

19 β ,28-Epoxy-18 α -olean-3 β -ol

R. C. Santos,^a R. M. A. Pinto,^a A. Matos Beja,^b
J. A. R. Salvador^a and J. A. Paixão^{b*}

^aLaboratório de Química Farmacêutica, Faculdade de Farmácia, Universidade de Coimbra, P-3000-548 Coimbra, Portugal, and ^bCEMDRX, Departamento de Física, Faculdade de Ciências e Tecnologia, Universidade de Coimbra, P-3004-516 Coimbra, Portugal

Correspondence e-mail: jap@pollux.fis.uc.pt

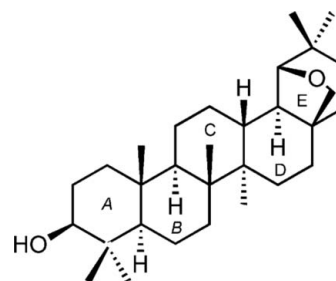
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.048; wR factor = 0.140; data-to-parameter ratio = 11.0.

The title triterpene, $\text{C}_{30}\text{H}_{50}\text{O}_2$, is an 18 α -oleanane derivative prepared by the Wagner–Meerwein rearrangement of betulin with $\text{Bi}(\text{OTf})_3 \cdot x\text{H}_2\text{O}$ (OTf is trifluoromethanesulfonate). There are two symmetry-independent molecules in the asymmetric unit that show no significant differences concerning bond lengths and angles. The conformation of the six-membered rings is close to a chair form, while the five-membered epoxide rings adopt envelope conformations. All rings are *trans*-fused. In the crystal, molecules are held together by $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds. A quantum-mechanical *ab initio* Roothan Hartree–Fock calculation on the isolated molecule gives values for bond lengths and valency angles close to the experimental values. The calculations also reproduce well the molecular conformation with calculated puckering parameters that match well the observed values.

Related literature

For terpene rearrangements, see: King *et al.* (1968). For Wagner–Meerwein rearrangements, see: Hanson (1991). For the synthesis of 18 α -oleanane derivatives, see: Salvador *et al.* (2009). For the cytotoxic activity of 18 α -oleanane derivatives, see: Urban *et al.* (2007); Thibeault *et al.* (2007). For puckering parameters, see: Cremer & Pople (1975) and for asymmetry parameters, see: Duax & Norton (1975). For the program *GAMMESS* used to perform the quantum chemical calculations, see: Schmidt *et al.* (1993).



Experimental

Crystal data

$\text{C}_{30}\text{H}_{50}\text{O}_2$
 $M_r = 442.70$
Monoclinic, $P2_1$
 $a = 13.2824$ (2) Å
 $b = 12.6702$ (2) Å
 $c = 15.5236$ (3) Å
 $\beta = 94.9990$ (10)°
 $V = 2602.54$ (8) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.07$ mm⁻¹
 $T = 293$ K
0.40 × 0.25 × 0.18 mm

Data collection

Bruker APEXII CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2000)
 $T_{\min} = 0.930$, $T_{\max} = 0.998$
63144 measured reflections
6510 independent reflections
5427 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.039$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.048$
 $wR(F^2) = 0.140$
 $S = 1.02$
6510 reflections
593 parameters
1 restraint
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.28$ e Å⁻³
 $\Delta\rho_{\min} = -0.19$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-\text{H} \cdots A$ | $D-\text{H}$ | $\text{H} \cdots A$ | $D \cdots A$ | $D-\text{H} \cdots A$ |
|---|--------------|---------------------|--------------|-----------------------|
| $\text{O3A}-\text{H3A1} \cdots \text{O19A}^i$ | 0.82 | 2.04 | 2.853 (3) | 171 |
| $\text{O3B}-\text{H3B1} \cdots \text{O3A}^{ii}$ | 0.82 | 2.12 | 2.920 (3) | 164 |

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

Data collection: *SMART* (Bruker, 2003); cell refinement: *SAINT* (Bruker, 2003); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5024).

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Comment

Terpenes were among those natural products wherein rearrangements were earliest detected and studied (King *et al.*, 1968). Rearrangements involving the formation of a carbocation and 1,2-migrations of hydrogen or alkyl groups from one carbon to a neighboring carbon are designated as Wagner–Meerwein rearrangements (Hanson, 1991). This group of transformations constitutes a particularly important field of research in organic chemistry and has been widely reported in triterpenoid chemistry. Recently, we have reported the use of bismuth(III) salts as catalysts for the Wagner–Meerwein rearrangement of lupane derivatives with expansion of ring E and formation of an additional O-containing ring (Salvador *et al.*, 2009). Using this procedure, the Wagner–Meerwein rearrangement of betulin afforded the title compound in high yield.

In this communication we report the molecular structure of the 19 β ,28-Epoxy-18 α -olean-3 β -ol (I), determined by single-crystal X-ray diffraction, and compare it with that of the free molecule as given by quantum mechanical *ab initio* calculation.

The structure of compound (I) with the corresponding atomic numbering scheme is shown in Fig. 1. All bond lengths and valency angles have typical values for this type of compounds. All rings are fused *trans* as shown by the angle between the least-squares planes of the rings [molecule A: rings A and B: 16.27 (12) $^\circ$, B and C: 9.60 (12) $^\circ$, C and D: 0.40 (12) $^\circ$, D and E: 15.59 (13) $^\circ$; molecule B: rings A and B: 13.68 (15) $^\circ$, B and C: 10.74 (14) $^\circ$, C and D: 2.78 (13) $^\circ$, D and E: 13.54 (16) $^\circ$]. Rings A to E have conformations close to chair as shown by the Cremer & Pople (1975) parameters [Molecule A: ring A: Q = 0.571 (3) Å, θ = 2.1 (3) $^\circ$ and φ = 79 (6) $^\circ$; B: Q = 0.575 (3) Å, θ = 9.2 (3) $^\circ$ and φ = 17.5 (16) $^\circ$; C: Q = 0.599 (3) Å, θ = 6.2 (3) $^\circ$ and φ = 336 (2) $^\circ$; D: Q = 0.521 (3) Å, θ = 168.4 (3) $^\circ$ and φ = 243.2 (14) $^\circ$; E: Q = 0.661 (3) Å, θ = 21.6 (3) $^\circ$ and φ = 59.1 (7) $^\circ$; Molecule B: ring A: Q = 0.553 (4) Å, θ = 4.6 (4) $^\circ$ and φ = 61 (5) $^\circ$; B: Q = 0.569 (3) Å, θ = 7.5 (3) $^\circ$ and φ = 11 (2) $^\circ$; C: Q = 0.585 (3) Å, θ = 4.2 (3) $^\circ$ and φ = 30 (3) $^\circ$; D: Q = 0.537 (3) Å, θ = 170.0 (3) $^\circ$ and φ = 236.5 (17) $^\circ$; E: Q = 0.656 (3) Å, θ = 18.9 (3) $^\circ$ and φ = 61.6 (10) $^\circ$; The epoxyde-ring has a C18-envelope conformation [molecule A: q_2 = 0.461 (3) Å and φ_2 = 255.3 (3) $^\circ$ and asymmetry parameters (Duax & Norton, 1975) $\Delta C_s(C18A) = \Delta C_s(C28A,O19A) = 2.1$ (2) $^\circ$ molecule B: q_2 = 0.468 (3) Å and φ_2 = 254.6 (4) $^\circ$ and asymmetry parameters (Duax & Norton, 1975) $\Delta C_s(C18B) = \Delta C_s(C28B,O19B) = 1.6$ (3) $^\circ$].

The molecules are hydrogen bonded involving the hydroxyl and epoxyde groups. The hydroxyl group of molecule A acts as a donor towards a neighbour hydroxyl group of a neighbour A molecule acting as an acceptor forming a chain of hydrogen bonds running along the *a* axis. The hydroxyl group of molecule B acts as a donor towards the epoxyde group of a neighbour A molecule.

In order to gain some insight on how the crystal packing of (I) might affect the molecular geometry we have performed a quantum chemical calculation on the equilibrium geometry of the free molecule. These calculations were performed with the computer program GAMMESS (Schmidt *et al.*, 1993). A molecular orbital Roothan Hartree-Fock method was used with an extended 6–31 G(d,p) basis set. Tight conditions for convergence of both the self-consistent field cycles and maximum

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density and energy gradient variations were imposed (10^{-6} atomic units). The program was run on the Milipeia cluster of UC-LCA (using 16 Opteron cores, 2.2 GHz running Linux).

The *ab-initio* calculations reproduce well the observed experimental bond length and valency angles of the molecule. Also, the calculated conformation of the rings are very close to the experimental values, with the exception of ring C for which the calculations gave a conformation closer to the ideal chair than experiment.

Experimental

The synthesis of the 19 β ,28-epoxy-18 α -olean-3 β -ol was efficiently accomplished by Wagner-Meerwein rearrangement of commercially available betulin with Bi(OTf)₃.xH₂O in CH₂Cl₂ at reflux (Salvador *et al.*, 2009). The product of this reaction was isolated in 95% yield and identified as the title compound from IR, ¹H and ¹³C NMR spectroscopy data (Salvador *et al.*, 2009). Recrystallization from ethanol at room temperature gave colourless single crystals suitable for X-ray diffraction analysis.

Refinement

All hydrogen atoms were refined as riding on their parent atoms using *SHELXL97* defaults except for that of the hydroxyl group which had its coordinates freely refined with $U_{\text{iso}} = 1.5 U_{\text{eq}}$ of the O atoms.

The absolute configuration was known from the synthetic route, but could not be determined from the X-ray data. Thus, Friedel pairs were merged for refinement.

