

Crystal structure of 11-(2,3-dimethoxyphenyl)-14-methyl-12-oxa-8,14-diazatetracyclo[8.3.3.0^{1,10}.0^{2,7}]hexadeca-2(7),3,5-triene-9,13-dione

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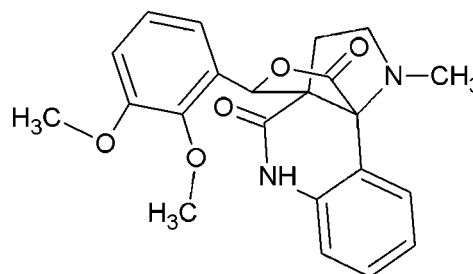
The title compound, C₂₂H₂₂N₂O₅, contains two conformationally similar molecules (*A* and *B*) in its the asymmetric unit (r.m.s. overlay fit for the 29 non-H atoms = 0.194 Å). In each molecule, the lactone ring has an envelope conformation with the spiro C atom as the flap. In the crystal, *A*+*A* and *B*+*B* inversion dimers linked by pairs of N—H···O hydrogen bonds occur; in both cases, R₂²(8) loops are generated. A weak C—H···O interaction is also observed, which links the dimers into [010] chains.

Keywords: crystal structure; diazatetracyclohexadecatrienedione; N—H···O hydrogen bonds.

CCDC reference: 1056691

1. Related literature

For general background and the biological and pharmacological properties of quinoline derivatives, see: Michael (1997). For a related structure, see: Vennila *et al.* (2011).



2. Experimental

2.1. Crystal data

C₂₂H₂₂N₂O₅
M_r = 394.42
 Triclinic, *P* $\bar{1}$
a = 10.1360 (4) Å
b = 10.3198 (4) Å
c = 18.8973 (7) Å
 α = 89.079 (2)°
 β = 74.955 (2)°
 γ = 89.406 (2)°
V = 1908.64 (13) Å³
Z = 4
 Mo *K* α radiation
 μ = 0.10 mm⁻¹
T = 293 K
 0.35 × 0.30 × 0.30 mm

2.2. Data collection

Bruker Kappa APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2004)
T_{min} = 0.967, *T_{max}* = 0.971
 35145 measured reflections
 6717 independent reflections
 5408 reflections with *I* > 2 σ (*I*)
R_{int} = 0.028

2.3. Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.047
 $wR(F^2)$ = 0.128
S = 1.02
 6717 reflections
 532 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max}$ = 0.65 e Å⁻³
 $\Delta\rho_{\min}$ = -0.54 e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| <i>D</i> —H··· <i>A</i> | <i>D</i> —H | H··· <i>A</i> | <i>D</i> ··· <i>A</i> | <i>D</i> —H··· <i>A</i> |
|---------------------------------|-------------|---------------|-----------------------|-------------------------|
| N8—H8···O9 ⁱ | 0.89 (3) | 2.01 (3) | 2.903 (2) | 177 (2) |
| N8A—H8A···O9A ⁱⁱ | 0.89 (2) | 2.07 (3) | 2.958 (2) | 175 (2) |
| C6A'—H6A'···O13A ⁱⁱⁱ | 0.93 | 2.43 | 3.322 (3) | 161 |

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT (Bruker, 2004); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

Acknowledgements

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7391).

References

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Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.
Vennila, K. N., Sankaran, M., Mohan, P. S. & Velmurugan, D. (2011). *Acta Cryst.* **E67**, o3376–o3377.

supporting information

Acta Cryst. (2015). E71, o293–o294 [doi:10.1107/S2056989015006386]

Crystal structure of 11-(2,3-dimethoxyphenyl)-14-methyl-12-oxa-8,14-diazatetracyclo[8.3.3.0^{1,10}.0^{2,7}]hexadeca-2(7),3,5-triene-9,13-dione

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S1. Comment

Quinolines exhibit physico-chemical activities which are useful in the field of pharmaceuticals and agrochemicals. Their derivatives are also present in a wide variety of natural products involved in several biological activities (Michael, 2006). The crystal structure of the title compound is presented here as a part of our on-going structural studies on quinoline derivatives.

The molecular structure of molecule (A) and molecule (B) is shown in Fig.1. The furan ring (O12A/C11A-C10A/C1A/C13A) of (A) exhibits an envelope conformation with C10A as the flap atom. The furan ring (O12/C11-C10/C1/C13) of (B) exhibits an envelope conformation with C10 as the flap atom. The quinoline ring (N8A/C1A-C10A) is almost coplanar showing a dihedral angle of 2.9 (8)° with the pyridine ring (N8A/C7A/C2A-C1A/C9A-C10A) of molecule (A) and is perpendicular with the pyridine ring (N8/C7/C2-C1/C9-C10) of molecule (B) inclined at an angle of 88.3 (8)°. The sum of angles at N8, N8A of the quinoline rings (360°) is in accordance with sp² hybridization.

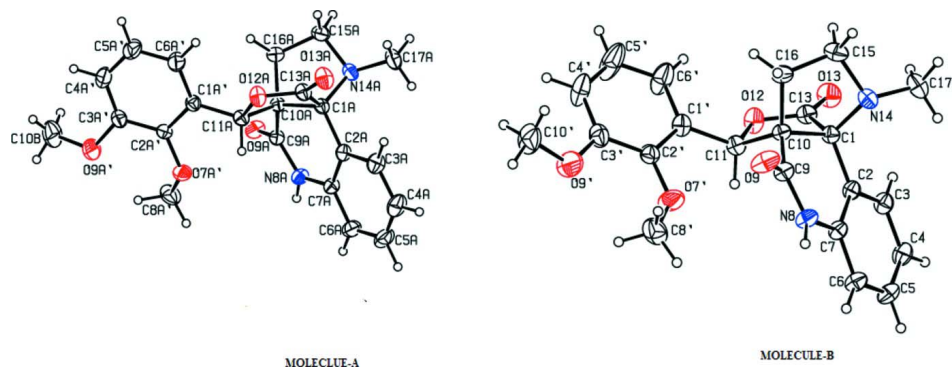
In the crystal of two independent molecules, hydrogen-bonded chains running along bc plane are generated by connecting neighbouring molecules via N-H...O, C-H...O hydrogen bonds forming a two dimensional structure (Fig.2.) The hydrogen bonds of N8-H8...O9 and N8A-H8A...O9A forming inversion dimers but enclosing smaller R₂²(8) loops and the hydrogen bond of C6A'-H6A'...O13A forms a one dimensional chain along [010] as shown in Fig.3.

S2. Experimental

A mixture of methyl 2-(hydroxy(m-tolyl)methyl)acrylate (252mgs, 1 mmol), isatin (161.7mgs, 1.1 mmol) and sarcosine (97.9mgs, 1.1 mmol) was placed in a round bottom flask and melted at 180°C until completion of the reaction was evidenced by TLC analysis. After completion of the reaction, the crude product was washed with 5ml of ethylacetate and hexane mixture (1:4 ratio) which successfully provided the pure product as colorless solid. The product was dissolved in ethyl acetate and heated for two minutes. The resulting solution was subjected to crystallization by slow evaporation of the solvent for 48 hours resulting in the formation of colourless blocks.

S3. Refinement

All H atoms were fixed geometrically and allowed to ride on their parent C atoms, with C-H distances fixed in the range 0.93-0.98 Å with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H $1.2U_{\text{eq}}(\text{C})$ for other H atoms.

**Figure 1**

The molecular structure of the title compound with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are omitted for clarity.

