

Crystal structure of anagryrine perchlorate

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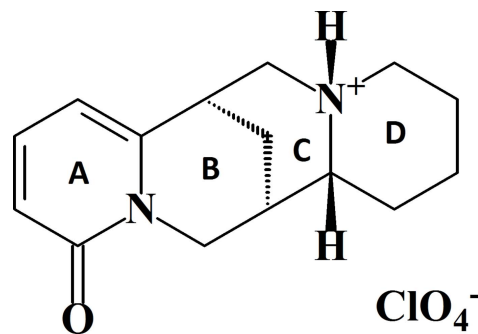
The title molecular salt, $C_{15}H_{21}N_2O^+ \cdot ClO_4^-$, crystallizes with four cations (*A*, *B*, *C* and *D*) and four anions in the chiral unit cell (space group $P2_1$). The alkaloid was isolated from the aerial parts of *Genista Hispanica* collected in the Samarkand region of Uzbekistan. Each cation is protonated at the N atom that bridges the alkaloid rings *C* and *D*. In each cation, ring *A* is almost planar and ring *B* adopts a sofa conformation with the methylene group bridging to the *C* ring as the flap. Rings *C* and *D* adopt chair conformations with a *cis* ring junction in all four cations. In the crystal, *A*+*B* and *C*+*D* dimeric pairs linked by pairs of $N-H \cdots O$ hydrogen bonds are observed, which generate $R_2^2(16)$ loops in each case. The dimers are consolidated by weak aromatic $\pi-\pi$ stacking interactions between the *A* rings [centroid-centroid distances = 3.913 (3) and 3.915 (3) Å].

Keywords: crystal structure; alkaloid; *Genista Hispanica*; anagryrine; perchlorate; $N-H \cdots O$ hydrogen bonds; $\pi-\pi$ stacking interactions.

CCDC reference: 1060546

1. Related literature

For the isolation of the title alkaloid, see: Orechoff *et al.* (1934); Sagen *et al.* (2002). For NMR spectra of the title alkaloid, see: Sagen *et al.* (2002). For theoretical studies of anagryrine and the crystal structure of anagryrine hydrochloride monohydrate, see: Galasso *et al.* (2006). For a related crystal structure, see: Atta-ur-Rahman *et al.* (1991).



2. Experimental

2.1. Crystal data

| | |
|------------------------------------|---|
| $C_{15}H_{21}N_2O^+ \cdot ClO_4^-$ | $V = 3125.41 (19) \text{ \AA}^3$ |
| $M_r = 344.79$ | $Z = 8$ |
| Monoclinic, $P2_1$ | Cu $K\alpha$ radiation |
| $a = 7.3550 (3) \text{ \AA}$ | $\mu = 2.42 \text{ mm}^{-1}$ |
| $b = 32.982 (1) \text{ \AA}$ | $T = 290 \text{ K}$ |
| $c = 12.8849 (4) \text{ \AA}$ | $0.65 \times 0.15 \times 0.04 \text{ mm}$ |
| $\beta = 90.709 (3)^\circ$ | |

2.2. Data collection

| | |
|---|--|
| Oxford Diffraction Xcalibur Ruby diffractometer | 53652 measured reflections |
| Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Oxford Diffraction, 2009) | 12780 independent reflections |
| $T_{\min} = 0.651$, $T_{\max} = 1.000$ | 9677 reflections with $I > 2\sigma(I)$ |
| | $R_{\text{int}} = 0.105$ |

2.3. Refinement

| | |
|--|--|
| $R[F^2 > 2\sigma(F^2)] = 0.064$ | $\Delta\rho_{\max} = 0.60 \text{ e \AA}^{-3}$ |
| $wR(F^2) = 0.189$ | $\Delta\rho_{\min} = -0.37 \text{ e \AA}^{-3}$ |
| $S = 1.02$ | Absolute structure: Flack x |
| 12780 reflections | determined using 3107 quotients |
| 846 parameters | $[(I^+) - (I^-)] / [(I^+) + (I^-)]$ (Parsons <i>et al.</i> , 2013) |
| 1 restraint | Absolute structure parameter: |
| H atoms treated by a mixture of independent and constrained refinement | $-0.024 (12)$ |

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

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|----------------------------|----------|--------------|--------------|----------------|
| $N2A-H2AN \cdots O1B$ | 1.03 (5) | 1.91 (6) | 2.741 (6) | 136 (5) |
| $N2B-H2CN \cdots O1A$ | 0.77 (7) | 2.00 (6) | 2.742 (5) | 163 (6) |
| $N2C-H2EN \cdots O1D^i$ | 0.90 (9) | 2.00 (9) | 2.735 (6) | 138 (8) |
| $N2D-H2GN \cdots O1C^{ii}$ | 1.05 (5) | 1.74 (5) | 2.754 (5) | 159 (5) |

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

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2009); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7409).

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supporting information

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Crystal structure of anagryne perchlorate

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S1. Comment

Quinolizidine alkaloids attracted the attention of researchers due to the structural characteristics and pharmacological activity. We have studied the aerial parts of *Genista Hispanica* collected in Samarkand region and isolated anagryne with R_f 0.5 (chloroform-methanol 6:1) along with other alkaloids. Anagryne a toxic alkaloid found in several species of *Lupinus* in the western United States. Acute poisoning produces nervousness, depression, loss of muscular control, convulsions, and coma.

Anagryne perchlorate crystallizes in the monoclinic space group $P2_1$ with a long unique b-axis of 32.982 (1) Å. The asymmetric unit consist of four protonated anagryne molecules and four perchlorate anions. The molecular structure of the alkaloid is shown in Fig. 1. The alkaloid molecule consists of four fused rings - planar ring A fused with the sofa ring B, and a twin-chair C/D fragment where C/D junction is *cis*. Conformation of all independent molecules matches. Anagryne molecule has three asymmetric centers at C6, C8, C10 and in addition by protonation of N2 it becomes as asymmetric senter. Configuration of chiral atoms are C-6R,C-8R,C-10R and N-2S. Crystal structure of thermopsine - a C-10-epimer of anagryne was investigated by Atta-ur-Rahman *et al.* (1991).

In the crystal, pairs of hydrogen bonds between protonated N atom of the base and the carbonyl O atom link molecules to form two molecular associates (Fig.2, Table 1.). In addition the associates are linked by weak π - π stacking interactions observed between aromatic rings of molecule [centroid—centroid distance = 3.913 (3) Å and 3.915 (3) Å for independent molecular pairs]

S2. Experimental

S2.1. Synthesis and crystallization

The powdered air-dried plant material were extracted with 80% ethanol. After distilling off the alcohol, the residue was acidified with H_2SO_4 and washed with chloroform, then the extract was basified with 25% aqueous ammonia, the sum of alkaloids (8.81 g) were extracted with chloroform. The resulting sum were dissolved in ethanol and acidified with HNO_3 to a weakly acidic medium, precipitated cytisine nitrate crystals (0.98g) were separated, and the mother liquor was evaporated. The resulting aqueous residue was basified with 25% aqueous ammonia and alkaloids was extracted with chloroform. The resulting sum of alkaloids was subjected to column chromatography on silica gel eluting with chloroform-methanol (100: 1) and isolated anagryne (0.16g). Obtained anagryne was dissolved in acetone and perchloric acid was added until acidic medium of pH 5-6. Precipitated anagryne perchlorate crystals were crystallized from methanol with m.p. 315 °C. Colourless prisms were obtained by re-crystallization from water at 50 °C .

S2.2. Refinement

Carbon-bound H atoms were placed geometrically and treated as riding on their parent atoms, with C—H distances of 0.93 Å (aromatic), 0.97 Å (methylene), 0.98 Å (tertiary carbon) and were refined with $U_{iso}(H)=1.2U_{eq}(C)$ for all hydrogen atoms. N-bound H atoms involved in the intermolecular hydrogen bonding were found by difference Fourier synthesis and refined isotropically [$N2A-H = 1.03(5)$ Å, $N2B-H = 0.77(7)$ Å, $N2C-H = 0.90(9)$ Å, $N2D-H = 1.05(5)$ Å].