Figures

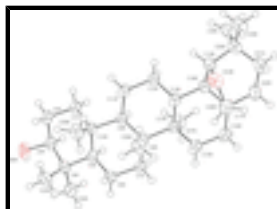


Fig. 1. *ORTEP* plot of one of the two molecules in the asymmetric unit. Displacement ellipsoids are drawn at the 50% level.

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Crystal data

C₃₀H₅₀O₂

$M_r = 442.70$

Monoclinic, $P2_1$

Hall symbol: P 2yb

$a = 13.2824$ (2) Å

$b = 12.6702$ (2) Å

$c = 15.5236$ (3) Å

$\beta = 94.9990$ (10)°

$V = 2602.54$ (8) Å³

$F_{000} = 984$

$D_x = 1.130$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 9126 reflections

$\theta = 2.5$ – 27.5 °

$\mu = 0.07$ mm⁻¹

$T = 293$ K

Block, clear colourless

$0.40 \times 0.25 \times 0.18$ mm

Z = 4

Data collection

| | |
|---|--|
| Bruker APEXII CCD area-detector diffractometer | 6510 independent reflections |
| Radiation source: fine-focus sealed tube | 5427 reflections with $I > 2\sigma(I)$ |
| Monochromator: graphite | $R_{\text{int}} = 0.039$ |
| $T = 293$ K | $\theta_{\text{max}} = 27.9^\circ$ |
| φ and ω scans | $\theta_{\text{min}} = 1.5^\circ$ |
| Absorption correction: multi-scan (SADABS; Sheldrick, 2000) | $h = -17 \rightarrow 17$ |
| $T_{\text{min}} = 0.930$, $T_{\text{max}} = 0.998$ | $k = -16 \rightarrow 16$ |
| 63144 measured reflections | $l = -20 \rightarrow 20$ |

Refinement

| | |
|--|--|
| Refinement on F^2 | Secondary atom site location: difference Fourier map |
| Least-squares matrix: full | Hydrogen site location: inferred from neighbouring sites |
| $R[F^2 > 2\sigma(F^2)] = 0.048$ | H-atom parameters constrained |
| $wR(F^2) = 0.140$ | $w = 1/[\sigma^2(F_o^2) + (0.0834P)^2 + 0.4115P]$ |
| $S = 1.02$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 6510 reflections | $(\Delta/\sigma)_{\text{max}} = 0.001$ |
| 593 parameters | $\Delta\rho_{\text{max}} = 0.28 \text{ e } \text{\AA}^{-3}$ |
| 1 restraint | $\Delta\rho_{\text{min}} = -0.19 \text{ e } \text{\AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | Extinction correction: none |

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)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|--------------|----------------------------------|
| C1A | 1.21538 (19) | 1.06069 (19) | 0.31109 (18) | 0.0404 (5) |
| H1A1 | 1.1745 | 1.1233 | 0.2993 | 0.049* |
| H1A2 | 1.2266 | 1.0531 | 0.3734 | 0.049* |

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|------|--------------|--------------|--------------|------------|
| C2A | 1.3177 (2) | 1.0762 (2) | 0.27390 (19) | 0.0436 (6) |
| H2A1 | 1.3071 | 1.0887 | 0.2121 | 0.052* |
| H2A2 | 1.3515 | 1.1374 | 0.3004 | 0.052* |
| C3A | 1.38282 (17) | 0.9802 (2) | 0.29068 (16) | 0.0380 (5) |
| H3A | 1.3919 | 0.9699 | 0.3534 | 0.046* |
| C4A | 1.33528 (17) | 0.8782 (2) | 0.25066 (16) | 0.0362 (5) |
| C23A | 1.3342 (2) | 0.8791 (3) | 0.15174 (18) | 0.0536 (7) |
| H23A | 1.4024 | 0.8802 | 0.1357 | 0.080* |
| H23B | 1.2991 | 0.9407 | 0.1291 | 0.080* |
| H23C | 1.3006 | 0.8170 | 0.1285 | 0.080* |
| C24A | 1.40181 (19) | 0.7852 (2) | 0.2846 (2) | 0.0509 (7) |
| H24A | 1.3767 | 0.7210 | 0.2578 | 0.076* |
| H24B | 1.4002 | 0.7798 | 0.3462 | 0.076* |
| H24C | 1.4701 | 0.7967 | 0.2710 | 0.076* |
| C5A | 1.22947 (16) | 0.86733 (18) | 0.28626 (14) | 0.0309 (4) |
| H5A | 1.2447 | 0.8626 | 0.3491 | 0.037* |
| C6A | 1.17428 (18) | 0.7645 (2) | 0.26226 (18) | 0.0408 (6) |
| H6A1 | 1.1449 | 0.7681 | 0.2029 | 0.049* |
| H6A2 | 1.2218 | 0.7062 | 0.2669 | 0.049* |
| C7A | 1.09128 (18) | 0.74588 (19) | 0.32233 (19) | 0.0410 (5) |
| H7A1 | 1.1221 | 0.7385 | 0.3810 | 0.049* |
| H7A2 | 1.0575 | 0.6799 | 0.3065 | 0.049* |
| C8A | 1.01189 (16) | 0.83426 (18) | 0.32031 (15) | 0.0331 (5) |
| C26A | 0.94773 (19) | 0.8264 (3) | 0.23252 (17) | 0.0479 (6) |
| H26A | 0.9915 | 0.8266 | 0.1865 | 0.072* |
| H26B | 0.9025 | 0.8855 | 0.2260 | 0.072* |
| H26C | 0.9093 | 0.7621 | 0.2306 | 0.072* |
| C9A | 1.06731 (16) | 0.94255 (17) | 0.33170 (15) | 0.0321 (4) |
| H9A | 1.0997 | 0.9408 | 0.3909 | 0.038* |
| C10A | 1.15646 (17) | 0.96339 (19) | 0.27366 (15) | 0.0331 (5) |
| C25A | 1.1193 (2) | 0.9870 (3) | 0.17901 (18) | 0.0508 (7) |
| H25A | 1.0552 | 1.0224 | 0.1769 | 0.076* |
| H25B | 1.1120 | 0.9221 | 0.1472 | 0.076* |
| H25C | 1.1675 | 1.0314 | 0.1538 | 0.076* |
| C11A | 0.99067 (19) | 1.0334 (2) | 0.3321 (2) | 0.0445 (6) |
| H11A | 1.0268 | 1.0994 | 0.3423 | 0.053* |
| H11B | 0.9530 | 1.0377 | 0.2758 | 0.053* |
| C12A | 0.91727 (19) | 1.0190 (2) | 0.4008 (2) | 0.0430 (6) |
| H12A | 0.8670 | 1.0747 | 0.3949 | 0.052* |
| H12B | 0.9537 | 1.0257 | 0.4574 | 0.052* |
| C13A | 0.86364 (16) | 0.91191 (19) | 0.39464 (15) | 0.0333 (5) |
| H13A | 0.8248 | 0.9089 | 0.3380 | 0.040* |
| C14A | 0.94152 (16) | 0.81979 (18) | 0.39779 (15) | 0.0330 (4) |
| C27A | 1.00401 (19) | 0.8173 (2) | 0.48636 (16) | 0.0434 (6) |
| H27A | 0.9595 | 0.8234 | 0.5316 | 0.065* |
| H27B | 1.0509 | 0.8751 | 0.4900 | 0.065* |
| H27C | 1.0404 | 0.7519 | 0.4925 | 0.065* |
| C15A | 0.88247 (19) | 0.7147 (2) | 0.38937 (19) | 0.0433 (6) |
| H15A | 0.8452 | 0.7119 | 0.3328 | 0.052* |