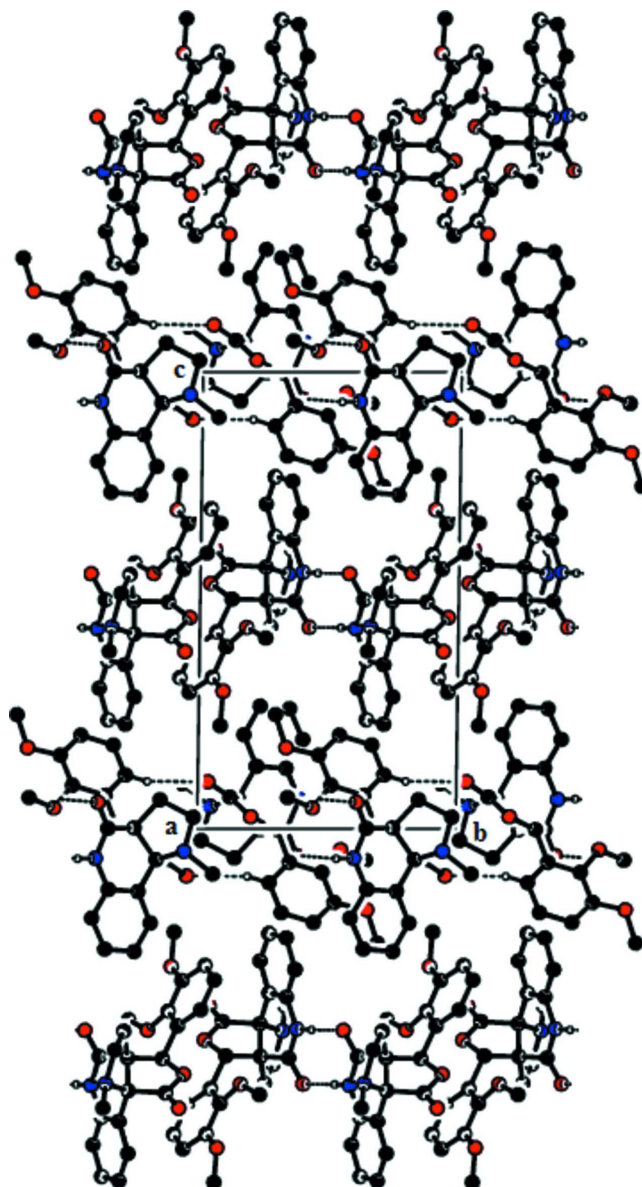
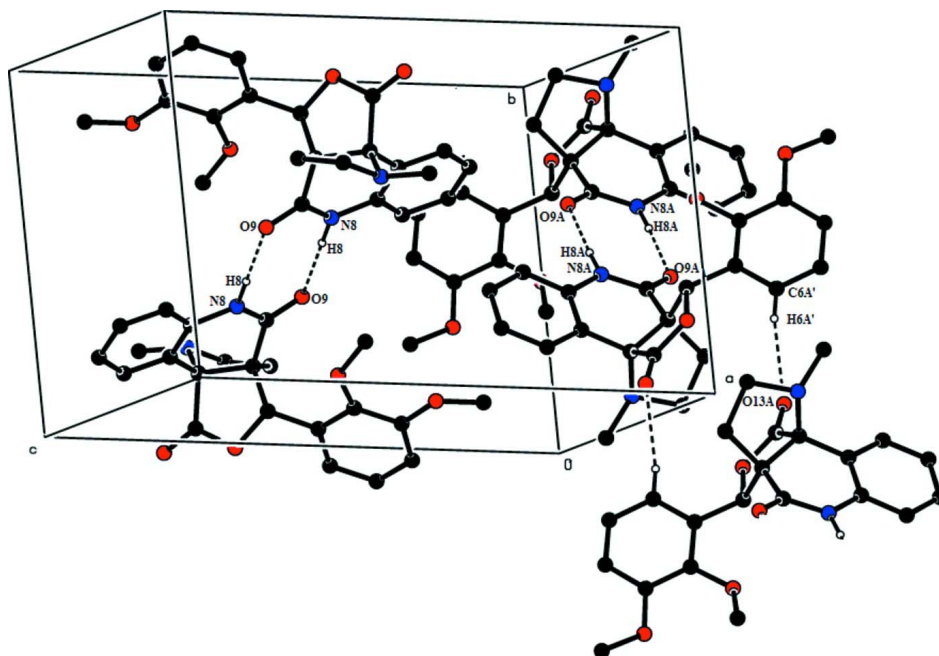


Figure 2

The molecular packing is viewed along the *a* axis. Dashed lines show the intermolecular N—H···O and C—H···O hydrogen bonds. H atoms not involved in hydrogen bonding have been omitted for clarity.

**Figure 3**

A partial view of the N8—H8...O9, N8A—H8A...O9A and C6A'—H6A'...O13A hydrogen-bonding interactions along the *c* axis.

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Crystal data

C₂₂H₂₂N₂O₅

M_r = 394.42

Triclinic, *P*1̄

Hall symbol: -P 1

a = 10.1360 (4) Å

b = 10.3198 (4) Å

c = 18.8973 (7) Å

α = 89.079 (2)°

β = 74.955 (2)°

γ = 89.406 (2)°

V = 1908.64 (13) Å³

Z = 4

F(000) = 832

D_x = 1.373 Mg m⁻³

Mo *K*α radiation, λ = 0.71073 Å

Cell parameters from 6717 reflections

θ = 1.1–25.0°

μ = 0.10 mm⁻¹

T = 293 K

Block, colourless

0.35 × 0.30 × 0.30 mm

Data collection

Bruker Kappa APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and φ scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2004)

T_{min} = 0.967, *T_{max}* = 0.971

35145 measured reflections

6717 independent reflections

5408 reflections with *I* > 2σ(*I*)

R_{int} = 0.028

θ_{max} = 25.0°, θ_{min} = 1.1°

h = -12→12

k = -12→12

l = -22→22

*Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.047$ $wR(F^2) = 0.128$ $S = 1.02$

6717 reflections

532 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sitesH atoms treated by a mixture of independent
and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0555P)^2 + 1.1105P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.65 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.54 \text{ e } \text{\AA}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0073 (9)

Special details

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

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > 2\sigma(F^2)$ 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)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|---------------|----------------------------------|
| C10B | 1.0615 (3) | 0.7030 (3) | -0.25020 (16) | 0.0810 (9) |
| H10A | 1.0438 | 0.7943 | -0.2528 | 0.121* |
| H10B | 1.0127 | 0.6574 | -0.2792 | 0.121* |
| H10C | 1.1577 | 0.6868 | -0.2686 | 0.121* |
| H8A | 0.517 (2) | 0.461 (2) | 0.0654 (12) | 0.049 (6)* |
| H8 | 0.057 (3) | 0.546 (2) | 0.4402 (13) | 0.061 (7)* |
| C1A | 0.71903 (18) | 0.15796 (17) | 0.06757 (10) | 0.0323 (4) |
| O12A | 0.95534 (13) | 0.20658 (13) | 0.02653 (8) | 0.0433 (4) |
| O9A | 0.61164 (15) | 0.38520 (14) | -0.05792 (7) | 0.0448 (4) |
| N8A | 0.57191 (17) | 0.39272 (17) | 0.06429 (9) | 0.0379 (4) |
| C9A | 0.63693 (18) | 0.34543 (18) | -0.00142 (10) | 0.0324 (4) |
| O13A | 0.90527 (15) | 0.03882 (14) | 0.10304 (9) | 0.0530 (4) |
| O7A' | 0.86006 (16) | 0.55593 (14) | -0.04580 (8) | 0.0513 (4) |
| C11A | 0.88282 (18) | 0.30883 (18) | -0.00169 (10) | 0.0337 (4) |
| H11A | 0.8642 | 0.3801 | 0.0333 | 0.040* |
| C7A | 0.5767 (2) | 0.34058 (19) | 0.13261 (10) | 0.0367 (4) |
| C10A | 0.74578 (18) | 0.24380 (17) | -0.00209 (10) | 0.0301 (4) |
| N14A | 0.64191 (16) | 0.04987 (15) | 0.04930 (9) | 0.0389 (4) |
| C1A' | 0.97041 (19) | 0.35816 (18) | -0.07344 (11) | 0.0363 (4) |
| C2A | 0.64674 (19) | 0.22556 (19) | 0.13708 (10) | 0.0355 (4) |
| C16A | 0.7560 (2) | 0.14502 (19) | -0.06377 (11) | 0.0400 (5) |
| H16A | 0.6961 | 0.1700 | -0.0943 | 0.048* |
| H16B | 0.8489 | 0.1397 | -0.0943 | 0.048* |