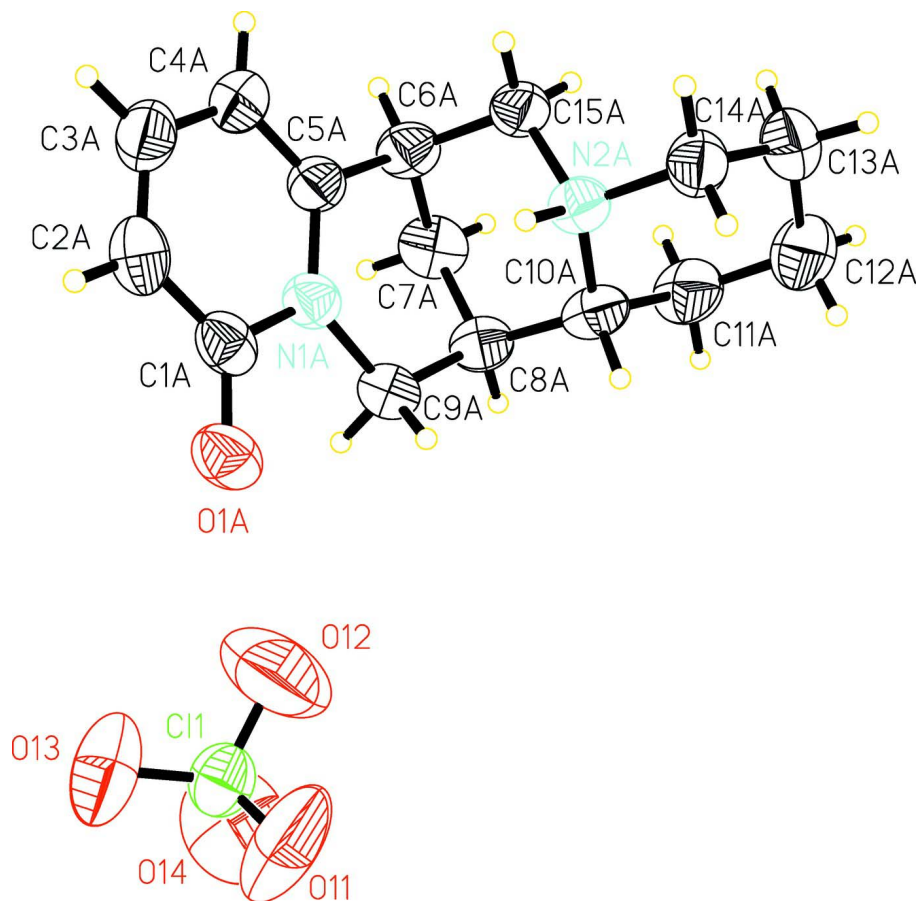


Figure 1

The molecular structure of cation A of the title compound, with displacement ellipsoids drawn at the 50% probability level.

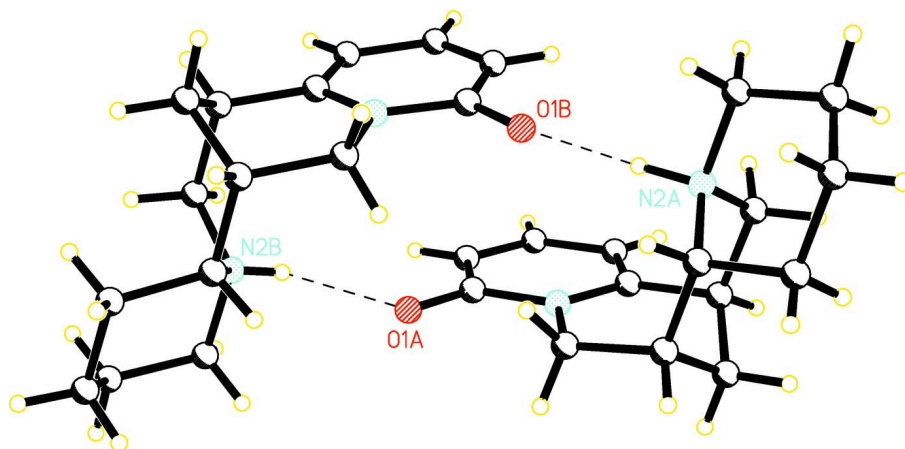


Figure 2

Hydrogen bonding between molecules.

Anagryne perchlorate

Crystal data

$C_{15}H_{21}N_2O^+ \cdot ClO_4^-$

$M_r = 344.79$

Monoclinic, $P2_1$

Hall symbol: $P\ 2y_b$

$a = 7.3550\ (3)\ \text{\AA}$

$b = 32.982\ (1)\ \text{\AA}$

$c = 12.8849\ (4)\ \text{\AA}$

$\beta = 90.709\ (3)^\circ$

$V = 3125.41\ (19)\ \text{\AA}^3$

$Z = 8$

$F(000) = 1456$

$D_x = 1.465\ \text{Mg m}^{-3}$

Melting point: $588(2)\ \text{K}$

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

Cell parameters from 6414 reflections

$\theta = 3.7\text{--}75.0^\circ$

$\mu = 2.42\ \text{mm}^{-1}$

$T = 290\ \text{K}$

Prism, colourless

$0.65 \times 0.15 \times 0.04\ \text{mm}$

Data collection

Oxford Diffraction Xcalibur Ruby
diffractometer

Radiation source: Enhance (Cu) X-ray Source

Graphite monochromator

Detector resolution: $10.2576\ \text{pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*CrysAlis PRO*; Oxford Diffraction, 2009)

$T_{\min} = 0.651$, $T_{\max} = 1.000$

53652 measured reflections

12780 independent reflections

9677 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.105$

$\theta_{\max} = 77.5^\circ$, $\theta_{\min} = 3.4^\circ$

$h = -9 \rightarrow 8$

$k = -41 \rightarrow 41$

$l = -16 \rightarrow 16$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.064$

$wR(F^2) = 0.189$

$S = 1.02$

12780 reflections

846 parameters

1 restraint

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H atoms treated by a mixture of independent
and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0663P)^2 + 0.7864P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.005$

$\Delta\rho_{\max} = 0.60\ \text{e \AA}^{-3}$

$\Delta\rho_{\min} = -0.37\ \text{e \AA}^{-3}$

Extinction correction: *SHELXL*,
 $F_c^* = kFc[1 + 0.001 \times Fc^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.00064 (10)

Absolute structure: Flack x determined using
 3107 quotients $[(F^+) - (F^-)] / [(F^+) + (F^-)]$ (Parsons *et al.*, 2013)
 Absolute structure parameter: -0.024 (12)

Special details

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

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

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|-------------|--------------|------------|----------------------------------|
| O1A | 0.4133 (6) | 0.31408 (11) | 0.7883 (3) | 0.0586 (9) |
| N1A | 0.5187 (5) | 0.35415 (11) | 0.9204 (3) | 0.0447 (8) |
| N2A | 0.2907 (6) | 0.38902 (11) | 1.1098 (3) | 0.0460 (8) |
| C1A | 0.4817 (7) | 0.34744 (15) | 0.8144 (4) | 0.0485 (10) |
| C2A | 0.5292 (7) | 0.37879 (17) | 0.7450 (4) | 0.0576 (12) |
| H2A | 0.5092 | 0.3752 | 0.6741 | 0.069* |
| C3A | 0.6032 (8) | 0.41377 (17) | 0.7797 (4) | 0.0593 (12) |
| H3A | 0.6344 | 0.4340 | 0.7328 | 0.071* |
| C4A | 0.6333 (8) | 0.41978 (15) | 0.8873 (4) | 0.0569 (12) |
| H4A | 0.6803 | 0.4443 | 0.9111 | 0.068* |
| C5A | 0.5937 (6) | 0.38989 (14) | 0.9558 (4) | 0.0468 (9) |
| C6A | 0.6195 (7) | 0.39559 (14) | 1.0709 (4) | 0.0509 (10) |
| H6A | 0.7237 | 0.4136 | 1.0823 | 0.061* |
| C7A | 0.6605 (8) | 0.35544 (16) | 1.1242 (4) | 0.0585 (12) |
| H7A | 0.6815 | 0.3595 | 1.1979 | 0.070* |
| H7B | 0.7678 | 0.3430 | 1.0948 | 0.070* |
| C8A | 0.4945 (7) | 0.32869 (14) | 1.1059 (4) | 0.0495 (10) |
| H8A | 0.5179 | 0.3029 | 1.1412 | 0.059* |
| C9A | 0.4759 (7) | 0.31962 (13) | 0.9901 (4) | 0.0500 (10) |
| H9A | 0.5558 | 0.2972 | 0.9735 | 0.060* |
| H9B | 0.3522 | 0.3109 | 0.9757 | 0.060* |
| C10A | 0.3225 (8) | 0.34697 (14) | 1.1534 (4) | 0.0525 (11) |
| H10A | 0.2194 | 0.3300 | 1.1316 | 0.063* |
| C11A | 0.3270 (10) | 0.34759 (18) | 1.2714 (4) | 0.0660 (14) |
| H11A | 0.4344 | 0.3621 | 1.2953 | 0.079* |
| H11B | 0.3345 | 0.3200 | 1.2973 | 0.079* |
| C12A | 0.1551 (12) | 0.3683 (2) | 1.3153 (5) | 0.0811 (19) |
| H12A | 0.0493 | 0.3519 | 1.2989 | 0.097* |
| H12B | 0.1663 | 0.3701 | 1.3903 | 0.097* |
| C13A | 0.1284 (10) | 0.41028 (19) | 1.2713 (4) | 0.0677 (14) |
| H13A | 0.0151 | 0.4215 | 1.2962 | 0.081* |