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|------|---------------|--------------|--------------|-------------|
| H15B | 0.9303 | 0.6567 | 0.3929 | 0.052* |
| C16A | 0.8090 (2) | 0.6995 (2) | 0.4580 (2) | 0.0475 (6) |
| H16A | 0.8473 | 0.6887 | 0.5134 | 0.057* |
| H16B | 0.7698 | 0.6361 | 0.4447 | 0.057* |
| C17A | 0.73689 (17) | 0.7918 (2) | 0.46578 (16) | 0.0391 (5) |
| C18A | 0.78804 (17) | 0.90099 (19) | 0.46337 (15) | 0.0348 (5) |
| H18A | 0.8214 | 0.9178 | 0.5205 | 0.042* |
| C19A | 0.69298 (17) | 0.9692 (2) | 0.44596 (16) | 0.0390 (5) |
| H19A | 0.7106 | 1.0387 | 0.4240 | 0.047* |
| C20A | 0.63477 (19) | 0.9810 (3) | 0.52715 (17) | 0.0464 (6) |
| C29A | 0.5409 (2) | 1.0501 (3) | 0.5060 (2) | 0.0584 (8) |
| H29A | 0.5615 | 1.1191 | 0.4889 | 0.088* |
| H29B | 0.5039 | 1.0558 | 0.5562 | 0.088* |
| H29C | 0.4988 | 1.0185 | 0.4596 | 0.088* |
| C30A | 0.7016 (3) | 1.0367 (3) | 0.5986 (2) | 0.0630 (9) |
| H30A | 0.7523 | 0.9885 | 0.6228 | 0.095* |
| H30B | 0.6608 | 1.0599 | 0.6430 | 0.095* |
| H30C | 0.7338 | 1.0965 | 0.5747 | 0.095* |
| C21A | 0.6028 (2) | 0.8707 (3) | 0.5547 (2) | 0.0594 (8) |
| H21A | 0.5843 | 0.8743 | 0.6137 | 0.071* |
| H21B | 0.5430 | 0.8498 | 0.5183 | 0.071* |
| C22A | 0.6834 (2) | 0.7858 (3) | 0.54953 (19) | 0.0522 (7) |
| H22A | 0.6522 | 0.7170 | 0.5533 | 0.063* |
| H22B | 0.7332 | 0.7932 | 0.5987 | 0.063* |
| C28A | 0.65841 (19) | 0.8003 (2) | 0.38829 (19) | 0.0474 (6) |
| H28A | 0.6862 | 0.7752 | 0.3363 | 0.057* |
| H28B | 0.5992 | 0.7585 | 0.3977 | 0.057* |
| O3A | 1.48053 (13) | 1.00257 (17) | 0.26184 (14) | 0.0501 (5) |
| H3A1 | 1.5240 | 0.9707 | 0.2923 | 0.075* |
| O19A | 0.63221 (13) | 0.91049 (17) | 0.38001 (12) | 0.0462 (4) |
| C1B | -0.2473 (3) | 0.2089 (3) | 0.1513 (3) | 0.0712 (11) |
| H1B1 | -0.2525 | 0.1906 | 0.0904 | 0.085* |
| H1B2 | -0.2003 | 0.1599 | 0.1813 | 0.085* |
| C2B | -0.3505 (3) | 0.1957 (3) | 0.1856 (3) | 0.0715 (10) |
| H2B1 | -0.3991 | 0.2409 | 0.1532 | 0.086* |
| H2B2 | -0.3729 | 0.1232 | 0.1776 | 0.086* |
| C3B | -0.3458 (2) | 0.2240 (3) | 0.2808 (2) | 0.0570 (8) |
| H3B | -0.2984 | 0.1756 | 0.3125 | 0.068* |
| C4B | -0.3095 (2) | 0.3373 (2) | 0.29996 (19) | 0.0486 (6) |
| C23B | -0.3896 (2) | 0.4190 (3) | 0.2711 (3) | 0.0698 (9) |
| H23D | -0.4115 | 0.4082 | 0.2111 | 0.105* |
| H23E | -0.3616 | 0.4885 | 0.2788 | 0.105* |
| H23F | -0.4462 | 0.4117 | 0.3050 | 0.105* |
| C24B | -0.2904 (3) | 0.3460 (4) | 0.3991 (2) | 0.0769 (11) |
| H24D | -0.2740 | 0.4177 | 0.4147 | 0.115* |
| H24E | -0.2352 | 0.3008 | 0.4190 | 0.115* |
| H24F | -0.3501 | 0.3250 | 0.4253 | 0.115* |
| C5B | -0.20754 (18) | 0.3516 (2) | 0.25790 (17) | 0.0408 (5) |
| H5B | -0.1617 | 0.3012 | 0.2890 | 0.049* |

supplementary materials

| | | | | |
|------|---------------|--------------|---------------|-------------|
| C6B | -0.1576 (2) | 0.4583 (3) | 0.2765 (2) | 0.0610 (8) |
| H6B1 | -0.1625 | 0.4771 | 0.3365 | 0.073* |
| H6B2 | -0.1924 | 0.5119 | 0.2406 | 0.073* |
| C7B | -0.0463 (2) | 0.4541 (3) | 0.2579 (2) | 0.0583 (8) |
| H7B1 | -0.0113 | 0.4045 | 0.2976 | 0.070* |
| H7B2 | -0.0165 | 0.5231 | 0.2693 | 0.070* |
| C8B | -0.02884 (19) | 0.4216 (2) | 0.16479 (19) | 0.0418 (5) |
| C26B | -0.0611 (3) | 0.5157 (3) | 0.1055 (3) | 0.0720 (11) |
| H26D | -0.0521 | 0.4977 | 0.0466 | 0.108* |
| H26E | -0.0203 | 0.5761 | 0.1223 | 0.108* |
| H26F | -0.1309 | 0.5318 | 0.1108 | 0.108* |
| C9B | -0.09107 (18) | 0.3195 (2) | 0.14203 (15) | 0.0388 (5) |
| H9B | -0.0595 | 0.2651 | 0.1803 | 0.047* |
| C10B | -0.20469 (19) | 0.3209 (2) | 0.16231 (17) | 0.0444 (6) |
| C25B | -0.2713 (2) | 0.3952 (4) | 0.0998 (2) | 0.0739 (11) |
| H25D | -0.2462 | 0.3944 | 0.0436 | 0.111* |
| H25E | -0.2685 | 0.4659 | 0.1222 | 0.111* |
| H25F | -0.3400 | 0.3708 | 0.0953 | 0.111* |
| C11B | -0.0740 (2) | 0.2824 (3) | 0.05069 (18) | 0.0598 (9) |
| H11C | -0.1128 | 0.2186 | 0.0376 | 0.072* |
| H11D | -0.0979 | 0.3361 | 0.0093 | 0.072* |
| C12B | 0.0367 (2) | 0.2607 (3) | 0.0419 (2) | 0.0569 (8) |
| H12C | 0.0450 | 0.2402 | -0.0172 | 0.068* |
| H12D | 0.0583 | 0.2018 | 0.0790 | 0.068* |
| C13B | 0.10390 (18) | 0.3549 (2) | 0.06550 (15) | 0.0394 (5) |
| H13B | 0.0829 | 0.4111 | 0.0244 | 0.047* |
| C14B | 0.08792 (18) | 0.39761 (19) | 0.15696 (16) | 0.0372 (5) |
| C27B | 0.1273 (2) | 0.3165 (3) | 0.22556 (17) | 0.0516 (7) |
| H27D | 0.1914 | 0.2897 | 0.2115 | 0.077* |
| H27E | 0.0799 | 0.2593 | 0.2267 | 0.077* |
| H27F | 0.1350 | 0.3497 | 0.2813 | 0.077* |
| C15B | 0.1532 (2) | 0.4977 (2) | 0.1723 (2) | 0.0571 (8) |
| H15C | 0.1288 | 0.5515 | 0.1312 | 0.069* |
| H15D | 0.1453 | 0.5244 | 0.2299 | 0.069* |
| C16B | 0.2653 (2) | 0.4786 (3) | 0.1634 (2) | 0.0636 (9) |
| H16C | 0.2925 | 0.4347 | 0.2111 | 0.076* |
| H16D | 0.3005 | 0.5457 | 0.1678 | 0.076* |
| C17B | 0.2866 (2) | 0.4255 (2) | 0.07837 (19) | 0.0492 (6) |
| C18B | 0.21616 (19) | 0.3323 (2) | 0.05480 (15) | 0.0396 (5) |
| H18B | 0.2384 | 0.2703 | 0.0891 | 0.048* |
| C19B | 0.2399 (2) | 0.3173 (3) | -0.03868 (18) | 0.0620 (9) |
| H19B | 0.1868 | 0.2753 | -0.0703 | 0.074* |
| C20B | 0.3440 (3) | 0.2659 (4) | -0.0460 (2) | 0.0688 (10) |
| C29B | 0.3596 (4) | 0.2538 (6) | -0.1423 (3) | 0.115 (2) |
| H29D | 0.3569 | 0.3220 | -0.1694 | 0.173* |
| H29E | 0.3073 | 0.2097 | -0.1695 | 0.173* |
| H29F | 0.4242 | 0.2221 | -0.1482 | 0.173* |
| C30B | 0.3482 (4) | 0.1572 (4) | -0.0041 (4) | 0.0938 (14) |
| H30D | 0.3997 | 0.1158 | -0.0277 | 0.141* |

