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1,3-Dibenzyl-1,2,3,4-tetrahydroquinazoline-2,4-dione

 Gavhar Karimova,^{a*} Jamshid Ashurov,^b Nasir Mukhamedov,^c Nusrat A. Parpiev^a and Khusniddin M. Shakhidoyatov^c

^aThe Mirzo Ulugbek National University of Uzbekistan, Faculty of Chemistry, University Str. 6, Tashkent 100779, Uzbekistan, ^bInstitute of Bioorganic Chemistry, Academy of Sciences of Uzbekistan, Mirzo Ulugbek Str. 83, Tashkent, 100125 Uzbekistan, and ^cS. Yunusov Institute of the Chemistry of Plant Substances, Academy of Sciences of Uzbekistan, Mirzo Ulugbek Str. 77, Tashkent 100170, Uzbekistan
Correspondence e-mail: gavhar1979.79@mail.ru

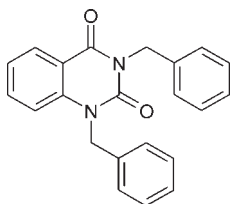
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.041; wR factor = 0.112; data-to-parameter ratio = 15.1.

The asymmetric unit of the title compound, $\text{C}_{22}\text{H}_{18}\text{N}_2\text{O}_2$, contains two independent molecules, which differ in the orientations of the benzyl groups with respect to the planar (r.m.s. deviations of 0.031 and 0.020 Å) quinazoline-2,4-dione skeletons [dihedral angles of 73.97 (4) and 70.07 (4)° in the first molecule and 75.63 (4) and 63.52 (3)° in the second]. The crystal structure is stabilized by weak intermolecular C—H \cdots O and C—H \cdots π interactions and aromatic π – π stacking interactions [centroid–centroid distance = 3.735 (2) Å].

Related literature

For the synthesis of the title compound, see: Hedayatullah (1981). For the synthesis of quinazoline-2,4-dione derivatives, see: Shi *et al.* (2007); Kuryazov *et al.* (2008). For the biological activity of quinazoline-2,4-dione derivatives, see: Colotaa *et al.* (2004); Yakhontov *et al.* (1977). For related structures, see: Mazza *et al.* (1988). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_{22}\text{H}_{18}\text{N}_2\text{O}_2$
 $M_r = 342.38$

 Orthorhombic, $Pbca$
 $a = 17.8989$ (4) Å

 $b = 14.0071$ (4) Å
 $c = 27.7222$ (6) Å
 $V = 6950.3$ (3) Å³
 $Z = 16$

 Cu $K\alpha$ radiation
 $\mu = 0.68$ mm⁻¹
 $T = 293$ K
 $0.5 \times 0.4 \times 0.35$ mm

Data collection

Oxford Diffraction Xcalibur Ruby diffractometer
 Absorption correction: multi-scan (*CrysAlis PRO*; Oxford Diffraction, 2009)
 $T_{\min} = 0.804$, $T_{\max} = 1.000$
 18947 measured reflections
 7088 independent reflections
 4141 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.112$
 $S = 0.90$
 7088 reflections
 470 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.28$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.14$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$Cg4$ and $Cg8$ are the centroids of the $C17A$ – $C22A$ and $C10B$ – $C15B$ rings, respectively.

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|------------------------------|-------|-------------|-------------|---------------|
| $C19A-H19A\cdots O2b^i$ | 0.93 | 2.70 | 3.419 (3) | 134 |
| $C6B-H6B\cdots C11a^{ii}$ | 0.93 | 2.89 | 3.604 (3) | 134 |
| $C21B-H21B\cdots C11a^{iii}$ | 0.93 | 2.80 | 3.600 (3) | 145 |
| $C19A-H19A\cdots O2b^i$ | 0.93 | 2.70 | 3.419 (3) | 134 |
| $C7B-H7B\cdots Cg4$ | 0.93 | 2.78 | 3.586 (2) | 146 |
| $C5A-H5A\cdots Cg8^{ii}$ | 0.93 | 2.90 | 3.641 (2) | 138 |

Symmetry codes: (i) $x + \frac{1}{2}, y, -z + \frac{3}{2}$; (ii) $-x + \frac{3}{2}, y, -\frac{1}{2}, z$; (iii) $x - \frac{1}{2}, y, -z + \frac{3}{2}$.

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

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FJ2301).

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supporting information

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1,3-Dibenzyl-1,2,3,4-tetrahydroquinazoline-2,4-dione

Gavhar Karimova, Jamshid Ashurov, Nasir Mukhamedov, Nusrat A. Parpiev and Khusniddin M. Shakhidoyatov

S1. Comment

Quinazoline-2,4-diones have been frequently used as intermediates and synthetic precursors for the preparation of a wide variety of heterocyclic compounds (Kuryazov *et al.*, 2008). In addition, they possess different biological activities (Colotaa *et al.*, 2004; Yakhontov *et al.*, 1977).