| | | | | |
|------|-------------|--------------|------------|-------------|
| H13B | 0.2268 | 0.4277 | 1.2946 | 0.081* |
| C14A | 0.1239 (7) | 0.40890 (16) | 1.1549 (4) | 0.0548 (11) |
| H14A | 0.1141 | 0.4363 | 1.1283 | 0.066* |
| H14B | 0.0167 | 0.3941 | 1.1321 | 0.066* |
| C15A | 0.4529 (7) | 0.41532 (14) | 1.1185 (3) | 0.0479 (10) |
| H15A | 0.4777 | 0.4211 | 1.1911 | 0.057* |
| H15B | 0.4285 | 0.4409 | 1.0836 | 0.057* |
| O1B | 0.0848 (6) | 0.36310 (12) | 0.9439 (3) | 0.0591 (9) |
| N1B | 0.0180 (6) | 0.34775 (12) | 0.7754 (3) | 0.0464 (8) |
| N2B | 0.2143 (5) | 0.28075 (11) | 0.6285 (3) | 0.0428 (8) |
| C1B | 0.0737 (7) | 0.37526 (15) | 0.8519 (4) | 0.0498 (10) |
| C2B | 0.1114 (8) | 0.41506 (17) | 0.8180 (5) | 0.0633 (13) |
| H2C | 0.1467 | 0.4345 | 0.8664 | 0.076* |
| C3B | 0.0975 (9) | 0.42544 (18) | 0.7174 (5) | 0.0682 (14) |
| H3C | 0.1227 | 0.4519 | 0.6974 | 0.082* |
| C4B | 0.0450 (9) | 0.39665 (17) | 0.6415 (5) | 0.0648 (14) |
| H4C | 0.0387 | 0.4038 | 0.5717 | 0.078* |
| C5B | 0.0042 (8) | 0.35841 (16) | 0.6721 (4) | 0.0538 (11) |
| C6B | -0.0512 (8) | 0.32671 (17) | 0.5943 (4) | 0.0559 (12) |
| H6C | -0.1171 | 0.3400 | 0.5372 | 0.067* |
| C7B | -0.1758 (8) | 0.2952 (2) | 0.6424 (4) | 0.0640 (14) |
| H7C | -0.2846 | 0.3081 | 0.6686 | 0.077* |
| H7D | -0.2117 | 0.2752 | 0.5910 | 0.077* |
| C8B | -0.0701 (7) | 0.27531 (16) | 0.7304 (4) | 0.0546 (11) |
| H8C | -0.1510 | 0.2552 | 0.7614 | 0.066* |
| C9B | -0.0252 (7) | 0.30642 (15) | 0.8148 (4) | 0.0501 (10) |
| H9C | -0.1278 | 0.3084 | 0.8611 | 0.060* |
| H9D | 0.0780 | 0.2966 | 0.8552 | 0.060* |
| C10B | 0.0950 (7) | 0.25221 (14) | 0.6903 (4) | 0.0495 (10) |
| H10C | 0.1657 | 0.2427 | 0.7505 | 0.059* |
| C11B | 0.0438 (9) | 0.21524 (18) | 0.6244 (5) | 0.0679 (15) |
| H11C | -0.0204 | 0.1959 | 0.6673 | 0.082* |
| H11D | -0.0384 | 0.2237 | 0.5691 | 0.082* |
| C12B | 0.2083 (10) | 0.19436 (17) | 0.5769 (5) | 0.0711 (16) |
| H12C | 0.2831 | 0.1827 | 0.6318 | 0.085* |
| H12D | 0.1673 | 0.1725 | 0.5321 | 0.085* |
| C13B | 0.3193 (9) | 0.22340 (17) | 0.5158 (5) | 0.0653 (14) |
| H13C | 0.2485 | 0.2332 | 0.4570 | 0.078* |
| H13D | 0.4260 | 0.2097 | 0.4896 | 0.078* |
| C14B | 0.3771 (7) | 0.25872 (16) | 0.5831 (4) | 0.0561 (11) |
| H14C | 0.4473 | 0.2776 | 0.5420 | 0.067* |
| H14D | 0.4546 | 0.2489 | 0.6391 | 0.067* |
| C15B | 0.1168 (8) | 0.30531 (15) | 0.5495 (3) | 0.0543 (11) |
| H15C | 0.0787 | 0.2879 | 0.4927 | 0.065* |
| H15D | 0.1991 | 0.3255 | 0.5221 | 0.065* |
| O1C | 0.8812 (5) | 0.12630 (11) | 0.2909 (3) | 0.0560 (8) |
| N1C | 0.7799 (5) | 0.08624 (11) | 0.4217 (3) | 0.0433 (7) |
| N2C | 1.0207 (6) | 0.05052 (11) | 0.6095 (3) | 0.0456 (8) |

| | | | | |
|------|-------------|--------------|------------|-------------|
| C1C | 0.8095 (6) | 0.09375 (14) | 0.3167 (3) | 0.0453 (9) |
| C2C | 0.7495 (7) | 0.06292 (16) | 0.2460 (4) | 0.0514 (10) |
| H2E | 0.7659 | 0.0666 | 0.1752 | 0.062* |
| C3C | 0.6694 (8) | 0.02861 (16) | 0.2800 (4) | 0.0574 (12) |
| H3E | 0.6292 | 0.0093 | 0.2324 | 0.069* |
| C4C | 0.6467 (8) | 0.02195 (15) | 0.3858 (4) | 0.0550 (11) |
| H4E | 0.5940 | -0.0020 | 0.4085 | 0.066* |
| C5C | 0.7012 (6) | 0.05031 (13) | 0.4561 (4) | 0.0449 (9) |
| C6C | 0.6872 (7) | 0.04424 (14) | 0.5715 (4) | 0.0498 (10) |
| H6E | 0.5835 | 0.0263 | 0.5838 | 0.060* |
| C7C | 0.6506 (8) | 0.08412 (16) | 0.6271 (4) | 0.0561 (11) |
| H7E | 0.5412 | 0.0968 | 0.5996 | 0.067* |
| H7F | 0.6354 | 0.0795 | 0.7009 | 0.067* |
| C8C | 0.8144 (8) | 0.11066 (13) | 0.6082 (4) | 0.0508 (11) |
| H8E | 0.7927 | 0.1364 | 0.6441 | 0.061* |
| C9C | 0.8253 (8) | 0.12038 (13) | 0.4926 (4) | 0.0500 (10) |
| H9E | 0.7433 | 0.1427 | 0.4774 | 0.060* |
| H9F | 0.9476 | 0.1295 | 0.4778 | 0.060* |
| C10C | 0.9917 (7) | 0.09288 (14) | 0.6542 (4) | 0.0509 (10) |
| H10E | 1.0924 | 0.1100 | 0.6312 | 0.061* |
| C11C | 0.9978 (10) | 0.09176 (17) | 0.7727 (4) | 0.0647 (14) |
| H11E | 0.9915 | 0.1193 | 0.7989 | 0.078* |
| H11F | 0.8920 | 0.0773 | 0.7973 | 0.078* |
| C12C | 1.1679 (11) | 0.0716 (2) | 0.8160 (5) | 0.0760 (17) |
| H12E | 1.2733 | 0.0879 | 0.7989 | 0.091* |
| H12F | 1.1605 | 0.0701 | 0.8910 | 0.091* |
| C13C | 1.1914 (10) | 0.0292 (2) | 0.7725 (4) | 0.0715 (15) |
| H13E | 1.0937 | 0.0119 | 0.7962 | 0.086* |
| H13F | 1.3057 | 0.0178 | 0.7970 | 0.086* |
| C14C | 1.1893 (8) | 0.03078 (18) | 0.6545 (4) | 0.0598 (12) |
| H14E | 1.2953 | 0.0457 | 0.6316 | 0.072* |
| H14F | 1.1983 | 0.0034 | 0.6277 | 0.072* |
| C15C | 0.8559 (7) | 0.02379 (14) | 0.6161 (3) | 0.0489 (10) |
| H15E | 0.8351 | 0.0168 | 0.6881 | 0.059* |
| H15F | 0.8781 | -0.0012 | 0.5785 | 0.059* |
| O1D | 0.2197 (6) | 0.07898 (11) | 0.4471 (3) | 0.0569 (8) |
| N1D | 0.2731 (5) | 0.09373 (11) | 0.2775 (3) | 0.0424 (7) |
| N2D | 0.0690 (5) | 0.16005 (10) | 0.1287 (3) | 0.0403 (7) |
| C1D | 0.2222 (7) | 0.06657 (14) | 0.3553 (3) | 0.0464 (10) |
| C2D | 0.1765 (7) | 0.02692 (15) | 0.3226 (4) | 0.0546 (11) |
| H2G | 0.1471 | 0.0074 | 0.3718 | 0.065* |
| C3D | 0.1750 (8) | 0.01683 (15) | 0.2199 (4) | 0.0569 (12) |
| H3G | 0.1380 | -0.0090 | 0.1996 | 0.068* |
| C4D | 0.2279 (8) | 0.04458 (15) | 0.1455 (4) | 0.0540 (11) |
| H4G | 0.2302 | 0.0371 | 0.0759 | 0.065* |
| C5D | 0.2762 (7) | 0.08259 (14) | 0.1743 (3) | 0.0461 (9) |
| C6D | 0.3330 (7) | 0.11426 (15) | 0.0955 (3) | 0.0492 (10) |
| H6G | 0.3962 | 0.1004 | 0.0391 | 0.059* |