| | | | | |
|------|--------------|--------------|---------------|------------|
| C13A | 0.8671 (2) | 0.12459 (19) | 0.06982 (11) | 0.0378 (5) |
| C2A' | 0.9520 (2) | 0.48460 (18) | -0.09543 (11) | 0.0362 (4) |
| C3A' | 1.0346 (2) | 0.5333 (2) | -0.16135 (12) | 0.0457 (5) |
| C3A | 0.6440 (2) | 0.1761 (2) | 0.20631 (12) | 0.0504 (6) |
| H3A | 0.6921 | 0.1004 | 0.2105 | 0.060* |
| C15A | 0.7124 (2) | 0.0159 (2) | -0.02557 (12) | 0.0449 (5) |
| H15A | 0.6518 | -0.0293 | -0.0489 | 0.054* |
| H15B | 0.7912 | -0.0388 | -0.0265 | 0.054* |
| C6A | 0.5070 (2) | 0.4038 (2) | 0.19557 (12) | 0.0512 (6) |
| H6A | 0.4620 | 0.4818 | 0.1921 | 0.061* |
| O9A' | 1.0186 (2) | 0.66034 (16) | -0.17763 (10) | 0.0789 (6) |
| C6A' | 1.0697 (2) | 0.2811 (2) | -0.11824 (14) | 0.0544 (6) |
| H6A' | 1.0829 | 0.1963 | -0.1039 | 0.065* |
| C4A' | 1.1314 (3) | 0.4546 (2) | -0.20506 (14) | 0.0599 (6) |
| H4A' | 1.1855 | 0.4864 | -0.2493 | 0.072* |
| C4A | 0.5714 (3) | 0.2371 (3) | 0.26883 (13) | 0.0615 (7) |
| H4A | 0.5680 | 0.2010 | 0.3147 | 0.074* |
| C17A | 0.6138 (2) | -0.0621 (2) | 0.09891 (14) | 0.0548 (6) |
| H17A | 0.5677 | -0.0341 | 0.1473 | 0.082* |
| H17B | 0.6983 | -0.1040 | 0.1000 | 0.082* |
| H17C | 0.5571 | -0.1219 | 0.0821 | 0.082* |
| C5A | 0.5043 (3) | 0.3513 (3) | 0.26324 (13) | 0.0610 (7) |
| H5A | 0.4568 | 0.3934 | 0.3054 | 0.073* |
| C8A' | 0.7778 (3) | 0.6521 (2) | -0.06863 (16) | 0.0621 (7) |
| H8A1 | 0.7195 | 0.6925 | -0.0264 | 0.093* |
| H8A2 | 0.7228 | 0.6124 | -0.0966 | 0.093* |
| H8A3 | 0.8354 | 0.7163 | -0.0984 | 0.093* |
| C5A' | 1.1486 (3) | 0.3291 (3) | -0.18348 (15) | 0.0672 (7) |
| H5A' | 1.2143 | 0.2764 | -0.2133 | 0.081* |
| N8 | 0.12270 (18) | 0.60382 (17) | 0.43819 (9) | 0.0413 (4) |
| O12 | 0.29646 (15) | 0.96066 (13) | 0.46254 (8) | 0.0480 (4) |
| C1 | 0.36492 (18) | 0.74700 (18) | 0.42665 (10) | 0.0334 (4) |
| C2 | 0.31784 (19) | 0.71338 (18) | 0.35954 (10) | 0.0348 (4) |
| C7 | 0.1971 (2) | 0.64514 (18) | 0.36828 (10) | 0.0360 (4) |
| O9 | 0.09671 (17) | 0.57648 (16) | 0.55871 (8) | 0.0579 (5) |
| O13 | 0.50407 (15) | 0.94355 (15) | 0.38830 (9) | 0.0551 (4) |
| C10 | 0.25101 (19) | 0.74324 (18) | 0.49834 (10) | 0.0322 (4) |
| C11 | 0.1820 (2) | 0.87718 (18) | 0.49699 (11) | 0.0374 (4) |
| H11 | 0.1197 | 0.8733 | 0.4652 | 0.045* |
| N14 | 0.46780 (16) | 0.65966 (17) | 0.44204 (10) | 0.0423 (4) |
| C9 | 0.1510 (2) | 0.63346 (19) | 0.50162 (10) | 0.0372 (4) |
| C13 | 0.4009 (2) | 0.89217 (19) | 0.42184 (11) | 0.0388 (5) |
| C1' | 0.1075 (2) | 0.93454 (19) | 0.56880 (12) | 0.0426 (5) |
| C3 | 0.3898 (2) | 0.7494 (2) | 0.28913 (11) | 0.0426 (5) |
| H3 | 0.4697 | 0.7969 | 0.2823 | 0.051* |
| C4 | 0.3449 (2) | 0.7160 (2) | 0.22934 (11) | 0.0471 (5) |
| H4 | 0.3957 | 0.7387 | 0.1824 | 0.057* |
| C2' | -0.0258 (2) | 0.8978 (2) | 0.60066 (11) | 0.0432 (5) |

