

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

Levocetirizinium dipicrate

 Jerry P. Jasinski,^{a*} Ray J. Butcher,^b M. S. Siddegowda,^c
 H. S. Yathirajan^c and A. R. Ramesha^d
^aDepartment of Chemistry, Keene State College, 229 Main Street, Keene, NH 03435-2001, USA, ^bDepartment of Chemistry, Howard University, 525 College Street NW, Washington, DC 20059, USA, ^cDepartment of Studies in Chemistry, University of Mysore, Manasagangotri, Mysore 570 006, India, and ^dRL Fine Chem., Bangalore 560 064, India

Correspondence e-mail: jjasinski@keene.edu

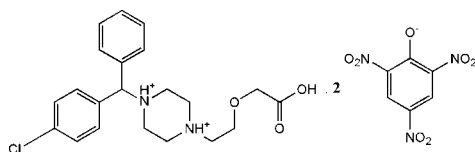
Received 29 October 2010; accepted 7 November 2010

 Key indicators: single-crystal X-ray study; $T = 123$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.044; wR factor = 0.122; data-to-parameter ratio = 10.5.

There are two cation–dianion pairs in the asymmetric unit of the title compound, $\text{C}_{21}\text{H}_{27}\text{ClN}_2\text{O}_3^{2+} \cdot 2\text{C}_6\text{H}_2\text{N}_3\text{O}_7^-$ [systematic name: 1-[2-(carboxymethoxy)ethyl]-4-[(*R*)-(4-chlorophenyl)phenylmethyl]piperazine-1,4-dium bis(2,4,6-trinitrophenolate)]. The piperazine group in the levocetirizinium cation is protonated at both N atoms. The acetyl end groups form $R_2^2(8)$ hydrogen-bonded motifs with adjacent cations. Each picrate anion interacts with the protonated N atom in the cation through a bifurcated $\text{N}-\text{H} \cdots \text{O}$ hydrogen bond, forming $R_1^1(6)$ ring motifs. Strong and weak intermolecular $\text{N}-\text{H} \cdots \text{O}$ and strong $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds, and weak π -ring and π - π stacking interactions [centroid–centroid distance = 3.7419 (14) Å] dominate the crystal packing, creating a three-dimensional supramolecular structure.

Related literature

For related background, see: Hair & Scott, (2006). For related structures, see: Jasinski *et al.* (2009, 2010*a,b*). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_{21}\text{H}_{27}\text{ClN}_2\text{O}_3^{2+} \cdot 2\text{C}_6\text{H}_2\text{N}_3\text{O}_7^-$
 $M_r = 847.11$

 Monoclinic, $P2_1$
 $a = 11.2444$ (1) Å

 $b = 15.7720$ (2) Å

 $c = 20.6204$ (2) Å

 $\beta = 95.998$ (1)°

 $V = 3636.94$ (7) Å³
 $Z = 4$

 Cu $K\alpha$ radiation

 $\mu = 1.74$ mm⁻¹
 $T = 123$ K

 $0.51 \times 0.47 \times 0.34$ mm

Data collection

Oxford Diffraction Xcalibur Ruby

Gemini diffractometer

Absorption correction: multi-scan

 (*CrysAlis RED*; Oxford

Diffraction, 2007)

 $T_{\min} = 0.533$, $T_{\max} = 1.000$

14383 measured reflections

11120 independent reflections

 10728 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.021$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.122$
 $S = 1.03$

11120 reflections

1063 parameters

1 restraint

H-atom parameters constrained

 $\Delta\rho_{\text{max}} = 0.96$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.64$ e Å⁻³

Absolute structure: Flack (1983),

3692 Friedel pairs

Flack parameter: 0.058 (13)

Table 1

Hydrogen-bond geometry (Å, °).

 Cg5 is the centroid of the $\text{C6B}-\text{C11B}$ ring.

| $D-\text{H} \cdots A$ | $D-\text{H}$ | $\text{H} \cdots A$ | $D \cdots A$ | $D-\text{H} \cdots A$ |
|---|--------------|---------------------|--------------|-----------------------|
| $\text{O2A}-\text{H2AD} \cdots \text{O2B}$ | 0.84 | 1.80 | 2.638 (4) | 180 |
| $\text{N1A}-\text{H1AC} \cdots \text{O1D}$ | 0.93 | 1.83 | 2.682 (3) | 152 |
| $\text{N1A}-\text{H1AC} \cdots \text{O7D}$ | 0.93 | 2.63 | 3.301 (3) | 129 |
| $\text{N2A}-\text{H2AC} \cdots \text{O1C}$ | 0.93 | 1.89 | 2.765 (3) | 155 |
| $\text{N2A}-\text{H2AC} \cdots \text{O7C}$ | 0.93 | 2.46 | 2.990 (3) | 116 |
| $\text{O3B}-\text{H3BC} \cdots \text{O3A}$ | 0.84 | 1.76 | 2.601 (4) | 180 |
| $\text{N1B}-\text{H1BC} \cdots \text{O1E}$ | 0.93 | 1.85 | 2.678 (3) | 147 |
| $\text{N1B}-\text{H1BC} \cdots \text{O7E}$ | 0.93 | 2.52 | 3.193 (3) | 130 |
| $\text{N2B}-\text{H2BC} \cdots \text{O1F}$ | 0.93 | 1.91 | 2.764 (3) | 153 |
| $\text{N2B}-\text{H2BC} \cdots \text{O7F}$ | 0.93 | 2.57 | 3.078 (4) | 115 |
| $\text{C19B}-\text{H19C} \cdots \text{Cg5}^i$ | 0.99 | 2.95 | 3.792 (4) | 144 |

 Symmetry code: (i) $x + 1, y, z$.

Data collection: *CrysAlis PRO* (Oxford Diffraction, 2007); cell refinement: *CrysAlis PRO* data reduction: *CrysAlis RED* (Oxford Diffraction, 2007); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

MSS thanks the University of Mysore (UOM) for research facilities and HSY thanks UOM for sabbatical leave. RJB acknowledges the NSF MRI program (grant No. CHE-0619278) for funds to purchase an X-ray diffractometer.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: OM2375).

References

- Allen, F. H., Kennard, O., Watson, D. G., Brammer, L., Orpen, A. G. & Taylor, R. (1987). *J. Chem. Soc. Perkin Trans. 2*, pp. S1–19.
- Flack, H. D. (1983). *Acta Cryst.* **A39**, 876–881.
- Hair, P. I. & Scott, L. J. (2006). *Drugs*, **66**, 973–996.
- Jasinski, J. P., Butcher, R. J., Hakim Al-Arique, Q. N. M., Yathirajan, H. S. & Narayana, B. (2009). *Acta Cryst.* **E65**, o1738–o1739.
- Jasinski, J. P., Butcher, R. J., Hakim Al-Arique, Q. N. M., Yathirajan, H. S. & Narayana, B. (2010*a*). *Acta Cryst.* **E66**, o347–o348.
- Jasinski, J. P., Butcher, R. J., Hakim Al-Arique, Q. N. M., Yathirajan, H. S. & Narayana, B. (2010*b*). *Acta Cryst.* **E66**, o411–o412.
- Oxford Diffraction (2007). *CrysAlis PRO* and *CrysAlis RED* Oxford Diffraction Ltd, Abingdon, England.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.

supporting information

Acta Cryst. (2010). E66, o3167 [https://doi.org/10.1107/S1600536810045733]

Levocetirizinium dipicrate

Jerry P. Jasinski, Ray J. Butcher, M. S. Siddegowda, H. S. Yathirajan and A. R. Ramesha

S1. Comment

Levocetirizine (as levocetirizine dihydrochloride) is a third-generation non-sedative antihistamine, developed from the second-generation antihistamine cetirizine. Chemically, levocetirizine is the active enantiomer of cetirizine. It is the *L*-enantiomer of the cetirizine racemate. Levocetirizine works by blocking histamine receptors. It does not prevent the actual release of histamine from mast cells, but prevents it from binding to its receptors. This in turn prevents the release of other allergy chemicals and increased blood supply to the area, and provides relief from the typical symptoms of hayfever. Levocetirizine is called a non-sedating antihistamine as it does not enter the brain in significant amounts, and is therefore unlikely to cause drowsiness. A review on the use of levocetirizine in the management of allergic rhinitis and skin allergies is described (Hair & Scott, 2006).

Recently, the crystal structures of propiverine picrate (Jasinski *et al.*, 2009), imatinibium dipicrate (Jasinski *et al.*, 2010*b*) and chlorimipraminium picrate (Jasinski *et al.*, 2010*a*) have been reported. The present work reports the crystal structure of the salt, $C_{21}H_{27}ClN_2O_3^{2+} \cdot 2C_6H_2N_3O_7^-$, formed by the interaction between 2-[2-[4-[(*R*)-(4-chlorophenyl)-phenyl-methyl] piperazin-1-yl]ethoxy]acetic acid and 2,4,6-trinitrophenol in aqueous medium.

In the crystal structure of the title compound the 6-membered piperazine groups (N1A/C1A/C2A/N2A/C3A/C4A & N1B/C1B/C2B/N2B/C3B/C4B) in the levocetirizinium cation are protonated at both N atoms (Fig. 1) and adopt slightly distorted chair conformations with puckering parameters Q , θ and φ of 0.591 (3) Å & 0.583 (3) Å, 171.6 (3)° & 170.8 (3)°, and 353.0 (17)° & 358.2 (19)°, for molecules A & B respectively (Figs. 1 & 2). For an ideal chair θ has a value of 0 or 180°. Bond distances (Allen *et al.*, 1987) and angles are in normal ranges. $R_2^1(6)$ graph-set motifs are formed between piperazine N1A—H1AC and N2A—H2AC groups and the picrate anions labeled D and C (Fig. 1) and piperazine N1B—H1BC and N2B—H2BC groups and the picrate anions labeled E and F (Fig. 2) through bifurcated N—H...O hydrogen bonds (Table 1). The acetyl end groups form an $R_2^2(8)$ hydrogen bonded motif with adjacent cations (Fig. 3). The dihedral angle between the mean planes of the anion benzene ring pairs is 31.9 (2)° (C—D) and 37.9 (6)° (E—F), respectively.

The mean plane of the two *o*-NO₂ groups in the two picrate anions are twisted by 15.8 (6)°, 53.7 (3)° (ring C), 25.9 (9)°, 38.5 (1)° (ring D), 24.5 (0)°, 38.7 (2)° (ring E) and 10.3 (3)°, 56.9 (9)° (ring F) with respect to the mean planes of the 6-membered benzene rings. The *p*-NO₂ groups in both picrate anions are nearly in the plane of the ring (torsion angles O4C/N2C/C4C/C3C = -8.8 (4)°; O4D/N2D/C4D/C3D = -175.8 (2)°; O4E/N2E/C4E/C3E = 2.6 (4)°; O4F/N2F/C4F/C3F = 3.4 (4)°). An extensive array of strong and weak N—H...O and strong O—H...O intermolecular hydrogen bonds (Table 1), weak π -ring (Table 2) and π - π (Table 3) stacking interactions dominate crystal packing in the unit cell creating a 3-D supramolecular structure (Fig. 4).

S2. Experimental

Levocetirizine (3.89 g, 0.01 mol) was dissolved in 20 ml of methanol and picric acid (2.4 g, 0.01 mol) was dissolved in 20 ml of methanol. Both the solutions were mixed and stirred in a beaker at room temperature for 1/2 half hour. The mixture was warmed for 10 min at 323 K & kept aside for two days at room temperature. The formed salt was filtered & dried in a vacuum desiccator over phosphorous pentoxide. The salt was recrystallized from dimethylsulphoxide by slow evaporation (m.p: 454–456 K).

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 1.00, 0.95 Å (CH), 0.99 Å (CH₂), 0.93 Å (NH), or 0.84 Å (OH). Isotropic displacement parameters for these atoms were set to 1.2 times (NH), 1.2 (CH, CH₂) or 1.5 (OH) times U_{eq} of the parent atom.

