation. The two conformers of the bicyclic spiro-lactam are connected to each other *via* intermolecular N—H···O hydrogen bonds forming a heterodimer. The asymmetric unit contains two such dimers.

Related literature

For related literature, see: Brimble & Trzoss (2004); Brimble *et al.* (2005); Ciminiello *et al.* (2007); Hu *et al.* (2001); MacKinnon *et al.* (2006); Schultz & Pettus (1997); Schultz *et al.* (1988).



Experimental

Crystal data

C₁₂H₂₁NO
 $M_r = 195.30$

Triclinic, P1
 $a = 8.5417(2)$ Å

Data collection

Bruker SMART diffractometer with APEX2 CCD detector
Absorption correction: none
25392 measured reflections

5160 independent reflections
4784 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.104$
 $S = 1.02$
5160 reflections
509 parameters

3 restraints
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.46$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1
Hydrogen-bond geometry (Å, °).

| D—H···A | D—H | H···A | D···A | D—H···A |
|---------------|------|-------|-----------|---------|
| N8—H8···O1B | 0.86 | 2.11 | 2.967 (2) | 173 |
| N8B—H8B···O1 | 0.86 | 2.03 | 2.868 (2) | 166 |
| N8A—H8A···O1C | 0.86 | 2.03 | 2.872 (2) | 165 |
| N8C—H8C···O1A | 0.86 | 2.10 | 2.959 (2) | 172 |

Data collection: SMART (Siemens, 1995); cell refinement: SAINT (Siemens, 1995); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEPIII (Burnett & Johnson, 1996); software used to prepare material for publication: SHELXTL (Sheldrick, 2008) and publCIF (Westrip, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: DN2350).

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supporting information

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(1*R*,6*R*)-1-Methyl-8-azaspiro[5.6]dodecan-7-one

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S1. Comment

The title spirolactam was prepared as part of a synthetic program directed towards the synthesis of spirolides A and C that were isolated from the culture of a toxic clone of the dinoflagellate *Alexandrium ostenfeldii* (Hu *et al.*, 2001, MacKinnon *et al.*, 2006, Ciminello *et al.*, 2007). The work demonstrates methodology to access enantiopure 7,6-spirolactams. The quaternary spirocyclic centre is generally considered a challenging stereocentre to be constructed in a stereoselective fashion in organic synthesis (Brimble *et al.*, 2005; Brimble & Trzoss, 2004). By employing (*S*)-methoxypyrrolidine as a chiral auxiliary, a highly diastereoselective Birch reductive alkylation (Schultz *et al.*, 1988 and Schultz & Pettus, 1997) furnished the alkylated product with the desired stereochemistry at the quaternary carbon which was then converted to the title spirolactam in several steps. Since the stereochemistry at C6 is known to be *R*, the absolute configuration at C1 has therefore been assigned as *R*.

The crystal structure indicates the presence of two conformers of the enantiopure bicyclic (1*R*,6*R*)-spirolactam. While in both conformers, the 7,6-bicyclic ring system adopts a chair-chair conformation, the methyl group (C13) adopts a differing position between the conformer. In one conformer, the methyl group (C13) adopts an axial position whereas in the other conformer, the same methyl group (C13B) adopts an equatorial position on their associated cyclohexane ring. In solution at room temperature, the two conformers are rapidly interconverting to each other as shown by the total lack of dynamic effects in the ¹H NMR spectrum at 400 MHz.

Each unit cell contains two heterodimers of the two chair-configured conformers of the bicyclic spirolactam. In each dimer, the axial and equatorial conformers are connected to each other by two adjacent intermolecular N—H···O hydrogen bonds (Figure 1).

S2. Experimental

To (2'S,1*R*,2*R*)-2-methyl-1-(4'-aminobutane)-1-[{(2'-ethoxymethyl)pyrrolidinyl}carbonyl]-2,5-cyclohexane (47.9 mg, 0.2 mmol) in water (1.6 ml) was added concentrated HCl (1.6 ml) and the mixture was heated under reflux overnight. After cooling to room temperature, the mixture was concentrated *in vacuo* and dried in a freeze-drier. The crude amino acid salt was dissolved in CH₂Cl₂/DMF (14.4 ml, 2:1) and DIPEA (0.15 ml, 0.9 mmol) was added. This resultant mixture was added dropwise to a solution of (benzotriazole-1-yloxy)tritypyrrolidinophosphonium hexafluorophosphate (229.0 mg, 0.4 mmol) and DMAP (53.8 mg, 0.4 mmol) in CH₂Cl₂/DMF (43.2 ml, 2:1) over 8.5 h using a syringe pump. After stirring for further 13 h, the mixture was concentrated *in vacuo*. The residual oil was dissolved in CH₂Cl₂ (50 ml) and washed with aqueous HCl solution (0.5 M, 2×50 ml). The combined aqueous layers were extracted with CH₂Cl₂ (60 ml). The combined organic layers were washed with saturated NaHCO₃ solution (60 ml), dried over anhydrous MgSO₄, filtered and concentrated *in vacuo*. Purification by flash chromatography (20:80→70:30 EtOAc-hexanes) afforded the *title compound* (7.8 mg, 26%) as a white solid. Recrystallization from CH₂Cl₂ afforded white prisms.

M. P. 392.4–393.4 K.

HRMS (+EI) calculated for $C_{12}H_{21}NO [M]^+$: 195.1623, found 195.1621.

IR (KBr plate neat) ν_{max} 3285, 2925, 2860, 1645, 1460, 1330, 1280, 1120 cm^{-1} .

^1H NMR (400 MHz, CDCl_3) δ 5.96 (1H, s, CONH), 3.34 (1 H, m, NHCH_aH_b), 3.08 (1 H, m, NHCH_aH_b), 2.15 (3 H, m, 1- CH and 5- CH_aH_b and 12- CH_aH_b), 1.74 (3H, m, 3- CH_aH_b , 5- CH_aH_b and 10- CH_aH_b), 1.50 (6 H, m, 2- CH_aH_b and 4- CH_aH_b and 10- CH_aH_b and 11- CH_2 and 12- CH_aH_b), 1.32 (2 H, m, 3- CH_aH_b and 4- CH_aH_b), 1.17 (1 H, m, 2- CH_aH_b), 1.00 (3H, d, $J = 7.1 \text{ Hz}$, 13- CH_3).

^{13}C NMR (100 MHz, CDCl_3) δ 181.0 (7-CO), 47.1 (6-C), 42.0 (9-NCH₂), 31.5 (1-CH), 29.6 (5- CH_2 and 2- CH_2), 29.4 (12- CH_2), 27.9 (3- CH_2), 23.3 (10- CH_2), 21.7 (11- CH_2), 20.6 (4- CH_2), 15.5 (13- CH_3).

m/z (+EI, 70 eV) 195 ([$M]^+$, 100), 180 (20), 166 (11), 140 (16%).