| | | | | |
|------|---------------|------------|--------------|-------------|
| C16 | 0.3306 (2) | 0.7269 (2) | 0.55711 (11) | 0.0437 (5) |
| H16C | 0.3282 | 0.8066 | 0.5840 | 0.052* |
| H16D | 0.2918 | 0.6580 | 0.5914 | 0.052* |
| C6 | 0.1502 (2) | 0.6138 (2) | 0.30803 (11) | 0.0461 (5) |
| H6 | 0.0687 | 0.5691 | 0.3145 | 0.055* |
| C5 | 0.2246 (2) | 0.6489 (2) | 0.23883 (11) | 0.0491 (5) |
| H5 | 0.1937 | 0.6273 | 0.1983 | 0.059* |
| O7' | -0.07771 (17) | 0.8186 (2) | 0.55922 (10) | 0.0873 (7) |
| C15 | 0.4763 (2) | 0.6935 (2) | 0.51562 (12) | 0.0511 (6) |
| H15C | 0.5360 | 0.7671 | 0.5137 | 0.061* |
| H15D | 0.5109 | 0.6209 | 0.5389 | 0.061* |
| C3' | -0.0965 (2) | 0.9489 (2) | 0.66811 (13) | 0.0518 (6) |
| C6' | 0.1686 (3) | 1.0237 (3) | 0.60371 (16) | 0.0825 (10) |
| H6' | 0.2582 | 1.0491 | 0.5828 | 0.099* |
| O9' | -0.22558 (19) | 0.9071 (3) | 0.69852 (11) | 0.0991 (8) |
| C17 | 0.5998 (2) | 0.6501 (2) | 0.38864 (14) | 0.0579 (6) |
| H17D | 0.6568 | 0.5892 | 0.4061 | 0.087* |
| H17E | 0.6426 | 0.7335 | 0.3817 | 0.087* |
| H17F | 0.5871 | 0.6212 | 0.3428 | 0.087* |
| C5' | 0.0973 (4) | 1.0747 (3) | 0.66906 (19) | 0.1108 (15) |
| H5' | 0.1388 | 1.1352 | 0.6920 | 0.133* |
| C4' | -0.0340 (3) | 1.0379 (3) | 0.70101 (16) | 0.0753 (9) |
| H4' | -0.0811 | 1.0736 | 0.7454 | 0.090* |
| C8' | -0.1935 (3) | 0.7448 (3) | 0.58216 (18) | 0.0735 (8) |
| H8'1 | -0.2080 | 0.6977 | 0.5415 | 0.110* |
| H8'2 | -0.2707 | 0.8003 | 0.6013 | 0.110* |
| H8'3 | -0.1827 | 0.6849 | 0.6197 | 0.110* |
| C10' | -0.2794 (3) | 0.9163 (3) | 0.77410 (16) | 0.0827 (9) |
| H10D | -0.3707 | 0.8829 | 0.7874 | 0.124* |
| H10E | -0.2809 | 1.0055 | 0.7881 | 0.124* |
| H10F | -0.2236 | 0.8670 | 0.7988 | 0.124* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|-------------|
| C10B | 0.101 (2) | 0.0716 (19) | 0.0727 (19) | -0.0136 (17) | -0.0279 (17) | 0.0330 (15) |
| C1A | 0.0312 (10) | 0.0281 (9) | 0.0380 (10) | 0.0034 (7) | -0.0103 (8) | 0.0037 (8) |
| O12A | 0.0312 (7) | 0.0457 (8) | 0.0549 (9) | 0.0008 (6) | -0.0157 (6) | 0.0150 (7) |
| O9A | 0.0504 (9) | 0.0500 (9) | 0.0365 (8) | 0.0156 (7) | -0.0163 (7) | 0.0022 (6) |
| N8A | 0.0395 (9) | 0.0367 (9) | 0.0354 (9) | 0.0147 (8) | -0.0066 (7) | 0.0016 (7) |
| C9A | 0.0305 (9) | 0.0323 (10) | 0.0344 (10) | 0.0026 (8) | -0.0086 (8) | 0.0022 (8) |
| O13A | 0.0475 (9) | 0.0455 (9) | 0.0692 (11) | 0.0064 (7) | -0.0221 (8) | 0.0192 (8) |
| O7A' | 0.0623 (10) | 0.0355 (8) | 0.0500 (9) | 0.0118 (7) | -0.0041 (8) | -0.0015 (7) |
| C11A | 0.0325 (10) | 0.0314 (10) | 0.0390 (11) | 0.0029 (8) | -0.0126 (8) | 0.0021 (8) |
| C7A | 0.0365 (10) | 0.0396 (11) | 0.0332 (10) | 0.0015 (8) | -0.0074 (8) | -0.0003 (8) |
| C10A | 0.0294 (9) | 0.0294 (9) | 0.0322 (10) | 0.0035 (7) | -0.0095 (8) | 0.0001 (7) |
| N14A | 0.0348 (9) | 0.0306 (8) | 0.0509 (10) | -0.0021 (7) | -0.0107 (8) | 0.0028 (7) |
| C1A' | 0.0311 (10) | 0.0357 (10) | 0.0417 (11) | -0.0013 (8) | -0.0089 (8) | 0.0011 (8) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C2A | 0.0347 (10) | 0.0389 (11) | 0.0337 (10) | -0.0007 (8) | -0.0106 (8) | 0.0039 (8) |
| C16A | 0.0434 (11) | 0.0382 (11) | 0.0385 (11) | 0.0031 (9) | -0.0106 (9) | -0.0069 (9) |
| C13A | 0.0367 (10) | 0.0343 (10) | 0.0442 (11) | 0.0033 (8) | -0.0141 (9) | 0.0051 (9) |
| C2A' | 0.0384 (10) | 0.0316 (10) | 0.0383 (11) | -0.0006 (8) | -0.0090 (9) | -0.0039 (8) |
| C3A' | 0.0532 (13) | 0.0374 (11) | 0.0450 (12) | -0.0072 (10) | -0.0099 (10) | 0.0053 (9) |
| C3A | 0.0544 (13) | 0.0573 (14) | 0.0417 (12) | 0.0022 (11) | -0.0171 (11) | 0.0096 (10) |
| C15A | 0.0451 (12) | 0.0357 (11) | 0.0536 (13) | 0.0020 (9) | -0.0117 (10) | -0.0092 (9) |
| C6A | 0.0553 (14) | 0.0508 (13) | 0.0428 (13) | 0.0082 (11) | -0.0045 (10) | -0.0073 (10) |
| O9A' | 0.1098 (16) | 0.0449 (10) | 0.0646 (12) | 0.0027 (10) | 0.0074 (11) | 0.0184 (9) |
| C6A' | 0.0436 (12) | 0.0444 (13) | 0.0654 (15) | 0.0104 (10) | 0.0025 (11) | 0.0081 (11) |
| C4A' | 0.0549 (14) | 0.0604 (15) | 0.0526 (14) | -0.0031 (12) | 0.0070 (12) | 0.0084 (12) |
| C4A | 0.0676 (16) | 0.0842 (19) | 0.0338 (12) | -0.0039 (14) | -0.0156 (11) | 0.0072 (12) |
| C17A | 0.0480 (13) | 0.0396 (12) | 0.0730 (16) | -0.0069 (10) | -0.0091 (12) | 0.0135 (11) |
| C5A | 0.0637 (16) | 0.0794 (18) | 0.0354 (12) | -0.0022 (14) | -0.0040 (11) | -0.0119 (12) |
| C8A' | 0.0543 (14) | 0.0403 (13) | 0.090 (2) | 0.0068 (11) | -0.0166 (14) | 0.0033 (12) |
| C5A' | 0.0538 (15) | 0.0607 (16) | 0.0698 (17) | 0.0129 (12) | 0.0144 (13) | 0.0020 (13) |
| N8 | 0.0456 (10) | 0.0458 (10) | 0.0310 (9) | -0.0206 (8) | -0.0067 (7) | 0.0022 (7) |
| O12 | 0.0506 (9) | 0.0326 (7) | 0.0507 (9) | -0.0067 (6) | 0.0046 (7) | 0.0040 (6) |
| C1 | 0.0315 (10) | 0.0352 (10) | 0.0326 (10) | -0.0048 (8) | -0.0066 (8) | 0.0031 (8) |
| C2 | 0.0374 (10) | 0.0339 (10) | 0.0307 (10) | -0.0024 (8) | -0.0046 (8) | 0.0006 (8) |
| C7 | 0.0418 (11) | 0.0346 (10) | 0.0305 (10) | -0.0045 (8) | -0.0071 (8) | 0.0015 (8) |
| O9 | 0.0753 (11) | 0.0630 (10) | 0.0348 (8) | -0.0380 (9) | -0.0126 (8) | 0.0138 (7) |
| O13 | 0.0449 (9) | 0.0522 (9) | 0.0619 (10) | -0.0192 (7) | -0.0023 (8) | 0.0087 (8) |
| C10 | 0.0333 (10) | 0.0345 (10) | 0.0293 (9) | -0.0060 (8) | -0.0088 (8) | 0.0019 (8) |
| C11 | 0.0358 (10) | 0.0372 (11) | 0.0365 (11) | -0.0057 (8) | -0.0049 (8) | 0.0047 (8) |
| N14 | 0.0338 (9) | 0.0466 (10) | 0.0458 (10) | 0.0019 (7) | -0.0091 (8) | 0.0045 (8) |
| C9 | 0.0406 (11) | 0.0379 (11) | 0.0318 (10) | -0.0099 (9) | -0.0070 (9) | 0.0035 (8) |
| C13 | 0.0380 (11) | 0.0403 (11) | 0.0372 (11) | -0.0088 (9) | -0.0081 (9) | 0.0046 (9) |
| C1' | 0.0447 (12) | 0.0347 (11) | 0.0440 (12) | -0.0009 (9) | -0.0041 (9) | 0.0012 (9) |
| C3 | 0.0424 (11) | 0.0434 (12) | 0.0374 (11) | -0.0054 (9) | -0.0021 (9) | 0.0021 (9) |
| C4 | 0.0589 (14) | 0.0477 (12) | 0.0298 (11) | -0.0007 (10) | -0.0028 (10) | 0.0024 (9) |
| C2' | 0.0395 (11) | 0.0517 (12) | 0.0386 (11) | 0.0032 (9) | -0.0106 (9) | -0.0002 (9) |
| C16 | 0.0473 (12) | 0.0494 (12) | 0.0385 (11) | -0.0036 (10) | -0.0184 (10) | 0.0034 (9) |
| C6 | 0.0535 (13) | 0.0474 (12) | 0.0387 (12) | -0.0120 (10) | -0.0137 (10) | -0.0025 (9) |
| C5 | 0.0679 (15) | 0.0490 (13) | 0.0326 (11) | -0.0033 (11) | -0.0167 (10) | -0.0033 (9) |
| O7' | 0.0427 (9) | 0.153 (2) | 0.0628 (12) | -0.0300 (11) | -0.0035 (8) | -0.0395 (12) |
| C15 | 0.0451 (12) | 0.0608 (14) | 0.0531 (14) | -0.0037 (11) | -0.0234 (11) | 0.0089 (11) |
| C3' | 0.0448 (12) | 0.0591 (14) | 0.0462 (13) | 0.0061 (11) | -0.0026 (10) | 0.0012 (11) |
| C6' | 0.0755 (19) | 0.0697 (18) | 0.081 (2) | -0.0368 (15) | 0.0209 (15) | -0.0341 (15) |
| O9' | 0.0486 (11) | 0.178 (2) | 0.0603 (12) | -0.0212 (13) | 0.0080 (9) | -0.0357 (14) |
| C17 | 0.0378 (12) | 0.0619 (15) | 0.0693 (16) | 0.0054 (11) | -0.0058 (11) | 0.0029 (12) |
| C5' | 0.113 (3) | 0.090 (2) | 0.098 (2) | -0.056 (2) | 0.034 (2) | -0.056 (2) |
| C4' | 0.087 (2) | 0.0551 (16) | 0.0642 (17) | -0.0073 (14) | 0.0179 (15) | -0.0229 (13) |
| C8' | 0.0470 (14) | 0.0657 (17) | 0.104 (2) | -0.0104 (12) | -0.0114 (14) | -0.0082 (16) |
| C10' | 0.0645 (18) | 0.103 (2) | 0.0633 (18) | -0.0090 (16) | 0.0159 (14) | -0.0132 (16) |