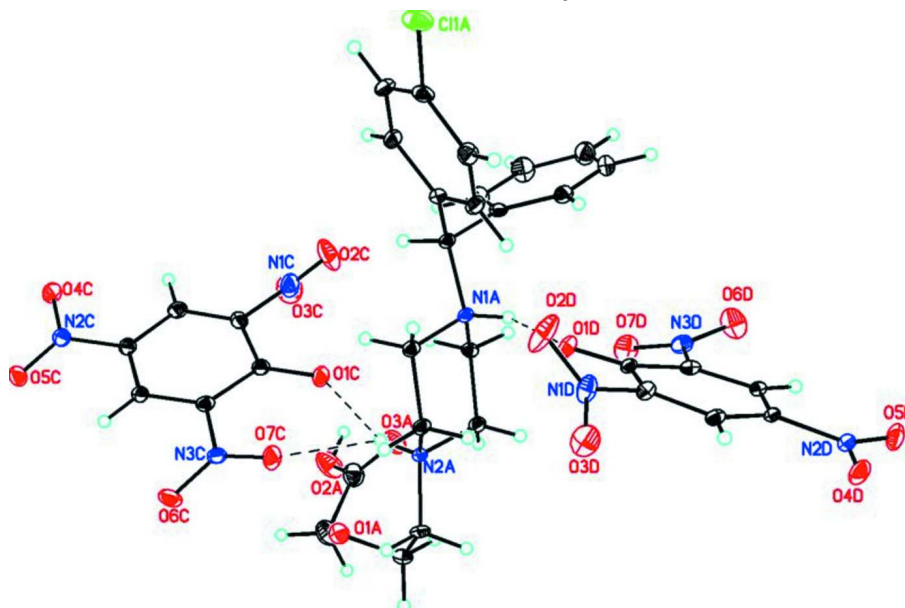


Figure 1

Molecular structure of the title compound showing the atom labeling scheme and 50% probability displacement ellipsoids. Dashed lines indicate strong and weak N—H...O intermolecular hydrogen bonds

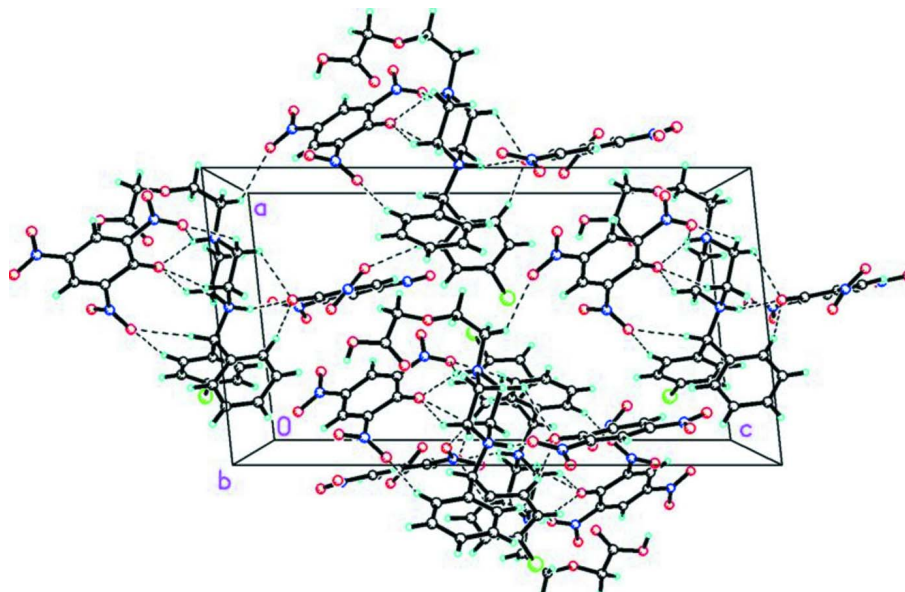


Figure 4

Packing diagram of the title compound viewed down the *b* axis. Dashed lines indicate strong and weak intermolecular N—H...O and O—H...O hydrogen bond interactions creating a 3-D supramolecular structure.

1-[2-(Carboxymethoxy)ethyl]-4-[(*R*)-(4-chlorophenyl)phenylmethyl]piperazine-1,4-dium bis(2,4,6-trinitrophenolate)

Crystal data

$C_{21}H_{27}ClN_2O_3^{2+} \cdot 2C_6H_2N_3O_7^-$

$M_r = 847.11$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 11.2444 (1) \text{ \AA}$

$b = 15.7720 (2) \text{ \AA}$

$c = 20.6204 (2) \text{ \AA}$

$\beta = 95.998 (1)^\circ$

$V = 3636.94 (7) \text{ \AA}^3$

$Z = 4$

$F(000) = 1752$

$D_x = 1.547 \text{ Mg m}^{-3}$

Cu $K\alpha$ radiation, $\lambda = 1.54178 \text{ \AA}$

Cell parameters from 12871 reflections

$\theta = 4.7\text{--}73.9^\circ$

$\mu = 1.74 \text{ mm}^{-1}$

$T = 123 \text{ K}$

Block, yellow

$0.51 \times 0.47 \times 0.34 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur Ruby Gemini diffractometer

Radiation source: Enhance (Cu) X-ray Source

Graphite monochromator

Detector resolution: $10.5081 \text{ pixels mm}^{-1}$

ω scans

Absorption correction: multi-scan

(*CrysAlis RED*; Oxford Diffraction, 2007)

$T_{\min} = 0.533$, $T_{\max} = 1.000$

14383 measured reflections

11120 independent reflections

10728 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.021$

$\theta_{\max} = 74.1^\circ$, $\theta_{\min} = 4.7^\circ$

$h = -9 \rightarrow 13$

$k = -19 \rightarrow 18$

$l = -25 \rightarrow 24$

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.122$ $S = 1.03$

11120 reflections

1063 parameters

1 restraint

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0826P)^2 + 1.9759P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.96 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.64 \text{ e } \text{\AA}^{-3}$ Absolute structure: Flack (1983), 3460 Friedel
pairs

Absolute structure parameter: 0.058 (13)

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 | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|--------------|----------------------------------|
| C11A | 1.43159 (6) | -0.07506 (5) | 0.46063 (4) | 0.03707 (19) |
| O1A | 0.51357 (19) | 0.14380 (15) | 0.62287 (11) | 0.0322 (5) |
| O2A | 0.5911 (3) | 0.2584 (2) | 0.76889 (13) | 0.0535 (7) |
| H2AD | 0.6533 | 0.2854 | 0.7821 | 0.080* |
| O3A | 0.6722 (2) | 0.26091 (17) | 0.67498 (11) | 0.0385 (6) |
| N1A | 0.95454 (18) | 0.17016 (15) | 0.53050 (10) | 0.0155 (4) |
| H1AC | 0.9645 | 0.1922 | 0.4896 | 0.019* |
| N2A | 0.69680 (19) | 0.15302 (15) | 0.53729 (11) | 0.0174 (4) |
| H2AC | 0.7070 | 0.1310 | 0.5793 | 0.021* |
| C1A | 0.8903 (2) | 0.08603 (17) | 0.52236 (12) | 0.0178 (5) |
| H1AA | 0.8923 | 0.0577 | 0.5653 | 0.021* |
| H1AB | 0.9322 | 0.0491 | 0.4933 | 0.021* |
| C2A | 0.7617 (2) | 0.09740 (18) | 0.49397 (12) | 0.0181 (5) |
| H2AA | 0.7593 | 0.1232 | 0.4501 | 0.022* |
| H2AB | 0.7218 | 0.0414 | 0.4894 | 0.022* |
| C3A | 0.7555 (2) | 0.23808 (18) | 0.53789 (13) | 0.0194 (5) |
| H3AA | 0.7120 | 0.2781 | 0.5639 | 0.023* |
| H3AB | 0.7521 | 0.2602 | 0.4928 | 0.023* |
| C4A | 0.8845 (2) | 0.23205 (17) | 0.56680 (12) | 0.0162 (5) |
| H4AA | 0.9221 | 0.2887 | 0.5658 | 0.019* |
| H4AB | 0.8872 | 0.2142 | 0.6130 | 0.019* |
| C5A | 1.0771 (2) | 0.15712 (18) | 0.56848 (12) | 0.0170 (5) |
| H5AA | 1.0633 | 0.1349 | 0.6125 | 0.020* |

| | | | | |
|------|--------------|---------------|--------------|------------|
| C6A | 1.1552 (2) | 0.09331 (18) | 0.53805 (13) | 0.0187 (5) |
| C7A | 1.2402 (2) | 0.05136 (19) | 0.58085 (14) | 0.0236 (6) |
| H7AA | 1.2406 | 0.0593 | 0.6265 | 0.028* |
| C8A | 1.3239 (3) | -0.00169 (19) | 0.55710 (15) | 0.0267 (6) |
| H8AA | 1.3825 | -0.0295 | 0.5861 | 0.032* |
| C9A | 1.3206 (2) | -0.01337 (18) | 0.49055 (16) | 0.0251 (6) |
| C10A | 1.2349 (3) | 0.0241 (2) | 0.44723 (15) | 0.0260 (6) |
| H10A | 1.2321 | 0.0128 | 0.4018 | 0.031* |
| C11A | 1.1524 (3) | 0.07878 (19) | 0.47100 (14) | 0.0240 (6) |
| H11A | 1.0941 | 0.1063 | 0.4416 | 0.029* |
| C12A | 1.1408 (2) | 0.24216 (18) | 0.57884 (13) | 0.0196 (5) |
| C13A | 1.1565 (3) | 0.2778 (2) | 0.64133 (14) | 0.0291 (7) |
| H13A | 1.1260 | 0.2496 | 0.6768 | 0.035* |
| C14A | 1.2168 (3) | 0.3543 (3) | 0.65159 (16) | 0.0390 (8) |
| H14A | 1.2260 | 0.3788 | 0.6939 | 0.047* |
| C15A | 1.2637 (3) | 0.3954 (2) | 0.60031 (17) | 0.0366 (8) |
| H15A | 1.3063 | 0.4471 | 0.6075 | 0.044* |
| C16A | 1.2474 (3) | 0.3596 (2) | 0.53822 (16) | 0.0308 (7) |
| H16A | 1.2787 | 0.3875 | 0.5029 | 0.037* |
| C17A | 1.1863 (3) | 0.28368 (19) | 0.52730 (14) | 0.0223 (6) |
| H17A | 1.1754 | 0.2601 | 0.4847 | 0.027* |
| C18A | 0.5646 (2) | 0.1559 (2) | 0.51504 (14) | 0.0260 (6) |
| H18A | 0.5514 | 0.1894 | 0.4743 | 0.031* |
| H18B | 0.5353 | 0.0976 | 0.5053 | 0.031* |
| C19A | 0.4949 (3) | 0.1943 (2) | 0.56542 (16) | 0.0302 (7) |
| H19A | 0.5221 | 0.2531 | 0.5749 | 0.036* |
| H19B | 0.4088 | 0.1957 | 0.5494 | 0.036* |
| C20A | 0.4897 (3) | 0.1860 (3) | 0.68060 (18) | 0.0403 (8) |
| H20A | 0.4714 | 0.1436 | 0.7135 | 0.048* |
| H20B | 0.4187 | 0.2228 | 0.6712 | 0.048* |
| C21A | 0.5937 (3) | 0.2386 (2) | 0.70758 (17) | 0.0370 (8) |
| Cl1B | 0.18104 (10) | 0.10273 (6) | 0.86966 (5) | 0.0527 (3) |
| O1B | 0.9688 (2) | 0.43787 (17) | 0.86713 (11) | 0.0376 (5) |
| O2B | 0.7870 (2) | 0.34294 (17) | 0.80996 (12) | 0.0391 (6) |
| O3B | 0.8678 (3) | 0.3414 (2) | 0.71568 (13) | 0.0575 (8) |
| H3BC | 0.8046 | 0.3154 | 0.7025 | 0.086* |
| N1B | 0.53581 (19) | 0.41964 (16) | 0.97035 (10) | 0.0190 (4) |
| H1BC | 0.5269 | 0.3949 | 1.0105 | 0.023* |
| N2B | 0.7932 (2) | 0.42637 (17) | 0.95643 (11) | 0.0224 (5) |
| H2BC | 0.7814 | 0.4518 | 0.9156 | 0.027* |
| C1B | 0.5976 (2) | 0.35768 (18) | 0.92903 (12) | 0.0198 (5) |
| H1BA | 0.5926 | 0.3792 | 0.8837 | 0.024* |
| H1BB | 0.5549 | 0.3027 | 0.9282 | 0.024* |
| C2B | 0.7275 (3) | 0.34358 (19) | 0.95376 (13) | 0.0224 (6) |
| H2BA | 0.7332 | 0.3180 | 0.9978 | 0.027* |
| H2BB | 0.7642 | 0.3039 | 0.9244 | 0.027* |
| C3B | 0.7382 (3) | 0.4805 (2) | 1.00382 (14) | 0.0250 (6) |
| H3BA | 0.7831 | 0.5344 | 1.0096 | 0.030* |