S3. Refinement

Hydrogen atoms were placed in calculated positions and refined using the riding model [C—H 0.93–0.97 \AA], with $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C})$.

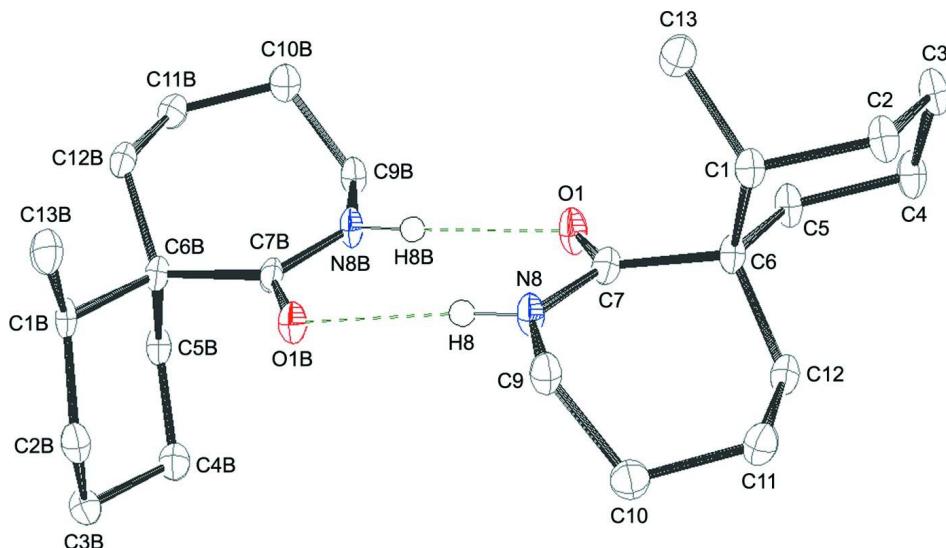


Figure 1

The molecular structure and atom numbering scheme of the heterodimer of (1*R*,6*R*)-1-methyl-8-azaspiro[5.6]dodecan-7-one. The respective methyl group (C13) adopts an axial or equatorial position on its associated cyclohexane ring as shown. Ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii. Hydrogen bonds are shown as dashed lines.

(1*R*,6*R*)-1-Methyl-8-azaspiro[5.6]dodecan-7-one

Crystal data

$C_{12}H_{21}NO$

$M_r = 195.30$

Triclinic, $P\bar{1}$

Hall symbol: P 1

$a = 8.5417(2) \text{ \AA}$

$b = 10.2807(2) \text{ \AA}$

$c = 12.6400(3) \text{ \AA}$

$\alpha = 102.850(1)^\circ$

$\beta = 90.091(1)^\circ$

$\gamma = 91.488(1)^\circ$

$V = 1081.78(4) \text{ \AA}^3$

$Z = 4$

$F(000) = 432$

$D_x = 1.199 \text{ Mg m}^{-3}$

Melting point: 392.4 K

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

Cell parameters from 9917 reflections

$\theta = 1.7\text{--}28.0^\circ$

$\mu = 0.08 \text{ mm}^{-1}$
 $T = 90 \text{ K}$

Prisms, white
 $0.34 \times 0.26 \times 0.23 \text{ mm}$

Data collection

Bruker SMART
diffractometer with APEX2 CCD detector
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
25392 measured reflections
5160 independent reflections

4784 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$
 $\theta_{\text{max}} = 28.0^\circ, \theta_{\text{min}} = 1.7^\circ$
 $h = -11 \rightarrow 10$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.104$
 $S = 1.02$
5160 reflections
509 parameters
3 restraints
Primary atom site location: structure-invariant direct methods

Secondary atom site location: difference Fourier map
Hydrogen site location: inferred from neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0667P)^2 + 0.214P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.46 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.21 \text{ e } \text{\AA}^{-3}$

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.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger. Friedel pairs were merged as recommended for light atom structures in the checkCIF program.

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|------------|------------|--------------|----------------------------------|
| C1 | 0.2813 (3) | 0.3685 (2) | 0.08818 (17) | 0.0169 (4) |
| H1 | 0.3372 | 0.3084 | 0.0303 | 0.020* |
| C2 | 0.2355 (3) | 0.2889 (2) | 0.17364 (18) | 0.0199 (4) |
| H2A | 0.1620 | 0.2176 | 0.1414 | 0.024* |
| H2B | 0.3282 | 0.2484 | 0.1953 | 0.024* |
| C3 | 0.1618 (3) | 0.3757 (2) | 0.27424 (18) | 0.0213 (5) |
| H3A | 0.0635 | 0.4092 | 0.2542 | 0.026* |
| H3B | 0.1398 | 0.3221 | 0.3270 | 0.026* |
| C4 | 0.2710 (3) | 0.4927 (2) | 0.32488 (18) | 0.0215 (5) |
| H4A | 0.3627 | 0.4594 | 0.3545 | 0.026* |
| H4B | 0.2178 | 0.5510 | 0.3841 | 0.026* |
| C5 | 0.3225 (3) | 0.5725 (2) | 0.24137 (17) | 0.0189 (4) |
| H5A | 0.3982 | 0.6411 | 0.2752 | 0.023* |
| H5B | 0.2323 | 0.6167 | 0.2205 | 0.023* |