Geometric parameters (Å, °)

| | | | |
|-----------|-----------|----------|-----------|
| C10B—O9A' | 1.392 (3) | N8—C9 | 1.344 (2) |
| C10B—H10A | 0.9600 | N8—C7 | 1.403 (2) |
| C10B—H10B | 0.9600 | N8—H8 | 0.89 (3) |
| C10B—H10C | 0.9600 | O12—C13 | 1.338 (2) |
| C1A—N14A | 1.465 (2) | O12—C11 | 1.457 (2) |
| C1A—C2A | 1.506 (3) | C1—N14 | 1.454 (2) |
| C1A—C10A | 1.541 (2) | C1—C2 | 1.513 (3) |
| C1A—C13A | 1.548 (3) | C1—C10 | 1.535 (3) |
| O12A—C13A | 1.341 (2) | C1—C13 | 1.541 (3) |
| O12A—C11A | 1.452 (2) | C2—C3 | 1.388 (3) |
| O9A—C9A | 1.225 (2) | C2—C7 | 1.389 (3) |
| N8A—C9A | 1.343 (2) | C7—C6 | 1.387 (3) |
| N8A—C7A | 1.403 (2) | O9—C9 | 1.221 (2) |
| N8A—H8A | 0.89 (2) | O13—C13 | 1.197 (2) |
| C9A—C10A | 1.512 (2) | C10—C9 | 1.518 (2) |
| O13A—C13A | 1.195 (2) | C10—C16 | 1.539 (3) |
| O7A'—C2A' | 1.358 (2) | C10—C11 | 1.546 (3) |
| O7A'—C8A' | 1.422 (3) | C11—C1' | 1.499 (3) |
| C11A—C1A' | 1.499 (3) | C11—H11 | 0.9800 |
| C11A—C10A | 1.551 (2) | N14—C17 | 1.455 (3) |
| C11A—H11A | 0.9800 | N14—C15 | 1.463 (3) |
| C7A—C6A | 1.386 (3) | C1'—C6' | 1.381 (3) |
| C7A—C2A | 1.390 (3) | C1'—C2' | 1.383 (3) |
| C10A—C16A | 1.543 (3) | C3—C4 | 1.373 (3) |
| N14A—C15A | 1.457 (3) | C3—H3 | 0.9300 |
| N14A—C17A | 1.459 (3) | C4—C5 | 1.378 (3) |
| C1A'—C6A' | 1.388 (3) | C4—H4 | 0.9300 |
| C1A'—C2A' | 1.388 (3) | C2'—O7' | 1.343 (3) |
| C2A—C3A | 1.390 (3) | C2'—C3' | 1.398 (3) |
| C16A—C15A | 1.519 (3) | C16—C15 | 1.521 (3) |
| C16A—H16A | 0.9700 | C16—H16C | 0.9700 |
| C16A—H16B | 0.9700 | C16—H16D | 0.9700 |
| C2A'—C3A' | 1.398 (3) | C6—C5 | 1.374 (3) |
| C3A'—O9A' | 1.359 (3) | C6—H6 | 0.9300 |
| C3A'—C4A' | 1.374 (3) | C5—H5 | 0.9300 |
| C3A—C4A | 1.377 (3) | O7'—C8' | 1.374 (3) |
| C3A—H3A | 0.9300 | C15—H15C | 0.9700 |
| C15A—H15A | 0.9700 | C15—H15D | 0.9700 |
| C15A—H15B | 0.9700 | C3'—O9' | 1.357 (3) |
| C6A—C5A | 1.375 (3) | C3'—C4' | 1.366 (4) |
| C6A—H6A | 0.9300 | C6'—C5' | 1.369 (4) |
| C6A'—C5A' | 1.372 (3) | C6'—H6' | 0.9300 |
| C6A'—H6A' | 0.9300 | O9'—C10' | 1.395 (3) |
| C4A'—C5A' | 1.375 (4) | C17—H17D | 0.9600 |
| C4A'—H4A' | 0.9300 | C17—H17E | 0.9600 |
| C4A—C5A | 1.370 (4) | C17—H17F | 0.9600 |