| | | | | |
|------|-------------|------------|---------------|-------------|
| H30E | 0.2840 | 0.1228 | -0.0155 | 0.141* |
| H30F | 0.3635 | 0.1644 | 0.0572 | 0.141* |
| C21B | 0.4246 (2) | 0.3413 (4) | -0.0031 (2) | 0.0659 (9) |
| H21C | 0.4882 | 0.3036 | 0.0068 | 0.079* |
| H21D | 0.4347 | 0.3992 | -0.0423 | 0.079* |
| C22B | 0.3958 (2) | 0.3857 (3) | 0.0823 (2) | 0.0608 (8) |
| H22C | 0.4411 | 0.4435 | 0.0995 | 0.073* |
| H22D | 0.4051 | 0.3313 | 0.1262 | 0.073* |
| C28B | 0.2655 (3) | 0.4934 (3) | -0.0014 (3) | 0.0750 (11) |
| H28C | 0.2104 | 0.5418 | 0.0062 | 0.090* |
| H28D | 0.3250 | 0.5341 | -0.0122 | 0.090* |
| O3B | -0.4427 (2) | 0.2113 (2) | 0.3131 (2) | 0.0794 (8) |
| H3B1 | -0.4599 | 0.1493 | 0.3090 | 0.119* |
| O19B | 0.2390 (2) | 0.4238 (3) | -0.07185 (16) | 0.0825 (9) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C1A | 0.0370 (12) | 0.0286 (11) | 0.0573 (15) | -0.0037 (9) | 0.0131 (11) | 0.0015 (10) |
| C2A | 0.0405 (13) | 0.0349 (12) | 0.0575 (15) | -0.0081 (10) | 0.0151 (11) | 0.0023 (11) |
| C3A | 0.0307 (11) | 0.0408 (13) | 0.0431 (12) | -0.0083 (10) | 0.0061 (9) | 0.0012 (10) |
| C4A | 0.0303 (10) | 0.0366 (12) | 0.0424 (12) | -0.0021 (9) | 0.0068 (9) | -0.0029 (10) |
| C23A | 0.0501 (15) | 0.0683 (19) | 0.0447 (14) | -0.0031 (14) | 0.0178 (12) | -0.0087 (13) |
| C24A | 0.0324 (12) | 0.0425 (14) | 0.079 (2) | 0.0015 (11) | 0.0106 (12) | 0.0039 (13) |
| C5A | 0.0277 (10) | 0.0303 (11) | 0.0350 (11) | -0.0034 (9) | 0.0039 (8) | -0.0024 (9) |
| C6A | 0.0332 (11) | 0.0328 (12) | 0.0571 (15) | -0.0037 (9) | 0.0087 (10) | -0.0124 (11) |
| C7A | 0.0317 (11) | 0.0273 (11) | 0.0655 (16) | -0.0036 (9) | 0.0124 (11) | -0.0051 (11) |
| C8A | 0.0259 (10) | 0.0328 (11) | 0.0406 (11) | -0.0017 (8) | 0.0022 (8) | -0.0032 (9) |
| C26A | 0.0348 (12) | 0.0666 (18) | 0.0418 (13) | -0.0087 (12) | 0.0005 (9) | -0.0081 (12) |
| C9A | 0.0286 (10) | 0.0286 (11) | 0.0392 (11) | -0.0003 (8) | 0.0041 (8) | 0.0003 (9) |
| C10A | 0.0316 (10) | 0.0313 (11) | 0.0368 (11) | 0.0005 (9) | 0.0043 (8) | 0.0042 (9) |
| C25A | 0.0412 (13) | 0.0647 (19) | 0.0463 (14) | 0.0044 (13) | 0.0030 (11) | 0.0171 (13) |
| C11A | 0.0365 (12) | 0.0308 (12) | 0.0681 (17) | 0.0038 (10) | 0.0145 (12) | 0.0092 (12) |
| C12A | 0.0349 (12) | 0.0298 (12) | 0.0662 (17) | -0.0004 (10) | 0.0155 (11) | -0.0025 (11) |
| C13A | 0.0276 (10) | 0.0326 (11) | 0.0399 (12) | -0.0002 (9) | 0.0044 (8) | 0.0012 (9) |
| C14A | 0.0276 (10) | 0.0288 (10) | 0.0426 (12) | -0.0007 (8) | 0.0038 (8) | 0.0003 (9) |
| C27A | 0.0361 (11) | 0.0497 (15) | 0.0440 (13) | 0.0024 (11) | 0.0018 (9) | 0.0052 (12) |
| C15A | 0.0370 (12) | 0.0317 (12) | 0.0622 (16) | -0.0041 (10) | 0.0096 (11) | -0.0029 (11) |
| C16A | 0.0417 (13) | 0.0362 (13) | 0.0656 (17) | -0.0055 (11) | 0.0098 (12) | 0.0070 (12) |
| C17A | 0.0315 (11) | 0.0414 (13) | 0.0448 (13) | -0.0054 (10) | 0.0059 (9) | 0.0024 (10) |
| C18A | 0.0293 (10) | 0.0360 (12) | 0.0392 (12) | -0.0017 (9) | 0.0039 (9) | -0.0002 (9) |
| C19A | 0.0319 (11) | 0.0438 (13) | 0.0414 (12) | 0.0010 (10) | 0.0043 (9) | 0.0001 (11) |
| C20A | 0.0374 (12) | 0.0576 (17) | 0.0452 (14) | 0.0042 (12) | 0.0104 (10) | -0.0046 (12) |
| C29A | 0.0428 (14) | 0.074 (2) | 0.0596 (18) | 0.0124 (15) | 0.0118 (13) | -0.0085 (16) |
| C30A | 0.0633 (19) | 0.079 (2) | 0.0469 (16) | 0.0099 (17) | 0.0034 (14) | -0.0166 (15) |
| C21A | 0.0505 (15) | 0.069 (2) | 0.0625 (18) | -0.0006 (15) | 0.0262 (13) | 0.0029 (16) |
| C22A | 0.0479 (14) | 0.0561 (17) | 0.0547 (16) | -0.0056 (13) | 0.0173 (12) | 0.0106 (13) |
| C28A | 0.0357 (12) | 0.0511 (15) | 0.0546 (15) | -0.0044 (11) | -0.0004 (10) | -0.0086 (12) |

supplementary materials

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|------|-------------|-------------|-------------|--------------|--------------|--------------|
| O3A | 0.0318 (8) | 0.0523 (11) | 0.0673 (12) | -0.0081 (8) | 0.0104 (8) | 0.0076 (10) |
| O19A | 0.0380 (9) | 0.0549 (11) | 0.0442 (10) | 0.0065 (8) | -0.0049 (7) | -0.0054 (8) |
| C1B | 0.064 (2) | 0.067 (2) | 0.086 (2) | -0.0353 (18) | 0.0331 (18) | -0.0368 (19) |
| C2B | 0.0626 (19) | 0.066 (2) | 0.089 (3) | -0.0335 (17) | 0.0255 (18) | -0.0284 (19) |
| C3B | 0.0506 (16) | 0.0530 (17) | 0.0701 (19) | -0.0124 (14) | 0.0211 (14) | -0.0017 (15) |
| C4B | 0.0413 (13) | 0.0477 (15) | 0.0575 (15) | -0.0065 (12) | 0.0084 (11) | -0.0075 (13) |
| C23B | 0.0454 (16) | 0.063 (2) | 0.104 (3) | 0.0024 (15) | 0.0206 (17) | 0.000 (2) |
| C24B | 0.0607 (19) | 0.112 (3) | 0.0605 (19) | -0.023 (2) | 0.0180 (15) | -0.017 (2) |
| C5B | 0.0352 (11) | 0.0368 (12) | 0.0504 (14) | -0.0047 (10) | 0.0035 (10) | -0.0068 (11) |
| C6B | 0.0501 (16) | 0.0507 (17) | 0.085 (2) | -0.0141 (14) | 0.0227 (15) | -0.0233 (16) |
| C7B | 0.0470 (15) | 0.0514 (16) | 0.079 (2) | -0.0181 (13) | 0.0180 (14) | -0.0327 (15) |
| C8B | 0.0379 (12) | 0.0282 (11) | 0.0590 (15) | -0.0054 (9) | 0.0026 (10) | 0.0031 (11) |
| C26B | 0.0490 (16) | 0.0510 (18) | 0.117 (3) | 0.0050 (14) | 0.0129 (18) | 0.037 (2) |
| C9B | 0.0401 (12) | 0.0371 (12) | 0.0388 (12) | -0.0108 (10) | 0.0009 (9) | -0.0040 (10) |
| C10B | 0.0393 (12) | 0.0494 (15) | 0.0443 (13) | -0.0125 (11) | 0.0024 (10) | -0.0038 (12) |
| C25B | 0.0444 (16) | 0.111 (3) | 0.065 (2) | -0.0022 (18) | -0.0034 (14) | 0.022 (2) |
| C11B | 0.0533 (16) | 0.083 (2) | 0.0436 (14) | -0.0297 (16) | 0.0066 (12) | -0.0196 (15) |
| C12B | 0.0573 (16) | 0.0645 (19) | 0.0505 (15) | -0.0237 (15) | 0.0146 (13) | -0.0230 (14) |
| C13B | 0.0390 (12) | 0.0444 (13) | 0.0340 (11) | -0.0114 (10) | -0.0013 (9) | 0.0048 (10) |
| C14B | 0.0364 (11) | 0.0314 (11) | 0.0430 (12) | -0.0055 (9) | -0.0002 (9) | -0.0040 (9) |
| C27B | 0.0461 (14) | 0.0678 (19) | 0.0398 (13) | 0.0066 (14) | -0.0015 (10) | 0.0051 (13) |
| C15B | 0.0489 (15) | 0.0438 (15) | 0.080 (2) | -0.0155 (12) | 0.0113 (14) | -0.0201 (15) |
| C16B | 0.0478 (15) | 0.062 (2) | 0.082 (2) | -0.0183 (15) | 0.0073 (14) | -0.0259 (17) |
| C17B | 0.0407 (13) | 0.0497 (16) | 0.0568 (16) | -0.0121 (12) | 0.0028 (11) | -0.0006 (13) |
| C18B | 0.0423 (12) | 0.0441 (14) | 0.0320 (11) | -0.0060 (11) | 0.0003 (9) | 0.0004 (10) |
| C19B | 0.0491 (15) | 0.099 (3) | 0.0386 (14) | -0.0242 (17) | 0.0072 (11) | -0.0054 (16) |
| C20B | 0.0576 (18) | 0.098 (3) | 0.0538 (17) | -0.0161 (19) | 0.0226 (14) | -0.0196 (18) |
| C29B | 0.083 (3) | 0.201 (7) | 0.067 (2) | -0.040 (4) | 0.036 (2) | -0.049 (3) |
| C30B | 0.090 (3) | 0.079 (3) | 0.120 (4) | 0.003 (2) | 0.045 (3) | -0.023 (3) |
| C21B | 0.0448 (15) | 0.095 (3) | 0.0585 (18) | -0.0074 (17) | 0.0080 (13) | 0.0002 (18) |
| C22B | 0.0371 (13) | 0.080 (2) | 0.0638 (18) | -0.0079 (15) | -0.0007 (12) | -0.0097 (17) |
| C28B | 0.0590 (19) | 0.069 (2) | 0.098 (3) | -0.0090 (17) | 0.0142 (19) | 0.035 (2) |
| O3B | 0.0670 (14) | 0.0738 (16) | 0.1036 (19) | -0.0283 (13) | 0.0428 (14) | -0.0159 (15) |
| O19B | 0.0687 (15) | 0.127 (3) | 0.0513 (13) | -0.0092 (16) | 0.0022 (11) | 0.0403 (15) |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|----------|-----------|
| C1A—C2A | 1.535 (3) | C1B—C2B | 1.522 (4) |
| C1A—C10A | 1.546 (3) | C1B—C10B | 1.532 (4) |
| C1A—H1A1 | 0.9700 | C1B—H1B1 | 0.9700 |
| C1A—H1A2 | 0.9700 | C1B—H1B2 | 0.9700 |
| C2A—C3A | 1.503 (4) | C2B—C3B | 1.518 (5) |
| C2A—H2A1 | 0.9700 | C2B—H2B1 | 0.9700 |
| C2A—H2A2 | 0.9700 | C2B—H2B2 | 0.9700 |
| C3A—O3A | 1.438 (3) | C3B—O3B | 1.431 (4) |
| C3A—C4A | 1.544 (3) | C3B—C4B | 1.535 (4) |
| C3A—H3A | 0.9800 | C3B—H3B | 0.9800 |
| C4A—C23A | 1.534 (4) | C4B—C23B | 1.523 (5) |
| C4A—C24A | 1.539 (4) | C4B—C24B | 1.542 (5) |