| | | | | |
|------|-------------|------------|---------------|------------|
| C6 | 0.3959 (3) | 0.4866 (2) | 0.13778 (16) | 0.0159 (4) |
| C7 | 0.4210 (3) | 0.5784 (2) | 0.05733 (17) | 0.0165 (4) |
| C9 | 0.5649 (3) | 0.4056 (2) | -0.08228 (18) | 0.0189 (4) |
| H9A | 0.6043 | 0.4063 | -0.1541 | 0.023* |
| H9B | 0.4854 | 0.3350 | -0.0901 | 0.023* |
| C10 | 0.6984 (3) | 0.3737 (2) | -0.01351 (19) | 0.0210 (4) |
| H10A | 0.7681 | 0.4514 | 0.0062 | 0.025* |
| H10B | 0.7576 | 0.3017 | -0.0565 | 0.025* |
| C11 | 0.6438 (3) | 0.3333 (2) | 0.08969 (19) | 0.0214 (4) |
| H11A | 0.7346 | 0.3083 | 0.1261 | 0.026* |
| H11B | 0.5756 | 0.2547 | 0.0693 | 0.026* |
| C12 | 0.5571 (3) | 0.4396 (2) | 0.17042 (18) | 0.0184 (4) |
| H12A | 0.6262 | 0.5178 | 0.1894 | 0.022* |
| H12B | 0.5417 | 0.4065 | 0.2359 | 0.022* |
| C13 | 0.1352 (3) | 0.4142 (2) | 0.0369 (2) | 0.0234 (5) |
| H13A | 0.0696 | 0.3379 | 0.0061 | 0.035* |
| H13B | 0.1664 | 0.4590 | -0.0190 | 0.035* |
| H13C | 0.0783 | 0.4742 | 0.0916 | 0.035* |
| C1A | 0.0229 (3) | 0.4909 (2) | 0.74816 (17) | 0.0182 (4) |
| H1A | 0.0141 | 0.4245 | 0.7928 | 0.022* |
| C2A | 0.1973 (3) | 0.5265 (2) | 0.74572 (18) | 0.0202 (4) |
| H2A1 | 0.2118 | 0.5969 | 0.7065 | 0.024* |
| H2A2 | 0.2346 | 0.5605 | 0.8195 | 0.024* |
| C3A | 0.2950 (3) | 0.4080 (2) | 0.69248 (19) | 0.0225 (5) |
| H3A1 | 0.4024 | 0.4379 | 0.6857 | 0.027* |
| H3A2 | 0.2947 | 0.3431 | 0.7376 | 0.027* |
| C4A | 0.2290 (3) | 0.3432 (2) | 0.58066 (19) | 0.0217 (4) |
| H4A1 | 0.2886 | 0.2650 | 0.5496 | 0.026* |
| H4A2 | 0.2397 | 0.4055 | 0.5334 | 0.026* |
| C5A | 0.0563 (3) | 0.3023 (2) | 0.58674 (18) | 0.0185 (4) |
| H5A1 | 0.0470 | 0.2366 | 0.6310 | 0.022* |
| H5A2 | 0.0188 | 0.2606 | 0.5144 | 0.022* |
| C6A | -0.0490 (3) | 0.4215 (2) | 0.63499 (17) | 0.0161 (4) |
| C7A | -0.0507 (3) | 0.5153 (2) | 0.55546 (17) | 0.0156 (4) |
| C9A | -0.2115 (3) | 0.3552 (2) | 0.40755 (18) | 0.0190 (4) |
| H9A1 | -0.1394 | 0.2822 | 0.3965 | 0.023* |
| H9A2 | -0.2472 | 0.3639 | 0.3366 | 0.023* |
| C10A | -0.3520 (3) | 0.3193 (2) | 0.47040 (18) | 0.0212 (4) |
| H10C | -0.4149 | 0.3976 | 0.4937 | 0.025* |
| H10D | -0.4162 | 0.2516 | 0.4228 | 0.025* |
| C11A | -0.3040 (3) | 0.2678 (2) | 0.56903 (19) | 0.0208 (4) |
| H11C | -0.2377 | 0.1918 | 0.5455 | 0.025* |
| H11D | -0.3972 | 0.2368 | 0.6008 | 0.025* |
| C12A | -0.2166 (3) | 0.3715 (2) | 0.65661 (18) | 0.0197 (4) |
| H12C | -0.2107 | 0.3350 | 0.7209 | 0.024* |
| H12D | -0.2816 | 0.4493 | 0.6747 | 0.024* |
| C13A | -0.0654 (3) | 0.6130 (2) | 0.81015 (18) | 0.0226 (5) |
| H13D | -0.0251 | 0.6394 | 0.8829 | 0.034* |

| | | | | |
|------|-------------|------------|---------------|------------|
| H13E | -0.1749 | 0.5903 | 0.8121 | 0.034* |
| H13F | -0.0513 | 0.6853 | 0.7741 | 0.034* |
| C1B | 0.5341 (3) | 0.8408 (2) | -0.36794 (16) | 0.0178 (4) |
| H1B | 0.5290 | 0.9070 | -0.4128 | 0.021* |
| C2B | 0.7072 (3) | 0.8067 (2) | -0.36470 (18) | 0.0199 (4) |
| H2B1 | 0.7181 | 0.7366 | -0.3251 | 0.024* |
| H2B2 | 0.7428 | 0.7725 | -0.4382 | 0.024* |
| C3B | 0.8108 (3) | 0.9260 (2) | -0.31153 (19) | 0.0209 (4) |
| H3B1 | 0.8137 | 0.9908 | -0.3568 | 0.025* |
| H3B2 | 0.9168 | 0.8971 | -0.3044 | 0.025* |
| C4B | 0.7482 (3) | 0.9906 (2) | -0.19965 (18) | 0.0203 (4) |
| H4B1 | 0.8119 | 1.0693 | -0.1685 | 0.024* |
| H4B2 | 0.7554 | 0.9286 | -0.1523 | 0.024* |
| C5B | 0.5778 (3) | 1.0301 (2) | -0.20684 (17) | 0.0180 (4) |
| H5B1 | 0.5723 | 1.0955 | -0.2514 | 0.022* |
| H5B2 | 0.5423 | 1.0720 | -0.1347 | 0.022* |
| C6B | 0.4660 (3) | 0.9104 (2) | -0.25522 (17) | 0.0153 (4) |
| C7B | 0.4590 (3) | 0.8169 (2) | -0.17528 (16) | 0.0152 (4) |
| C9B | 0.3049 (3) | 0.9764 (2) | -0.02876 (18) | 0.0185 (4) |
| H9B1 | 0.3806 | 1.0502 | -0.0171 | 0.022* |
| H9B2 | 0.2677 | 0.9675 | 0.0418 | 0.022* |
| C10B | 0.1677 (3) | 1.0105 (2) | -0.09283 (19) | 0.0207 (4) |
| H10E | 0.1063 | 1.0777 | -0.0460 | 0.025* |
| H10F | 0.1011 | 0.9314 | -0.1166 | 0.025* |
| C11B | 0.2195 (3) | 1.0621 (2) | -0.19119 (18) | 0.0191 (4) |
| H11E | 0.2901 | 1.1385 | -0.1673 | 0.023* |
| H11F | 0.1284 | 1.0925 | -0.2237 | 0.023* |
| C12B | 0.3016 (3) | 0.9586 (2) | -0.27761 (18) | 0.0184 (4) |
| H12E | 0.3094 | 0.9945 | -0.3422 | 0.022* |
| H12F | 0.2325 | 0.8803 | -0.2955 | 0.022* |
| C13B | 0.4399 (3) | 0.7182 (2) | -0.43024 (18) | 0.0226 (5) |
| H13G | 0.4805 | 0.6909 | -0.5023 | 0.034* |
| H13H | 0.3319 | 0.7404 | -0.4340 | 0.034* |
| H13I | 0.4485 | 0.6466 | -0.3933 | 0.034* |
| C1C | -0.2016 (3) | 0.9626 (2) | 0.29179 (17) | 0.0170 (4) |
| H1C | -0.1435 | 1.0228 | 0.3506 | 0.020* |
| C2C | -0.2420 (3) | 1.0427 (2) | 0.20650 (17) | 0.0196 (4) |
| H2C1 | -0.3126 | 1.1129 | 0.2384 | 0.023* |
| H2C2 | -0.1470 | 1.0846 | 0.1864 | 0.023* |
| C3C | -0.3183 (3) | 0.9565 (2) | 0.10453 (18) | 0.0210 (4) |
| H3C1 | -0.4190 | 0.9219 | 0.1228 | 0.025* |
| H3C2 | -0.3361 | 1.0107 | 0.0521 | 0.025* |
| C4C | -0.2146 (3) | 0.8405 (2) | 0.05437 (18) | 0.0206 (4) |
| H4C1 | -0.1205 | 0.8749 | 0.0260 | 0.025* |
| H4C2 | -0.2700 | 0.7823 | -0.0057 | 0.025* |
| C5C | -0.1689 (3) | 0.7601 (2) | 0.13766 (17) | 0.0192 (4) |
| H5C1 | -0.0964 | 0.6922 | 0.1042 | 0.023* |
| H5C2 | -0.2618 | 0.7150 | 0.1573 | 0.023* |