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|------|-------------|--------------|--------------|-------------|
| C7D | 0.4602 (7) | 0.14484 (17) | 0.1420 (4) | 0.0554 (12) |
| H7G | 0.4957 | 0.1644 | 0.0900 | 0.066* |
| H7H | 0.5688 | 0.1315 | 0.1685 | 0.066* |
| C8D | 0.3609 (7) | 0.16602 (14) | 0.2301 (4) | 0.0508 (10) |
| H8G | 0.4437 | 0.1863 | 0.2599 | 0.061* |
| C9D | 0.3199 (7) | 0.13482 (14) | 0.3155 (3) | 0.0480 (10) |
| H9G | 0.4254 | 0.1328 | 0.3611 | 0.058* |
| H9H | 0.2198 | 0.1449 | 0.3565 | 0.058* |
| C10D | 0.1933 (7) | 0.18822 (13) | 0.1895 (3) | 0.0474 (10) |
| H10G | 0.1259 | 0.1980 | 0.2496 | 0.057* |
| C11D | 0.2425 (9) | 0.22534 (15) | 0.1232 (5) | 0.0623 (14) |
| H11G | 0.3119 | 0.2443 | 0.1656 | 0.075* |
| H11H | 0.3192 | 0.2167 | 0.0667 | 0.075* |
| C12D | 0.0761 (9) | 0.24698 (16) | 0.0785 (5) | 0.0662 (14) |
| H12G | 0.0047 | 0.2582 | 0.1344 | 0.079* |
| H12H | 0.1144 | 0.2692 | 0.0344 | 0.079* |
| C13D | -0.0384 (9) | 0.21742 (18) | 0.0160 (5) | 0.0664 (14) |
| H13G | 0.0306 | 0.2077 | -0.0427 | 0.080* |
| H13H | -0.1460 | 0.2311 | -0.0106 | 0.080* |
| C14D | -0.0952 (7) | 0.18134 (16) | 0.0832 (4) | 0.0540 (11) |
| H14G | -0.1719 | 0.1908 | 0.1389 | 0.065* |
| H14H | -0.1652 | 0.1624 | 0.0414 | 0.065* |
| C15D | 0.1650 (7) | 0.13525 (14) | 0.0495 (3) | 0.0498 (10) |
| H15G | 0.2014 | 0.1526 | -0.0073 | 0.060* |
| H15H | 0.0822 | 0.1149 | 0.0218 | 0.060* |
| Cl1 | 0.5283 (2) | 0.18310 (5) | 0.83543 (11) | 0.0683 (3) |
| O11 | 0.3429 (11) | 0.1756 (3) | 0.8283 (6) | 0.150 (4) |
| O12 | 0.5416 (16) | 0.2170 (3) | 0.8995 (9) | 0.181 (5) |
| O13 | 0.6026 (10) | 0.1909 (2) | 0.7395 (5) | 0.120 (2) |
| O14 | 0.6176 (15) | 0.1521 (3) | 0.8875 (7) | 0.161 (4) |
| Cl2 | 0.6107 (3) | 0.42676 (4) | 0.45172 (10) | 0.0705 (4) |
| O21 | 0.452 (2) | 0.4201 (8) | 0.4894 (10) | 0.355 (15) |
| O22 | 0.590 (2) | 0.4423 (4) | 0.3594 (7) | 0.244 (8) |
| O23 | 0.670 (2) | 0.3903 (3) | 0.4433 (15) | 0.298 (11) |
| O24 | 0.7186 (14) | 0.4502 (2) | 0.5180 (6) | 0.153 (4) |
| Cl3 | 0.7597 (2) | 0.25649 (4) | 0.33650 (10) | 0.0654 (3) |
| O31 | 0.6728 (15) | 0.2873 (3) | 0.3883 (7) | 0.167 (4) |
| O32 | 0.7512 (19) | 0.2224 (3) | 0.3989 (9) | 0.195 (5) |
| O33 | 0.6803 (8) | 0.24878 (19) | 0.2401 (4) | 0.1007 (18) |
| O34 | 0.9439 (11) | 0.2649 (4) | 0.3267 (6) | 0.182 (5) |
| Cl4 | 0.7284 (3) | 0.02077 (4) | 0.96609 (10) | 0.0700 (4) |
| O41 | 0.8759 (18) | -0.0026 (3) | 1.0007 (9) | 0.196 (6) |
| O42 | 0.5883 (17) | 0.0062 (4) | 1.0232 (7) | 0.198 (6) |
| O43 | 0.7115 (10) | 0.01230 (17) | 0.8606 (3) | 0.0973 (18) |
| O44 | 0.7781 (14) | 0.06101 (17) | 0.9878 (4) | 0.135 (3) |
| H2AN | 0.259 (8) | 0.3892 (17) | 1.032 (4) | 0.051 (14)* |
| H2CN | 0.264 (9) | 0.2945 (18) | 0.668 (5) | 0.052 (15)* |
| H2EN | 1.034 (13) | 0.055 (3) | 0.541 (7) | 0.11 (3)* |