| | | | |
|----------------|-------------|-------------|-------------|
| C4A—H4A | 0.9300 | C5'—C4' | 1.366 (4) |
| C17A—H17A | 0.9600 | C5'—H5' | 0.9300 |
| C17A—H17B | 0.9600 | C4'—H4' | 0.9300 |
| C17A—H17C | 0.9600 | C8'—H8'1 | 0.9600 |
| C5A—H5A | 0.9300 | C8'—H8'2 | 0.9600 |
| C8A'—H8A1 | 0.9600 | C8'—H8'3 | 0.9600 |
| C8A'—H8A2 | 0.9600 | C10'—H10D | 0.9600 |
| C8A'—H8A3 | 0.9600 | C10'—H10E | 0.9600 |
| C5A'—H5A' | 0.9300 | C10'—H10F | 0.9600 |
| O9A'—C10B—H10A | 109.5 | C9—N8—C7 | 125.51 (16) |
| O9A'—C10B—H10B | 109.5 | C9—N8—H8 | 117.3 (16) |
| H10A—C10B—H10B | 109.5 | C7—N8—H8 | 116.9 (16) |
| O9A'—C10B—H10C | 109.5 | C13—O12—C11 | 111.27 (15) |
| H10A—C10B—H10C | 109.5 | N14—C1—C2 | 114.61 (16) |
| H10B—C10B—H10C | 109.5 | N14—C1—C10 | 103.07 (15) |
| N14A—C1A—C2A | 113.64 (15) | C2—C1—C10 | 113.98 (15) |
| N14A—C1A—C10A | 102.66 (14) | N14—C1—C13 | 115.71 (15) |
| C2A—C1A—C10A | 114.65 (15) | C2—C1—C13 | 108.15 (15) |
| N14A—C1A—C13A | 115.07 (15) | C10—C1—C13 | 100.52 (15) |
| C2A—C1A—C13A | 109.39 (15) | C3—C2—C7 | 118.31 (18) |
| C10A—C1A—C13A | 100.71 (14) | C3—C2—C1 | 122.57 (17) |
| C13A—O12A—C11A | 110.64 (14) | C7—C2—C1 | 119.12 (16) |
| C9A—N8A—C7A | 126.00 (17) | C6—C7—C2 | 120.67 (18) |
| C9A—N8A—H8A | 117.9 (15) | C6—C7—N8 | 118.95 (17) |
| C7A—N8A—H8A | 116.0 (15) | C2—C7—N8 | 120.36 (17) |
| O9A—C9A—N8A | 121.57 (17) | C9—C10—C1 | 113.04 (15) |
| O9A—C9A—C10A | 121.93 (17) | C9—C10—C16 | 111.75 (15) |
| N8A—C9A—C10A | 116.47 (16) | C1—C10—C16 | 102.91 (15) |
| C2A'—O7A'—C8A' | 121.12 (18) | C9—C10—C11 | 111.71 (15) |
| O12A—C11A—C1A' | 109.37 (15) | C1—C10—C11 | 102.04 (14) |
| O12A—C11A—C10A | 103.00 (14) | C16—C10—C11 | 114.72 (16) |
| C1A'—C11A—C10A | 117.58 (15) | O12—C11—C1' | 108.71 (15) |
| O12A—C11A—H11A | 108.8 | O12—C11—C10 | 102.87 (15) |
| C1A'—C11A—H11A | 108.8 | C1'—C11—C10 | 117.93 (16) |
| C10A—C11A—H11A | 108.8 | O12—C11—H11 | 109.0 |
| C6A—C7A—C2A | 120.61 (18) | C1'—C11—H11 | 109.0 |
| C6A—C7A—N8A | 118.81 (18) | C10—C11—H11 | 109.0 |
| C2A—C7A—N8A | 120.56 (17) | C1—N14—C17 | 118.80 (17) |
| C9A—C10A—C1A | 114.39 (15) | C1—N14—C15 | 104.91 (16) |
| C9A—C10A—C16A | 111.90 (15) | C17—N14—C15 | 113.96 (18) |
| C1A—C10A—C16A | 103.25 (14) | O9—C9—N8 | 121.40 (17) |
| C9A—C10A—C11A | 110.44 (14) | O9—C9—C10 | 122.32 (17) |
| C1A—C10A—C11A | 101.78 (14) | N8—C9—C10 | 116.26 (16) |
| C16A—C10A—C11A | 114.60 (15) | O13—C13—O12 | 121.44 (19) |
| C15A—N14A—C17A | 113.06 (17) | O13—C13—C1 | 128.59 (19) |
| C15A—N14A—C1A | 105.39 (15) | O12—C13—C1 | 109.97 (15) |
| C17A—N14A—C1A | 118.68 (17) | C6'—C1'—C2' | 119.2 (2) |

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|----------------|-------------|---------------|-------------|
| C6A'—C1A'—C2A' | 119.18 (19) | C6'—C1'—C11 | 121.6 (2) |
| C6A'—C1A'—C11A | 121.77 (18) | C2'—C1'—C11 | 119.20 (18) |
| C2A'—C1A'—C11A | 119.05 (17) | C4—C3—C2 | 121.04 (19) |
| C7A—C2A—C3A | 118.03 (19) | C4—C3—H3 | 119.5 |
| C7A—C2A—C1A | 119.21 (16) | C2—C3—H3 | 119.5 |
| C3A—C2A—C1A | 122.74 (18) | C3—C4—C5 | 120.03 (19) |
| C15A—C16A—C10A | 105.86 (16) | C3—C4—H4 | 120.0 |
| C15A—C16A—H16A | 110.6 | C5—C4—H4 | 120.0 |
| C10A—C16A—H16A | 110.6 | O7'—C2'—C1' | 113.90 (19) |
| C15A—C16A—H16B | 110.6 | O7'—C2'—C3' | 125.9 (2) |
| C10A—C16A—H16B | 110.6 | C1'—C2'—C3' | 120.1 (2) |
| H16A—C16A—H16B | 108.7 | C15—C16—C10 | 105.64 (16) |
| O13A—C13A—O12A | 121.52 (18) | C15—C16—H16C | 110.6 |
| O13A—C13A—C1A | 128.10 (18) | C10—C16—H16C | 110.6 |
| O12A—C13A—C1A | 110.37 (15) | C15—C16—H16D | 110.6 |
| O7A'—C2A'—C1A' | 115.13 (17) | C10—C16—H16D | 110.6 |
| O7A'—C2A'—C3A' | 124.65 (18) | H16C—C16—H16D | 108.7 |
| C1A'—C2A'—C3A' | 119.92 (19) | C5—C6—C7 | 119.8 (2) |
| O9A'—C3A'—C4A' | 122.8 (2) | C5—C6—H6 | 120.1 |
| O9A'—C3A'—C2A' | 117.4 (2) | C7—C6—H6 | 120.1 |
| C4A'—C3A'—C2A' | 119.8 (2) | C6—C5—C4 | 120.1 (2) |
| C4A—C3A—C2A | 121.3 (2) | C6—C5—H5 | 120.0 |
| C4A—C3A—H3A | 119.4 | C4—C5—H5 | 120.0 |
| C2A—C3A—H3A | 119.4 | C2'—O7'—C8' | 126.5 (2) |
| N14A—C15A—C16A | 104.53 (16) | N14—C15—C16 | 104.96 (16) |
| N14A—C15A—H15A | 110.8 | N14—C15—H15C | 110.8 |
| C16A—C15A—H15A | 110.8 | C16—C15—H15C | 110.8 |
| N14A—C15A—H15B | 110.8 | N14—C15—H15D | 110.8 |
| C16A—C15A—H15B | 110.8 | C16—C15—H15D | 110.8 |
| H15A—C15A—H15B | 108.9 | H15C—C15—H15D | 108.8 |
| C5A—C6A—C7A | 119.9 (2) | O9'—C3'—C4' | 122.5 (2) |
| C5A—C6A—H6A | 120.0 | O9'—C3'—C2' | 118.2 (2) |
| C7A—C6A—H6A | 120.0 | C4'—C3'—C2' | 119.4 (2) |
| C3A'—O9A'—C10B | 119.3 (2) | C5'—C6'—C1' | 120.0 (2) |
| C5A'—C6A'—C1A' | 120.4 (2) | C5'—C6'—H6' | 120.0 |
| C5A'—C6A'—H6A' | 119.8 | C1'—C6'—H6' | 120.0 |
| C1A'—C6A'—H6A' | 119.8 | C3'—O9'—C10' | 119.0 (2) |
| C3A'—C4A'—C5A' | 120.2 (2) | N14—C17—H17D | 109.5 |
| C3A'—C4A'—H4A' | 119.9 | N14—C17—H17E | 109.5 |
| C5A'—C4A'—H4A' | 119.9 | H17D—C17—H17E | 109.5 |
| C5A—C4A—C3A | 119.8 (2) | N14—C17—H17F | 109.5 |
| C5A—C4A—H4A | 120.1 | H17D—C17—H17F | 109.5 |
| C3A—C4A—H4A | 120.1 | H17E—C17—H17F | 109.5 |
| N14A—C17A—H17A | 109.5 | C4'—C5'—C6' | 120.9 (3) |
| N14A—C17A—H17B | 109.5 | C4'—C5'—H5' | 119.6 |
| H17A—C17A—H17B | 109.5 | C6'—C5'—H5' | 119.6 |
| N14A—C17A—H17C | 109.5 | C3'—C4'—C5' | 120.3 (2) |
| H17A—C17A—H17C | 109.5 | C3'—C4'—H4' | 119.8 |