| | | | | |
|------|--------------|--------------|---------------|------------|
| C6C | -0.0923 (3) | 0.8458 (2) | 0.24222 (16) | 0.0152 (4) |
| C7C | -0.0736 (3) | 0.7534 (2) | 0.32233 (17) | 0.0172 (4) |
| C9C | 0.0785 (3) | 0.9260 (2) | 0.46415 (18) | 0.0195 (4) |
| H9C1 | 0.1166 | 0.9250 | 0.5363 | 0.023* |
| H9C2 | 0.0033 | 0.9964 | 0.4713 | 0.023* |
| C10C | 0.2153 (3) | 0.9581 (2) | 0.39660 (19) | 0.0223 (5) |
| H10G | 0.2795 | 0.8801 | 0.3766 | 0.027* |
| H10H | 0.2793 | 1.0293 | 0.4405 | 0.027* |
| C11C | 0.1644 (3) | 1.0004 (2) | 0.29360 (19) | 0.0226 (5) |
| H11G | 0.1011 | 1.0788 | 0.3143 | 0.027* |
| H11H | 0.2572 | 1.0259 | 0.2581 | 0.027* |
| C12C | 0.0718 (3) | 0.8950 (2) | 0.21178 (18) | 0.0188 (4) |
| H12G | 0.1366 | 0.8173 | 0.1921 | 0.023* |
| H12H | 0.0586 | 0.9294 | 0.1469 | 0.023* |
| C13C | -0.3507 (3) | 0.9146 (2) | 0.3408 (2) | 0.0236 (5) |
| H13J | -0.4126 | 0.9900 | 0.3719 | 0.035* |
| H13K | -0.3226 | 0.8688 | 0.3962 | 0.035* |
| H13L | -0.4101 | 0.8549 | 0.2850 | 0.035* |
| N8 | 0.4924 (2) | 0.53304 (18) | -0.03820 (15) | 0.0186 (4) |
| H8 | 0.4959 | 0.5884 | -0.0802 | 0.022* |
| N8A | -0.1270 (2) | 0.47750 (18) | 0.45948 (15) | 0.0187 (4) |
| H8A | -0.1254 | 0.5375 | 0.4216 | 0.022* |
| N8B | 0.3840 (2) | 0.85451 (18) | -0.07988 (15) | 0.0180 (4) |
| H8B | 0.3826 | 0.7949 | -0.0417 | 0.022* |
| N8C | -0.0011 (2) | 0.79845 (18) | 0.41869 (15) | 0.0184 (4) |
| H8C | -0.0018 | 0.7428 | 0.4604 | 0.022* |
| O1 | 0.3740 (2) | 0.69436 (15) | 0.07917 (13) | 0.0225 (3) |
| O1A | 0.01941 (19) | 0.62531 (15) | 0.57620 (12) | 0.0188 (3) |
| O1B | 0.52257 (19) | 0.70723 (15) | -0.19575 (12) | 0.0182 (3) |
| O1C | -0.1266 (2) | 0.63695 (15) | 0.29949 (13) | 0.0222 (3) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|-------------|------------|
| C1 | 0.0211 (10) | 0.0169 (10) | 0.0134 (10) | 0.0004 (8) | 0.0009 (8) | 0.0048 (8) |
| C2 | 0.0296 (12) | 0.0158 (10) | 0.0151 (10) | -0.0021 (8) | 0.0021 (9) | 0.0056 (8) |
| C3 | 0.0303 (12) | 0.0187 (10) | 0.0165 (10) | -0.0002 (9) | 0.0053 (9) | 0.0077 (8) |
| C4 | 0.0348 (13) | 0.0183 (10) | 0.0120 (10) | 0.0007 (9) | 0.0030 (9) | 0.0045 (8) |
| C5 | 0.0298 (12) | 0.0143 (9) | 0.0134 (10) | 0.0015 (8) | 0.0031 (8) | 0.0046 (8) |
| C6 | 0.0228 (11) | 0.0149 (9) | 0.0109 (9) | 0.0000 (8) | -0.0001 (8) | 0.0047 (8) |
| C7 | 0.0210 (11) | 0.0158 (9) | 0.0130 (10) | -0.0003 (8) | -0.0018 (8) | 0.0041 (8) |
| C9 | 0.0245 (11) | 0.0192 (10) | 0.0140 (10) | 0.0025 (8) | 0.0026 (8) | 0.0053 (8) |
| C10 | 0.0214 (11) | 0.0204 (11) | 0.0216 (11) | 0.0037 (8) | 0.0022 (9) | 0.0053 (9) |
| C11 | 0.0247 (11) | 0.0209 (11) | 0.0205 (11) | 0.0030 (9) | -0.0015 (9) | 0.0082 (9) |
| C12 | 0.0235 (11) | 0.0189 (10) | 0.0144 (10) | 0.0000 (8) | -0.0018 (8) | 0.0070 (8) |
| C13 | 0.0250 (12) | 0.0262 (11) | 0.0206 (11) | 0.0000 (9) | -0.0027 (9) | 0.0087 (9) |
| C1A | 0.0250 (11) | 0.0197 (10) | 0.0115 (9) | 0.0010 (8) | 0.0000 (8) | 0.0066 (8) |
| C2A | 0.0288 (12) | 0.0206 (10) | 0.0127 (9) | -0.0014 (9) | -0.0036 (8) | 0.0072 (8) |