The title compound consists of a quinazoline-2,4-dione skeleton with two benzyl groups. The asymmetric unit contains two molecules of 1,2,3,4-tetrahydro-1,3-dibenzylquinazoline-2,4-dione (Fig. 1). Orientation of benzyl groups with respect to the planar quinazoline-2,4-dione skeletons are different for independent molecules. Dihedral angles between planar quinazoline-2,4-dione system and benzyl group planes are 73.97 (4)° and 70.07 (4)° (for molecule A) and 75.63 (4)° and 63.52 (3)° (for molecule B). Torsion angles responsible for orientation of benzyl groups are shown in table 1. (In order compare torsions between A and B independent molecules must be taken absolute values of torsion angles).

Quinazoline-2,4-dione system of the molecules are packed into sheets along *b* axis by a aromatic π - π stacking interaction. The benzene rings in the quinazoline-2,4-dione system standing nearly parallel (molecules A and B) are separated with distance of 3.477 (2) Å and benzene ring-centroid separation is 3.735 (2) Å with ring offset of 1.364 (2) Å. This distances for molecules of A at (*x*,*y*,*z*) and B at (1.5 - *x*, 1/2 + *y*, *z*) are 3.493 (2) Å, 3.791 (2) Å and 1.473 (2) Å.

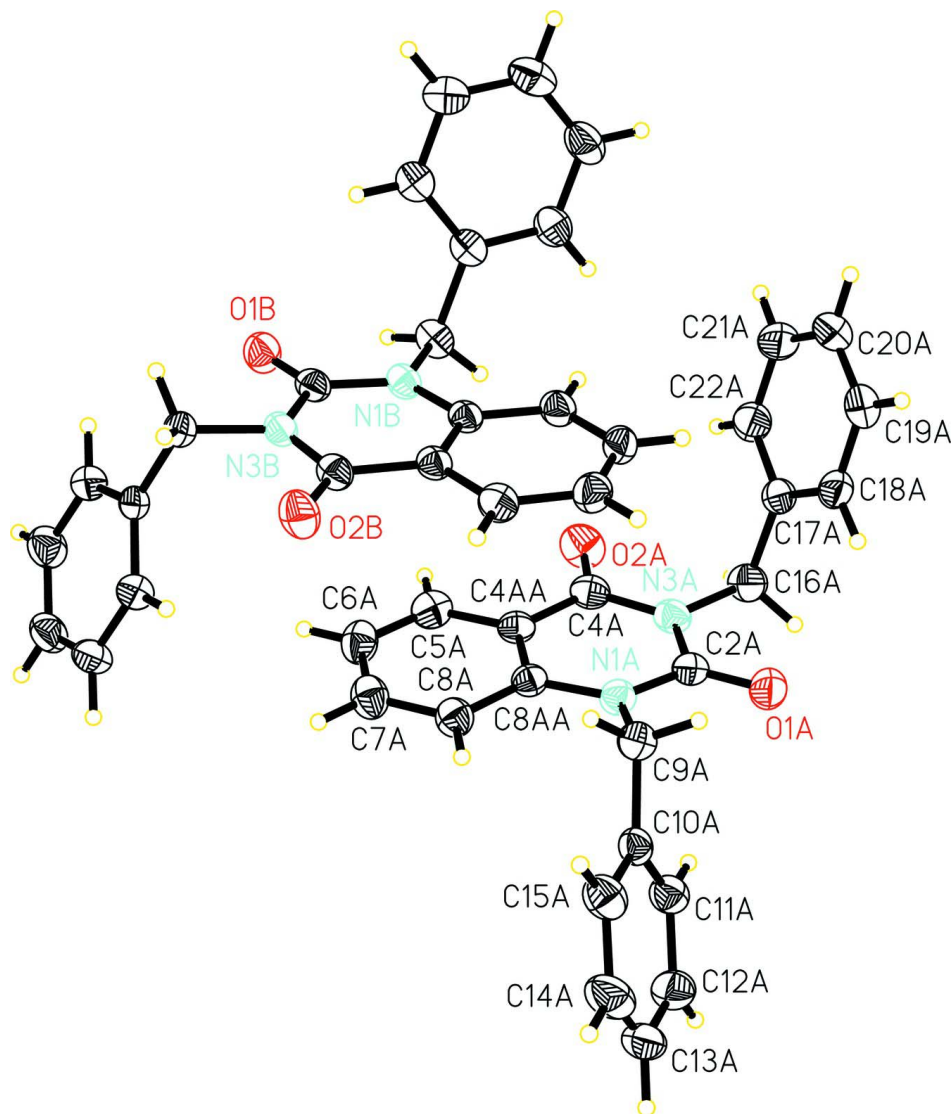
The observed structure is stabilized by weak C—H \cdots O and C—H \cdots C_{ar} type hydrogen bonds (Table 2). The bond distances and angles in organic compound molecules are in normal ranges (Allen *et al.*, 1987).