| | | | | |
|------|--------------|---------------|--------------|-------------|
| H3BB | 0.7433 | 0.4515 | 1.0466 | 0.030* |
| C4B | 0.6090 (3) | 0.49921 (18) | 0.98106 (14) | 0.0223 (6) |
| H4BA | 0.5748 | 0.5349 | 1.0140 | 0.027* |
| H4BB | 0.6047 | 0.5317 | 0.9398 | 0.027* |
| C5B | 0.4118 (2) | 0.43970 (19) | 0.93531 (13) | 0.0221 (6) |
| H5BA | 0.4251 | 0.4663 | 0.8927 | 0.026* |
| C6B | 0.3450 (2) | 0.3572 (2) | 0.91958 (13) | 0.0233 (6) |
| C7B | 0.3046 (3) | 0.3084 (2) | 0.96865 (15) | 0.0263 (6) |
| H7BA | 0.3131 | 0.3283 | 1.0124 | 0.032* |
| C8B | 0.2515 (3) | 0.2303 (2) | 0.95391 (17) | 0.0339 (7) |
| H8BA | 0.2244 | 0.1961 | 0.9873 | 0.041* |
| C9B | 0.2388 (3) | 0.2028 (2) | 0.88932 (17) | 0.0365 (8) |
| C10B | 0.2738 (4) | 0.2517 (3) | 0.84000 (16) | 0.0408 (8) |
| H10B | 0.2618 | 0.2328 | 0.7961 | 0.049* |
| C11B | 0.3272 (3) | 0.3296 (2) | 0.85527 (16) | 0.0343 (7) |
| H11B | 0.3517 | 0.3643 | 0.8215 | 0.041* |
| C12B | 0.3390 (3) | 0.50213 (19) | 0.97057 (15) | 0.0244 (6) |
| C13B | 0.3556 (3) | 0.5196 (2) | 1.03748 (15) | 0.0292 (7) |
| H13B | 0.4170 | 0.4916 | 1.0645 | 0.035* |
| C14B | 0.2819 (3) | 0.5781 (2) | 1.06445 (18) | 0.0353 (7) |
| H14B | 0.2957 | 0.5916 | 1.1095 | 0.042* |
| C15B | 0.1889 (3) | 0.6167 (2) | 1.0263 (2) | 0.0417 (8) |
| H15B | 0.1369 | 0.6548 | 1.0452 | 0.050* |
| C16B | 0.1725 (3) | 0.5990 (2) | 0.9599 (2) | 0.0427 (9) |
| H16B | 0.1091 | 0.6255 | 0.9332 | 0.051* |
| C17B | 0.2473 (3) | 0.5434 (2) | 0.93261 (17) | 0.0338 (7) |
| H17B | 0.2361 | 0.5331 | 0.8870 | 0.041* |
| C18B | 0.9256 (3) | 0.4166 (2) | 0.97499 (15) | 0.0327 (7) |
| H18C | 0.9391 | 0.3790 | 1.0135 | 0.039* |
| H18D | 0.9607 | 0.4727 | 0.9871 | 0.039* |
| C19B | 0.9876 (3) | 0.3799 (3) | 0.91983 (19) | 0.0396 (8) |
| H19C | 1.0741 | 0.3730 | 0.9333 | 0.047* |
| H19D | 0.9535 | 0.3238 | 0.9070 | 0.047* |
| C20B | 0.9815 (4) | 0.4014 (3) | 0.80583 (19) | 0.0483 (10) |
| H20C | 1.0471 | 0.3593 | 0.8105 | 0.058* |
| H20D | 1.0034 | 0.4461 | 0.7755 | 0.058* |
| C21B | 0.8681 (4) | 0.3587 (3) | 0.77753 (17) | 0.0415 (8) |
| O1C | 0.79680 (19) | 0.09712 (15) | 0.65794 (10) | 0.0274 (4) |
| O2C | 1.0086 (3) | 0.1450 (2) | 0.73266 (12) | 0.0616 (10) |
| O3C | 0.9234 (3) | 0.17726 (19) | 0.81843 (16) | 0.0577 (8) |
| O4C | 0.8609 (2) | -0.11502 (16) | 0.91389 (10) | 0.0331 (5) |
| O5C | 0.7121 (2) | -0.18315 (15) | 0.86224 (11) | 0.0313 (5) |
| O6C | 0.56936 (19) | -0.10817 (17) | 0.64531 (11) | 0.0361 (6) |
| O7C | 0.6664 (2) | -0.02135 (15) | 0.59006 (10) | 0.0309 (5) |
| N1C | 0.9382 (3) | 0.1313 (2) | 0.77214 (14) | 0.0401 (7) |
| N2C | 0.7885 (2) | -0.12627 (17) | 0.86580 (12) | 0.0255 (5) |
| N3C | 0.6489 (2) | -0.05485 (17) | 0.64152 (11) | 0.0224 (5) |
| C1C | 0.7961 (2) | 0.04300 (19) | 0.70205 (13) | 0.0205 (5) |

| | | | | |
|------|--------------|---------------|--------------|-------------|
| C2C | 0.8661 (3) | 0.0537 (2) | 0.76457 (14) | 0.0254 (6) |
| C3C | 0.8660 (3) | 0.0012 (2) | 0.81686 (14) | 0.0249 (6) |
| H3CA | 0.9145 | 0.0127 | 0.8564 | 0.030* |
| C4C | 0.7922 (2) | -0.0702 (2) | 0.81035 (13) | 0.0223 (6) |
| C5C | 0.7203 (2) | -0.08658 (19) | 0.75341 (13) | 0.0212 (5) |
| H5CA | 0.6686 | -0.1344 | 0.7505 | 0.025* |
| C6C | 0.7240 (2) | -0.03285 (19) | 0.70048 (13) | 0.0207 (6) |
| O1D | 0.93928 (19) | 0.18889 (14) | 0.40063 (9) | 0.0249 (4) |
| O2D | 0.9747 (3) | 0.05081 (17) | 0.32418 (14) | 0.0548 (8) |
| O3D | 0.8281 (2) | 0.05186 (16) | 0.24704 (12) | 0.0424 (6) |
| O4D | 0.8374 (2) | 0.30033 (16) | 0.11042 (10) | 0.0309 (5) |
| O5D | 0.8824 (2) | 0.42219 (16) | 0.15462 (11) | 0.0380 (5) |
| O6D | 1.0061 (3) | 0.44020 (16) | 0.38295 (13) | 0.0443 (6) |
| O7D | 0.9339 (2) | 0.34861 (17) | 0.44645 (10) | 0.0364 (5) |
| N1D | 0.8989 (3) | 0.08785 (16) | 0.28649 (12) | 0.0285 (6) |
| N2D | 0.8686 (2) | 0.34472 (17) | 0.15838 (11) | 0.0231 (5) |
| N3D | 0.9598 (2) | 0.37156 (16) | 0.39307 (12) | 0.0241 (5) |
| C1D | 0.9264 (2) | 0.22591 (18) | 0.34697 (12) | 0.0160 (5) |
| C2D | 0.9020 (2) | 0.18040 (18) | 0.28569 (13) | 0.0198 (5) |
| C3D | 0.8806 (2) | 0.21806 (19) | 0.22587 (12) | 0.0179 (5) |
| H3DA | 0.8595 | 0.1850 | 0.1879 | 0.022* |
| C4D | 0.8903 (2) | 0.30535 (18) | 0.22165 (13) | 0.0180 (5) |
| C5D | 0.9179 (2) | 0.35465 (18) | 0.27694 (13) | 0.0186 (5) |
| H5DA | 0.9264 | 0.4143 | 0.2733 | 0.022* |
| C6D | 0.9327 (2) | 0.31620 (18) | 0.33673 (13) | 0.0184 (5) |
| O1E | 0.5620 (2) | 0.39892 (13) | 1.09998 (9) | 0.0253 (4) |
| O2E | 0.5266 (3) | 0.53058 (18) | 1.18380 (13) | 0.0507 (7) |
| O3E | 0.6761 (3) | 0.52363 (18) | 1.25888 (13) | 0.0496 (7) |
| O4E | 0.6578 (2) | 0.2660 (2) | 1.38424 (11) | 0.0452 (7) |
| O5E | 0.6351 (2) | 0.14505 (19) | 1.33419 (12) | 0.0447 (7) |
| O6E | 0.4915 (4) | 0.14663 (19) | 1.10720 (15) | 0.0659 (10) |
| O7E | 0.5555 (2) | 0.24316 (15) | 1.04664 (10) | 0.0331 (5) |
| N1E | 0.6011 (3) | 0.49029 (19) | 1.21902 (13) | 0.0342 (6) |
| N2E | 0.6358 (2) | 0.2233 (2) | 1.33420 (13) | 0.0336 (7) |
| N3E | 0.5367 (2) | 0.21628 (17) | 1.09969 (13) | 0.0277 (5) |
| C1E | 0.5733 (2) | 0.35792 (19) | 1.15211 (13) | 0.0194 (5) |
| C2E | 0.5990 (2) | 0.39845 (19) | 1.21537 (14) | 0.0218 (6) |
| C3E | 0.6207 (2) | 0.3559 (2) | 1.27318 (13) | 0.0248 (6) |
| H3EA | 0.6411 | 0.3861 | 1.3127 | 0.030* |
| C4E | 0.6126 (2) | 0.2685 (2) | 1.27344 (14) | 0.0259 (6) |
| C5E | 0.5844 (2) | 0.2231 (2) | 1.21629 (14) | 0.0239 (6) |
| H5EA | 0.5775 | 0.1631 | 1.2172 | 0.029* |
| C6E | 0.5666 (2) | 0.26696 (19) | 1.15802 (13) | 0.0195 (5) |
| O1F | 0.68330 (18) | 0.50280 (14) | 0.84530 (10) | 0.0261 (4) |
| O2F | 0.4523 (2) | 0.4860 (2) | 0.77442 (12) | 0.0477 (7) |
| O3F | 0.5271 (3) | 0.4344 (2) | 0.69206 (16) | 0.0716 (10) |
| O4F | 0.6482 (2) | 0.71341 (16) | 0.58982 (10) | 0.0324 (5) |
| O5F | 0.79813 (19) | 0.77792 (16) | 0.64310 (10) | 0.0305 (5) |