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|-----------|-----------|-----------|-----------|
| C4A—C5A | 1.560 (3) | C4B—C5B | 1.565 (4) |
| C23A—H23A | 0.9600 | C23B—H23D | 0.9600 |
| C23A—H23B | 0.9600 | C23B—H23E | 0.9600 |
| C23A—H23C | 0.9600 | C23B—H23F | 0.9600 |
| C24A—H24A | 0.9600 | C24B—H24D | 0.9600 |
| C24A—H24B | 0.9600 | C24B—H24E | 0.9600 |
| C24A—H24C | 0.9600 | C24B—H24F | 0.9600 |
| C5A—C6A | 1.525 (3) | C5B—C6B | 1.521 (4) |
| C5A—C10A | 1.558 (3) | C5B—C10B | 1.538 (4) |
| C5A—H5A | 0.9800 | C5B—H5B | 0.9800 |
| C6A—C7A | 1.523 (3) | C6B—C7B | 1.532 (4) |
| C6A—H6A1 | 0.9700 | C6B—H6B1 | 0.9700 |
| C6A—H6A2 | 0.9700 | C6B—H6B2 | 0.9700 |
| C7A—C8A | 1.537 (3) | C7B—C8B | 1.540 (4) |
| C7A—H7A1 | 0.9700 | C7B—H7B1 | 0.9700 |
| C7A—H7A2 | 0.9700 | C7B—H7B2 | 0.9700 |
| C8A—C26A | 1.546 (3) | C8B—C26B | 1.543 (4) |
| C8A—C9A | 1.560 (3) | C8B—C9B | 1.559 (3) |
| C8A—C14A | 1.597 (3) | C8B—C14B | 1.595 (3) |
| C26A—H26A | 0.9600 | C26B—H26D | 0.9600 |
| C26A—H26B | 0.9600 | C26B—H26E | 0.9600 |
| C26A—H26C | 0.9600 | C26B—H26F | 0.9600 |
| C9A—C11A | 1.537 (3) | C9B—C11B | 1.529 (4) |
| C9A—C10A | 1.572 (3) | C9B—C10B | 1.568 (3) |
| C9A—H9A | 0.9800 | C9B—H9B | 0.9800 |
| C10A—C25A | 1.538 (3) | C10B—C25B | 1.569 (5) |
| C25A—H25A | 0.9600 | C25B—H25D | 0.9600 |
| C25A—H25B | 0.9600 | C25B—H25E | 0.9600 |
| C25A—H25C | 0.9600 | C25B—H25F | 0.9600 |
| C11A—C12A | 1.517 (4) | C11B—C12B | 1.514 (4) |
| C11A—H11A | 0.9700 | C11B—H11C | 0.9700 |
| C11A—H11B | 0.9700 | C11B—H11D | 0.9700 |
| C12A—C13A | 1.532 (3) | C12B—C13B | 1.516 (4) |
| C12A—H12A | 0.9700 | C12B—H12C | 0.9700 |
| C12A—H12B | 0.9700 | C12B—H12D | 0.9700 |
| C13A—C18A | 1.534 (3) | C13B—C18B | 1.542 (4) |
| C13A—C14A | 1.558 (3) | C13B—C14B | 1.551 (3) |
| C13A—H13A | 0.9800 | C13B—H13B | 0.9800 |
| C14A—C27A | 1.543 (3) | C14B—C27B | 1.539 (4) |
| C14A—C15A | 1.545 (3) | C14B—C15B | 1.543 (3) |
| C27A—H27A | 0.9600 | C27B—H27D | 0.9600 |
| C27A—H27B | 0.9600 | C27B—H27E | 0.9600 |
| C27A—H27C | 0.9600 | C27B—H27F | 0.9600 |
| C15A—C16A | 1.518 (4) | C15B—C16B | 1.527 (4) |
| C15A—H15A | 0.9700 | C15B—H15C | 0.9700 |
| C15A—H15B | 0.9700 | C15B—H15D | 0.9700 |
| C16A—C17A | 1.523 (4) | C16B—C17B | 1.529 (4) |
| C16A—H16A | 0.9700 | C16B—H16C | 0.9700 |
| C16A—H16B | 0.9700 | C16B—H16D | 0.9700 |

supplementary materials

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|---------------|-------------|---------------|-----------|
| C17A—C28A | 1.525 (4) | C17B—C28B | 1.514 (5) |
| C17A—C22A | 1.537 (4) | C17B—C18B | 1.532 (4) |
| C17A—C18A | 1.544 (3) | C17B—C22B | 1.533 (4) |
| C18A—C19A | 1.534 (3) | C18B—C19B | 1.523 (4) |
| C18A—H18A | 0.9800 | C18B—H18B | 0.9800 |
| C19A—O19A | 1.452 (3) | C19B—O19B | 1.444 (5) |
| C19A—C20A | 1.542 (3) | C19B—C20B | 1.541 (5) |
| C19A—H19A | 0.9800 | C19B—H19B | 0.9800 |
| C20A—C30A | 1.531 (4) | C20B—C30B | 1.522 (7) |
| C20A—C21A | 1.533 (5) | C20B—C29B | 1.535 (5) |
| C20A—C29A | 1.535 (4) | C20B—C21B | 1.542 (5) |
| C29A—H29A | 0.9600 | C29B—H29D | 0.9600 |
| C29A—H29B | 0.9600 | C29B—H29E | 0.9600 |
| C29A—H29C | 0.9600 | C29B—H29F | 0.9600 |
| C30A—H30A | 0.9600 | C30B—H30D | 0.9600 |
| C30A—H30B | 0.9600 | C30B—H30E | 0.9600 |
| C30A—H30C | 0.9600 | C30B—H30F | 0.9600 |
| C21A—C22A | 1.524 (5) | C21B—C22B | 1.519 (5) |
| C21A—H21A | 0.9700 | C21B—H21C | 0.9700 |
| C21A—H21B | 0.9700 | C21B—H21D | 0.9700 |
| C22A—H22A | 0.9700 | C22B—H22C | 0.9700 |
| C22A—H22B | 0.9700 | C22B—H22D | 0.9700 |
| C28A—O19A | 1.442 (4) | C28B—O19B | 1.424 (5) |
| C28A—H28A | 0.9700 | C28B—H28C | 0.9700 |
| C28A—H28B | 0.9700 | C28B—H28D | 0.9700 |
| O3A—H3A1 | 0.8200 | O3B—H3B1 | 0.8200 |
| C2A—C1A—C10A | 113.4 (2) | C2B—C1B—C10B | 113.4 (3) |
| C2A—C1A—H1A1 | 108.9 | C2B—C1B—H1B1 | 108.9 |
| C10A—C1A—H1A1 | 108.9 | C10B—C1B—H1B1 | 108.9 |
| C2A—C1A—H1A2 | 108.9 | C2B—C1B—H1B2 | 108.9 |
| C10A—C1A—H1A2 | 108.9 | C10B—C1B—H1B2 | 108.9 |
| H1A1—C1A—H1A2 | 107.7 | H1B1—C1B—H1B2 | 107.7 |
| C3A—C2A—C1A | 110.3 (2) | C3B—C2B—C1B | 110.6 (3) |
| C3A—C2A—H2A1 | 109.6 | C3B—C2B—H2B1 | 109.5 |
| C1A—C2A—H2A1 | 109.6 | C1B—C2B—H2B1 | 109.5 |
| C3A—C2A—H2A2 | 109.6 | C3B—C2B—H2B2 | 109.5 |
| C1A—C2A—H2A2 | 109.6 | C1B—C2B—H2B2 | 109.5 |
| H2A1—C2A—H2A2 | 108.1 | H2B1—C2B—H2B2 | 108.1 |
| O3A—C3A—C2A | 107.9 (2) | O3B—C3B—C2B | 110.7 (3) |
| O3A—C3A—C4A | 112.9 (2) | O3B—C3B—C4B | 108.3 (3) |
| C2A—C3A—C4A | 113.49 (19) | C2B—C3B—C4B | 113.3 (3) |
| O3A—C3A—H3A | 107.4 | O3B—C3B—H3B | 108.2 |
| C2A—C3A—H3A | 107.4 | C2B—C3B—H3B | 108.2 |
| C4A—C3A—H3A | 107.4 | C4B—C3B—H3B | 108.2 |
| C23A—C4A—C24A | 107.7 (2) | C23B—C4B—C3B | 112.3 (3) |
| C23A—C4A—C3A | 111.4 (2) | C23B—C4B—C24B | 107.2 (3) |
| C24A—C4A—C3A | 107.50 (19) | C3B—C4B—C24B | 106.4 (3) |
| C23A—C4A—C5A | 115.2 (2) | C23B—C4B—C5B | 114.0 (3) |
| C24A—C4A—C5A | 108.6 (2) | C3B—C4B—C5B | 107.2 (2) |