S2. Experimental

To suspension of 1*H*-quinazoline-2,4-dione (1.62 g) in 40 ml benzene was added 10% aqueous solution of sodium hydroxide (40 ml), tetrabutylammonium bromide (1.29 g, 4 mmol) and benzyl chloride (3.80 g, 30 mmol). The mixture was heated until 60° C and helded out for 6 h (Hedayatullah, 1981). The organic layer was separated, washed with water until neutral reaction and dried with Na₂SO₄, benzene was evaporated. Residue was recrystallized from benzene and obtained in 88% yield (3.02 g) of title compound. Colorless crystals suitable for X-ray analysis were obtained from dimethylformamide by slow evaporation.

S3. Refinement

Carbon-bound H atoms were positioned geometrically and treated as riding on their C atoms, with C—H distances of 0.93 Å (aromatic) and 0.97 Å (CH₂) and were refined with $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C})$.

**Figure 1**

Asymmetric unit of cell with atom labels and 30% probability displacement ellipsoids for non-H atoms.

1,3-Dibenzyl-1,2,3,4-tetrahydroquinazoline-2,4-dione

Crystal data

$C_{22}H_{18}N_2O_2$

$M_r = 342.38$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 17.8989(4) \text{ \AA}$

$b = 14.0071(4) \text{ \AA}$

$c = 27.7222(6) \text{ \AA}$

$V = 6950.3(3) \text{ \AA}^3$

$Z = 16$

$F(000) = 2880$

$D_x = 1.309 \text{ Mg m}^{-3}$

Melting point: 398(2) K

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

Cell parameters from 2463 reflections

$\theta = 3.5\text{--}35.8^\circ$

$\mu = 0.68 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Prism, colourless

$0.5 \times 0.4 \times 0.35 \text{ mm}$

Data collection

Oxford Diffraction Xcalibur Ruby
diffractometer
Radiation source: Enhance (Cu) X-ray Source
Graphite monochromator
Detector resolution: 10.2576 pixels mm⁻¹
 ω scans
Absorption correction: multi-scan
(*CrysAlis PRO*; Oxford Diffraction, 2009)
 $T_{\min} = 0.804$, $T_{\max} = 1.000$

18947 measured reflections
7088 independent reflections
4141 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$
 $\theta_{\max} = 75.8^\circ$, $\theta_{\min} = 4.0^\circ$
 $h = -22 \rightarrow 18$
 $k = -11 \rightarrow 17$
 $l = -31 \rightarrow 34$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.112$
 $S = 0.90$
7088 reflections
470 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0587P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.28 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.14 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.00081 (5)

Special details

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

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

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

| | x | y | z | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|--------------|--------------|-------------|----------------------------------|
| O1A | 0.90039 (7) | 0.27786 (11) | 0.72998 (5) | 0.0745 (4) |
| O2A | 0.85328 (8) | 0.30319 (11) | 0.89063 (5) | 0.0770 (4) |
| N1A | 0.77945 (8) | 0.27259 (10) | 0.75417 (5) | 0.0498 (4) |
| N3A | 0.87693 (8) | 0.29234 (10) | 0.81052 (6) | 0.0531 (4) |
| C2A | 0.85450 (10) | 0.28112 (13) | 0.76252 (7) | 0.0538 (5) |
| C4A | 0.82872 (11) | 0.29654 (13) | 0.84973 (7) | 0.0542 (5) |
| C4AA | 0.74920 (10) | 0.29394 (11) | 0.83793 (6) | 0.0476 (4) |
| C5A | 0.69582 (11) | 0.30185 (13) | 0.87449 (7) | 0.0596 (5) |
| H5A | 0.7113 | 0.3091 | 0.9063 | 0.072* |
| C6A | 0.62102 (11) | 0.29916 (14) | 0.86422 (8) | 0.0639 (5) |
| H6A | 0.5859 | 0.3033 | 0.8888 | 0.077* |
| C7A | 0.59835 (11) | 0.29017 (13) | 0.81658 (8) | 0.0627 (5) |
| H7A | 0.5476 | 0.2894 | 0.8094 | 0.075* |
| C8A | 0.64939 (10) | 0.28232 (13) | 0.77977 (7) | 0.0563 (5) |