| | | | | |
|------|--------------|--------------|--------------|------------|
| O6F | 0.92948 (19) | 0.69978 (19) | 0.86099 (11) | 0.0389 (6) |
| O7F | 0.8319 (2) | 0.61047 (17) | 0.91374 (10) | 0.0344 (5) |
| N1F | 0.5248 (2) | 0.48557 (18) | 0.73525 (12) | 0.0302 (6) |
| N2F | 0.7192 (2) | 0.72331 (17) | 0.63870 (11) | 0.0230 (5) |
| N3F | 0.8502 (2) | 0.64698 (18) | 0.86315 (11) | 0.0251 (5) |
| C1F | 0.6912 (2) | 0.55608 (19) | 0.80234 (13) | 0.0201 (5) |
| C2F | 0.6167 (2) | 0.55174 (19) | 0.74045 (13) | 0.0219 (6) |
| C3F | 0.6259 (2) | 0.6021 (2) | 0.68762 (13) | 0.0210 (5) |
| H3FA | 0.5766 | 0.5930 | 0.6480 | 0.025* |
| C4F | 0.7102 (2) | 0.66759 (19) | 0.69346 (13) | 0.0204 (5) |
| C5F | 0.7830 (2) | 0.6799 (2) | 0.75121 (13) | 0.0219 (6) |
| H5FA | 0.8398 | 0.7246 | 0.7546 | 0.026* |
| C6F | 0.7730 (2) | 0.6275 (2) | 0.80350 (13) | 0.0220 (6) |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C11A | 0.0230 (3) | 0.0272 (3) | 0.0621 (5) | 0.0041 (3) | 0.0098 (3) | -0.0105 (3) |
| O1A | 0.0289 (11) | 0.0372 (12) | 0.0308 (11) | -0.0023 (9) | 0.0039 (9) | 0.0049 (10) |
| O2A | 0.0541 (16) | 0.069 (2) | 0.0402 (14) | -0.0227 (15) | 0.0193 (12) | -0.0114 (14) |
| O3A | 0.0374 (13) | 0.0474 (14) | 0.0326 (12) | -0.0101 (11) | 0.0119 (10) | -0.0043 (11) |
| N1A | 0.0137 (9) | 0.0221 (11) | 0.0104 (9) | 0.0022 (9) | 0.0003 (8) | 0.0029 (8) |
| N2A | 0.0118 (9) | 0.0228 (11) | 0.0170 (10) | -0.0004 (8) | -0.0011 (8) | 0.0052 (9) |
| C1A | 0.0172 (12) | 0.0205 (13) | 0.0155 (11) | 0.0013 (10) | 0.0005 (9) | 0.0004 (10) |
| C2A | 0.0178 (12) | 0.0230 (13) | 0.0124 (11) | 0.0008 (10) | -0.0029 (9) | 0.0004 (10) |
| C3A | 0.0174 (12) | 0.0214 (14) | 0.0196 (12) | 0.0036 (10) | 0.0023 (10) | 0.0030 (11) |
| C4A | 0.0138 (11) | 0.0211 (13) | 0.0140 (11) | 0.0000 (10) | 0.0030 (9) | -0.0025 (10) |
| C5A | 0.0150 (11) | 0.0236 (13) | 0.0122 (11) | 0.0026 (10) | 0.0003 (9) | 0.0030 (10) |
| C6A | 0.0138 (11) | 0.0215 (13) | 0.0204 (12) | 0.0013 (10) | 0.0001 (10) | 0.0044 (11) |
| C7A | 0.0225 (13) | 0.0262 (14) | 0.0206 (13) | -0.0008 (11) | -0.0046 (10) | 0.0059 (11) |
| C8A | 0.0188 (13) | 0.0245 (15) | 0.0352 (16) | 0.0037 (11) | -0.0048 (11) | 0.0078 (13) |
| C9A | 0.0147 (12) | 0.0172 (13) | 0.0445 (17) | 0.0031 (10) | 0.0091 (11) | -0.0003 (12) |
| C10A | 0.0275 (14) | 0.0285 (15) | 0.0230 (14) | 0.0031 (12) | 0.0078 (11) | 0.0018 (12) |
| C11A | 0.0214 (13) | 0.0271 (15) | 0.0235 (13) | 0.0081 (11) | 0.0027 (11) | 0.0065 (12) |
| C12A | 0.0122 (11) | 0.0267 (14) | 0.0194 (12) | 0.0003 (10) | -0.0008 (9) | 0.0016 (11) |
| C13A | 0.0307 (15) | 0.0394 (18) | 0.0164 (13) | -0.0086 (13) | -0.0012 (11) | 0.0021 (12) |
| C14A | 0.047 (2) | 0.043 (2) | 0.0244 (15) | -0.0132 (16) | -0.0075 (13) | -0.0055 (15) |
| C15A | 0.0389 (18) | 0.0336 (17) | 0.0357 (17) | -0.0098 (14) | -0.0045 (14) | 0.0048 (14) |
| C16A | 0.0283 (15) | 0.0328 (17) | 0.0316 (15) | -0.0017 (13) | 0.0046 (12) | 0.0107 (14) |
| C17A | 0.0217 (13) | 0.0237 (14) | 0.0215 (13) | 0.0043 (11) | 0.0019 (10) | 0.0002 (11) |
| C18A | 0.0127 (12) | 0.0367 (16) | 0.0267 (14) | -0.0022 (11) | -0.0072 (10) | 0.0072 (13) |
| C19A | 0.0143 (12) | 0.0363 (17) | 0.0397 (17) | 0.0017 (11) | 0.0021 (11) | 0.0104 (14) |
| C20A | 0.0407 (18) | 0.045 (2) | 0.0373 (18) | -0.0047 (16) | 0.0161 (15) | -0.0020 (16) |
| C21A | 0.0402 (18) | 0.0380 (19) | 0.0339 (17) | 0.0000 (15) | 0.0097 (14) | 0.0020 (15) |
| Cl1B | 0.0762 (7) | 0.0329 (4) | 0.0430 (5) | -0.0139 (4) | -0.0218 (4) | 0.0062 (4) |
| O1B | 0.0362 (12) | 0.0415 (14) | 0.0356 (12) | -0.0070 (10) | 0.0058 (10) | -0.0004 (11) |
| O2B | 0.0378 (12) | 0.0464 (14) | 0.0349 (12) | -0.0041 (11) | 0.0125 (10) | -0.0074 (11) |
| O3B | 0.0541 (16) | 0.081 (2) | 0.0408 (14) | -0.0284 (16) | 0.0203 (12) | -0.0149 (15) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| N1B | 0.0194 (10) | 0.0241 (11) | 0.0129 (10) | -0.0010 (9) | -0.0013 (8) | 0.0018 (9) |
| N2B | 0.0213 (11) | 0.0276 (12) | 0.0166 (10) | -0.0040 (10) | -0.0060 (8) | 0.0065 (10) |
| C1B | 0.0229 (13) | 0.0238 (13) | 0.0121 (11) | -0.0016 (11) | -0.0008 (10) | 0.0016 (10) |
| C2B | 0.0225 (13) | 0.0246 (14) | 0.0192 (12) | -0.0029 (11) | -0.0022 (10) | 0.0031 (11) |
| C3B | 0.0272 (14) | 0.0265 (14) | 0.0201 (13) | -0.0085 (12) | -0.0031 (11) | 0.0027 (12) |
| C4B | 0.0303 (15) | 0.0195 (13) | 0.0168 (12) | -0.0020 (11) | 0.0012 (10) | 0.0028 (11) |
| C5B | 0.0200 (12) | 0.0309 (15) | 0.0139 (12) | -0.0017 (11) | -0.0052 (10) | 0.0085 (11) |
| C6B | 0.0196 (13) | 0.0294 (15) | 0.0194 (13) | 0.0022 (11) | -0.0052 (10) | 0.0040 (12) |
| C7B | 0.0237 (14) | 0.0311 (15) | 0.0232 (14) | 0.0026 (12) | -0.0013 (11) | 0.0059 (12) |
| C8B | 0.0373 (17) | 0.0284 (16) | 0.0348 (17) | -0.0031 (14) | -0.0021 (13) | 0.0093 (14) |
| C9B | 0.0387 (18) | 0.0281 (16) | 0.0386 (18) | -0.0073 (13) | -0.0152 (14) | 0.0030 (14) |
| C10B | 0.055 (2) | 0.042 (2) | 0.0224 (15) | -0.0088 (17) | -0.0138 (14) | -0.0012 (15) |
| C11B | 0.0391 (17) | 0.0393 (18) | 0.0221 (14) | -0.0101 (15) | -0.0080 (12) | 0.0072 (14) |
| C12B | 0.0222 (13) | 0.0244 (14) | 0.0268 (14) | -0.0006 (11) | 0.0034 (11) | 0.0080 (12) |
| C13B | 0.0307 (15) | 0.0326 (16) | 0.0247 (15) | 0.0033 (13) | 0.0051 (12) | 0.0032 (13) |
| C14B | 0.0361 (17) | 0.0317 (17) | 0.0400 (17) | 0.0029 (14) | 0.0129 (14) | 0.0000 (14) |
| C15B | 0.0371 (18) | 0.0255 (16) | 0.064 (2) | 0.0038 (13) | 0.0119 (17) | 0.0029 (16) |
| C16B | 0.0293 (17) | 0.0344 (18) | 0.062 (2) | 0.0083 (15) | -0.0050 (16) | 0.0123 (18) |
| C17B | 0.0352 (17) | 0.0322 (16) | 0.0320 (16) | -0.0005 (14) | -0.0057 (13) | 0.0085 (14) |
| C18B | 0.0203 (13) | 0.0451 (19) | 0.0305 (15) | -0.0038 (13) | -0.0087 (12) | 0.0066 (14) |
| C19B | 0.0205 (14) | 0.047 (2) | 0.051 (2) | 0.0004 (14) | 0.0002 (14) | 0.0025 (17) |
| C20B | 0.048 (2) | 0.057 (3) | 0.042 (2) | -0.0171 (18) | 0.0182 (17) | -0.0066 (18) |
| C21B | 0.050 (2) | 0.043 (2) | 0.0324 (17) | -0.0087 (17) | 0.0099 (15) | -0.0010 (15) |
| O1C | 0.0310 (10) | 0.0296 (11) | 0.0204 (10) | -0.0041 (9) | -0.0035 (8) | 0.0087 (9) |
| O2C | 0.0673 (18) | 0.088 (2) | 0.0270 (12) | -0.0519 (18) | -0.0058 (12) | 0.0159 (14) |
| O3C | 0.0692 (19) | 0.0376 (15) | 0.0648 (19) | -0.0111 (14) | -0.0002 (15) | -0.0087 (14) |
| O4C | 0.0332 (11) | 0.0427 (13) | 0.0207 (10) | -0.0072 (10) | -0.0099 (8) | 0.0117 (10) |
| O5C | 0.0292 (11) | 0.0374 (12) | 0.0267 (10) | -0.0088 (9) | 0.0004 (9) | 0.0107 (10) |
| O6C | 0.0218 (10) | 0.0573 (16) | 0.0278 (11) | -0.0154 (10) | -0.0049 (8) | 0.0090 (11) |
| O7C | 0.0406 (12) | 0.0345 (12) | 0.0165 (10) | -0.0071 (10) | -0.0020 (8) | 0.0073 (9) |
| N1C | 0.0459 (17) | 0.0417 (17) | 0.0293 (14) | -0.0153 (14) | -0.0128 (12) | 0.0133 (13) |
| N2C | 0.0233 (11) | 0.0312 (13) | 0.0216 (12) | -0.0020 (10) | 0.0000 (9) | 0.0068 (10) |
| N3C | 0.0153 (10) | 0.0310 (13) | 0.0205 (11) | 0.0015 (9) | 0.0004 (9) | 0.0040 (10) |
| C1C | 0.0171 (12) | 0.0275 (14) | 0.0170 (12) | 0.0016 (11) | 0.0022 (10) | 0.0032 (11) |
| C2C | 0.0213 (13) | 0.0305 (15) | 0.0239 (14) | -0.0048 (12) | 0.0002 (11) | 0.0070 (12) |
| C3C | 0.0211 (13) | 0.0322 (16) | 0.0200 (13) | -0.0013 (12) | -0.0050 (10) | 0.0023 (12) |
| C4C | 0.0201 (12) | 0.0295 (15) | 0.0173 (13) | -0.0002 (11) | 0.0017 (10) | 0.0060 (12) |
| C5C | 0.0177 (12) | 0.0261 (14) | 0.0198 (13) | 0.0001 (11) | 0.0019 (10) | 0.0052 (11) |
| C6C | 0.0154 (12) | 0.0303 (15) | 0.0165 (12) | 0.0031 (11) | 0.0027 (10) | 0.0029 (11) |
| O1D | 0.0319 (11) | 0.0273 (11) | 0.0150 (9) | 0.0069 (9) | -0.0009 (8) | 0.0046 (8) |
| O2D | 0.087 (2) | 0.0286 (13) | 0.0426 (14) | 0.0205 (14) | -0.0201 (14) | 0.0040 (12) |
| O3D | 0.0560 (15) | 0.0280 (12) | 0.0408 (13) | -0.0048 (11) | -0.0066 (12) | -0.0063 (11) |
| O4D | 0.0341 (11) | 0.0426 (13) | 0.0150 (9) | 0.0040 (10) | -0.0027 (8) | 0.0058 (9) |
| O5D | 0.0449 (13) | 0.0324 (12) | 0.0346 (12) | -0.0054 (11) | -0.0050 (10) | 0.0190 (10) |
| O6D | 0.0595 (16) | 0.0320 (13) | 0.0415 (13) | -0.0186 (12) | 0.0063 (12) | -0.0136 (11) |
| O7D | 0.0487 (14) | 0.0398 (13) | 0.0205 (10) | -0.0008 (11) | 0.0022 (9) | -0.0082 (10) |
| N1D | 0.0437 (15) | 0.0195 (13) | 0.0218 (12) | 0.0033 (11) | 0.0013 (11) | 0.0009 (10) |
| N2D | 0.0164 (11) | 0.0327 (13) | 0.0197 (11) | 0.0018 (9) | 0.0003 (9) | 0.0087 (10) |