H2GN 0.015 (7) 0.1414 (16) 0.187 (4) 0.043 (13)*

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-----------|-------------|-------------|--------------|--------------|--------------|
| O1A | 0.060 (2) | 0.0617 (19) | 0.0538 (19) | -0.0031 (16) | -0.0063 (16) | -0.0159 (15) |
| N1A | 0.040 (2) | 0.0453 (17) | 0.0491 (19) | -0.0026 (15) | 0.0007 (15) | -0.0076 (14) |
| N2A | 0.050 (2) | 0.0452 (17) | 0.0423 (18) | -0.0046 (16) | -0.0050 (16) | 0.0012 (14) |
| C1A | 0.038 (2) | 0.055 (2) | 0.053 (2) | 0.0049 (19) | -0.0001 (18) | -0.0126 (19) |
| C2A | 0.048 (3) | 0.074 (3) | 0.051 (3) | 0.010 (2) | 0.007 (2) | -0.002 (2) |
| C3A | 0.058 (3) | 0.062 (3) | 0.059 (3) | 0.001 (2) | 0.015 (2) | 0.006 (2) |
| C4A | 0.058 (3) | 0.055 (3) | 0.058 (3) | -0.010 (2) | 0.013 (2) | -0.002 (2) |
| C5A | 0.040 (2) | 0.045 (2) | 0.055 (2) | -0.0056 (18) | 0.0040 (18) | -0.0063 (18) |
| C6A | 0.045 (3) | 0.052 (2) | 0.056 (3) | -0.0108 (19) | -0.0066 (19) | -0.0060 (19) |
| C7A | 0.053 (3) | 0.062 (3) | 0.061 (3) | 0.003 (2) | -0.017 (2) | -0.001 (2) |
| C8A | 0.051 (3) | 0.045 (2) | 0.053 (2) | -0.0014 (18) | -0.008 (2) | 0.0045 (17) |
| C9A | 0.054 (3) | 0.044 (2) | 0.052 (2) | 0.0011 (19) | -0.002 (2) | -0.0052 (17) |
| C10A | 0.061 (3) | 0.046 (2) | 0.051 (2) | -0.014 (2) | -0.008 (2) | 0.0050 (18) |
| C11A | 0.090 (4) | 0.062 (3) | 0.045 (3) | -0.007 (3) | -0.003 (3) | 0.014 (2) |
| C12A | 0.107 (6) | 0.081 (4) | 0.055 (3) | -0.011 (4) | 0.023 (3) | 0.011 (3) |
| C13A | 0.076 (4) | 0.071 (3) | 0.057 (3) | 0.003 (3) | 0.016 (3) | -0.007 (2) |
| C14A | 0.054 (3) | 0.059 (3) | 0.051 (2) | -0.003 (2) | 0.006 (2) | -0.001 (2) |
| C15A | 0.051 (3) | 0.047 (2) | 0.045 (2) | -0.0101 (19) | -0.0018 (18) | -0.0063 (16) |
| O1B | 0.061 (2) | 0.070 (2) | 0.0461 (18) | 0.0027 (17) | -0.0119 (15) | -0.0126 (15) |
| N1B | 0.044 (2) | 0.0549 (19) | 0.0403 (18) | 0.0049 (16) | -0.0033 (15) | -0.0072 (15) |
| N2B | 0.045 (2) | 0.0442 (17) | 0.0396 (17) | -0.0071 (15) | -0.0041 (15) | -0.0049 (14) |
| C1B | 0.039 (2) | 0.060 (2) | 0.050 (2) | 0.0102 (19) | -0.0046 (18) | -0.0109 (19) |
| C2B | 0.053 (3) | 0.055 (3) | 0.081 (4) | 0.003 (2) | -0.001 (3) | -0.018 (2) |
| C3B | 0.060 (4) | 0.056 (3) | 0.088 (4) | 0.005 (2) | 0.011 (3) | 0.009 (3) |
| C4B | 0.072 (4) | 0.064 (3) | 0.058 (3) | 0.020 (3) | 0.001 (3) | 0.007 (2) |
| C5B | 0.056 (3) | 0.063 (3) | 0.042 (2) | 0.011 (2) | -0.002 (2) | 0.0016 (19) |
| C6B | 0.056 (3) | 0.071 (3) | 0.041 (2) | 0.013 (2) | -0.013 (2) | -0.005 (2) |
| C7B | 0.039 (3) | 0.090 (4) | 0.063 (3) | -0.001 (2) | -0.001 (2) | -0.027 (3) |
| C8B | 0.052 (3) | 0.062 (3) | 0.051 (2) | -0.018 (2) | 0.007 (2) | -0.010 (2) |
| C9B | 0.050 (3) | 0.060 (3) | 0.040 (2) | -0.006 (2) | 0.0037 (18) | -0.0037 (18) |
| C10B | 0.053 (3) | 0.047 (2) | 0.049 (2) | -0.0091 (19) | 0.0043 (19) | -0.0035 (17) |
| C11B | 0.067 (4) | 0.062 (3) | 0.076 (4) | -0.025 (3) | 0.008 (3) | -0.015 (3) |
| C12B | 0.083 (4) | 0.055 (3) | 0.075 (4) | -0.005 (3) | -0.004 (3) | -0.026 (3) |
| C13B | 0.061 (3) | 0.062 (3) | 0.073 (3) | -0.001 (2) | 0.003 (3) | -0.025 (3) |
| C14B | 0.050 (3) | 0.057 (3) | 0.062 (3) | 0.006 (2) | -0.003 (2) | -0.013 (2) |
| C15B | 0.067 (3) | 0.061 (3) | 0.035 (2) | 0.001 (2) | 0.002 (2) | 0.0004 (18) |
| O1C | 0.057 (2) | 0.0607 (19) | 0.0500 (17) | -0.0013 (16) | 0.0074 (15) | 0.0148 (14) |
| N1C | 0.043 (2) | 0.0449 (17) | 0.0420 (18) | -0.0007 (15) | 0.0033 (14) | 0.0020 (14) |
| N2C | 0.053 (2) | 0.0494 (19) | 0.0346 (17) | -0.0036 (16) | 0.0057 (15) | 0.0040 (14) |
| C1C | 0.041 (2) | 0.054 (2) | 0.041 (2) | 0.0054 (19) | -0.0002 (17) | 0.0071 (17) |
| C2C | 0.046 (3) | 0.067 (3) | 0.041 (2) | 0.007 (2) | -0.0032 (18) | -0.0056 (19) |
| C3C | 0.056 (3) | 0.059 (3) | 0.056 (3) | 0.009 (2) | -0.010 (2) | -0.014 (2) |
| C4C | 0.057 (3) | 0.048 (2) | 0.061 (3) | -0.007 (2) | 0.000 (2) | -0.007 (2) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C5C | 0.036 (2) | 0.050 (2) | 0.049 (2) | -0.0040 (17) | 0.0008 (17) | 0.0022 (17) |
| C6C | 0.051 (3) | 0.048 (2) | 0.050 (2) | -0.0107 (19) | 0.013 (2) | 0.0030 (18) |
| C7C | 0.054 (3) | 0.063 (3) | 0.052 (3) | -0.004 (2) | 0.014 (2) | 0.001 (2) |
| C8C | 0.061 (3) | 0.042 (2) | 0.050 (2) | -0.0019 (19) | 0.011 (2) | -0.0077 (17) |
| C9C | 0.059 (3) | 0.045 (2) | 0.047 (2) | -0.0036 (19) | -0.001 (2) | 0.0022 (17) |
| C10C | 0.060 (3) | 0.050 (2) | 0.043 (2) | -0.011 (2) | 0.0060 (19) | 0.0022 (17) |
| C11C | 0.092 (4) | 0.061 (3) | 0.042 (2) | -0.005 (3) | 0.003 (2) | -0.007 (2) |
| C12C | 0.089 (5) | 0.088 (4) | 0.050 (3) | -0.013 (4) | -0.013 (3) | -0.002 (3) |
| C13C | 0.077 (4) | 0.087 (4) | 0.050 (3) | 0.004 (3) | -0.004 (3) | 0.011 (3) |
| C14C | 0.057 (3) | 0.068 (3) | 0.055 (3) | 0.009 (2) | 0.004 (2) | 0.011 (2) |
| C15C | 0.057 (3) | 0.046 (2) | 0.044 (2) | -0.004 (2) | 0.0043 (19) | 0.0037 (17) |
| O1D | 0.066 (2) | 0.0650 (19) | 0.0397 (16) | 0.0003 (17) | 0.0116 (15) | 0.0094 (14) |
| N1D | 0.042 (2) | 0.0485 (17) | 0.0369 (16) | 0.0005 (15) | 0.0040 (14) | 0.0026 (13) |
| N2D | 0.0384 (19) | 0.0408 (16) | 0.0419 (17) | -0.0041 (14) | 0.0030 (14) | 0.0061 (13) |
| C1D | 0.046 (3) | 0.053 (2) | 0.041 (2) | 0.0062 (19) | 0.0088 (17) | 0.0103 (17) |
| C2D | 0.051 (3) | 0.051 (2) | 0.061 (3) | 0.004 (2) | 0.005 (2) | 0.011 (2) |
| C3D | 0.062 (3) | 0.044 (2) | 0.065 (3) | 0.006 (2) | 0.007 (2) | 0.000 (2) |
| C4D | 0.060 (3) | 0.056 (2) | 0.045 (2) | 0.010 (2) | 0.003 (2) | -0.0046 (19) |
| C5D | 0.045 (2) | 0.055 (2) | 0.038 (2) | 0.0077 (19) | 0.0029 (17) | 0.0017 (17) |
| C6D | 0.051 (3) | 0.060 (2) | 0.0372 (19) | 0.002 (2) | 0.0140 (18) | 0.0031 (17) |
| C7D | 0.031 (2) | 0.075 (3) | 0.059 (3) | -0.007 (2) | 0.0002 (19) | 0.028 (2) |
| C8D | 0.047 (3) | 0.053 (2) | 0.051 (2) | -0.0144 (19) | -0.0042 (19) | 0.0082 (19) |
| C9D | 0.049 (3) | 0.058 (2) | 0.037 (2) | -0.006 (2) | -0.0053 (17) | 0.0024 (17) |
| C10D | 0.057 (3) | 0.042 (2) | 0.043 (2) | -0.0100 (19) | 0.0025 (18) | 0.0010 (16) |
| C11D | 0.073 (4) | 0.049 (2) | 0.065 (3) | -0.018 (2) | -0.008 (3) | 0.014 (2) |
| C12D | 0.080 (4) | 0.047 (2) | 0.071 (3) | -0.008 (2) | 0.004 (3) | 0.016 (2) |
| C13D | 0.067 (4) | 0.068 (3) | 0.065 (3) | -0.002 (3) | -0.006 (3) | 0.025 (3) |
| C14D | 0.047 (3) | 0.056 (2) | 0.059 (3) | -0.001 (2) | 0.002 (2) | 0.018 (2) |
| C15D | 0.059 (3) | 0.056 (2) | 0.0351 (19) | 0.004 (2) | 0.0044 (18) | 0.0027 (17) |
| C11 | 0.0701 (9) | 0.0747 (8) | 0.0602 (7) | -0.0005 (7) | 0.0010 (6) | 0.0107 (6) |
| O11 | 0.099 (5) | 0.237 (10) | 0.114 (5) | -0.075 (6) | 0.002 (4) | 0.053 (6) |
| O12 | 0.197 (11) | 0.136 (7) | 0.211 (10) | 0.006 (7) | 0.029 (8) | -0.090 (7) |
| O13 | 0.106 (5) | 0.149 (6) | 0.105 (4) | 0.017 (4) | 0.039 (4) | 0.050 (4) |
| O14 | 0.202 (10) | 0.155 (7) | 0.124 (6) | 0.066 (7) | -0.028 (6) | 0.041 (5) |
| C12 | 0.1003 (12) | 0.0585 (6) | 0.0522 (6) | 0.0064 (7) | -0.0139 (7) | -0.0059 (5) |
| O21 | 0.232 (16) | 0.68 (4) | 0.151 (10) | -0.22 (2) | 0.081 (10) | -0.093 (17) |
| O22 | 0.350 (16) | 0.269 (13) | 0.110 (6) | -0.210 (13) | -0.120 (8) | 0.074 (7) |
| O23 | 0.331 (19) | 0.117 (7) | 0.44 (2) | 0.107 (10) | -0.243 (18) | -0.132 (10) |
| O24 | 0.243 (10) | 0.091 (4) | 0.121 (5) | -0.004 (5) | -0.107 (6) | -0.023 (4) |
| C13 | 0.0663 (8) | 0.0713 (7) | 0.0584 (7) | 0.0034 (6) | -0.0049 (6) | -0.0103 (5) |
| O31 | 0.206 (10) | 0.174 (8) | 0.122 (6) | 0.079 (7) | -0.004 (6) | -0.070 (6) |
| O32 | 0.274 (15) | 0.119 (6) | 0.192 (10) | 0.028 (8) | -0.049 (9) | 0.059 (6) |
| O33 | 0.092 (4) | 0.119 (4) | 0.090 (3) | 0.020 (3) | -0.037 (3) | -0.027 (3) |
| O34 | 0.088 (5) | 0.338 (15) | 0.119 (5) | -0.068 (7) | 0.002 (4) | -0.092 (7) |
| C14 | 0.1067 (12) | 0.0591 (6) | 0.0445 (6) | -0.0147 (7) | 0.0163 (6) | -0.0037 (5) |
| O41 | 0.260 (14) | 0.134 (7) | 0.192 (10) | 0.046 (8) | -0.121 (10) | -0.031 (7) |
| O42 | 0.242 (12) | 0.241 (11) | 0.113 (5) | -0.130 (10) | 0.107 (7) | -0.060 (6) |
| O43 | 0.148 (5) | 0.097 (3) | 0.047 (2) | -0.033 (3) | 0.012 (3) | -0.003 (2) |

| | | | | | | |
|-----|------------|-----------|-----------|------------|-----------|------------|
| O44 | 0.268 (11) | 0.069 (3) | 0.068 (3) | -0.033 (4) | 0.011 (4) | -0.012 (2) |
|-----|------------|-----------|-----------|------------|-----------|------------|