| | | | | |
|------|--------------|--------------|--------------|------------|
| H8A | 0.6332 | 0.2767 | 0.7480 | 0.068* |
| C8AA | 0.72575 (9) | 0.28287 (11) | 0.79034 (6) | 0.0454 (4) |
| C9A | 0.75706 (10) | 0.24198 (13) | 0.70537 (6) | 0.0531 (4) |
| H9AA | 0.7164 | 0.1965 | 0.7084 | 0.064* |
| H9AB | 0.7988 | 0.2089 | 0.6906 | 0.064* |
| C10A | 0.73253 (9) | 0.32101 (13) | 0.67192 (6) | 0.0480 (4) |
| C11A | 0.76693 (11) | 0.40913 (13) | 0.67155 (7) | 0.0583 (5) |
| H11A | 0.8039 | 0.4228 | 0.6941 | 0.070* |
| C12A | 0.74673 (12) | 0.47754 (15) | 0.63777 (8) | 0.0722 (6) |
| H12A | 0.7701 | 0.5368 | 0.6377 | 0.087* |
| C13A | 0.69229 (13) | 0.45750 (18) | 0.60452 (8) | 0.0775 (7) |
| H13A | 0.6785 | 0.5035 | 0.5820 | 0.093* |
| C14A | 0.65844 (13) | 0.37067 (19) | 0.60429 (8) | 0.0801 (7) |
| H14A | 0.6220 | 0.3572 | 0.5814 | 0.096* |
| C15A | 0.67784 (11) | 0.30240 (16) | 0.63795 (7) | 0.0671 (5) |
| H15A | 0.6539 | 0.2434 | 0.6378 | 0.081* |
| C16A | 0.95826 (10) | 0.29432 (13) | 0.81973 (8) | 0.0627 (5) |
| H16A | 0.9681 | 0.3332 | 0.8480 | 0.075* |
| H16C | 0.9834 | 0.3235 | 0.7925 | 0.075* |
| C17A | 0.98951 (9) | 0.19523 (13) | 0.82771 (7) | 0.0530 (4) |
| C18A | 1.00645 (10) | 0.13649 (14) | 0.78927 (8) | 0.0617 (5) |
| H18A | 1.0011 | 0.1593 | 0.7579 | 0.074* |
| C19A | 1.03111 (11) | 0.04455 (16) | 0.79674 (9) | 0.0748 (6) |
| H19A | 1.0409 | 0.0049 | 0.7706 | 0.090* |
| C20A | 1.04120 (12) | 0.01171 (18) | 0.84297 (10) | 0.0873 (7) |
| H20A | 1.0579 | -0.0503 | 0.8481 | 0.105* |
| C21A | 1.02667 (12) | 0.07020 (18) | 0.88159 (9) | 0.0853 (7) |
| H21A | 1.0346 | 0.0482 | 0.9128 | 0.102* |
| C22A | 1.00042 (10) | 0.16140 (16) | 0.87410 (7) | 0.0684 (6) |
| H22A | 0.9900 | 0.2005 | 0.9004 | 0.082* |
| O1B | 0.61180 (7) | 0.06448 (9) | 0.98324 (5) | 0.0633 (4) |
| O2B | 0.54505 (7) | 0.03102 (10) | 0.82607 (5) | 0.0668 (4) |
| N1B | 0.70512 (7) | 0.06323 (10) | 0.92718 (5) | 0.0467 (3) |
| N3B | 0.57889 (7) | 0.04627 (9) | 0.90467 (5) | 0.0455 (3) |
| C2B | 0.63124 (9) | 0.05873 (12) | 0.94108 (7) | 0.0480 (4) |
| C4AB | 0.67396 (9) | 0.04264 (11) | 0.84349 (6) | 0.0444 (4) |
| C4B | 0.59496 (9) | 0.03999 (12) | 0.85581 (6) | 0.0476 (4) |
| C5B | 0.69580 (10) | 0.03363 (12) | 0.79517 (6) | 0.0542 (5) |
| H5B | 0.6597 | 0.0278 | 0.7712 | 0.065* |
| C6B | 0.76984 (11) | 0.03330 (13) | 0.78293 (7) | 0.0599 (5) |
| H6B | 0.7842 | 0.0264 | 0.7509 | 0.072* |
| C7B | 0.82289 (10) | 0.04332 (13) | 0.81853 (7) | 0.0576 (5) |
| H7B | 0.8732 | 0.0429 | 0.8102 | 0.069* |
| C8AB | 0.72776 (9) | 0.05321 (11) | 0.87922 (6) | 0.0435 (4) |
| C8B | 0.80291 (9) | 0.05388 (12) | 0.86611 (7) | 0.0533 (4) |
| H8B | 0.8396 | 0.0615 | 0.8896 | 0.064* |
| C9B | 0.76085 (10) | 0.07851 (12) | 0.96557 (6) | 0.0545 (5) |
| H9B | 0.7359 | 0.1041 | 0.9938 | 0.065* |