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|---------------------|--------------|-----------------|--------------|
| H17B—C17A—H17C | 109.5 | C5'—C4'—H4' | 119.8 |
| C4A—C5A—C6A | 120.3 (2) | O7'—C8'—H8'1 | 109.5 |
| C4A—C5A—H5A | 119.8 | O7'—C8'—H8'2 | 109.5 |
| C6A—C5A—H5A | 119.8 | H8'1—C8'—H8'2 | 109.5 |
| O7A'—C8A'—H8A1 | 109.5 | O7'—C8'—H8'3 | 109.5 |
| O7A'—C8A'—H8A2 | 109.5 | H8'1—C8'—H8'3 | 109.5 |
| H8A1—C8A'—H8A2 | 109.5 | H8'2—C8'—H8'3 | 109.5 |
| O7A'—C8A'—H8A3 | 109.5 | O9'—C10'—H10D | 109.5 |
| H8A1—C8A'—H8A3 | 109.5 | O9'—C10'—H10E | 109.5 |
| H8A2—C8A'—H8A3 | 109.5 | H10D—C10'—H10E | 109.5 |
| C6A'—C5A'—C4A' | 120.5 (2) | O9'—C10'—H10F | 109.5 |
| C6A'—C5A'—H5A' | 119.7 | H10D—C10'—H10F | 109.5 |
| C4A'—C5A'—H5A' | 119.7 | H10E—C10'—H10F | 109.5 |
| | | | |
| C7A—N8A—C9A—O9A | 169.07 (18) | N14—C1—C2—C3 | -82.4 (2) |
| C7A—N8A—C9A—C10A | -12.8 (3) | C10—C1—C2—C3 | 159.17 (18) |
| C13A—O12A—C11A—C1A' | -153.09 (16) | C13—C1—C2—C3 | 48.3 (2) |
| C13A—O12A—C11A—C10A | -27.3 (2) | N14—C1—C2—C7 | 98.4 (2) |
| C9A—N8A—C7A—C6A | 177.86 (19) | C10—C1—C2—C7 | -20.0 (3) |
| C9A—N8A—C7A—C2A | -4.0 (3) | C13—C1—C2—C7 | -130.91 (18) |
| O9A—C9A—C10A—C1A | -150.95 (18) | C3—C2—C7—C6 | 0.1 (3) |
| N8A—C9A—C10A—C1A | 30.9 (2) | C1—C2—C7—C6 | 179.38 (19) |
| O9A—C9A—C10A—C16A | -34.0 (2) | C3—C2—C7—N8 | 178.38 (18) |
| N8A—C9A—C10A—C16A | 147.90 (17) | C1—C2—C7—N8 | -2.4 (3) |
| O9A—C9A—C10A—C11A | 95.0 (2) | C9—N8—C7—C6 | -175.7 (2) |
| N8A—C9A—C10A—C11A | -83.2 (2) | C9—N8—C7—C2 | 6.0 (3) |
| N14A—C1A—C10A—C9A | 90.48 (17) | N14—C1—C10—C9 | -86.85 (18) |
| C2A—C1A—C10A—C9A | -33.2 (2) | C2—C1—C10—C9 | 38.0 (2) |
| C13A—C1A—C10A—C9A | -150.54 (15) | C13—C1—C10—C9 | 153.41 (16) |
| N14A—C1A—C10A—C16A | -31.36 (17) | N14—C1—C10—C16 | 33.86 (18) |
| C2A—C1A—C10A—C16A | -155.09 (15) | C2—C1—C10—C16 | 158.69 (16) |
| C13A—C1A—C10A—C16A | 87.62 (16) | C13—C1—C10—C16 | -85.88 (17) |
| N14A—C1A—C10A—C11A | -150.43 (14) | N14—C1—C10—C11 | 153.04 (14) |
| C2A—C1A—C10A—C11A | 85.84 (17) | C2—C1—C10—C11 | -82.13 (18) |
| C13A—C1A—C10A—C11A | -31.45 (17) | C13—C1—C10—C11 | 33.30 (17) |
| O12A—C11A—C10A—C9A | 158.13 (15) | C13—O12—C11—C1' | 148.84 (17) |
| C1A'—C11A—C10A—C9A | -81.6 (2) | C13—O12—C11—C10 | 23.1 (2) |
| O12A—C11A—C10A—C1A | 36.27 (17) | C9—C10—C11—O12 | -156.02 (15) |
| C1A'—C11A—C10A—C1A | 156.57 (16) | C1—C10—C11—O12 | -34.98 (17) |
| O12A—C11A—C10A—C16A | -74.39 (18) | C16—C10—C11—O12 | 75.49 (19) |
| C1A'—C11A—C10A—C16A | 45.9 (2) | C9—C10—C11—C1' | 84.4 (2) |
| C2A—C1A—N14A—C15A | 168.15 (16) | C1—C10—C11—C1' | -154.55 (16) |
| C10A—C1A—N14A—C15A | 43.74 (18) | C16—C10—C11—C1' | -44.1 (2) |
| C13A—C1A—N14A—C15A | -64.6 (2) | C2—C1—N14—C17 | 62.7 (2) |
| C2A—C1A—N14A—C17A | -64.0 (2) | C10—C1—N14—C17 | -172.91 (18) |
| C10A—C1A—N14A—C17A | 171.59 (17) | C13—C1—N14—C17 | -64.3 (2) |
| C13A—C1A—N14A—C17A | 63.2 (2) | C2—C1—N14—C15 | -168.57 (16) |
| O12A—C11A—C1A'—C6A' | 26.3 (3) | C10—C1—N14—C15 | -44.15 (18) |