Geometric parameters (Å, °)

| | | | |
|-----------|------------|-----------|------------|
| O1A—C1A | 1.254 (6) | C1C—C2C | 1.432 (7) |
| N1A—C5A | 1.377 (6) | C2C—C3C | 1.351 (8) |
| N1A—C1A | 1.407 (6) | C2C—H2E | 0.9300 |
| N1A—C9A | 1.487 (6) | C3C—C4C | 1.393 (8) |
| N2A—C15A | 1.478 (6) | C3C—H3E | 0.9300 |
| N2A—C14A | 1.513 (7) | C4C—C5C | 1.359 (7) |
| N2A—C10A | 1.514 (6) | C4C—H4E | 0.9300 |
| N2A—H2AN | 1.03 (6) | C5C—C6C | 1.504 (6) |
| C1A—C2A | 1.414 (8) | C6C—C15C | 1.520 (7) |
| C2A—C3A | 1.349 (8) | C6C—C7C | 1.524 (7) |
| C2A—H2A | 0.9300 | C6C—H6E | 0.9800 |
| C3A—C4A | 1.414 (8) | C7C—C8C | 1.512 (7) |
| C3A—H3A | 0.9300 | C7C—H7E | 0.9700 |
| C4A—C5A | 1.357 (7) | C7C—H7F | 0.9700 |
| C4A—H4A | 0.9300 | C8C—C9C | 1.526 (6) |
| C5A—C6A | 1.505 (7) | C8C—C10C | 1.541 (8) |
| C6A—C7A | 1.520 (7) | C8C—H8E | 0.9800 |
| C6A—C15A | 1.523 (7) | C9C—H9E | 0.9700 |
| C6A—H6A | 0.9800 | C9C—H9F | 0.9700 |
| C7A—C8A | 1.522 (7) | C10C—C11C | 1.527 (6) |
| C7A—H7A | 0.9700 | C10C—H10E | 0.9800 |
| C7A—H7B | 0.9700 | C11C—C12C | 1.518 (10) |
| C8A—C9A | 1.526 (7) | C11C—H11E | 0.9700 |
| C8A—C10A | 1.536 (8) | C11C—H11F | 0.9700 |
| C8A—H8A | 0.9800 | C12C—C13C | 1.517 (10) |
| C9A—H9A | 0.9700 | C12C—H12E | 0.9700 |
| C9A—H9B | 0.9700 | C12C—H12F | 0.9700 |
| C10A—C11A | 1.521 (7) | C13C—C14C | 1.521 (7) |
| C10A—H10A | 0.9800 | C13C—H13E | 0.9700 |
| C11A—C12A | 1.550 (10) | C13C—H13F | 0.9700 |
| C11A—H11A | 0.9700 | C14C—H14E | 0.9700 |
| C11A—H11B | 0.9700 | C14C—H14F | 0.9700 |
| C12A—C13A | 1.507 (9) | C15C—H15E | 0.9700 |
| C12A—H12A | 0.9700 | C15C—H15F | 0.9700 |
| C12A—H12B | 0.9700 | O1D—C1D | 1.252 (6) |
| C13A—C14A | 1.500 (7) | N1D—C5D | 1.380 (6) |
| C13A—H13A | 0.9700 | N1D—C1D | 1.399 (5) |
| C13A—H13B | 0.9700 | N1D—C9D | 1.480 (6) |
| C14A—H14A | 0.9700 | N2D—C15D | 1.492 (6) |
| C14A—H14B | 0.9700 | N2D—C14D | 1.509 (6) |
| C15A—H15A | 0.9700 | N2D—C10D | 1.515 (6) |
| C15A—H15B | 0.9700 | N2D—H2GN | 1.05 (5) |
| O1B—C1B | 1.253 (6) | C1D—C2D | 1.413 (7) |
| N1B—C5B | 1.379 (6) | C2D—C3D | 1.365 (8) |

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| N1B—C1B | 1.397 (6) | C2D—H2G | 0.9300 |
| N1B—C9B | 1.490 (6) | C3D—C4D | 1.384 (7) |
| N2B—C15B | 1.479 (6) | C3D—H3G | 0.9300 |
| N2B—C10B | 1.519 (6) | C4D—C5D | 1.353 (7) |
| N2B—C14B | 1.524 (6) | C4D—H4G | 0.9300 |
| N2B—H2CN | 0.77 (6) | C5D—C6D | 1.519 (6) |
| C1B—C2B | 1.412 (8) | C6D—C7D | 1.496 (7) |
| C2B—C3B | 1.343 (9) | C6D—C15D | 1.529 (7) |
| C2B—H2C | 0.9300 | C6D—H6G | 0.9800 |
| C3B—C4B | 1.414 (9) | C7D—C8D | 1.526 (8) |
| C3B—H3C | 0.9300 | C7D—H7G | 0.9700 |
| C4B—C5B | 1.356 (8) | C7D—H7H | 0.9700 |
| C4B—H4C | 0.9300 | C8D—C10D | 1.521 (7) |
| C5B—C6B | 1.502 (7) | C8D—C9D | 1.539 (6) |
| C6B—C7B | 1.522 (9) | C8D—H8G | 0.9800 |
| C6B—C15B | 1.541 (8) | C9D—H9G | 0.9700 |
| C6B—H6C | 0.9800 | C9D—H9H | 0.9700 |
| C7B—C8B | 1.516 (8) | C10D—C11D | 1.538 (6) |
| C7B—H7C | 0.9700 | C10D—H10G | 0.9800 |
| C7B—H7D | 0.9700 | C11D—C12D | 1.524 (9) |
| C8B—C10B | 1.528 (8) | C11D—H11G | 0.9700 |
| C8B—C9B | 1.529 (6) | C11D—H11H | 0.9700 |
| C8B—H8C | 0.9800 | C12D—C13D | 1.514 (9) |
| C9B—H9C | 0.9700 | C12D—H12G | 0.9700 |
| C9B—H9D | 0.9700 | C12D—H12H | 0.9700 |
| C10B—C11B | 1.530 (6) | C13D—C14D | 1.533 (7) |
| C10B—H10C | 0.9800 | C13D—H13G | 0.9700 |
| C11B—C12B | 1.527 (9) | C13D—H13H | 0.9700 |
| C11B—H11C | 0.9700 | C14D—H14G | 0.9700 |
| C11B—H11D | 0.9700 | C14D—H14H | 0.9700 |
| C12B—C13B | 1.490 (9) | C15D—H15G | 0.9700 |
| C12B—H12C | 0.9700 | C15D—H15H | 0.9700 |
| C12B—H12D | 0.9700 | C11—O13 | 1.381 (6) |
| C13B—C14B | 1.510 (7) | C11—O14 | 1.385 (7) |
| C13B—H13C | 0.9700 | C11—O11 | 1.387 (7) |
| C13B—H13D | 0.9700 | C11—O12 | 1.393 (8) |
| C14B—H14C | 0.9700 | C12—O23 | 1.285 (8) |
| C14B—H14D | 0.9700 | C12—O21 | 1.290 (13) |
| C15B—H15C | 0.9700 | C12—O22 | 1.302 (8) |
| C15B—H15D | 0.9700 | C12—O24 | 1.392 (6) |
| O1C—C1C | 1.243 (6) | C13—O31 | 1.377 (7) |
| N1C—C5C | 1.394 (6) | C13—O32 | 1.385 (9) |
| N1C—C1C | 1.395 (6) | C13—O33 | 1.389 (5) |
| N1C—C9C | 1.486 (6) | C13—O34 | 1.390 (8) |
| N2C—C15C | 1.502 (6) | C14—O42 | 1.362 (8) |
| N2C—C14C | 1.510 (7) | C14—O43 | 1.391 (5) |
| N2C—C10C | 1.527 (6) | C14—O41 | 1.400 (10) |
| N2C—H2EN | 0.91 (9) | C14—O44 | 1.404 (5) |