| | | | | | | |
|------|-------------|-------------|-------------|-------------|-------------|------------|
| C3A | 0.0242 (11) | 0.0255 (11) | 0.0196 (11) | 0.0019 (9) | -0.0014 (9) | 0.0086 (9) |
| C4A | 0.0252 (11) | 0.0207 (10) | 0.0195 (11) | 0.0047 (8) | 0.0017 (9) | 0.0049 (8) |
| C5A | 0.0261 (12) | 0.0167 (10) | 0.0139 (10) | 0.0025 (8) | -0.0009 (8) | 0.0054 (8) |
| C6A | 0.0219 (11) | 0.0169 (9) | 0.0107 (9) | 0.0026 (8) | 0.0012 (8) | 0.0055 (7) |
| C7A | 0.0197 (10) | 0.0164 (9) | 0.0120 (9) | 0.0043 (8) | 0.0034 (8) | 0.0053 (7) |
| C9A | 0.0271 (11) | 0.0175 (10) | 0.0130 (10) | -0.0014 (8) | -0.0031 (8) | 0.0048 (8) |
| C10A | 0.0244 (11) | 0.0225 (11) | 0.0171 (10) | -0.0009 (9) | -0.0021 (9) | 0.0055 (8) |
| C11A | 0.0251 (12) | 0.0201 (10) | 0.0184 (11) | -0.0018 (8) | 0.0009 (9) | 0.0070 (8) |
| C12A | 0.0247 (11) | 0.0208 (10) | 0.0149 (10) | -0.0005 (9) | 0.0023 (8) | 0.0072 (8) |
| C13A | 0.0303 (12) | 0.0239 (11) | 0.0131 (10) | 0.0017 (9) | 0.0002 (9) | 0.0030 (8) |
| C1B | 0.0265 (11) | 0.0197 (10) | 0.0085 (9) | 0.0005 (8) | 0.0013 (8) | 0.0061 (8) |
| C2B | 0.0257 (11) | 0.0211 (10) | 0.0138 (10) | 0.0017 (9) | 0.0036 (8) | 0.0055 (8) |
| C3B | 0.0211 (11) | 0.0239 (11) | 0.0192 (11) | 0.0004 (9) | 0.0029 (8) | 0.0083 (9) |
| C4B | 0.0241 (11) | 0.0201 (10) | 0.0174 (10) | -0.0017 (8) | -0.0007 (9) | 0.0057 (8) |
| C5B | 0.0267 (11) | 0.0162 (10) | 0.0117 (9) | -0.0013 (8) | 0.0006 (8) | 0.0049 (8) |
| C6B | 0.0208 (10) | 0.0152 (9) | 0.0109 (9) | -0.0002 (8) | -0.0014 (8) | 0.0049 (7) |
| C7B | 0.0192 (10) | 0.0174 (9) | 0.0099 (9) | -0.0023 (8) | -0.0027 (7) | 0.0054 (7) |
| C9B | 0.0255 (11) | 0.0180 (10) | 0.0129 (10) | 0.0020 (8) | 0.0025 (8) | 0.0053 (8) |
| C10B | 0.0253 (12) | 0.0194 (10) | 0.0178 (10) | 0.0025 (9) | 0.0015 (9) | 0.0046 (8) |
| C11B | 0.0218 (11) | 0.0188 (10) | 0.0181 (10) | 0.0015 (8) | -0.0021 (8) | 0.0068 (8) |
| C12B | 0.0219 (11) | 0.0212 (11) | 0.0138 (10) | 0.0011 (8) | -0.0025 (8) | 0.0074 (8) |
| C13B | 0.0300 (12) | 0.0239 (11) | 0.0130 (10) | -0.0011 (9) | 0.0000 (9) | 0.0023 (8) |
| C1C | 0.0225 (11) | 0.0165 (10) | 0.0128 (10) | 0.0018 (8) | 0.0006 (8) | 0.0050 (8) |
| C2C | 0.0279 (12) | 0.0175 (10) | 0.0142 (10) | 0.0034 (8) | -0.0019 (8) | 0.0053 (8) |
| C3C | 0.0325 (12) | 0.0188 (10) | 0.0129 (10) | 0.0033 (9) | -0.0045 (8) | 0.0054 (8) |
| C4C | 0.0340 (12) | 0.0170 (10) | 0.0113 (9) | 0.0018 (9) | -0.0027 (9) | 0.0040 (8) |
| C5C | 0.0310 (12) | 0.0142 (9) | 0.0125 (10) | 0.0008 (8) | -0.0031 (8) | 0.0033 (7) |
| C6C | 0.0220 (10) | 0.0148 (9) | 0.0101 (9) | -0.0002 (8) | 0.0004 (8) | 0.0056 (8) |
| C7C | 0.0207 (11) | 0.0188 (10) | 0.0133 (10) | 0.0005 (8) | 0.0006 (8) | 0.0062 (8) |
| C9C | 0.0267 (12) | 0.0193 (10) | 0.0133 (10) | -0.0021 (8) | -0.0034 (8) | 0.0056 (8) |
| C10C | 0.0261 (12) | 0.0204 (11) | 0.0213 (11) | -0.0027 (9) | -0.0020 (9) | 0.0069 (9) |
| C11C | 0.0263 (12) | 0.0214 (11) | 0.0218 (11) | -0.0029 (9) | 0.0004 (9) | 0.0091 (9) |
| C12C | 0.0234 (11) | 0.0199 (10) | 0.0154 (10) | 0.0018 (8) | 0.0037 (8) | 0.0086 (8) |
| C13C | 0.0254 (12) | 0.0278 (12) | 0.0191 (11) | 0.0016 (9) | 0.0039 (9) | 0.0086 (9) |
| N8 | 0.0268 (10) | 0.0172 (8) | 0.0138 (8) | 0.0030 (7) | 0.0031 (7) | 0.0073 (7) |
| N8A | 0.0293 (10) | 0.0160 (8) | 0.0124 (8) | -0.0018 (7) | -0.0021 (7) | 0.0072 (7) |
| N8B | 0.0279 (10) | 0.0161 (8) | 0.0122 (8) | 0.0025 (7) | 0.0024 (7) | 0.0075 (7) |
| N8C | 0.0277 (10) | 0.0167 (8) | 0.0126 (8) | -0.0017 (7) | -0.0013 (7) | 0.0078 (7) |
| O1 | 0.0352 (9) | 0.0169 (7) | 0.0174 (8) | 0.0043 (7) | 0.0055 (7) | 0.0079 (6) |
| O1A | 0.0277 (8) | 0.0152 (7) | 0.0147 (7) | -0.0004 (6) | -0.0012 (6) | 0.0061 (6) |
| O1B | 0.0263 (8) | 0.0156 (7) | 0.0140 (7) | 0.0012 (6) | 0.0014 (6) | 0.0057 (6) |
| O1C | 0.0348 (9) | 0.0169 (7) | 0.0165 (8) | -0.0032 (6) | -0.0051 (7) | 0.0074 (6) |