| | | | | | | |
|-----|-------------|-------------|-------------|--------------|--------------|--------------|
| N3D | 0.0227 (11) | 0.0260 (13) | 0.0225 (12) | 0.0014 (10) | -0.0033 (9) | -0.0054 (10) |
| C1D | 0.0131 (11) | 0.0218 (13) | 0.0127 (11) | 0.0025 (10) | -0.0003 (9) | 0.0024 (10) |
| C2D | 0.0204 (12) | 0.0190 (13) | 0.0197 (13) | 0.0033 (10) | 0.0000 (10) | 0.0009 (11) |
| C3D | 0.0147 (11) | 0.0250 (13) | 0.0136 (12) | 0.0034 (10) | -0.0006 (9) | -0.0012 (10) |
| C4D | 0.0117 (11) | 0.0259 (14) | 0.0162 (12) | 0.0020 (10) | -0.0002 (9) | 0.0067 (11) |
| C5D | 0.0124 (11) | 0.0199 (13) | 0.0231 (13) | 0.0015 (10) | 0.0003 (9) | 0.0059 (11) |
| C6D | 0.0125 (11) | 0.0229 (13) | 0.0194 (13) | 0.0016 (10) | -0.0007 (9) | -0.0043 (11) |
| O1E | 0.0384 (11) | 0.0251 (11) | 0.0117 (9) | 0.0039 (9) | 0.0000 (8) | 0.0020 (8) |
| O2E | 0.0682 (18) | 0.0368 (14) | 0.0428 (14) | 0.0218 (13) | -0.0140 (13) | -0.0115 (12) |
| O3E | 0.0667 (18) | 0.0423 (15) | 0.0358 (13) | -0.0032 (13) | -0.0139 (12) | -0.0175 (12) |
| O4E | 0.0369 (13) | 0.081 (2) | 0.0165 (11) | 0.0089 (13) | -0.0002 (9) | 0.0123 (12) |
| O5E | 0.0317 (12) | 0.0609 (18) | 0.0396 (13) | -0.0068 (12) | -0.0052 (10) | 0.0312 (13) |
| O6E | 0.112 (3) | 0.0382 (16) | 0.0483 (16) | -0.0372 (18) | 0.0095 (17) | -0.0060 (13) |
| O7E | 0.0512 (14) | 0.0322 (12) | 0.0151 (9) | -0.0055 (10) | -0.0006 (9) | -0.0010 (9) |
| N1E | 0.0437 (16) | 0.0342 (15) | 0.0232 (13) | 0.0069 (12) | -0.0033 (11) | -0.0115 (12) |
| N2E | 0.0125 (11) | 0.065 (2) | 0.0231 (13) | 0.0012 (12) | 0.0016 (9) | 0.0171 (14) |
| N3E | 0.0319 (13) | 0.0232 (13) | 0.0275 (13) | -0.0049 (11) | -0.0001 (10) | -0.0021 (11) |
| C1E | 0.0149 (12) | 0.0272 (14) | 0.0158 (12) | 0.0035 (10) | 0.0002 (9) | 0.0007 (11) |
| C2E | 0.0188 (12) | 0.0268 (15) | 0.0196 (13) | 0.0040 (10) | 0.0014 (10) | -0.0028 (11) |
| C3E | 0.0139 (12) | 0.0486 (19) | 0.0121 (12) | 0.0033 (12) | 0.0019 (9) | -0.0049 (12) |
| C4E | 0.0131 (12) | 0.0466 (19) | 0.0184 (13) | 0.0025 (12) | 0.0025 (10) | 0.0110 (13) |
| C5E | 0.0154 (12) | 0.0302 (15) | 0.0258 (14) | -0.0018 (11) | 0.0009 (10) | 0.0093 (12) |
| C6E | 0.0154 (12) | 0.0252 (14) | 0.0177 (12) | -0.0016 (10) | 0.0001 (10) | 0.0013 (11) |
| O1F | 0.0262 (10) | 0.0294 (11) | 0.0217 (10) | 0.0024 (8) | -0.0022 (8) | 0.0094 (9) |
| O2F | 0.0445 (14) | 0.0596 (18) | 0.0400 (14) | -0.0253 (13) | 0.0084 (11) | 0.0029 (13) |
| O3F | 0.086 (2) | 0.067 (2) | 0.064 (2) | -0.0362 (19) | 0.0176 (17) | -0.0337 (18) |
| O4F | 0.0310 (11) | 0.0461 (13) | 0.0183 (10) | -0.0084 (10) | -0.0061 (8) | 0.0079 (9) |
| O5F | 0.0220 (10) | 0.0427 (13) | 0.0267 (10) | -0.0079 (9) | 0.0027 (8) | 0.0118 (10) |
| O6F | 0.0215 (11) | 0.0641 (17) | 0.0295 (11) | -0.0136 (11) | -0.0049 (9) | 0.0105 (12) |
| O7F | 0.0427 (13) | 0.0410 (13) | 0.0178 (10) | -0.0085 (11) | -0.0045 (9) | 0.0087 (10) |
| N1F | 0.0380 (14) | 0.0319 (14) | 0.0192 (12) | -0.0092 (12) | -0.0045 (10) | 0.0053 (11) |
| N2F | 0.0176 (11) | 0.0341 (13) | 0.0176 (11) | -0.0002 (10) | 0.0032 (9) | 0.0041 (10) |
| N3F | 0.0193 (11) | 0.0366 (14) | 0.0188 (11) | 0.0008 (10) | -0.0008 (9) | 0.0031 (11) |
| C1F | 0.0178 (12) | 0.0282 (14) | 0.0150 (12) | 0.0058 (11) | 0.0053 (10) | 0.0028 (11) |
| C2F | 0.0218 (13) | 0.0252 (14) | 0.0189 (12) | -0.0002 (11) | 0.0029 (10) | -0.0022 (11) |
| C3F | 0.0194 (12) | 0.0297 (14) | 0.0137 (11) | 0.0020 (11) | 0.0011 (9) | 0.0000 (11) |
| C4F | 0.0142 (11) | 0.0311 (15) | 0.0162 (12) | 0.0044 (11) | 0.0037 (9) | 0.0035 (11) |
| C5F | 0.0143 (11) | 0.0328 (15) | 0.0189 (13) | 0.0021 (11) | 0.0034 (10) | 0.0019 (12) |
| C6F | 0.0145 (12) | 0.0311 (15) | 0.0202 (13) | 0.0043 (11) | 0.0003 (10) | 0.0020 (12) |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|-----------|-----------|
| C11A—C9A | 1.745 (3) | C11B—H11B | 0.9500 |
| O1A—C20A | 1.413 (4) | C12B—C17B | 1.389 (4) |
| O1A—C19A | 1.425 (4) | C12B—C13B | 1.400 (4) |
| O2A—C21A | 1.306 (4) | C13B—C14B | 1.394 (5) |
| O2A—H2AD | 0.8400 | C13B—H13B | 0.9500 |
| O3A—C21A | 1.216 (4) | C14B—C15B | 1.382 (5) |

| | | | |
|-----------|-----------|-----------|-----------|
| N1A—C4A | 1.502 (3) | C14B—H14B | 0.9500 |
| N1A—C1A | 1.512 (3) | C15B—C16B | 1.390 (6) |
| N1A—C5A | 1.526 (3) | C15B—H15B | 0.9500 |
| N1A—H1AC | 0.9300 | C16B—C17B | 1.376 (5) |
| N2A—C3A | 1.495 (4) | C16B—H16B | 0.9500 |
| N2A—C2A | 1.495 (3) | C17B—H17B | 0.9500 |
| N2A—C18A | 1.510 (3) | C18B—C19B | 1.510 (5) |
| N2A—H2AC | 0.9300 | C18B—H18C | 0.9900 |
| C1A—C2A | 1.513 (3) | C18B—H18D | 0.9900 |
| C1A—H1AA | 0.9900 | C19B—H19C | 0.9900 |
| C1A—H1AB | 0.9900 | C19B—H19D | 0.9900 |
| C2A—H2AA | 0.9900 | C20B—C21B | 1.505 (5) |
| C2A—H2AB | 0.9900 | C20B—H20C | 0.9900 |
| C3A—C4A | 1.512 (3) | C20B—H20D | 0.9900 |
| C3A—H3AA | 0.9900 | O1C—C1C | 1.248 (4) |
| C3A—H3AB | 0.9900 | O2C—N1C | 1.213 (4) |
| C4A—H4AA | 0.9900 | O3C—N1C | 1.224 (4) |
| C4A—H4AB | 0.9900 | O4C—N2C | 1.228 (3) |
| C5A—C6A | 1.514 (4) | O5C—N2C | 1.239 (3) |
| C5A—C12A | 1.525 (4) | O6C—N3C | 1.236 (3) |
| C5A—H5AA | 1.0000 | O7C—N3C | 1.220 (3) |
| C6A—C7A | 1.398 (4) | N1C—C2C | 1.467 (4) |
| C6A—C11A | 1.399 (4) | N2C—C4C | 1.449 (4) |
| C7A—C8A | 1.386 (4) | N3C—C6C | 1.448 (4) |
| C7A—H7AA | 0.9500 | C1C—C6C | 1.444 (4) |
| C8A—C9A | 1.381 (5) | C1C—C2C | 1.448 (4) |
| C8A—H8AA | 0.9500 | C2C—C3C | 1.360 (4) |
| C9A—C10A | 1.376 (4) | C3C—C4C | 1.398 (4) |
| C10A—C11A | 1.393 (4) | C3C—H3CA | 0.9500 |
| C10A—H10A | 0.9500 | C4C—C5C | 1.378 (4) |
| C11A—H11A | 0.9500 | C5C—C6C | 1.386 (4) |
| C12A—C17A | 1.390 (4) | C5C—H5CA | 0.9500 |
| C12A—C13A | 1.400 (4) | O1D—C1D | 1.246 (3) |
| C13A—C14A | 1.390 (5) | O2D—N1D | 1.238 (4) |
| C13A—H13A | 0.9500 | O3D—N1D | 1.218 (4) |
| C14A—C15A | 1.390 (5) | O4D—N2D | 1.232 (3) |
| C14A—H14A | 0.9500 | O5D—N2D | 1.235 (4) |
| C15A—C16A | 1.394 (5) | O6D—N3D | 1.228 (4) |
| C15A—H15A | 0.9500 | O7D—N3D | 1.222 (3) |
| C16A—C17A | 1.387 (5) | N1D—C2D | 1.460 (4) |
| C16A—H16A | 0.9500 | N2D—C4D | 1.443 (3) |
| C17A—H17A | 0.9500 | N3D—C6D | 1.460 (4) |
| C18A—C19A | 1.494 (4) | C1D—C6D | 1.442 (4) |
| C18A—H18A | 0.9900 | C1D—C2D | 1.454 (4) |
| C18A—H18B | 0.9900 | C2D—C3D | 1.368 (4) |
| C19A—H19A | 0.9900 | C3D—C4D | 1.384 (4) |
| C19A—H19B | 0.9900 | C3D—H3DA | 0.9500 |
| C20A—C21A | 1.493 (5) | C4D—C5D | 1.388 (4) |