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| C5A—N1A—C1A | 121.9 (4) | C3C—C2C—C1C | 121.4 (4) |
| C5A—N1A—C9A | 122.9 (4) | C3C—C2C—H2E | 119.3 |
| C1A—N1A—C9A | 115.2 (4) | C1C—C2C—H2E | 119.3 |
| C15A—N2A—C14A | 112.0 (4) | C2C—C3C—C4C | 120.5 (5) |
| C15A—N2A—C10A | 112.9 (4) | C2C—C3C—H3E | 119.8 |
| C14A—N2A—C10A | 112.1 (4) | C4C—C3C—H3E | 119.8 |
| C15A—N2A—H2AN | 104 (3) | C5C—C4C—C3C | 120.4 (5) |
| C14A—N2A—H2AN | 101 (3) | C5C—C4C—H4E | 119.8 |
| C10A—N2A—H2AN | 114 (3) | C3C—C4C—H4E | 119.8 |
| O1A—C1A—N1A | 118.1 (5) | C4C—C5C—N1C | 119.5 (4) |
| O1A—C1A—C2A | 125.0 (5) | C4C—C5C—C6C | 122.9 (4) |
| N1A—C1A—C2A | 116.9 (4) | N1C—C5C—C6C | 117.5 (4) |
| C3A—C2A—C1A | 121.1 (5) | C5C—C6C—C15C | 111.5 (4) |
| C3A—C2A—H2A | 119.4 | C5C—C6C—C7C | 111.4 (4) |
| C1A—C2A—H2A | 119.4 | C15C—C6C—C7C | 110.7 (4) |
| C2A—C3A—C4A | 120.2 (5) | C5C—C6C—H6E | 107.7 |
| C2A—C3A—H3A | 119.9 | C15C—C6C—H6E | 107.7 |
| C4A—C3A—H3A | 119.9 | C7C—C6C—H6E | 107.7 |
| C5A—C4A—C3A | 120.2 (5) | C8C—C7C—C6C | 106.1 (4) |
| C5A—C4A—H4A | 119.9 | C8C—C7C—H7E | 110.5 |
| C3A—C4A—H4A | 119.9 | C6C—C7C—H7E | 110.5 |
| C4A—C5A—N1A | 119.6 (4) | C8C—C7C—H7F | 110.5 |
| C4A—C5A—C6A | 121.7 (4) | C6C—C7C—H7F | 110.5 |
| N1A—C5A—C6A | 118.6 (4) | H7E—C7C—H7F | 108.7 |
| C5A—C6A—C7A | 111.0 (4) | C7C—C8C—C9C | 109.3 (4) |
| C5A—C6A—C15A | 111.0 (4) | C7C—C8C—C10C | 112.9 (4) |
| C7A—C6A—C15A | 110.3 (4) | C9C—C8C—C10C | 113.6 (4) |
| C5A—C6A—H6A | 108.2 | C7C—C8C—H8E | 106.8 |
| C7A—C6A—H6A | 108.2 | C9C—C8C—H8E | 106.8 |
| C15A—C6A—H6A | 108.2 | C10C—C8C—H8E | 106.8 |
| C6A—C7A—C8A | 106.3 (4) | N1C—C9C—C8C | 115.3 (4) |
| C6A—C7A—H7A | 110.5 | N1C—C9C—H9E | 108.5 |
| C8A—C7A—H7A | 110.5 | C8C—C9C—H9E | 108.5 |
| C6A—C7A—H7B | 110.5 | N1C—C9C—H9F | 108.5 |
| C8A—C7A—H7B | 110.5 | C8C—C9C—H9F | 108.5 |
| H7A—C7A—H7B | 108.7 | H9E—C9C—H9F | 107.5 |
| C7A—C8A—C9A | 109.1 (4) | N2C—C10C—C11C | 110.6 (4) |
| C7A—C8A—C10A | 111.9 (4) | N2C—C10C—C8C | 108.9 (4) |
| C9A—C8A—C10A | 113.7 (4) | C11C—C10C—C8C | 114.1 (5) |
| C7A—C8A—H8A | 107.3 | N2C—C10C—H10E | 107.7 |
| C9A—C8A—H8A | 107.3 | C11C—C10C—H10E | 107.7 |
| C10A—C8A—H8A | 107.3 | C8C—C10C—H10E | 107.7 |
| N1A—C9A—C8A | 115.0 (4) | C12C—C11C—C10C | 113.1 (5) |
| N1A—C9A—H9A | 108.5 | C12C—C11C—H11E | 109.0 |
| C8A—C9A—H9A | 108.5 | C10C—C11C—H11E | 109.0 |
| N1A—C9A—H9B | 108.5 | C12C—C11C—H11F | 109.0 |
| C8A—C9A—H9B | 108.5 | C10C—C11C—H11F | 109.0 |

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| H9A—C9A—H9B | 107.5 | H11E—C11C—H11F | 107.8 |
| N2A—C10A—C11A | 111.1 (4) | C13C—C12C—C11C | 111.4 (5) |
| N2A—C10A—C8A | 109.6 (4) | C13C—C12C—H12E | 109.3 |
| C11A—C10A—C8A | 113.4 (5) | C11C—C12C—H12E | 109.3 |
| N2A—C10A—H10A | 107.5 | C13C—C12C—H12F | 109.3 |
| C11A—C10A—H10A | 107.5 | C11C—C12C—H12F | 109.3 |
| C8A—C10A—H10A | 107.5 | H12E—C12C—H12F | 108.0 |
| C10A—C11A—C12A | 111.3 (5) | C12C—C13C—C14C | 109.7 (5) |
| C10A—C11A—H11A | 109.4 | C12C—C13C—H13E | 109.7 |
| C12A—C11A—H11A | 109.4 | C14C—C13C—H13E | 109.7 |
| C10A—C11A—H11B | 109.4 | C12C—C13C—H13F | 109.7 |
| C12A—C11A—H11B | 109.4 | C14C—C13C—H13F | 109.7 |
| H11A—C11A—H11B | 108.0 | H13E—C13C—H13F | 108.2 |
| C13A—C12A—C11A | 111.8 (5) | N2C—C14C—C13C | 113.3 (5) |
| C13A—C12A—H12A | 109.3 | N2C—C14C—H14E | 108.9 |
| C11A—C12A—H12A | 109.3 | C13C—C14C—H14E | 108.9 |
| C13A—C12A—H12B | 109.3 | N2C—C14C—H14F | 108.9 |
| C11A—C12A—H12B | 109.3 | C13C—C14C—H14F | 108.9 |
| H12A—C12A—H12B | 107.9 | H14E—C14C—H14F | 107.7 |
| C14A—C13A—C12A | 110.5 (5) | N2C—C15C—C6C | 112.0 (4) |
| C14A—C13A—H13A | 109.6 | N2C—C15C—H15E | 109.2 |
| C12A—C13A—H13A | 109.6 | C6C—C15C—H15E | 109.2 |
| C14A—C13A—H13B | 109.6 | N2C—C15C—H15F | 109.2 |
| C12A—C13A—H13B | 109.6 | C6C—C15C—H15F | 109.2 |
| H13A—C13A—H13B | 108.1 | H15E—C15C—H15F | 107.9 |
| C13A—C14A—N2A | 112.9 (5) | C5D—N1D—C1D | 121.8 (4) |
| C13A—C14A—H14A | 109.0 | C5D—N1D—C9D | 123.8 (4) |
| N2A—C14A—H14A | 109.0 | C1D—N1D—C9D | 114.4 (4) |
| C13A—C14A—H14B | 109.0 | C15D—N2D—C14D | 111.9 (4) |
| N2A—C14A—H14B | 109.0 | C15D—N2D—C10D | 113.7 (4) |
| H14A—C14A—H14B | 107.8 | C14D—N2D—C10D | 113.0 (4) |
| N2A—C15A—C6A | 111.8 (4) | C15D—N2D—H2GN | 111 (3) |
| N2A—C15A—H15A | 109.3 | C14D—N2D—H2GN | 104 (3) |
| C6A—C15A—H15A | 109.3 | C10D—N2D—H2GN | 103 (3) |
| N2A—C15A—H15B | 109.3 | O1D—C1D—N1D | 118.4 (4) |
| C6A—C15A—H15B | 109.3 | O1D—C1D—C2D | 125.3 (4) |
| H15A—C15A—H15B | 107.9 | N1D—C1D—C2D | 116.4 (4) |
| C5B—N1B—C1B | 122.2 (4) | C3D—C2D—C1D | 120.9 (5) |
| C5B—N1B—C9B | 123.3 (4) | C3D—C2D—H2G | 119.6 |
| C1B—N1B—C9B | 114.6 (4) | C1D—C2D—H2G | 119.6 |
| C15B—N2B—C10B | 114.9 (4) | C2D—C3D—C4D | 120.7 (5) |
| C15B—N2B—C14B | 111.9 (4) | C2D—C3D—H3G | 119.6 |
| C10B—N2B—C14B | 111.6 (4) | C4D—C3D—H3G | 119.6 |
| C15B—N2B—H2CN | 111 (4) | C5D—C4D—C3D | 119.9 (5) |
| C10B—N2B—H2CN | 107 (4) | C5D—C4D—H4G | 120.1 |
| C14B—N2B—H2CN | 100 (5) | C3D—C4D—H4G | 120.1 |
| O1B—C1B—N1B | 118.3 (5) | C4D—C5D—N1D | 120.2 (4) |
| O1B—C1B—C2B | 125.4 (5) | C4D—C5D—C6D | 121.8 (4) |