| | | | | |
|------|--------------|---------------|-------------|------------|
| H9D | 0.7968 | 0.1257 | 0.9547 | 0.065* |
| C10B | 0.80190 (10) | -0.01121 (13) | 0.97963 (6) | 0.0498 (4) |
| C11B | 0.76367 (11) | -0.08904 (13) | 0.99740 (7) | 0.0588 (5) |
| H11B | 0.7120 | -0.0862 | 1.0005 | 0.071* |
| C12B | 0.80122 (13) | -0.17137 (15) | 1.01063 (8) | 0.0711 (6) |
| H12B | 0.7749 | -0.2229 | 1.0233 | 0.085* |
| C13B | 0.87748 (13) | -0.17699 (17) | 1.00508 (8) | 0.0769 (6) |
| H13B | 0.9026 | -0.2329 | 1.0131 | 0.092* |
| C14B | 0.91636 (12) | -0.09989 (17) | 0.98765 (8) | 0.0736 (6) |
| H14B | 0.9679 | -0.1035 | 0.9840 | 0.088* |
| C15B | 0.87924 (10) | -0.01730 (15) | 0.97548 (7) | 0.0617 (5) |
| H15B | 0.9061 | 0.0351 | 0.9644 | 0.074* |
| C16B | 0.50000 (9) | 0.03632 (12) | 0.91967 (7) | 0.0511 (4) |
| H16B | 0.4981 | -0.0001 | 0.9494 | 0.061* |
| H16D | 0.4733 | 0.0004 | 0.8952 | 0.061* |
| C17B | 0.46085 (8) | 0.13002 (12) | 0.92738 (6) | 0.0458 (4) |
| C18B | 0.44536 (10) | 0.16354 (14) | 0.97336 (7) | 0.0588 (5) |
| H18B | 0.4611 | 0.1286 | 1.0000 | 0.071* |
| C19B | 0.40708 (11) | 0.24777 (15) | 0.98023 (8) | 0.0698 (6) |
| H19B | 0.3966 | 0.2689 | 1.0113 | 0.084* |
| C20B | 0.38446 (11) | 0.30048 (14) | 0.94116 (8) | 0.0667 (6) |
| H20B | 0.3591 | 0.3577 | 0.9456 | 0.080* |
| C21B | 0.39945 (10) | 0.26832 (14) | 0.89542 (8) | 0.0636 (5) |
| H21B | 0.3841 | 0.3040 | 0.8689 | 0.076* |
| C22B | 0.43716 (10) | 0.18334 (13) | 0.88838 (7) | 0.0546 (4) |
| H22B | 0.4466 | 0.1620 | 0.8572 | 0.066* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| O1A | 0.0545 (8) | 0.1060 (12) | 0.0629 (9) | 0.0081 (8) | 0.0042 (7) | 0.0195 (8) |
| O2A | 0.0782 (10) | 0.0927 (11) | 0.0600 (8) | 0.0061 (8) | -0.0186 (8) | -0.0133 (8) |
| N1A | 0.0497 (8) | 0.0526 (8) | 0.0470 (8) | -0.0007 (7) | -0.0019 (7) | 0.0021 (7) |
| N3A | 0.0485 (8) | 0.0514 (9) | 0.0592 (9) | -0.0013 (7) | -0.0091 (7) | 0.0050 (7) |
| C2A | 0.0508 (10) | 0.0543 (11) | 0.0564 (11) | 0.0043 (8) | -0.0011 (9) | 0.0129 (9) |
| C4A | 0.0613 (11) | 0.0454 (10) | 0.0558 (11) | 0.0007 (9) | -0.0080 (10) | -0.0016 (9) |
| C4AA | 0.0562 (10) | 0.0357 (8) | 0.0509 (10) | 0.0001 (8) | -0.0025 (8) | -0.0001 (8) |
| C5A | 0.0755 (13) | 0.0490 (11) | 0.0544 (11) | -0.0009 (10) | 0.0017 (10) | -0.0074 (9) |
| C6A | 0.0621 (12) | 0.0599 (12) | 0.0699 (13) | -0.0041 (10) | 0.0134 (11) | -0.0085 (10) |
| C7A | 0.0521 (11) | 0.0594 (12) | 0.0765 (14) | -0.0048 (9) | 0.0008 (10) | -0.0087 (11) |
| C8A | 0.0519 (10) | 0.0581 (11) | 0.0589 (11) | -0.0027 (9) | -0.0024 (9) | -0.0011 (10) |
| C8AA | 0.0496 (9) | 0.0364 (8) | 0.0501 (10) | -0.0015 (7) | 0.0003 (8) | 0.0025 (8) |
| C9A | 0.0567 (10) | 0.0526 (10) | 0.0500 (10) | 0.0017 (9) | -0.0012 (9) | -0.0030 (9) |
| C10A | 0.0447 (9) | 0.0556 (10) | 0.0436 (9) | 0.0055 (8) | 0.0024 (8) | -0.0033 (8) |
| C11A | 0.0571 (11) | 0.0617 (12) | 0.0561 (11) | 0.0052 (9) | -0.0041 (9) | -0.0008 (10) |
| C12A | 0.0811 (15) | 0.0588 (12) | 0.0766 (14) | 0.0096 (11) | 0.0074 (12) | 0.0109 (11) |
| C13A | 0.0798 (15) | 0.0928 (17) | 0.0599 (13) | 0.0336 (14) | 0.0013 (12) | 0.0178 (13) |
| C14A | 0.0747 (15) | 0.1002 (18) | 0.0653 (14) | 0.0219 (14) | -0.0215 (12) | -0.0031 (14) |