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

| | | | |
|--------|-----------|---------|-----------|
| C1—C13 | 1.537 (3) | C1B—C6B | 1.565 (3) |
| C1—C2 | 1.539 (3) | C1B—H1B | 0.9800 |
| C1—C6 | 1.556 (3) | C2B—C3B | 1.521 (3) |

| | | | |
|----------|-----------|-----------|-----------|
| C1—H1 | 0.9800 | C2B—H2B1 | 0.9700 |
| C2—C3 | 1.529 (3) | C2B—H2B2 | 0.9700 |
| C2—H2A | 0.9700 | C3B—C4B | 1.527 (3) |
| C2—H2B | 0.9700 | C3B—H3B1 | 0.9700 |
| C3—C4 | 1.525 (3) | C3B—H3B2 | 0.9700 |
| C3—H3A | 0.9700 | C4B—C5B | 1.529 (3) |
| C3—H3B | 0.9700 | C4B—H4B1 | 0.9700 |
| C4—C5 | 1.532 (3) | C4B—H4B2 | 0.9700 |
| C4—H4A | 0.9700 | C5B—C6B | 1.550 (3) |
| C4—H4B | 0.9700 | C5B—H5B1 | 0.9700 |
| C5—C6 | 1.551 (3) | C5B—H5B2 | 0.9700 |
| C5—H5A | 0.9700 | C6B—C7B | 1.542 (3) |
| C5—H5B | 0.9700 | C6B—C12B | 1.548 (3) |
| C6—C7 | 1.545 (3) | C7B—O1B | 1.239 (3) |
| C6—C12 | 1.556 (3) | C7B—N8B | 1.350 (3) |
| C7—O1 | 1.240 (3) | C9B—N8B | 1.460 (3) |
| C7—N8 | 1.347 (3) | C9B—C10B | 1.515 (3) |
| C9—N8 | 1.459 (3) | C9B—H9B1 | 0.9700 |
| C9—C10 | 1.518 (3) | C9B—H9B2 | 0.9700 |
| C9—H9A | 0.9700 | C10B—C11B | 1.519 (3) |
| C9—H9B | 0.9700 | C10B—H10E | 0.9700 |
| C10—C11 | 1.525 (3) | C10B—H10F | 0.9700 |
| C10—H10A | 0.9700 | C11B—C12B | 1.530 (3) |
| C10—H10B | 0.9700 | C11B—H11E | 0.9700 |
| C11—C12 | 1.527 (3) | C11B—H11F | 0.9700 |
| C11—H11A | 0.9700 | C12B—H12E | 0.9700 |
| C11—H11B | 0.9700 | C12B—H12F | 0.9700 |
| C12—H12A | 0.9700 | C13B—H13G | 0.9600 |
| C12—H12B | 0.9700 | C13B—H13H | 0.9600 |
| C13—H13A | 0.9600 | C13B—H13I | 0.9600 |
| C13—H13B | 0.9600 | C1C—C13C | 1.534 (3) |
| C13—H13C | 0.9600 | C1C—C2C | 1.539 (3) |
| C1A—C2A | 1.527 (3) | C1C—C6C | 1.560 (3) |
| C1A—C13A | 1.540 (3) | C1C—H1C | 0.9800 |
| C1A—C6A | 1.566 (3) | C2C—C3C | 1.526 (3) |
| C1A—H1A | 0.9800 | C2C—H2C1 | 0.9700 |
| C2A—C3A | 1.524 (3) | C2C—H2C2 | 0.9700 |
| C2A—H2A1 | 0.9700 | C3C—C4C | 1.526 (3) |
| C2A—H2A2 | 0.9700 | C3C—H3C1 | 0.9700 |
| C3A—C4A | 1.523 (3) | C3C—H3C2 | 0.9700 |
| C3A—H3A1 | 0.9700 | C4C—C5C | 1.532 (3) |
| C3A—H3A2 | 0.9700 | C4C—H4C1 | 0.9700 |
| C4A—C5A | 1.531 (3) | C4C—H4C2 | 0.9700 |
| C4A—H4A1 | 0.9700 | C5C—C6C | 1.549 (3) |
| C4A—H4A2 | 0.9700 | C5C—H5C1 | 0.9700 |
| C5A—C6A | 1.552 (3) | C5C—H5C2 | 0.9700 |
| C5A—H5A1 | 0.9700 | C6C—C7C | 1.545 (3) |
| C5A—H5A2 | 0.9700 | C6C—C12C | 1.555 (3) |

| | | | |
|------------|-------------|---------------|-------------|
| C6A—C7A | 1.540 (3) | C7C—O1C | 1.241 (3) |
| C6A—C12A | 1.554 (3) | C7C—N8C | 1.347 (3) |
| C7A—O1A | 1.241 (3) | C9C—N8C | 1.459 (3) |
| C7A—N8A | 1.350 (3) | C9C—C10C | 1.520 (3) |
| C9A—N8A | 1.454 (3) | C9C—H9C1 | 0.9700 |
| C9A—C10A | 1.523 (3) | C9C—H9C2 | 0.9700 |
| C9A—H9A1 | 0.9700 | C10C—C11C | 1.527 (3) |
| C9A—H9A2 | 0.9700 | C10C—H10G | 0.9700 |
| C10A—C11A | 1.518 (3) | C10C—H10H | 0.9700 |
| C10A—H10C | 0.9700 | C11C—C12C | 1.524 (3) |
| C10A—H10D | 0.9700 | C11C—H11G | 0.9700 |
| C11A—C12A | 1.533 (3) | C11C—H11H | 0.9700 |
| C11A—H11C | 0.9700 | C12C—H12G | 0.9700 |
| C11A—H11D | 0.9700 | C12C—H12H | 0.9700 |
| C12A—H12C | 0.9700 | C13C—H13J | 0.9600 |
| C12A—H12D | 0.9700 | C13C—H13K | 0.9600 |
| C13A—H13D | 0.9600 | C13C—H13L | 0.9600 |
| C13A—H13E | 0.9600 | N8—H8 | 0.8600 |
| C13A—H13F | 0.9600 | N8A—H8A | 0.8600 |
| C1B—C2B | 1.531 (3) | N8B—H8B | 0.8600 |
| C1B—C13B | 1.537 (3) | N8C—H8C | 0.8600 |
| | | | |
| C13—C1—C2 | 110.99 (19) | C3B—C2B—C1B | 112.89 (18) |
| C13—C1—C6 | 112.59 (17) | C3B—C2B—H2B1 | 109.0 |
| C2—C1—C6 | 110.66 (17) | C1B—C2B—H2B1 | 109.0 |
| C13—C1—H1 | 107.4 | C3B—C2B—H2B2 | 109.0 |
| C2—C1—H1 | 107.4 | C1B—C2B—H2B2 | 109.0 |
| C6—C1—H1 | 107.4 | H2B1—C2B—H2B2 | 107.8 |
| C3—C2—C1 | 112.60 (17) | C2B—C3B—C4B | 110.34 (18) |
| C3—C2—H2A | 109.1 | C2B—C3B—H3B1 | 109.6 |
| C1—C2—H2A | 109.1 | C4B—C3B—H3B1 | 109.6 |
| C3—C2—H2B | 109.1 | C2B—C3B—H3B2 | 109.6 |
| C1—C2—H2B | 109.1 | C4B—C3B—H3B2 | 109.6 |
| H2A—C2—H2B | 107.8 | H3B1—C3B—H3B2 | 108.1 |
| C4—C3—C2 | 110.84 (19) | C3B—C4B—C5B | 111.08 (18) |
| C4—C3—H3A | 109.5 | C3B—C4B—H4B1 | 109.4 |
| C2—C3—H3A | 109.5 | C5B—C4B—H4B1 | 109.4 |
| C4—C3—H3B | 109.5 | C3B—C4B—H4B2 | 109.4 |
| C2—C3—H3B | 109.5 | C5B—C4B—H4B2 | 109.4 |
| H3A—C3—H3B | 108.1 | H4B1—C4B—H4B2 | 108.0 |
| C3—C4—C5 | 111.46 (18) | C4B—C5B—C6B | 113.30 (18) |
| C3—C4—H4A | 109.3 | C4B—C5B—H5B1 | 108.9 |
| C5—C4—H4A | 109.3 | C6B—C5B—H5B1 | 108.9 |
| C3—C4—H4B | 109.3 | C4B—C5B—H5B2 | 108.9 |
| C5—C4—H4B | 109.3 | C6B—C5B—H5B2 | 108.9 |
| H4A—C4—H4B | 108.0 | H5B1—C5B—H5B2 | 107.7 |
| C4—C5—C6 | 113.92 (17) | C7B—C6B—C12B | 111.70 (17) |
| C4—C5—H5A | 108.8 | C7B—C6B—C5B | 108.14 (16) |