| | | | |
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| C10A—C11A—C1A'—C6A' | -90.6 (2) | C13—C1—N14—C15 | 64.5 (2) |
| O12A—C11A—C1A'—C2A' | -153.38 (17) | C7—N8—C9—O9 | -167.2 (2) |
| C10A—C11A—C1A'—C2A' | 89.7 (2) | C7—N8—C9—C10 | 14.3 (3) |
| C6A—C7A—C2A—C3A | 0.3 (3) | C1—C10—C9—O9 | 145.7 (2) |
| N8A—C7A—C2A—C3A | -177.79 (19) | C16—C10—C9—O9 | 30.2 (3) |
| C6A—C7A—C2A—C1A | 178.77 (18) | C11—C10—C9—O9 | -99.9 (2) |
| N8A—C7A—C2A—C1A | 0.6 (3) | C1—C10—C9—N8 | -35.7 (2) |
| N14A—C1A—C2A—C7A | -99.4 (2) | C16—C10—C9—N8 | -151.28 (19) |
| C10A—C1A—C2A—C7A | 18.2 (2) | C11—C10—C9—N8 | 78.7 (2) |
| C13A—C1A—C2A—C7A | 130.44 (18) | C11—O12—C13—O13 | 179.06 (19) |
| N14A—C1A—C2A—C3A | 78.9 (2) | C11—O12—C13—C1 | -1.1 (2) |
| C10A—C1A—C2A—C3A | -163.43 (18) | N14—C1—C13—O13 | 48.4 (3) |
| C13A—C1A—C2A—C3A | -51.2 (2) | C2—C1—C13—O13 | -81.7 (3) |
| C9A—C10A—C16A—C15A | -114.59 (18) | C10—C1—C13—O13 | 158.5 (2) |
| C1A—C10A—C16A—C15A | 8.91 (19) | N14—C1—C13—O12 | -131.46 (18) |
| C11A—C10A—C16A—C15A | 118.69 (17) | C2—C1—C13—O12 | 98.43 (18) |
| C11A—O12A—C13A—O13A | -174.39 (19) | C10—C1—C13—O12 | -21.30 (19) |
| C11A—O12A—C13A—C1A | 6.6 (2) | O12—C11—C1'—C6' | -19.7 (3) |
| N14A—C1A—C13A—O13A | -52.4 (3) | C10—C11—C1'—C6' | 96.8 (3) |
| C2A—C1A—C13A—O13A | 76.9 (3) | O12—C11—C1'—C2' | 159.99 (18) |
| C10A—C1A—C13A—O13A | -162.0 (2) | C10—C11—C1'—C2' | -83.5 (2) |
| N14A—C1A—C13A—O12A | 126.50 (17) | C7—C2—C3—C4 | -1.5 (3) |
| C2A—C1A—C13A—O12A | -104.16 (18) | C1—C2—C3—C4 | 179.28 (19) |
| C10A—C1A—C13A—O12A | 16.9 (2) | C2—C3—C4—C5 | 1.9 (3) |
| C8A'—O7A'—C2A'—C1A' | -146.66 (19) | C6'—C1'—C2'—O7' | 175.2 (3) |
| C8A'—O7A'—C2A'—C3A' | 39.6 (3) | C11—C1'—C2'—O7' | -4.5 (3) |
| C6A'—C1A'—C2A'—O7A' | -175.19 (19) | C6'—C1'—C2'—C3' | -1.2 (4) |
| C11A—C1A'—C2A'—O7A' | 4.5 (3) | C11—C1'—C2'—C3' | 179.14 (19) |
| C6A'—C1A'—C2A'—C3A' | -1.1 (3) | C9—C10—C16—C15 | 109.63 (19) |
| C11A—C1A'—C2A'—C3A' | 178.60 (18) | C1—C10—C16—C15 | -12.0 (2) |
| O7A'—C2A'—C3A'—O9A' | -2.0 (3) | C11—C10—C16—C15 | -121.91 (18) |
| C1A'—C2A'—C3A'—O9A' | -175.4 (2) | C2—C7—C6—C5 | 0.9 (3) |
| O7A'—C2A'—C3A'—C4A' | 175.2 (2) | N8—C7—C6—C5 | -177.4 (2) |
| C1A'—C2A'—C3A'—C4A' | 1.7 (3) | C7—C6—C5—C4 | -0.5 (3) |
| C7A—C2A—C3A—C4A | 1.5 (3) | C3—C4—C5—C6 | -0.8 (3) |
| C1A—C2A—C3A—C4A | -176.8 (2) | C1'—C2'—O7'—C8' | 164.7 (3) |
| C17A—N14A—C15A—C16A | -169.27 (17) | C3'—C2'—O7'—C8' | -19.2 (4) |
| C1A—N14A—C15A—C16A | -38.11 (19) | C1—N14—C15—C16 | 36.4 (2) |
| C10A—C16A—C15A—N14A | 16.9 (2) | C17—N14—C15—C16 | 167.98 (18) |
| C2A—C7A—C6A—C5A | -1.4 (3) | C10—C16—C15—N14 | -14.0 (2) |
| N8A—C7A—C6A—C5A | 176.7 (2) | O7'—C2'—C3'—O9' | 6.2 (4) |
| C4A'—C3A'—O9A'—C10B | 24.3 (4) | C1'—C2'—C3'—O9' | -177.9 (2) |
| C2A'—C3A'—O9A'—C10B | -158.7 (2) | O7'—C2'—C3'—C4' | -174.0 (3) |
| C2A'—C1A'—C6A'—C5A' | -0.1 (4) | C1'—C2'—C3'—C4' | 1.9 (4) |
| C11A—C1A'—C6A'—C5A' | -179.8 (2) | C2'—C1'—C6'—C5' | 0.0 (5) |
| O9A'—C3A'—C4A'—C5A' | 175.9 (3) | C11—C1'—C6'—C5' | 179.6 (3) |
| C2A'—C3A'—C4A'—C5A' | -1.1 (4) | C4'—C3'—O9'—C10' | -23.3 (4) |
| C2A—C3A—C4A—C5A | -2.3 (4) | C2'—C3'—O9'—C10' | 156.6 (3) |

| | | | |
|---------------------|----------|-----------------|-----------|
| C3A—C4A—C5A—C6A | 1.2 (4) | C1'—C6'—C5'—C4' | 0.6 (6) |
| C7A—C6A—C5A—C4A | 0.6 (4) | O9'—C3'—C4'—C5' | 178.5 (3) |
| C1A'—C6A'—C5A'—C4A' | 0.7 (4) | C2'—C3'—C4'—C5' | -1.4 (5) |
| C3A'—C4A'—C5A'—C6A' | -0.1 (4) | C6'—C5'—C4'—C3' | 0.1 (6) |

Hydrogen-bond geometry (Å, °)

| <i>D—H...A</i> | <i>D—H</i> | <i>H...A</i> | <i>D...A</i> | <i>D—H...A</i> |
|---------------------------------|------------|--------------|--------------|----------------|
| N8—H8...O9 ⁱ | 0.89 (3) | 2.01 (3) | 2.903 (2) | 177 (2) |
| N8A—H8A...O9A ⁱⁱ | 0.89 (2) | 2.07 (3) | 2.958 (2) | 175 (2) |
| C6A'—H6A'...O13A ⁱⁱⁱ | 0.93 | 2.43 | 3.322 (3) | 161 |

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