| | | | |
|---------------|-------------|----------------|-----------|
| C20A—H20A | 0.9900 | C5D—C6D | 1.368 (4) |
| C20A—H20B | 0.9900 | C5D—H5DA | 0.9500 |
| C11B—C9B | 1.739 (3) | O1E—C1E | 1.249 (3) |
| O1B—C20B | 1.410 (4) | O2E—N1E | 1.227 (4) |
| O1B—C19B | 1.419 (4) | O3E—N1E | 1.232 (4) |
| O2B—C21B | 1.211 (4) | O4E—N2E | 1.235 (4) |
| O3B—C21B | 1.304 (4) | O5E—N2E | 1.235 (4) |
| O3B—H3BC | 0.8400 | O6E—N3E | 1.227 (4) |
| N1B—C4B | 1.505 (4) | O7E—N3E | 1.212 (3) |
| N1B—C1B | 1.512 (4) | N1E—C2E | 1.451 (4) |
| N1B—C5B | 1.535 (3) | N2E—C4E | 1.441 (4) |
| N1B—H1BC | 0.9300 | N3E—C6E | 1.454 (4) |
| N2B—C3B | 1.481 (4) | C1E—C6E | 1.442 (4) |
| N2B—C2B | 1.499 (4) | C1E—C2E | 1.454 (4) |
| N2B—C18B | 1.505 (4) | C2E—C3E | 1.368 (4) |
| N2B—H2BC | 0.9300 | C3E—C4E | 1.382 (5) |
| C1B—C2B | 1.512 (4) | C3E—H3EA | 0.9500 |
| C1B—H1BA | 0.9900 | C4E—C5E | 1.387 (5) |
| C1B—H1BB | 0.9900 | C5E—C6E | 1.383 (4) |
| C2B—H2BA | 0.9900 | C5E—H5EA | 0.9500 |
| C2B—H2BB | 0.9900 | O1F—C1F | 1.231 (4) |
| C3B—C4B | 1.508 (4) | O2F—N1F | 1.206 (4) |
| C3B—H3BA | 0.9900 | O3F—N1F | 1.205 (4) |
| C3B—H3BB | 0.9900 | O4F—N2F | 1.228 (3) |
| C4B—H4BA | 0.9900 | O5F—N2F | 1.233 (3) |
| C4B—H4BB | 0.9900 | O6F—N3F | 1.224 (4) |
| C5B—C12B | 1.514 (4) | O7F—N3F | 1.228 (3) |
| C5B—C6B | 1.520 (4) | N1F—C2F | 1.465 (4) |
| C5B—H5BA | 1.0000 | N2F—C4F | 1.443 (4) |
| C6B—C7B | 1.386 (4) | N3F—C6F | 1.462 (4) |
| C6B—C11B | 1.390 (4) | C1F—C2F | 1.453 (4) |
| C7B—C8B | 1.388 (5) | C1F—C6F | 1.453 (4) |
| C7B—H7BA | 0.9500 | C2F—C3F | 1.361 (4) |
| C8B—C9B | 1.394 (5) | C3F—C4F | 1.399 (4) |
| C8B—H8BA | 0.9500 | C3F—H3FA | 0.9500 |
| C9B—C10B | 1.367 (5) | C4F—C5F | 1.386 (4) |
| C10B—C11B | 1.388 (5) | C5F—C6F | 1.372 (4) |
| C10B—H10B | 0.9500 | C5F—H5FA | 0.9500 |
| C20A—O1A—C19A | 114.4 (3) | C10B—C9B—C11B | 117.9 (3) |
| C21A—O2A—H2AD | 109.5 | C8B—C9B—C11B | 120.2 (3) |
| C4A—N1A—C1A | 110.86 (19) | C9B—C10B—C11B | 118.8 (3) |
| C4A—N1A—C5A | 108.93 (19) | C9B—C10B—H10B | 120.6 |
| C1A—N1A—C5A | 109.52 (19) | C11B—C10B—H10B | 120.6 |
| C4A—N1A—H1AC | 109.2 | C10B—C11B—C6B | 120.6 (3) |
| C1A—N1A—H1AC | 109.2 | C10B—C11B—H11B | 119.7 |
| C5A—N1A—H1AC | 109.2 | C6B—C11B—H11B | 119.7 |
| C3A—N2A—C2A | 106.79 (19) | C17B—C12B—C13B | 118.5 (3) |

| | | | |
|---------------|-----------|----------------|-----------|
| C3A—N2A—C18A | 113.3 (2) | C17B—C12B—C5B | 116.1 (3) |
| C2A—N2A—C18A | 111.2 (2) | C13B—C12B—C5B | 125.3 (3) |
| C3A—N2A—H2AC | 108.5 | C14B—C13B—C12B | 120.0 (3) |
| C2A—N2A—H2AC | 108.5 | C14B—C13B—H13B | 120.0 |
| C18A—N2A—H2AC | 108.5 | C12B—C13B—H13B | 120.0 |
| N1A—C1A—C2A | 111.4 (2) | C15B—C14B—C13B | 120.7 (3) |
| N1A—C1A—H1AA | 109.3 | C15B—C14B—H14B | 119.7 |
| C2A—C1A—H1AA | 109.3 | C13B—C14B—H14B | 119.7 |
| N1A—C1A—H1AB | 109.3 | C14B—C15B—C16B | 119.2 (3) |
| C2A—C1A—H1AB | 109.3 | C14B—C15B—H15B | 120.4 |
| H1AA—C1A—H1AB | 108.0 | C16B—C15B—H15B | 120.4 |
| N2A—C2A—C1A | 110.2 (2) | C17B—C16B—C15B | 120.5 (3) |
| N2A—C2A—H2AA | 109.6 | C17B—C16B—H16B | 119.8 |
| C1A—C2A—H2AA | 109.6 | C15B—C16B—H16B | 119.8 |
| N2A—C2A—H2AB | 109.6 | C16B—C17B—C12B | 121.2 (3) |
| C1A—C2A—H2AB | 109.6 | C16B—C17B—H17B | 119.4 |
| H2AA—C2A—H2AB | 108.1 | C12B—C17B—H17B | 119.4 |
| N2A—C3A—C4A | 110.6 (2) | N2B—C18B—C19B | 111.7 (3) |
| N2A—C3A—H3AA | 109.5 | N2B—C18B—H18C | 109.3 |
| C4A—C3A—H3AA | 109.5 | C19B—C18B—H18C | 109.3 |
| N2A—C3A—H3AB | 109.5 | N2B—C18B—H18D | 109.3 |
| C4A—C3A—H3AB | 109.5 | C19B—C18B—H18D | 109.3 |
| H3AA—C3A—H3AB | 108.1 | H18C—C18B—H18D | 107.9 |
| N1A—C4A—C3A | 112.1 (2) | O1B—C19B—C18B | 106.6 (3) |
| N1A—C4A—H4AA | 109.2 | O1B—C19B—H19C | 110.4 |
| C3A—C4A—H4AA | 109.2 | C18B—C19B—H19C | 110.4 |
| N1A—C4A—H4AB | 109.2 | O1B—C19B—H19D | 110.4 |
| C3A—C4A—H4AB | 109.2 | C18B—C19B—H19D | 110.4 |
| H4AA—C4A—H4AB | 107.9 | H19C—C19B—H19D | 108.6 |
| C6A—C5A—C12A | 111.0 (2) | O1B—C20B—C21B | 111.6 (3) |
| C6A—C5A—N1A | 114.1 (2) | O1B—C20B—H20C | 109.3 |
| C12A—C5A—N1A | 109.8 (2) | C21B—C20B—H20C | 109.3 |
| C6A—C5A—H5AA | 107.2 | O1B—C20B—H20D | 109.3 |
| C12A—C5A—H5AA | 107.2 | C21B—C20B—H20D | 109.3 |
| N1A—C5A—H5AA | 107.2 | H20C—C20B—H20D | 108.0 |
| C7A—C6A—C11A | 119.3 (3) | O2B—C21B—O3B | 124.9 (4) |
| C7A—C6A—C5A | 116.2 (2) | O2B—C21B—C20B | 122.3 (3) |
| C11A—C6A—C5A | 124.4 (2) | O3B—C21B—C20B | 112.8 (3) |
| C8A—C7A—C6A | 120.5 (3) | O2C—N1C—O3C | 124.7 (3) |
| C8A—C7A—H7AA | 119.8 | O2C—N1C—C2C | 118.0 (3) |
| C6A—C7A—H7AA | 119.8 | O3C—N1C—C2C | 117.3 (3) |
| C9A—C8A—C7A | 118.9 (3) | O4C—N2C—O5C | 123.3 (2) |
| C9A—C8A—H8AA | 120.6 | O4C—N2C—C4C | 118.6 (2) |
| C7A—C8A—H8AA | 120.6 | O5C—N2C—C4C | 118.1 (2) |
| C10A—C9A—C8A | 122.1 (3) | O7C—N3C—O6C | 122.2 (2) |
| C10A—C9A—C11A | 119.2 (2) | O7C—N3C—C6C | 119.6 (2) |
| C8A—C9A—C11A | 118.7 (2) | O6C—N3C—C6C | 118.2 (2) |
| C9A—C10A—C11A | 118.9 (3) | O1C—C1C—C6C | 126.6 (2) |