| | | | |
|---------------|-------------|----------------|-------------|
| C6—C5—H5A | 108.8 | C12B—C6B—C5B | 110.97 (17) |
| C4—C5—H5B | 108.8 | C7B—C6B—C1B | 112.38 (17) |
| C6—C5—H5B | 108.8 | C12B—C6B—C1B | 106.61 (17) |
| H5A—C5—H5B | 107.7 | C5B—C6B—C1B | 106.96 (17) |
| C7—C6—C5 | 106.96 (17) | O1B—C7B—N8B | 118.46 (18) |
| C7—C6—C1 | 110.23 (16) | O1B—C7B—C6B | 121.51 (18) |
| C5—C6—C1 | 109.32 (17) | N8B—C7B—C6B | 120.03 (18) |
| C7—C6—C12 | 109.54 (17) | N8B—C9B—C10B | 114.80 (18) |
| C5—C6—C12 | 107.66 (17) | N8B—C9B—H9B1 | 108.6 |
| C1—C6—C12 | 112.92 (16) | C10B—C9B—H9B1 | 108.6 |
| O1—C7—N8 | 119.01 (19) | N8B—C9B—H9B2 | 108.6 |
| O1—C7—C6 | 120.49 (19) | C10B—C9B—H9B2 | 108.6 |
| N8—C7—C6 | 120.49 (18) | H9B1—C9B—H9B2 | 107.5 |
| N8—C9—C10 | 114.21 (19) | C9B—C10B—C11B | 112.40 (19) |
| N8—C9—H9A | 108.7 | C9B—C10B—H10E | 109.1 |
| C10—C9—H9A | 108.7 | C11B—C10B—H10E | 109.1 |
| N8—C9—H9B | 108.7 | C9B—C10B—H10F | 109.1 |
| C10—C9—H9B | 108.7 | C11B—C10B—H10F | 109.1 |
| H9A—C9—H9B | 107.6 | H10E—C10B—H10F | 107.9 |
| C9—C10—C11 | 113.44 (19) | C10B—C11B—C12B | 113.93 (18) |
| C9—C10—H10A | 108.9 | C10B—C11B—H11E | 108.8 |
| C11—C10—H10A | 108.9 | C12B—C11B—H11E | 108.8 |
| C9—C10—H10B | 108.9 | C10B—C11B—H11F | 108.8 |
| C11—C10—H10B | 108.9 | C12B—C11B—H11F | 108.8 |
| H10A—C10—H10B | 107.7 | H11E—C11B—H11F | 107.7 |
| C10—C11—C12 | 115.62 (18) | C11B—C12B—C6B | 120.43 (19) |
| C10—C11—H11A | 108.4 | C11B—C12B—H12E | 107.2 |
| C12—C11—H11A | 108.4 | C6B—C12B—H12E | 107.2 |
| C10—C11—H11B | 108.4 | C11B—C12B—H12F | 107.2 |
| C12—C11—H11B | 108.4 | C6B—C12B—H12F | 107.2 |
| H11A—C11—H11B | 107.4 | H12E—C12B—H12F | 106.9 |
| C11—C12—C6 | 119.42 (18) | C1B—C13B—H13G | 109.5 |
| C11—C12—H12A | 107.5 | C1B—C13B—H13H | 109.5 |
| C6—C12—H12A | 107.5 | H13G—C13B—H13H | 109.5 |
| C11—C12—H12B | 107.5 | C1B—C13B—H13I | 109.5 |
| C6—C12—H12B | 107.5 | H13G—C13B—H13I | 109.5 |
| H12A—C12—H12B | 107.0 | H13H—C13B—H13I | 109.5 |
| C1—C13—H13A | 109.5 | C13C—C1C—C2C | 110.98 (19) |
| C1—C13—H13B | 109.5 | C13C—C1C—C6C | 112.58 (17) |
| H13A—C13—H13B | 109.5 | C2C—C1C—C6C | 110.45 (17) |
| C1—C13—H13C | 109.5 | C13C—C1C—H1C | 107.5 |
| H13A—C13—H13C | 109.5 | C2C—C1C—H1C | 107.5 |
| H13B—C13—H13C | 109.5 | C6C—C1C—H1C | 107.5 |
| C2A—C1A—C13A | 109.58 (18) | C3C—C2C—C1C | 112.85 (18) |
| C2A—C1A—C6A | 114.28 (17) | C3C—C2C—H2C1 | 109.0 |
| C13A—C1A—C6A | 115.17 (18) | C1C—C2C—H2C1 | 109.0 |
| C2A—C1A—H1A | 105.6 | C3C—C2C—H2C2 | 109.0 |
| C13A—C1A—H1A | 105.6 | C1C—C2C—H2C2 | 109.0 |