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| N1B—C1B—C2B | 116.3 (5) | N1D—C5D—C6D | 118.0 (4) |
| C3B—C2B—C1B | 121.5 (5) | C7D—C6D—C5D | 111.8 (4) |
| C3B—C2B—H2C | 119.3 | C7D—C6D—C15D | 110.4 (4) |
| C1B—C2B—H2C | 119.3 | C5D—C6D—C15D | 110.0 (4) |
| C2B—C3B—C4B | 120.9 (6) | C7D—C6D—H6G | 108.2 |
| C2B—C3B—H3C | 119.6 | C5D—C6D—H6G | 108.2 |
| C4B—C3B—H3C | 119.6 | C15D—C6D—H6G | 108.2 |
| C5B—C4B—C3B | 118.8 (5) | C6D—C7D—C8D | 107.7 (4) |
| C5B—C4B—H4C | 120.6 | C6D—C7D—H7G | 110.2 |
| C3B—C4B—H4C | 120.6 | C8D—C7D—H7G | 110.2 |
| C4B—C5B—N1B | 120.3 (5) | C6D—C7D—H7H | 110.2 |
| C4B—C5B—C6B | 120.8 (5) | C8D—C7D—H7H | 110.2 |
| N1B—C5B—C6B | 118.9 (5) | H7G—C7D—H7H | 108.5 |
| C5B—C6B—C7B | 111.2 (4) | C10D—C8D—C7D | 111.0 (4) |
| C5B—C6B—C15B | 110.9 (4) | C10D—C8D—C9D | 113.8 (4) |
| C7B—C6B—C15B | 109.3 (5) | C7D—C8D—C9D | 109.0 (4) |
| C5B—C6B—H6C | 108.5 | C10D—C8D—H8G | 107.6 |
| C7B—C6B—H6C | 108.5 | C7D—C8D—H8G | 107.6 |
| C15B—C6B—H6C | 108.5 | C9D—C8D—H8G | 107.6 |
| C8B—C7B—C6B | 107.1 (4) | N1D—C9D—C8D | 115.0 (4) |
| C8B—C7B—H7C | 110.3 | N1D—C9D—H9G | 108.5 |
| C6B—C7B—H7C | 110.3 | C8D—C9D—H9G | 108.5 |
| C8B—C7B—H7D | 110.3 | N1D—C9D—H9H | 108.5 |
| C6B—C7B—H7D | 110.3 | C8D—C9D—H9H | 108.5 |
| H7C—C7B—H7D | 108.5 | H9G—C9D—H9H | 107.5 |
| C7B—C8B—C10B | 111.5 (4) | N2D—C10D—C8D | 111.3 (4) |
| C7B—C8B—C9B | 110.2 (4) | N2D—C10D—C11D | 110.2 (4) |
| C10B—C8B—C9B | 114.2 (4) | C8D—C10D—C11D | 112.3 (4) |
| C7B—C8B—H8C | 106.9 | N2D—C10D—H10G | 107.6 |
| C10B—C8B—H8C | 106.9 | C8D—C10D—H10G | 107.6 |
| C9B—C8B—H8C | 106.9 | C11D—C10D—H10G | 107.6 |
| N1B—C9B—C8B | 114.6 (4) | C12D—C11D—C10D | 112.9 (5) |
| N1B—C9B—H9C | 108.6 | C12D—C11D—H11G | 109.0 |
| C8B—C9B—H9C | 108.6 | C10D—C11D—H11G | 109.0 |
| N1B—C9B—H9D | 108.6 | C12D—C11D—H11H | 109.0 |
| C8B—C9B—H9D | 108.6 | C10D—C11D—H11H | 109.0 |
| H9C—C9B—H9D | 107.6 | H11G—C11D—H11H | 107.8 |
| N2B—C10B—C8B | 109.5 (4) | C13D—C12D—C11D | 109.8 (5) |
| N2B—C10B—C11B | 110.0 (4) | C13D—C12D—H12G | 109.7 |
| C8B—C10B—C11B | 113.2 (4) | C11D—C12D—H12G | 109.7 |
| N2B—C10B—H10C | 108.0 | C13D—C12D—H12H | 109.7 |
| C8B—C10B—H10C | 108.0 | C11D—C12D—H12H | 109.7 |
| C11B—C10B—H10C | 108.0 | H12G—C12D—H12H | 108.2 |
| C12B—C11B—C10B | 113.0 (5) | C12D—C13D—C14D | 110.7 (5) |
| C12B—C11B—H11C | 109.0 | C12D—C13D—H13G | 109.5 |
| C10B—C11B—H11C | 109.0 | C14D—C13D—H13G | 109.5 |
| C12B—C11B—H11D | 109.0 | C12D—C13D—H13H | 109.5 |
| C10B—C11B—H11D | 109.0 | C14D—C13D—H13H | 109.5 |

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| H11C—C11B—H11D | 107.8 | H13G—C13D—H13H | 108.1 |
| C13B—C12B—C11B | 111.3 (5) | N2D—C14D—C13D | 111.0 (4) |
| C13B—C12B—H12C | 109.4 | N2D—C14D—H14G | 109.4 |
| C11B—C12B—H12C | 109.4 | C13D—C14D—H14G | 109.4 |
| C13B—C12B—H12D | 109.4 | N2D—C14D—H14H | 109.4 |
| C11B—C12B—H12D | 109.4 | C13D—C14D—H14H | 109.4 |
| H12C—C12B—H12D | 108.0 | H14G—C14D—H14H | 108.0 |
| C12B—C13B—C14B | 110.2 (5) | N2D—C15D—C6D | 111.7 (4) |
| C12B—C13B—H13C | 109.6 | N2D—C15D—H15G | 109.3 |
| C14B—C13B—H13C | 109.6 | C6D—C15D—H15G | 109.3 |
| C12B—C13B—H13D | 109.6 | N2D—C15D—H15H | 109.3 |
| C14B—C13B—H13D | 109.6 | C6D—C15D—H15H | 109.3 |
| H13C—C13B—H13D | 108.1 | H15G—C15D—H15H | 107.9 |
| C13B—C14B—N2B | 111.8 (4) | O13—C11—O14 | 112.3 (5) |
| C13B—C14B—H14C | 109.3 | O13—C11—O11 | 112.0 (5) |
| N2B—C14B—H14C | 109.3 | O14—C11—O11 | 111.1 (6) |
| C13B—C14B—H14D | 109.3 | O13—C11—O12 | 110.8 (6) |
| N2B—C14B—H14D | 109.3 | O14—C11—O12 | 106.1 (7) |
| H14C—C14B—H14D | 107.9 | O11—C11—O12 | 104.1 (7) |
| N2B—C15B—C6B | 112.2 (4) | O23—C12—O21 | 100.5 (13) |
| N2B—C15B—H15C | 109.2 | O23—C12—O22 | 109.0 (11) |
| C6B—C15B—H15C | 109.2 | O21—C12—O22 | 108.4 (11) |
| N2B—C15B—H15D | 109.2 | O23—C12—O24 | 112.3 (6) |
| C6B—C15B—H15D | 109.2 | O21—C12—O24 | 112.1 (8) |
| H15C—C15B—H15D | 107.9 | O22—C12—O24 | 113.6 (5) |
| C5C—N1C—C1C | 122.0 (4) | O31—C13—O32 | 107.0 (7) |
| C5C—N1C—C9C | 122.7 (4) | O31—C13—O33 | 112.1 (5) |
| C1C—N1C—C9C | 115.2 (4) | O32—C13—O33 | 110.5 (6) |
| C15C—N2C—C14C | 112.6 (4) | O31—C13—O34 | 110.8 (7) |
| C15C—N2C—C10C | 113.5 (4) | O32—C13—O34 | 105.4 (8) |
| C14C—N2C—C10C | 111.6 (4) | O33—C13—O34 | 110.8 (4) |
| C15C—N2C—H2EN | 105 (6) | O42—C14—O43 | 113.4 (5) |
| C14C—N2C—H2EN | 111 (6) | O42—C14—O41 | 102.8 (8) |
| C10C—N2C—H2EN | 103 (6) | O43—C14—O41 | 105.1 (6) |
| O1C—C1C—N1C | 119.0 (4) | O42—C14—O44 | 115.1 (6) |
| O1C—C1C—C2C | 124.9 (4) | O43—C14—O44 | 113.9 (3) |
| N1C—C1C—C2C | 116.1 (4) | O41—C14—O44 | 105.0 (6) |

Hydrogen-bond geometry (Å, °)

| <i>D</i> —H \cdots <i>A</i> | <i>D</i> —H | H \cdots <i>A</i> | <i>D</i> \cdots <i>A</i> | <i>D</i> —H \cdots <i>A</i> |
|--|-------------|---------------------|----------------------------|-------------------------------|
| N2 <i>A</i> —H2 <i>AN</i> \cdots O1 <i>B</i> | 1.03 (5) | 1.91 (6) | 2.741 (6) | 136 (5) |
| N2 <i>B</i> —H2 <i>CN</i> \cdots O1 <i>A</i> | 0.77 (7) | 2.00 (6) | 2.742 (5) | 163 (6) |
| N2 <i>C</i> —H2 <i>EN</i> \cdots O1 <i>D</i> ⁱ | 0.90 (9) | 2.00 (9) | 2.735 (6) | 138 (8) |
| N2 <i>D</i> —H2 <i>GN</i> \cdots O1 <i>C</i> ⁱⁱ | 1.05 (5) | 1.74 (5) | 2.754 (5) | 159 (5) |

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