| | | | |
|----------------|-----------|--------------|-----------|
| C9A—C10A—H10A | 120.5 | O1C—C1C—C2C | 121.6 (3) |
| C11A—C10A—H10A | 120.5 | C6C—C1C—C2C | 111.6 (2) |
| C10A—C11A—C6A | 120.2 (3) | C3C—C2C—C1C | 126.1 (3) |
| C10A—C11A—H11A | 119.9 | C3C—C2C—N1C | 118.0 (3) |
| C6A—C11A—H11A | 119.9 | C1C—C2C—N1C | 115.8 (3) |
| C17A—C12A—C13A | 119.5 (3) | C2C—C3C—C4C | 117.7 (3) |
| C17A—C12A—C5A | 120.8 (2) | C2C—C3C—H3CA | 121.2 |
| C13A—C12A—C5A | 119.6 (2) | C4C—C3C—H3CA | 121.2 |
| C14A—C13A—C12A | 120.1 (3) | C5C—C4C—C3C | 121.5 (3) |
| C14A—C13A—H13A | 119.9 | C5C—C4C—N2C | 119.7 (3) |
| C12A—C13A—H13A | 119.9 | C3C—C4C—N2C | 118.8 (2) |
| C13A—C14A—C15A | 120.4 (3) | C4C—C5C—C6C | 119.6 (3) |
| C13A—C14A—H14A | 119.8 | C4C—C5C—H5CA | 120.2 |
| C15A—C14A—H14A | 119.8 | C6C—C5C—H5CA | 120.2 |
| C14A—C15A—C16A | 119.1 (3) | C5C—C6C—C1C | 123.5 (2) |
| C14A—C15A—H15A | 120.4 | C5C—C6C—N3C | 116.9 (3) |
| C16A—C15A—H15A | 120.4 | C1C—C6C—N3C | 119.6 (2) |
| C17A—C16A—C15A | 120.9 (3) | O3D—N1D—O2D | 123.9 (3) |
| C17A—C16A—H16A | 119.5 | O3D—N1D—C2D | 118.2 (3) |
| C15A—C16A—H16A | 119.5 | O2D—N1D—C2D | 117.6 (3) |
| C16A—C17A—C12A | 119.9 (3) | O4D—N2D—O5D | 122.6 (2) |
| C16A—C17A—H17A | 120.1 | O4D—N2D—C4D | 119.2 (3) |
| C12A—C17A—H17A | 120.1 | O5D—N2D—C4D | 118.2 (3) |
| C19A—C18A—N2A | 111.9 (2) | O7D—N3D—O6D | 124.0 (3) |
| C19A—C18A—H18A | 109.2 | O7D—N3D—C6D | 119.3 (3) |
| N2A—C18A—H18A | 109.2 | O6D—N3D—C6D | 116.7 (2) |
| C19A—C18A—H18B | 109.2 | O1D—C1D—C6D | 126.1 (2) |
| N2A—C18A—H18B | 109.2 | O1D—C1D—C2D | 122.3 (3) |
| H18A—C18A—H18B | 107.9 | C6D—C1D—C2D | 111.6 (2) |
| O1A—C19A—C18A | 108.0 (3) | C3D—C2D—C1D | 124.7 (3) |
| O1A—C19A—H19A | 110.1 | C3D—C2D—N1D | 116.3 (3) |
| C18A—C19A—H19A | 110.1 | C1D—C2D—N1D | 119.1 (2) |
| O1A—C19A—H19B | 110.1 | C2D—C3D—C4D | 118.8 (3) |
| C18A—C19A—H19B | 110.1 | C2D—C3D—H3DA | 120.6 |
| H19A—C19A—H19B | 108.4 | C4D—C3D—H3DA | 120.6 |
| O1A—C20A—C21A | 111.4 (3) | C3D—C4D—C5D | 121.2 (2) |
| O1A—C20A—H20A | 109.3 | C3D—C4D—N2D | 118.5 (3) |
| C21A—C20A—H20A | 109.3 | C5D—C4D—N2D | 120.3 (3) |
| O1A—C20A—H20B | 109.3 | C6D—C5D—C4D | 119.2 (3) |
| C21A—C20A—H20B | 109.3 | C6D—C5D—H5DA | 120.4 |
| H20A—C20A—H20B | 108.0 | C4D—C5D—H5DA | 120.4 |
| O3A—C21A—O2A | 123.8 (3) | C5D—C6D—C1D | 124.5 (3) |
| O3A—C21A—C20A | 122.7 (3) | C5D—C6D—N3D | 116.6 (3) |
| O2A—C21A—C20A | 113.4 (3) | C1D—C6D—N3D | 118.9 (2) |
| C20B—O1B—C19B | 113.8 (3) | O2E—N1E—O3E | 123.4 (3) |
| C21B—O3B—H3BC | 109.5 | O2E—N1E—C2E | 118.7 (3) |
| C4B—N1B—C1B | 110.2 (2) | O3E—N1E—C2E | 117.8 (3) |
| C4B—N1B—C5B | 110.7 (2) | O5E—N2E—O4E | 123.1 (3) |

| | | | |
|---------------|-----------|--------------|-----------|
| C1B—N1B—C5B | 108.4 (2) | O5E—N2E—C4E | 119.6 (3) |
| C4B—N1B—H1BC | 109.2 | O4E—N2E—C4E | 117.4 (3) |
| C1B—N1B—H1BC | 109.2 | O7E—N3E—O6E | 122.7 (3) |
| C5B—N1B—H1BC | 109.2 | O7E—N3E—C6E | 120.5 (2) |
| C3B—N2B—C2B | 106.7 (2) | O6E—N3E—C6E | 116.8 (3) |
| C3B—N2B—C18B | 111.2 (2) | O1E—C1E—C6E | 125.9 (3) |
| C2B—N2B—C18B | 113.0 (2) | O1E—C1E—C2E | 122.5 (3) |
| C3B—N2B—H2BC | 108.6 | C6E—C1E—C2E | 111.6 (2) |
| C2B—N2B—H2BC | 108.6 | C3E—C2E—N1E | 116.4 (3) |
| C18B—N2B—H2BC | 108.6 | C3E—C2E—C1E | 124.5 (3) |
| N1B—C1B—C2B | 112.9 (2) | N1E—C2E—C1E | 119.1 (3) |
| N1B—C1B—H1BA | 109.0 | C2E—C3E—C4E | 119.1 (3) |
| C2B—C1B—H1BA | 109.0 | C2E—C3E—H3EA | 120.4 |
| N1B—C1B—H1BB | 109.0 | C4E—C3E—H3EA | 120.4 |
| C2B—C1B—H1BB | 109.0 | C3E—C4E—C5E | 121.4 (3) |
| H1BA—C1B—H1BB | 107.8 | C3E—C4E—N2E | 119.4 (3) |
| N2B—C2B—C1B | 109.9 (2) | C5E—C4E—N2E | 119.2 (3) |
| N2B—C2B—H2BA | 109.7 | C6E—C5E—C4E | 118.6 (3) |
| C1B—C2B—H2BA | 109.7 | C6E—C5E—H5EA | 120.7 |
| N2B—C2B—H2BB | 109.7 | C4E—C5E—H5EA | 120.7 |
| C1B—C2B—H2BB | 109.7 | C5E—C6E—C1E | 124.5 (3) |
| H2BA—C2B—H2BB | 108.2 | C5E—C6E—N3E | 116.3 (3) |
| N2B—C3B—C4B | 111.3 (2) | C1E—C6E—N3E | 119.1 (2) |
| N2B—C3B—H3BA | 109.4 | O3F—N1F—O2F | 124.5 (3) |
| C4B—C3B—H3BA | 109.4 | O3F—N1F—C2F | 117.5 (3) |
| N2B—C3B—H3BB | 109.4 | O2F—N1F—C2F | 117.9 (3) |
| C4B—C3B—H3BB | 109.4 | O4F—N2F—O5F | 123.4 (2) |
| H3BA—C3B—H3BB | 108.0 | O4F—N2F—C4F | 118.2 (2) |
| N1B—C4B—C3B | 112.1 (2) | O5F—N2F—C4F | 118.5 (2) |
| N1B—C4B—H4BA | 109.2 | O6F—N3F—O7F | 122.5 (2) |
| C3B—C4B—H4BA | 109.2 | O6F—N3F—C6F | 118.9 (2) |
| N1B—C4B—H4BB | 109.2 | O7F—N3F—C6F | 118.6 (2) |
| C3B—C4B—H4BB | 109.2 | O1F—C1F—C2F | 121.4 (3) |
| H4BA—C4B—H4BB | 107.9 | O1F—C1F—C6F | 127.6 (3) |
| C12B—C5B—C6B | 112.3 (2) | C2F—C1F—C6F | 110.9 (2) |
| C12B—C5B—N1B | 114.7 (2) | C3F—C2F—C1F | 126.2 (3) |
| C6B—C5B—N1B | 109.2 (2) | C3F—C2F—N1F | 118.0 (2) |
| C12B—C5B—H5BA | 106.7 | C1F—C2F—N1F | 115.8 (2) |
| C6B—C5B—H5BA | 106.7 | C2F—C3F—C4F | 118.0 (2) |
| N1B—C5B—H5BA | 106.7 | C2F—C3F—H3FA | 121.0 |
| C7B—C6B—C11B | 119.8 (3) | C4F—C3F—H3FA | 121.0 |
| C7B—C6B—C5B | 120.8 (3) | C5F—C4F—C3F | 120.7 (3) |
| C11B—C6B—C5B | 119.4 (3) | C5F—C4F—N2F | 120.3 (3) |
| C6B—C7B—C8B | 120.0 (3) | C3F—C4F—N2F | 119.0 (2) |
| C6B—C7B—H7BA | 120.0 | C6F—C5F—C4F | 120.1 (3) |
| C8B—C7B—H7BA | 120.0 | C6F—C5F—H5FA | 119.9 |
| C7B—C8B—C9B | 118.9 (3) | C4F—C5F—H5FA | 119.9 |
| C7B—C8B—H8BA | 120.5 | C5F—C6F—C1F | 123.9 (3) |

| | | | |
|---------------------|------------|-----------------|------------|
| C9B—C8B—H8BA | 120.5 | C5F—C6F—N3F | 116.4 (3) |
| C10B—C9B—C8B | 121.8 (3) | C1F—C6F—N3F | 119.7 (2) |
| C4A—N1A—C1A—C2A | 51.3 (3) | N1C—C2C—C3C—C4C | 176.6 (3) |
| C5A—N1A—C1A—C2A | 171.5 (2) | C2C—C3C—C4C—C5C | -1.0 (4) |
| C3A—N2A—C2A—C1A | 63.6 (3) | C2C—C3C—C4C—N2C | -178.7 (3) |
| C18A—N2A—C2A—C1A | -172.3 (2) | O4C—N2C—C4C—C5C | 173.5 (3) |
| N1A—C1A—C2A—N2A | -59.0 (3) | O5C—N2C—C4C—C5C | -6.7 (4) |
| C2A—N2A—C3A—C4A | -62.8 (3) | O4C—N2C—C4C—C3C | -8.7 (4) |
| C18A—N2A—C3A—C4A | 174.4 (2) | O5C—N2C—C4C—C3C | 171.1 (3) |
| C1A—N1A—C4A—C3A | -50.6 (3) | C3C—C4C—C5C—C6C | 2.3 (4) |
| C5A—N1A—C4A—C3A | -171.2 (2) | N2C—C4C—C5C—C6C | 180.0 (3) |
| N2A—C3A—C4A—N1A | 57.6 (3) | C4C—C5C—C6C—C1C | -2.8 (4) |
| C4A—N1A—C5A—C6A | 177.9 (2) | C4C—C5C—C6C—N3C | 178.3 (3) |
| C1A—N1A—C5A—C6A | 56.5 (3) | O1C—C1C—C6C—C5C | -174.3 (3) |
| C4A—N1A—C5A—C12A | -56.7 (3) | C2C—C1C—C6C—C5C | 2.0 (4) |
| C1A—N1A—C5A—C12A | -178.1 (2) | O1C—C1C—C6C—N3C | 4.5 (4) |
| C12A—C5A—C6A—C7A | 82.4 (3) | C2C—C1C—C6C—N3C | -179.2 (2) |
| N1A—C5A—C6A—C7A | -152.8 (2) | O7C—N3C—C6C—C5C | -164.6 (3) |
| C12A—C5A—C6A—C11A | -93.1 (3) | O6C—N3C—C6C—C5C | 14.7 (4) |
| N1A—C5A—C6A—C11A | 31.6 (4) | O7C—N3C—C6C—C1C | 16.5 (4) |
| C11A—C6A—C7A—C8A | 2.4 (4) | O6C—N3C—C6C—C1C | -164.2 (3) |
| C5A—C6A—C7A—C8A | -173.4 (3) | O1D—C1D—C2D—C3D | 176.3 (3) |
| C6A—C7A—C8A—C9A | -1.0 (4) | C6D—C1D—C2D—C3D | -3.6 (4) |
| C7A—C8A—C9A—C10A | -1.8 (4) | O1D—C1D—C2D—N1D | -3.5 (4) |
| C7A—C8A—C9A—C11A | 176.2 (2) | C6D—C1D—C2D—N1D | 176.7 (3) |
| C8A—C9A—C10A—C11A | 3.2 (5) | O3D—N1D—C2D—C3D | -34.6 (4) |
| C11A—C9A—C10A—C11A | -174.8 (2) | O2D—N1D—C2D—C3D | 140.2 (3) |
| C9A—C10A—C11A—C6A | -1.7 (4) | O3D—N1D—C2D—C1D | 145.1 (3) |
| C7A—C6A—C11A—C10A | -1.0 (4) | O2D—N1D—C2D—C1D | -40.0 (4) |
| C5A—C6A—C11A—C10A | 174.4 (3) | C1D—C2D—C3D—C4D | 4.6 (4) |
| C6A—C5A—C12A—C17A | 54.8 (3) | N1D—C2D—C3D—C4D | -175.7 (3) |
| N1A—C5A—C12A—C17A | -72.3 (3) | C2D—C3D—C4D—C5D | -1.7 (4) |
| C6A—C5A—C12A—C13A | -123.5 (3) | C2D—C3D—C4D—N2D | 179.5 (2) |
| N1A—C5A—C12A—C13A | 109.4 (3) | O4D—N2D—C4D—C3D | 3.0 (4) |
| C17A—C12A—C13A—C14A | 0.3 (5) | O5D—N2D—C4D—C3D | -177.0 (3) |
| C5A—C12A—C13A—C14A | 178.6 (3) | O4D—N2D—C4D—C5D | -175.8 (2) |
| C12A—C13A—C14A—C15A | -1.1 (5) | O5D—N2D—C4D—C5D | 4.3 (4) |
| C13A—C14A—C15A—C16A | 1.3 (6) | C3D—C4D—C5D—C6D | -1.7 (4) |
| C14A—C15A—C16A—C17A | -0.5 (5) | N2D—C4D—C5D—C6D | 177.0 (2) |
| C15A—C16A—C17A—C12A | -0.3 (5) | C4D—C5D—C6D—C1D | 2.6 (4) |
| C13A—C12A—C17A—C16A | 0.5 (4) | C4D—C5D—C6D—N3D | -179.1 (2) |
| C5A—C12A—C17A—C16A | -177.8 (2) | O1D—C1D—C6D—C5D | -179.9 (2) |
| C3A—N2A—C18A—C19A | -72.7 (3) | C2D—C1D—C6D—C5D | -0.1 (4) |
| C2A—N2A—C18A—C19A | 167.0 (2) | O1D—C1D—C6D—N3D | 1.8 (4) |
| C20A—O1A—C19A—C18A | 159.0 (3) | C2D—C1D—C6D—N3D | -178.4 (2) |
| N2A—C18A—C19A—O1A | -60.1 (3) | O7D—N3D—C6D—C5D | 154.8 (3) |
| C19A—O1A—C20A—C21A | -82.5 (4) | O6D—N3D—C6D—C5D | -23.9 (4) |