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| C3A—C4A—C5A | 106.31 (18) | C24B—C4B—C5B | 109.6 (2) |
| C4A—C23A—H23A | 109.5 | C4B—C23B—H23D | 109.5 |
| C4A—C23A—H23B | 109.5 | C4B—C23B—H23E | 109.5 |
| H23A—C23A—H23B | 109.5 | H23D—C23B—H23E | 109.5 |
| C4A—C23A—H23C | 109.5 | C4B—C23B—H23F | 109.5 |
| H23A—C23A—H23C | 109.5 | H23D—C23B—H23F | 109.5 |
| H23B—C23A—H23C | 109.5 | H23E—C23B—H23F | 109.5 |
| C4A—C24A—H24A | 109.5 | C4B—C24B—H24D | 109.5 |
| C4A—C24A—H24B | 109.5 | C4B—C24B—H24E | 109.5 |
| H24A—C24A—H24B | 109.5 | H24D—C24B—H24E | 109.5 |
| C4A—C24A—H24C | 109.5 | C4B—C24B—H24F | 109.5 |
| H24A—C24A—H24C | 109.5 | H24D—C24B—H24F | 109.5 |
| H24B—C24A—H24C | 109.5 | H24E—C24B—H24F | 109.5 |
| C6A—C5A—C10A | 110.85 (17) | C6B—C5B—C10B | 111.1 (2) |
| C6A—C5A—C4A | 114.69 (19) | C6B—C5B—C4B | 113.8 (2) |
| C10A—C5A—C4A | 117.16 (19) | C10B—C5B—C4B | 117.9 (2) |
| C6A—C5A—H5A | 104.1 | C6B—C5B—H5B | 104.1 |
| C10A—C5A—H5A | 104.1 | C10B—C5B—H5B | 104.1 |
| C4A—C5A—H5A | 104.1 | C4B—C5B—H5B | 104.1 |
| C7A—C6A—C5A | 109.95 (19) | C5B—C6B—C7B | 110.2 (2) |
| C7A—C6A—H6A1 | 109.7 | C5B—C6B—H6B1 | 109.6 |
| C5A—C6A—H6A1 | 109.7 | C7B—C6B—H6B1 | 109.6 |
| C7A—C6A—H6A2 | 109.7 | C5B—C6B—H6B2 | 109.6 |
| C5A—C6A—H6A2 | 109.7 | C7B—C6B—H6B2 | 109.6 |
| H6A1—C6A—H6A2 | 108.2 | H6B1—C6B—H6B2 | 108.1 |
| C6A—C7A—C8A | 114.0 (2) | C6B—C7B—C8B | 114.4 (2) |
| C6A—C7A—H7A1 | 108.8 | C6B—C7B—H7B1 | 108.7 |
| C8A—C7A—H7A1 | 108.8 | C8B—C7B—H7B1 | 108.7 |
| C6A—C7A—H7A2 | 108.8 | C6B—C7B—H7B2 | 108.7 |
| C8A—C7A—H7A2 | 108.8 | C8B—C7B—H7B2 | 108.7 |
| H7A1—C7A—H7A2 | 107.7 | H7B1—C7B—H7B2 | 107.6 |
| C7A—C8A—C26A | 107.2 (2) | C7B—C8B—C26B | 107.2 (3) |
| C7A—C8A—C9A | 108.81 (17) | C7B—C8B—C9B | 108.3 (2) |
| C26A—C8A—C9A | 112.0 (2) | C26B—C8B—C9B | 113.1 (2) |
| C7A—C8A—C14A | 110.28 (19) | C7B—C8B—C14B | 110.5 (2) |
| C26A—C8A—C14A | 110.04 (18) | C26B—C8B—C14B | 108.8 (2) |
| C9A—C8A—C14A | 108.49 (17) | C9B—C8B—C14B | 108.9 (2) |
| C8A—C26A—H26A | 109.5 | C8B—C26B—H26D | 109.5 |
| C8A—C26A—H26B | 109.5 | C8B—C26B—H26E | 109.5 |
| H26A—C26A—H26B | 109.5 | H26D—C26B—H26E | 109.5 |
| C8A—C26A—H26C | 109.5 | C8B—C26B—H26F | 109.5 |
| H26A—C26A—H26C | 109.5 | H26D—C26B—H26F | 109.5 |
| H26B—C26A—H26C | 109.5 | H26E—C26B—H26F | 109.5 |
| C11A—C9A—C8A | 110.67 (18) | C11B—C9B—C8B | 110.3 (2) |
| C11A—C9A—C10A | 114.15 (19) | C11B—C9B—C10B | 114.3 (2) |
| C8A—C9A—C10A | 117.04 (19) | C8B—C9B—C10B | 116.5 (2) |
| C11A—C9A—H9A | 104.5 | C11B—C9B—H9B | 104.8 |
| C8A—C9A—H9A | 104.5 | C8B—C9B—H9B | 104.8 |
| C10A—C9A—H9A | 104.5 | C10B—C9B—H9B | 104.8 |

supplementary materials

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|----------------|-------------|----------------|-------------|
| C25A—C10A—C1A | 108.3 (2) | C1B—C10B—C5B | 107.6 (2) |
| C25A—C10A—C5A | 114.6 (2) | C1B—C10B—C9B | 108.6 (2) |
| C1A—C10A—C5A | 106.66 (17) | C5B—C10B—C9B | 107.56 (19) |
| C25A—C10A—C9A | 112.66 (19) | C1B—C10B—C25B | 107.7 (3) |
| C1A—C10A—C9A | 107.49 (19) | C5B—C10B—C25B | 112.6 (3) |
| C5A—C10A—C9A | 106.78 (18) | C9B—C10B—C25B | 112.5 (2) |
| C10A—C25A—H25A | 109.5 | C10B—C25B—H25D | 109.5 |
| C10A—C25A—H25B | 109.5 | C10B—C25B—H25E | 109.5 |
| H25A—C25A—H25B | 109.5 | H25D—C25B—H25E | 109.5 |
| C10A—C25A—H25C | 109.5 | C10B—C25B—H25F | 109.5 |
| H25A—C25A—H25C | 109.5 | H25D—C25B—H25F | 109.5 |
| H25B—C25A—H25C | 109.5 | H25E—C25B—H25F | 109.5 |
| C12A—C11A—C9A | 112.3 (2) | C12B—C11B—C9B | 111.2 (2) |
| C12A—C11A—H11A | 109.1 | C12B—C11B—H11C | 109.4 |
| C9A—C11A—H11A | 109.1 | C9B—C11B—H11C | 109.4 |
| C12A—C11A—H11B | 109.1 | C12B—C11B—H11D | 109.4 |
| C9A—C11A—H11B | 109.1 | C9B—C11B—H11D | 109.4 |
| H11A—C11A—H11B | 107.9 | H11C—C11B—H11D | 108.0 |
| C11A—C12A—C13A | 112.7 (2) | C11B—C12B—C13B | 113.0 (3) |
| C11A—C12A—H12A | 109.0 | C11B—C12B—H12C | 109.0 |
| C13A—C12A—H12A | 109.0 | C13B—C12B—H12C | 109.0 |
| C11A—C12A—H12B | 109.0 | C11B—C12B—H12D | 109.0 |
| C13A—C12A—H12B | 109.0 | C13B—C12B—H12D | 109.0 |
| H12A—C12A—H12B | 107.8 | H12C—C12B—H12D | 107.8 |
| C12A—C13A—C18A | 111.5 (2) | C12B—C13B—C18B | 112.5 (2) |
| C12A—C13A—C14A | 110.98 (17) | C12B—C13B—C14B | 111.8 (2) |
| C18A—C13A—C14A | 112.58 (19) | C18B—C13B—C14B | 111.98 (18) |
| C12A—C13A—H13A | 107.2 | C12B—C13B—H13B | 106.7 |
| C18A—C13A—H13A | 107.2 | C18B—C13B—H13B | 106.7 |
| C14A—C13A—H13A | 107.2 | C14B—C13B—H13B | 106.7 |
| C27A—C14A—C15A | 106.7 (2) | C27B—C14B—C15B | 106.8 (2) |
| C27A—C14A—C13A | 110.32 (19) | C27B—C14B—C13B | 109.6 (2) |
| C15A—C14A—C13A | 108.20 (17) | C15B—C14B—C13B | 108.0 (2) |
| C27A—C14A—C8A | 111.64 (18) | C27B—C14B—C8B | 110.4 (2) |
| C15A—C14A—C8A | 111.18 (19) | C15B—C14B—C8B | 111.6 (2) |
| C13A—C14A—C8A | 108.73 (18) | C13B—C14B—C8B | 110.35 (19) |
| C14A—C27A—H27A | 109.5 | C14B—C27B—H27D | 109.5 |
| C14A—C27A—H27B | 109.5 | C14B—C27B—H27E | 109.5 |
| H27A—C27A—H27B | 109.5 | H27D—C27B—H27E | 109.5 |
| C14A—C27A—H27C | 109.5 | C14B—C27B—H27F | 109.5 |
| H27A—C27A—H27C | 109.5 | H27D—C27B—H27F | 109.5 |
| H27B—C27A—H27C | 109.5 | H27E—C27B—H27F | 109.5 |
| C16A—C15A—C14A | 113.7 (2) | C16B—C15B—C14B | 113.2 (2) |
| C16A—C15A—H15A | 108.8 | C16B—C15B—H15C | 108.9 |
| C14A—C15A—H15A | 108.8 | C14B—C15B—H15C | 108.9 |
| C16A—C15A—H15B | 108.8 | C16B—C15B—H15D | 108.9 |
| C14A—C15A—H15B | 108.8 | C14B—C15B—H15D | 108.9 |
| H15A—C15A—H15B | 107.7 | H15C—C15B—H15D | 107.7 |
| C15A—C16A—C17A | 113.9 (2) | C15B—C16B—C17B | 113.8 (2) |