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|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C15A | 0.0579 (11) | 0.0750 (14) | 0.0684 (13) | 0.0040 (10) | -0.0115 (10) | -0.0061 (11) |
| C16A | 0.0493 (10) | 0.0628 (12) | 0.0758 (14) | -0.0107 (9) | -0.0113 (10) | 0.0083 (10) |
| C17A | 0.0356 (8) | 0.0619 (11) | 0.0616 (11) | -0.0065 (8) | -0.0052 (8) | 0.0112 (10) |
| C18A | 0.0470 (10) | 0.0761 (14) | 0.0621 (12) | -0.0012 (10) | 0.0051 (9) | 0.0156 (11) |
| C19A | 0.0567 (12) | 0.0811 (15) | 0.0867 (16) | 0.0119 (11) | 0.0168 (11) | 0.0081 (13) |
| C20A | 0.0631 (14) | 0.0857 (17) | 0.113 (2) | 0.0246 (12) | 0.0145 (14) | 0.0336 (16) |
| C21A | 0.0695 (14) | 0.110 (2) | 0.0764 (16) | 0.0181 (14) | 0.0016 (12) | 0.0386 (15) |
| C22A | 0.0568 (12) | 0.0878 (15) | 0.0607 (12) | 0.0031 (11) | -0.0043 (10) | 0.0106 (12) |
| O1B | 0.0576 (8) | 0.0803 (9) | 0.0520 (8) | 0.0075 (7) | 0.0021 (6) | 0.0003 (7) |
| O2B | 0.0490 (7) | 0.0854 (10) | 0.0661 (8) | -0.0013 (7) | -0.0096 (7) | -0.0101 (7) |
| N1B | 0.0412 (7) | 0.0463 (8) | 0.0525 (8) | 0.0014 (6) | -0.0047 (6) | -0.0038 (7) |
| N3B | 0.0388 (7) | 0.0430 (8) | 0.0545 (9) | 0.0020 (6) | -0.0008 (6) | 0.0011 (7) |
| C2B | 0.0468 (9) | 0.0414 (9) | 0.0558 (11) | 0.0049 (8) | -0.0025 (8) | 0.0012 (8) |
| C4AB | 0.0451 (9) | 0.0342 (8) | 0.0539 (10) | 0.0011 (7) | -0.0002 (8) | -0.0014 (8) |
| C4B | 0.0430 (9) | 0.0424 (9) | 0.0573 (11) | 0.0001 (8) | -0.0052 (8) | -0.0026 (8) |
| C5B | 0.0563 (11) | 0.0509 (10) | 0.0556 (11) | 0.0017 (9) | -0.0021 (9) | -0.0032 (9) |
| C6B | 0.0647 (12) | 0.0540 (11) | 0.0610 (12) | 0.0016 (9) | 0.0103 (10) | 0.0006 (10) |
| C7B | 0.0469 (10) | 0.0493 (10) | 0.0766 (14) | -0.0005 (8) | 0.0122 (9) | 0.0075 (10) |
| C8AB | 0.0410 (9) | 0.0329 (8) | 0.0567 (10) | -0.0001 (7) | -0.0021 (8) | 0.0006 (8) |
| C8B | 0.0426 (9) | 0.0503 (10) | 0.0670 (12) | -0.0011 (8) | -0.0036 (9) | 0.0051 (9) |
| C9B | 0.0515 (10) | 0.0515 (10) | 0.0605 (11) | 0.0012 (8) | -0.0098 (9) | -0.0124 (9) |
| C10B | 0.0493 (10) | 0.0514 (10) | 0.0486 (10) | 0.0019 (8) | -0.0101 (8) | -0.0098 (8) |
| C11B | 0.0569 (11) | 0.0610 (12) | 0.0584 (11) | 0.0023 (9) | -0.0048 (9) | -0.0066 (10) |
| C12B | 0.0818 (15) | 0.0609 (13) | 0.0704 (13) | 0.0033 (11) | -0.0094 (12) | 0.0060 (11) |
| C13B | 0.0835 (16) | 0.0728 (15) | 0.0745 (15) | 0.0234 (13) | -0.0169 (13) | 0.0020 (12) |
| C14B | 0.0555 (12) | 0.0895 (16) | 0.0759 (14) | 0.0186 (12) | -0.0117 (11) | 0.0002 (13) |
| C15B | 0.0499 (11) | 0.0696 (13) | 0.0654 (12) | 0.0003 (10) | -0.0100 (9) | -0.0030 (10) |
| C16B | 0.0393 (9) | 0.0478 (10) | 0.0662 (12) | -0.0032 (8) | 0.0020 (8) | 0.0087 (9) |
| C17B | 0.0338 (8) | 0.0465 (10) | 0.0570 (10) | -0.0023 (7) | 0.0011 (8) | 0.0081 (8) |
| C18B | 0.0556 (11) | 0.0633 (12) | 0.0574 (12) | 0.0069 (9) | 0.0038 (9) | 0.0122 (10) |
| C19B | 0.0674 (13) | 0.0741 (14) | 0.0679 (14) | 0.0142 (11) | 0.0075 (11) | -0.0061 (12) |
| C20B | 0.0549 (11) | 0.0543 (12) | 0.0908 (16) | 0.0124 (9) | 0.0031 (11) | -0.0005 (11) |
| C21B | 0.0558 (11) | 0.0586 (12) | 0.0764 (14) | 0.0083 (9) | -0.0088 (10) | 0.0154 (11) |
| C22B | 0.0502 (10) | 0.0575 (11) | 0.0561 (11) | 0.0037 (9) | -0.0032 (9) | 0.0078 (9) |