| | | | |
|---------------------|------------|-----------------|------------|
| O1A—C20A—C21A—O3A | 19.3 (5) | O7D—N3D—C6D—C1D | -26.8 (4) |
| O1A—C20A—C21A—O2A | -161.1 (3) | O6D—N3D—C6D—C1D | 154.5 (3) |
| C4B—N1B—C1B—C2B | 50.0 (3) | O2E—N1E—C2E—C3E | -140.4 (3) |
| C5B—N1B—C1B—C2B | 171.3 (2) | O3E—N1E—C2E—C3E | 37.7 (4) |
| C3B—N2B—C2B—C1B | 62.5 (3) | O2E—N1E—C2E—C1E | 39.0 (4) |
| C18B—N2B—C2B—C1B | -174.9 (2) | O3E—N1E—C2E—C1E | -143.0 (3) |
| N1B—C1B—C2B—N2B | -57.8 (3) | O1E—C1E—C2E—C3E | -175.0 (3) |
| C2B—N2B—C3B—C4B | -63.3 (3) | C6E—C1E—C2E—C3E | 3.3 (4) |
| C18B—N2B—C3B—C4B | 173.0 (2) | O1E—C1E—C2E—N1E | 5.7 (4) |
| C1B—N1B—C4B—C3B | -49.4 (3) | C6E—C1E—C2E—N1E | -175.9 (3) |
| C5B—N1B—C4B—C3B | -169.4 (2) | N1E—C2E—C3E—C4E | 176.0 (3) |
| N2B—C3B—C4B—N1B | 58.2 (3) | C1E—C2E—C3E—C4E | -3.3 (4) |
| C4B—N1B—C5B—C12B | -58.4 (3) | C2E—C3E—C4E—C5E | 0.8 (4) |
| C1B—N1B—C5B—C12B | -179.4 (2) | C2E—C3E—C4E—N2E | 179.8 (2) |
| C4B—N1B—C5B—C6B | 174.6 (2) | O5E—N2E—C4E—C3E | -176.0 (3) |
| C1B—N1B—C5B—C6B | 53.6 (3) | O4E—N2E—C4E—C3E | 2.6 (4) |
| C12B—C5B—C6B—C7B | -58.2 (3) | O5E—N2E—C4E—C5E | 3.0 (4) |
| N1B—C5B—C6B—C7B | 70.2 (3) | O4E—N2E—C4E—C5E | -178.3 (3) |
| C12B—C5B—C6B—C11B | 123.4 (3) | C3E—C4E—C5E—C6E | 1.3 (4) |
| N1B—C5B—C6B—C11B | -108.2 (3) | N2E—C4E—C5E—C6E | -177.7 (2) |
| C11B—C6B—C7B—C8B | 3.0 (5) | C4E—C5E—C6E—C1E | -1.1 (4) |
| C5B—C6B—C7B—C8B | -175.3 (3) | C4E—C5E—C6E—N3E | -179.7 (2) |
| C6B—C7B—C8B—C9B | -0.7 (5) | O1E—C1E—C6E—C5E | 177.2 (3) |
| C7B—C8B—C9B—C10B | -1.9 (5) | C2E—C1E—C6E—C5E | -1.1 (4) |
| C7B—C8B—C9B—C11B | 176.1 (3) | O1E—C1E—C6E—N3E | -4.2 (4) |
| C8B—C9B—C10B—C11B | 2.2 (6) | C2E—C1E—C6E—N3E | 177.5 (2) |
| C11B—C9B—C10B—C11B | -175.9 (3) | O7E—N3E—C6E—C5E | -157.0 (3) |
| C9B—C10B—C11B—C6B | 0.2 (6) | O6E—N3E—C6E—C5E | 23.5 (4) |
| C7B—C6B—C11B—C10B | -2.8 (5) | O7E—N3E—C6E—C1E | 24.3 (4) |
| C5B—C6B—C11B—C10B | 175.6 (3) | O6E—N3E—C6E—C1E | -155.2 (3) |
| C6B—C5B—C12B—C17B | -75.6 (3) | O1F—C1F—C2F—C3F | -173.2 (3) |
| N1B—C5B—C12B—C17B | 159.0 (3) | C6F—C1F—C2F—C3F | 5.0 (4) |
| C6B—C5B—C12B—C13B | 103.1 (3) | O1F—C1F—C2F—N1F | 5.8 (4) |
| N1B—C5B—C12B—C13B | -22.3 (4) | C6F—C1F—C2F—N1F | -175.9 (2) |
| C17B—C12B—C13B—C14B | -0.8 (5) | O3F—N1F—C2F—C3F | 57.0 (4) |
| C5B—C12B—C13B—C14B | -179.5 (3) | O2F—N1F—C2F—C3F | -123.1 (3) |
| C12B—C13B—C14B—C15B | 2.8 (5) | O3F—N1F—C2F—C1F | -122.1 (3) |
| C13B—C14B—C15B—C16B | -2.6 (5) | O2F—N1F—C2F—C1F | 57.7 (4) |
| C14B—C15B—C16B—C17B | 0.4 (6) | C1F—C2F—C3F—C4F | -3.7 (4) |
| C15B—C16B—C17B—C12B | 1.6 (6) | N1F—C2F—C3F—C4F | 177.3 (3) |
| C13B—C12B—C17B—C16B | -1.4 (5) | C2F—C3F—C4F—C5F | 0.8 (4) |
| C5B—C12B—C17B—C16B | 177.4 (3) | C2F—C3F—C4F—N2F | -178.2 (2) |
| C3B—N2B—C18B—C19B | -166.4 (3) | O4F—N2F—C4F—C5F | -175.7 (3) |
| C2B—N2B—C18B—C19B | 73.6 (3) | O5F—N2F—C4F—C5F | 4.9 (4) |
| C20B—O1B—C19B—C18B | -158.3 (3) | O4F—N2F—C4F—C3F | 3.4 (4) |
| N2B—C18B—C19B—O1B | 61.1 (4) | O5F—N2F—C4F—C3F | -176.1 (3) |
| C19B—O1B—C20B—C21B | 84.1 (4) | C3F—C4F—C5F—C6F | 0.1 (4) |
| O1B—C20B—C21B—O2B | -14.5 (6) | N2F—C4F—C5F—C6F | 179.1 (2) |

| | | | |
|-------------------|------------|-----------------|------------|
| O1B—C20B—C21B—O3B | 165.5 (4) | C4F—C5F—C6F—C1F | 1.8 (4) |
| O1C—C1C—C2C—C3C | 175.9 (3) | C4F—C5F—C6F—N3F | -177.8 (3) |
| C6C—C1C—C2C—C3C | -0.7 (4) | O1F—C1F—C6F—C5F | 174.2 (3) |
| O1C—C1C—C2C—N1C | -0.6 (4) | C2F—C1F—C6F—C5F | -4.0 (4) |
| C6C—C1C—C2C—N1C | -177.2 (3) | O1F—C1F—C6F—N3F | -6.2 (4) |
| O2C—N1C—C2C—C3C | 127.6 (3) | C2F—C1F—C6F—N3F | 175.6 (2) |
| O3C—N1C—C2C—C3C | -51.8 (4) | O6F—N3F—C6F—C5F | -10.1 (4) |
| O2C—N1C—C2C—C1C | -55.6 (4) | O7F—N3F—C6F—C5F | 169.0 (3) |
| O3C—N1C—C2C—C1C | 124.9 (3) | O6F—N3F—C6F—C1F | 170.3 (3) |
| C1C—C2C—C3C—C4C | 0.2 (5) | O7F—N3F—C6F—C1F | -10.6 (4) |

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

Cg5 is the centroid of the C6B—C11B ring.

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-------------------------------------|-------|-------------|-------------|---------------|
| O2A—H2AD \cdots O2B | 0.84 | 1.80 | 2.638 (4) | 180 |
| N1A—H1AC \cdots O1D | 0.93 | 1.83 | 2.682 (3) | 152 |
| N1A—H1AC \cdots O7D | 0.93 | 2.63 | 3.301 (3) | 129 |
| N2A—H2AC \cdots O1C | 0.93 | 1.89 | 2.765 (3) | 155 |
| N2A—H2AC \cdots O7C | 0.93 | 2.46 | 2.990 (3) | 116 |
| O3B—H3BC \cdots O3A | 0.84 | 1.76 | 2.601 (4) | 180 |
| N1B—H1BC \cdots O1E | 0.93 | 1.85 | 2.678 (3) | 147 |
| N1B—H1BC \cdots O7E | 0.93 | 2.52 | 3.193 (3) | 130 |
| N2B—H2BC \cdots O1F | 0.93 | 1.91 | 2.764 (3) | 153 |
| N2B—H2BC \cdots O7F | 0.93 | 2.57 | 3.078 (4) | 115 |
| C19B—H19C \cdots Cg5 ⁱ | 0.99 | 2.95 | 3.792 (4) | 144 |

Symmetry code: (i) $x+1, y, z$.