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| C15A—C16A—H16A | 108.8 | C15B—C16B—H16C | 108.8 |
| C17A—C16A—H16A | 108.8 | C17B—C16B—H16C | 108.8 |
| C15A—C16A—H16B | 108.8 | C15B—C16B—H16D | 108.8 |
| C17A—C16A—H16B | 108.8 | C17B—C16B—H16D | 108.8 |
| H16A—C16A—H16B | 107.7 | H16C—C16B—H16D | 107.7 |
| C16A—C17A—C28A | 112.5 (2) | C28B—C17B—C16B | 114.9 (3) |
| C16A—C17A—C22A | 111.7 (2) | C28B—C17B—C18B | 100.2 (2) |
| C28A—C17A—C22A | 109.6 (2) | C16B—C17B—C18B | 113.0 (2) |
| C16A—C17A—C18A | 113.90 (18) | C28B—C17B—C22B | 109.1 (3) |
| C28A—C17A—C18A | 100.9 (2) | C16B—C17B—C22B | 110.9 (2) |
| C22A—C17A—C18A | 107.5 (2) | C18B—C17B—C22B | 108.2 (3) |
| C13A—C18A—C19A | 113.9 (2) | C19B—C18B—C17B | 98.7 (2) |
| C13A—C18A—C17A | 114.4 (2) | C19B—C18B—C13B | 114.1 (2) |
| C19A—C18A—C17A | 98.75 (18) | C17B—C18B—C13B | 114.0 (2) |
| C13A—C18A—H18A | 109.8 | C19B—C18B—H18B | 109.9 |
| C19A—C18A—H18A | 109.8 | C17B—C18B—H18B | 109.9 |
| C17A—C18A—H18A | 109.8 | C13B—C18B—H18B | 109.9 |
| O19A—C19A—C18A | 103.5 (2) | O19B—C19B—C18B | 103.1 (3) |
| O19A—C19A—C20A | 109.8 (2) | O19B—C19B—C20B | 110.3 (3) |
| C18A—C19A—C20A | 112.0 (2) | C18B—C19B—C20B | 112.6 (3) |
| O19A—C19A—H19A | 110.5 | O19B—C19B—H19B | 110.2 |
| C18A—C19A—H19A | 110.5 | C18B—C19B—H19B | 110.2 |
| C20A—C19A—H19A | 110.5 | C20B—C19B—H19B | 110.2 |
| C30A—C20A—C21A | 112.1 (3) | C30B—C20B—C29B | 108.7 (4) |
| C30A—C20A—C29A | 107.6 (3) | C30B—C20B—C19B | 110.5 (3) |
| C21A—C20A—C29A | 109.9 (2) | C29B—C20B—C19B | 108.2 (3) |
| C30A—C20A—C19A | 109.6 (2) | C30B—C20B—C21B | 112.2 (4) |
| C21A—C20A—C19A | 108.1 (2) | C29B—C20B—C21B | 109.6 (3) |
| C29A—C20A—C19A | 109.6 (2) | C19B—C20B—C21B | 107.5 (3) |
| C20A—C29A—H29A | 109.5 | C20B—C29B—H29D | 109.5 |
| C20A—C29A—H29B | 109.5 | C20B—C29B—H29E | 109.5 |
| H29A—C29A—H29B | 109.5 | H29D—C29B—H29E | 109.5 |
| C20A—C29A—H29C | 109.5 | C20B—C29B—H29F | 109.5 |
| H29A—C29A—H29C | 109.5 | H29D—C29B—H29F | 109.5 |
| H29B—C29A—H29C | 109.5 | H29E—C29B—H29F | 109.5 |
| C20A—C30A—H30A | 109.5 | C20B—C30B—H30D | 109.5 |
| C20A—C30A—H30B | 109.5 | C20B—C30B—H30E | 109.5 |
| H30A—C30A—H30B | 109.5 | H30D—C30B—H30E | 109.5 |
| C20A—C30A—H30C | 109.5 | C20B—C30B—H30F | 109.5 |
| H30A—C30A—H30C | 109.5 | H30D—C30B—H30F | 109.5 |
| H30B—C30A—H30C | 109.5 | H30E—C30B—H30F | 109.5 |
| C22A—C21A—C20A | 114.5 (2) | C22B—C21B—C20B | 112.8 (3) |
| C22A—C21A—H21A | 108.6 | C22B—C21B—H21C | 109.0 |
| C20A—C21A—H21A | 108.6 | C20B—C21B—H21C | 109.0 |
| C22A—C21A—H21B | 108.6 | C22B—C21B—H21D | 109.0 |
| C20A—C21A—H21B | 108.6 | C20B—C21B—H21D | 109.0 |
| H21A—C21A—H21B | 107.6 | H21C—C21B—H21D | 107.8 |
| C21A—C22A—C17A | 112.9 (2) | C21B—C22B—C17B | 113.4 (3) |
| C21A—C22A—H22A | 109.0 | C21B—C22B—H22C | 108.9 |

supplementary materials

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| C17A—C22A—H22A | 109.0 | C17B—C22B—H22C | 108.9 |
| C21A—C22A—H22B | 109.0 | C21B—C22B—H22D | 108.9 |
| C17A—C22A—H22B | 109.0 | C17B—C22B—H22D | 108.9 |
| H22A—C22A—H22B | 107.8 | H22C—C22B—H22D | 107.7 |
| O19A—C28A—C17A | 106.4 (2) | O19B—C28B—C17B | 107.0 (3) |
| O19A—C28A—H28A | 110.5 | O19B—C28B—H28C | 110.3 |
| C17A—C28A—H28A | 110.5 | C17B—C28B—H28C | 110.3 |
| O19A—C28A—H28B | 110.5 | O19B—C28B—H28D | 110.3 |
| C17A—C28A—H28B | 110.5 | C17B—C28B—H28D | 110.3 |
| H28A—C28A—H28B | 108.6 | H28C—C28B—H28D | 108.6 |
| C3A—O3A—H3A1 | 109.5 | C3B—O3B—H3B1 | 109.5 |
| C28A—O19A—C19A | 108.55 (19) | C28B—O19B—C19B | 108.1 (2) |
| C10A—C1A—C2A—C3A | -58.3 (3) | C10B—C1B—C2B—C3B | -58.4 (5) |
| C1A—C2A—C3A—O3A | -174.8 (2) | C1B—C2B—C3B—O3B | -179.8 (3) |
| C1A—C2A—C3A—C4A | 59.4 (3) | C1B—C2B—C3B—C4B | 58.4 (4) |
| O3A—C3A—C4A—C23A | -52.3 (3) | O3B—C3B—C4B—C23B | -49.9 (4) |
| C2A—C3A—C4A—C23A | 70.9 (3) | C2B—C3B—C4B—C23B | 73.2 (4) |
| O3A—C3A—C4A—C24A | 65.4 (3) | O3B—C3B—C4B—C24B | 67.1 (3) |
| C2A—C3A—C4A—C24A | -171.4 (2) | C2B—C3B—C4B—C24B | -169.8 (3) |
| O3A—C3A—C4A—C5A | -178.45 (19) | O3B—C3B—C4B—C5B | -175.8 (3) |
| C2A—C3A—C4A—C5A | -55.3 (3) | C2B—C3B—C4B—C5B | -52.7 (3) |
| C23A—C4A—C5A—C6A | 62.6 (3) | C23B—C4B—C5B—C6B | 59.1 (3) |
| C24A—C4A—C5A—C6A | -58.1 (3) | C3B—C4B—C5B—C6B | -176.0 (3) |
| C3A—C4A—C5A—C6A | -173.5 (2) | C24B—C4B—C5B—C6B | -61.0 (4) |
| C23A—C4A—C5A—C10A | -69.9 (3) | C23B—C4B—C5B—C10B | -73.6 (3) |
| C24A—C4A—C5A—C10A | 169.3 (2) | C3B—C4B—C5B—C10B | 51.3 (3) |
| C3A—C4A—C5A—C10A | 53.9 (2) | C24B—C4B—C5B—C10B | 166.3 (3) |
| C10A—C5A—C6A—C7A | -63.0 (3) | C10B—C5B—C6B—C7B | -61.3 (3) |
| C4A—C5A—C6A—C7A | 161.6 (2) | C4B—C5B—C6B—C7B | 162.8 (3) |
| C5A—C6A—C7A—C8A | 59.0 (3) | C5B—C6B—C7B—C8B | 57.5 (4) |
| C6A—C7A—C8A—C26A | 71.5 (3) | C6B—C7B—C8B—C26B | 72.7 (3) |
| C6A—C7A—C8A—C9A | -49.8 (3) | C6B—C7B—C8B—C9B | -49.6 (3) |
| C6A—C7A—C8A—C14A | -168.72 (19) | C6B—C7B—C8B—C14B | -168.8 (3) |
| C7A—C8A—C9A—C11A | -178.7 (2) | C7B—C8B—C9B—C11B | -178.6 (2) |
| C26A—C8A—C9A—C11A | 62.9 (3) | C26B—C8B—C9B—C11B | 62.7 (3) |
| C14A—C8A—C9A—C11A | -58.7 (2) | C14B—C8B—C9B—C11B | -58.4 (3) |
| C7A—C8A—C9A—C10A | 48.2 (3) | C7B—C8B—C9B—C10B | 49.0 (3) |
| C26A—C8A—C9A—C10A | -70.2 (2) | C26B—C8B—C9B—C10B | -69.7 (3) |
| C14A—C8A—C9A—C10A | 168.15 (18) | C14B—C8B—C9B—C10B | 169.2 (2) |
| C2A—C1A—C10A—C25A | -70.7 (3) | C2B—C1B—C10B—C5B | 52.9 (4) |
| C2A—C1A—C10A—C5A | 53.1 (3) | C2B—C1B—C10B—C9B | 169.0 (3) |
| C2A—C1A—C10A—C9A | 167.3 (2) | C2B—C1B—C10B—C25B | -68.8 (4) |
| C6A—C5A—C10A—C25A | -67.7 (3) | C6B—C5B—C10B—C1B | 174.8 (3) |
| C4A—C5A—C10A—C25A | 66.6 (3) | C4B—C5B—C10B—C1B | -51.3 (3) |
| C6A—C5A—C10A—C1A | 172.5 (2) | C6B—C5B—C10B—C9B | 58.0 (3) |
| C4A—C5A—C10A—C1A | -53.2 (2) | C4B—C5B—C10B—C9B | -168.1 (2) |
| C6A—C5A—C10A—C9A | 57.8 (2) | C6B—C5B—C10B—C25B | -66.6 (3) |
| C4A—C5A—C10A—C9A | -167.93 (18) | C4B—C5B—C10B—C25B | 67.3 (3) |
| C11A—C9A—C10A—C25A | -57.1 (3) | C11B—C9B—C10B—C1B | 59.6 (3) |