Geometric parameters (Å, °)

| | | | |
|-----------|-----------|-----------|-------------|
| O1A—C2A | 1.221 (2) | O1B—C2B | 1.222 (2) |
| O2A—C4A | 1.220 (2) | O2B—C4B | 1.2222 (19) |
| N1A—C2A | 1.368 (2) | N1B—C2B | 1.379 (2) |
| N1A—C8AA | 1.397 (2) | N1B—C8AB | 1.397 (2) |
| N1A—C9A | 1.474 (2) | N1B—C9B | 1.474 (2) |
| N3A—C4A | 1.389 (2) | N3B—C4B | 1.387 (2) |
| N3A—C2A | 1.399 (2) | N3B—C2B | 1.388 (2) |
| N3A—C16A | 1.478 (2) | N3B—C16B | 1.479 (2) |
| C4A—C4AA | 1.461 (2) | C4AB—C8AB | 1.389 (2) |
| C4AA—C8AA | 1.393 (2) | C4AB—C5B | 1.401 (2) |
| C4AA—C5A | 1.397 (2) | C4AB—C4B | 1.455 (2) |

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|--------------|-------------|---------------|-------------|
| C5A—C6A | 1.369 (3) | C5B—C6B | 1.368 (2) |
| C5A—H5A | 0.9300 | C5B—H5B | 0.9300 |
| C6A—C7A | 1.387 (3) | C6B—C7B | 1.377 (3) |
| C6A—H6A | 0.9300 | C6B—H6B | 0.9300 |
| C7A—C8A | 1.374 (2) | C7B—C8B | 1.375 (2) |
| C7A—H7A | 0.9300 | C7B—H7B | 0.9300 |
| C8A—C8AA | 1.398 (2) | C8AB—C8B | 1.393 (2) |
| C8A—H8A | 0.9300 | C8B—H8B | 0.9300 |
| C9A—C10A | 1.510 (2) | C9B—C10B | 1.507 (2) |
| C9A—H9AA | 0.9700 | C9B—H9B | 0.9700 |
| C9A—H9AB | 0.9700 | C9B—H9D | 0.9700 |
| C10A—C11A | 1.379 (2) | C10B—C11B | 1.378 (2) |
| C10A—C15A | 1.383 (2) | C10B—C15B | 1.392 (2) |
| C11A—C12A | 1.388 (3) | C11B—C12B | 1.384 (3) |
| C11A—H11A | 0.9300 | C11B—H11B | 0.9300 |
| C12A—C13A | 1.370 (3) | C12B—C13B | 1.376 (3) |
| C12A—H12A | 0.9300 | C12B—H12B | 0.9300 |
| C13A—C14A | 1.359 (3) | C13B—C14B | 1.373 (3) |
| C13A—H13A | 0.9300 | C13B—H13B | 0.9300 |
| C14A—C15A | 1.380 (3) | C14B—C15B | 1.376 (3) |
| C14A—H14A | 0.9300 | C14B—H14B | 0.9300 |
| C15A—H15A | 0.9300 | C15B—H15B | 0.9300 |
| C16A—C17A | 1.513 (2) | C16B—C17B | 1.503 (2) |
| C16A—H16A | 0.9700 | C16B—H16B | 0.9700 |
| C16A—H16C | 0.9700 | C16B—H16D | 0.9700 |
| C17A—C18A | 1.380 (3) | C17B—C22B | 1.381 (2) |
| C17A—C22A | 1.384 (3) | C17B—C18B | 1.386 (2) |
| C18A—C19A | 1.377 (3) | C18B—C19B | 1.378 (2) |