| | | | |
|----------------|-------------|----------------|-------------|
| C6A—C1A—H1A | 105.6 | H2C1—C2C—H2C2 | 107.8 |
| C3A—C2A—C1A | 112.93 (19) | C4C—C3C—C2C | 110.91 (19) |
| C3A—C2A—H2A1 | 109.0 | C4C—C3C—H3C1 | 109.5 |
| C1A—C2A—H2A1 | 109.0 | C2C—C3C—H3C1 | 109.5 |
| C3A—C2A—H2A2 | 109.0 | C4C—C3C—H3C2 | 109.5 |
| C1A—C2A—H2A2 | 109.0 | C2C—C3C—H3C2 | 109.5 |
| H2A1—C2A—H2A2 | 107.8 | H3C1—C3C—H3C2 | 108.0 |
| C4A—C3A—C2A | 110.37 (19) | C3C—C4C—C5C | 111.50 (18) |
| C4A—C3A—H3A1 | 109.6 | C3C—C4C—H4C1 | 109.3 |
| C2A—C3A—H3A1 | 109.6 | C5C—C4C—H4C1 | 109.3 |
| C4A—C3A—H3A2 | 109.6 | C3C—C4C—H4C2 | 109.3 |
| C2A—C3A—H3A2 | 109.6 | C5C—C4C—H4C2 | 109.3 |
| H3A1—C3A—H3A2 | 108.1 | H4C1—C4C—H4C2 | 108.0 |
| C3A—C4A—C5A | 111.31 (19) | C4C—C5C—C6C | 113.84 (17) |
| C3A—C4A—H4A1 | 109.4 | C4C—C5C—H5C1 | 108.8 |
| C5A—C4A—H4A1 | 109.4 | C6C—C5C—H5C1 | 108.8 |
| C3A—C4A—H4A2 | 109.4 | C4C—C5C—H5C2 | 108.8 |
| C5A—C4A—H4A2 | 109.4 | C6C—C5C—H5C2 | 108.8 |
| H4A1—C4A—H4A2 | 108.0 | H5C1—C5C—H5C2 | 107.7 |
| C4A—C5A—C6A | 113.14 (18) | C7C—C6C—C5C | 106.96 (17) |
| C4A—C5A—H5A1 | 109.0 | C7C—C6C—C12C | 109.73 (18) |
| C6A—C5A—H5A1 | 109.0 | C5C—C6C—C12C | 107.82 (17) |
| C4A—C5A—H5A2 | 109.0 | C7C—C6C—C1C | 109.88 (16) |
| C6A—C5A—H5A2 | 109.0 | C5C—C6C—C1C | 109.36 (18) |
| H5A1—C5A—H5A2 | 107.8 | C12C—C6C—C1C | 112.90 (16) |
| C7A—C6A—C5A | 108.18 (16) | O1C—C7C—N8C | 118.87 (19) |
| C7A—C6A—C12A | 111.81 (18) | O1C—C7C—C6C | 120.59 (19) |
| C5A—C6A—C12A | 110.80 (17) | N8C—C7C—C6C | 120.53 (18) |
| C7A—C6A—C1A | 112.39 (17) | N8C—C9C—C10C | 113.87 (18) |
| C5A—C6A—C1A | 106.92 (17) | N8C—C9C—H9C1 | 108.8 |
| C12A—C6A—C1A | 106.65 (17) | C10C—C9C—H9C1 | 108.8 |
| O1A—C7A—N8A | 118.28 (19) | N8C—C9C—H9C2 | 108.8 |
| O1A—C7A—C6A | 121.6 (2) | C10C—C9C—H9C2 | 108.8 |
| N8A—C7A—C6A | 120.09 (19) | H9C1—C9C—H9C2 | 107.7 |
| N8A—C9A—C10A | 114.84 (18) | C9C—C10C—C11C | 113.3 (2) |
| N8A—C9A—H9A1 | 108.6 | C9C—C10C—H10G | 108.9 |
| C10A—C9A—H9A1 | 108.6 | C11C—C10C—H10G | 108.9 |
| N8A—C9A—H9A2 | 108.6 | C9C—C10C—H10H | 108.9 |
| C10A—C9A—H9A2 | 108.6 | C11C—C10C—H10H | 108.9 |
| H9A1—C9A—H9A2 | 107.5 | H10G—C10C—H10H | 107.7 |
| C11A—C10A—C9A | 112.36 (19) | C12C—C11C—C10C | 115.43 (19) |
| C11A—C10A—H10C | 109.1 | C12C—C11C—H11G | 108.4 |
| C9A—C10A—H10C | 109.1 | C10C—C11C—H11G | 108.4 |
| C11A—C10A—H10D | 109.1 | C12C—C11C—H11H | 108.4 |
| C9A—C10A—H10D | 109.1 | C10C—C11C—H11H | 108.4 |
| H10C—C10A—H10D | 107.9 | H11G—C11C—H11H | 107.5 |
| C10A—C11A—C12A | 114.16 (19) | C11C—C12C—C6C | 119.95 (18) |
| C10A—C11A—H11C | 108.7 | C11C—C12C—H12G | 107.3 |

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| C12A—C11A—H11C | 108.7 | C6C—C12C—H12G | 107.3 |
| C10A—C11A—H11D | 108.7 | C11C—C12C—H12H | 107.3 |
| C12A—C11A—H11D | 108.7 | C6C—C12C—H12H | 107.3 |
| H11C—C11A—H11D | 107.6 | H12G—C12C—H12H | 106.9 |
| C11A—C12A—C6A | 120.26 (18) | C1C—C13C—H13J | 109.5 |
| C11A—C12A—H12C | 107.3 | C1C—C13C—H13K | 109.5 |
| C6A—C12A—H12C | 107.3 | H13J—C13C—H13K | 109.5 |
| C11A—C12A—H12D | 107.3 | C1C—C13C—H13L | 109.5 |
| C6A—C12A—H12D | 107.3 | H13J—C13C—H13L | 109.5 |
| H12C—C12A—H12D | 106.9 | H13K—C13C—H13L | 109.5 |
| C1A—C13A—H13D | 109.5 | C7—N8—C9 | 130.71 (18) |
| C1A—C13A—H13E | 109.5 | C7—N8—H8 | 114.6 |
| H13D—C13A—H13E | 109.5 | C9—N8—H8 | 114.6 |
| C1A—C13A—H13F | 109.5 | C7A—N8A—C9A | 132.58 (18) |
| H13D—C13A—H13F | 109.5 | C7A—N8A—H8A | 113.7 |
| H13E—C13A—H13F | 109.5 | C9A—N8A—H8A | 113.7 |
| C2B—C1B—C13B | 109.51 (18) | C7B—N8B—C9B | 132.66 (18) |
| C2B—C1B—C6B | 114.23 (17) | C7B—N8B—H8B | 113.7 |
| C13B—C1B—C6B | 115.45 (18) | C9B—N8B—H8B | 113.7 |
| C2B—C1B—H1B | 105.6 | C7C—N8C—C9C | 130.77 (18) |
| C13B—C1B—H1B | 105.6 | C7C—N8C—H8C | 114.6 |
| C6B—C1B—H1B | 105.6 | C9C—N8C—H8C | 114.6 |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|---------------|------|-------|-----------|---------|
| N8—H8···O1B | 0.86 | 2.11 | 2.967 (2) | 173 |
| N8B—H8B···O1 | 0.86 | 2.03 | 2.868 (2) | 166 |
| N8A—H8A···O1C | 0.86 | 2.03 | 2.872 (2) | 165 |
| N8C—H8C···O1A | 0.86 | 2.10 | 2.959 (2) | 172 |