| | | | |
|---------------------|--------------|---------------------|------------|
| C8A—C9A—C10A—C25A | 74.4 (3) | C8B—C9B—C10B—C1B | -169.8 (3) |
| C11A—C9A—C10A—C1A | 62.0 (3) | C11B—C9B—C10B—C5B | 175.8 (3) |
| C8A—C9A—C10A—C1A | -166.42 (19) | C8B—C9B—C10B—C5B | -53.7 (3) |
| C11A—C9A—C10A—C5A | 176.2 (2) | C11B—C9B—C10B—C25B | -59.6 (4) |
| C8A—C9A—C10A—C5A | -52.3 (2) | C8B—C9B—C10B—C25B | 71.0 (3) |
| C8A—C9A—C11A—C12A | 55.8 (3) | C8B—C9B—C11B—C12B | 58.7 (3) |
| C10A—C9A—C11A—C12A | -169.6 (2) | C10B—C9B—C11B—C12B | -167.8 (3) |
| C9A—C11A—C12A—C13A | -53.6 (3) | C9B—C11B—C12B—C13B | -56.3 (4) |
| C11A—C12A—C13A—C18A | -178.5 (2) | C11B—C12B—C13B—C18B | -179.0 (2) |
| C11A—C12A—C13A—C14A | 55.1 (3) | C11B—C12B—C13B—C14B | 54.0 (3) |
| C12A—C13A—C14A—C27A | 64.8 (3) | C12B—C13B—C14B—C27B | 68.1 (3) |
| C18A—C13A—C14A—C27A | -60.9 (2) | C18B—C13B—C14B—C27B | -59.1 (3) |
| C12A—C13A—C14A—C15A | -178.8 (2) | C12B—C13B—C14B—C15B | -175.9 (2) |
| C18A—C13A—C14A—C15A | 55.5 (2) | C18B—C13B—C14B—C15B | 56.9 (3) |
| C12A—C13A—C14A—C8A | -57.9 (2) | C12B—C13B—C14B—C8B | -53.6 (3) |
| C18A—C13A—C14A—C8A | 176.36 (18) | C18B—C13B—C14B—C8B | 179.1 (2) |
| C7A—C8A—C14A—C27A | 56.9 (3) | C7B—C8B—C14B—C27B | 53.5 (3) |
| C26A—C8A—C14A—C27A | 175.0 (2) | C26B—C8B—C14B—C27B | 170.9 (3) |
| C9A—C8A—C14A—C27A | -62.1 (2) | C9B—C8B—C14B—C27B | -65.4 (3) |
| C7A—C8A—C14A—C15A | -62.1 (2) | C7B—C8B—C14B—C15B | -65.2 (3) |
| C26A—C8A—C14A—C15A | 55.9 (3) | C26B—C8B—C14B—C15B | 52.2 (3) |
| C9A—C8A—C14A—C15A | 178.82 (18) | C9B—C8B—C14B—C15B | 175.9 (2) |
| C7A—C8A—C14A—C13A | 178.88 (18) | C7B—C8B—C14B—C13B | 174.7 (2) |
| C26A—C8A—C14A—C13A | -63.1 (2) | C26B—C8B—C14B—C13B | -67.8 (3) |
| C9A—C8A—C14A—C13A | 59.8 (2) | C9B—C8B—C14B—C13B | 55.9 (3) |
| C27A—C14A—C15A—C16A | 61.8 (3) | C27B—C14B—C15B—C16B | 60.5 (3) |
| C13A—C14A—C15A—C16A | -57.0 (3) | C13B—C14B—C15B—C16B | -57.2 (3) |
| C8A—C14A—C15A—C16A | -176.3 (2) | C8B—C14B—C15B—C16B | -178.7 (3) |
| C14A—C15A—C16A—C17A | 52.2 (3) | C14B—C15B—C16B—C17B | 52.6 (4) |
| C15A—C16A—C17A—C28A | 70.5 (3) | C15B—C16B—C17B—C28B | 68.9 (4) |
| C15A—C16A—C17A—C22A | -165.6 (2) | C15B—C16B—C17B—C18B | -45.2 (4) |
| C15A—C16A—C17A—C18A | -43.5 (3) | C15B—C16B—C17B—C22B | -166.8 (3) |
| C12A—C13A—C18A—C19A | 71.8 (3) | C28B—C17B—C18B—C19B | 44.3 (3) |
| C14A—C13A—C18A—C19A | -162.71 (19) | C16B—C17B—C18B—C19B | 167.1 (3) |
| C12A—C13A—C18A—C17A | -175.6 (2) | C22B—C17B—C18B—C19B | -69.8 (3) |
| C14A—C13A—C18A—C17A | -50.1 (3) | C28B—C17B—C18B—C13B | -77.1 (3) |
| C16A—C17A—C18A—C13A | 43.1 (3) | C16B—C17B—C18B—C13B | 45.7 (3) |
| C28A—C17A—C18A—C13A | -77.8 (2) | C22B—C17B—C18B—C13B | 168.8 (2) |
| C22A—C17A—C18A—C13A | 167.4 (2) | C12B—C13B—C18B—C19B | 67.9 (3) |
| C16A—C17A—C18A—C19A | 164.4 (2) | C14B—C13B—C18B—C19B | -165.3 (2) |
| C28A—C17A—C18A—C19A | 43.6 (2) | C12B—C13B—C18B—C17B | -179.8 (2) |
| C22A—C17A—C18A—C19A | -71.2 (2) | C14B—C13B—C18B—C17B | -52.9 (3) |
| C13A—C18A—C19A—O19A | 78.2 (2) | C17B—C18B—C19B—O19B | -44.7 (3) |
| C17A—C18A—C19A—O19A | -43.5 (2) | C13B—C18B—C19B—O19B | 76.6 (3) |
| C13A—C18A—C19A—C20A | -163.6 (2) | C17B—C18B—C19B—C20B | 74.2 (3) |
| C17A—C18A—C19A—C20A | 74.6 (2) | C13B—C18B—C19B—C20B | -164.5 (3) |
| O19A—C19A—C20A—C30A | 176.4 (3) | O19B—C19B—C20B—C30B | 175.2 (3) |
| C18A—C19A—C20A—C30A | 62.0 (3) | C18B—C19B—C20B—C30B | 60.6 (4) |
| O19A—C19A—C20A—C21A | 54.0 (3) | O19B—C19B—C20B—C29B | -65.9 (4) |

supplementary materials

| | | | |
|---------------------|------------|---------------------|------------|
| C18A—C19A—C20A—C21A | -60.4 (3) | C18B—C19B—C20B—C29B | 179.5 (4) |
| O19A—C19A—C20A—C29A | -65.8 (3) | O19B—C19B—C20B—C21B | 52.4 (3) |
| C18A—C19A—C20A—C29A | 179.8 (2) | C18B—C19B—C20B—C21B | -62.2 (4) |
| C30A—C20A—C21A—C22A | -79.1 (3) | C30B—C20B—C21B—C22B | -77.8 (4) |
| C29A—C20A—C21A—C22A | 161.3 (3) | C29B—C20B—C21B—C22B | 161.3 (4) |
| C19A—C20A—C21A—C22A | 41.8 (3) | C19B—C20B—C21B—C22B | 43.9 (4) |
| C20A—C21A—C22A—C17A | -44.1 (4) | C20B—C21B—C22B—C17B | -46.1 (5) |
| C16A—C17A—C22A—C21A | -174.1 (2) | C28B—C17B—C22B—C21B | -47.1 (4) |
| C28A—C17A—C22A—C21A | -48.6 (3) | C16B—C17B—C22B—C21B | -174.6 (3) |
| C18A—C17A—C22A—C21A | 60.3 (3) | C18B—C17B—C22B—C21B | 61.0 (4) |
| C16A—C17A—C28A—O19A | -151.2 (2) | C16B—C17B—C28B—O19B | -151.1 (3) |
| C22A—C17A—C28A—O19A | 83.8 (3) | C18B—C17B—C28B—O19B | -29.7 (3) |
| C18A—C17A—C28A—O19A | -29.4 (3) | C22B—C17B—C28B—O19B | 83.7 (3) |
| C17A—C28A—O19A—C19A | 2.1 (3) | C17B—C28B—O19B—C19B | 1.7 (4) |
| C18A—C19A—O19A—C28A | 26.6 (3) | C18B—C19B—O19B—C28B | 27.5 (3) |
| C20A—C19A—O19A—C28A | -93.1 (2) | C20B—C19B—O19B—C28B | -93.0 (3) |

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-------------------------------------|-------|-------------|-------------|---------------|
| O3A—H3A1 \cdots O19A ⁱ | 0.82 | 2.04 | 2.853 (3) | 171 |
| O3B—H3B1 \cdots O3A ⁱⁱ | 0.82 | 2.12 | 2.920 (3) | 164 |

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

Fig. 1