| C18A—H18A | 0.9300 | C18B—H18B | 0.9300 |
| C19A—C20A | 1.374 (3) | C19B—C20B | 1.372 (3) |
| C19A—H19A | 0.9300 | C19B—H19B | 0.9300 |
| C20A—C21A | 1.373 (3) | C20B—C21B | 1.372 (3) |
| C20A—H20A | 0.9300 | C20B—H20B | 0.9300 |
| C21A—C22A | 1.377 (3) | C21B—C22B | 1.382 (2) |
| C21A—H21A | 0.9300 | C21B—H21B | 0.9300 |
| C22A—H22A | 0.9300 | C22B—H22B | 0.9300 |
| | | | |
| C2A—N1A—C8AA | 123.03 (15) | C2B—N1B—C8AB | 122.66 (14) |
| C2A—N1A—C9A | 116.58 (15) | C2B—N1B—C9B | 116.98 (14) |
| C8AA—N1A—C9A | 120.11 (14) | C8AB—N1B—C9B | 120.36 (14) |
| C4A—N3A—C2A | 124.81 (15) | C4B—N3B—C2B | 125.30 (14) |
| C4A—N3A—C16A | 118.42 (15) | C4B—N3B—C16B | 117.80 (14) |
| C2A—N3A—C16A | 116.68 (16) | C2B—N3B—C16B | 116.87 (14) |
| O1A—C2A—N1A | 122.15 (17) | O1B—C2B—N1B | 122.51 (16) |
| O1A—C2A—N3A | 120.93 (17) | O1B—C2B—N3B | 120.76 (16) |
| N1A—C2A—N3A | 116.91 (16) | N1B—C2B—N3B | 116.73 (15) |
| O2A—C4A—N3A | 120.45 (18) | C8AB—C4AB—C5B | 119.85 (16) |
| O2A—C4A—C4AA | 124.14 (18) | C8AB—C4AB—C4B | 120.59 (16) |

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|----------------|-------------|----------------|-------------|
| N3A—C4A—C4AA | 115.41 (16) | C5B—C4AB—C4B | 119.56 (16) |
| C8AA—C4AA—C5A | 119.31 (16) | O2B—C4B—N3B | 120.91 (15) |
| C8AA—C4AA—C4A | 120.56 (16) | O2B—C4B—C4AB | 123.68 (17) |
| C5A—C4AA—C4A | 120.13 (16) | N3B—C4B—C4AB | 115.40 (15) |
| C6A—C5A—C4AA | 121.04 (18) | C6B—C5B—C4AB | 120.52 (17) |
| C6A—C5A—H5A | 119.5 | C6B—C5B—H5B | 119.7 |
| C4AA—C5A—H5A | 119.5 | C4AB—C5B—H5B | 119.7 |
| C5A—C6A—C7A | 119.11 (19) | C5B—C6B—C7B | 119.34 (18) |
| C5A—C6A—H6A | 120.4 | C5B—C6B—H6B | 120.3 |
| C7A—C6A—H6A | 120.4 | C7B—C6B—H6B | 120.3 |
| C8A—C7A—C6A | 121.32 (18) | C8B—C7B—C6B | 121.30 (18) |
| C8A—C7A—H7A | 119.3 | C8B—C7B—H7B | 119.4 |
| C6A—C7A—H7A | 119.3 | C6B—C7B—H7B | 119.4 |
| C7A—C8A—C8AA | 119.60 (18) | C4AB—C8AB—C8B | 118.93 (16) |
| C7A—C8A—H8A | 120.2 | C4AB—C8AB—N1B | 119.22 (15) |
| C8AA—C8A—H8A | 120.2 | C8B—C8AB—N1B | 121.84 (15) |
| C4AA—C8AA—N1A | 118.95 (15) | C7B—C8B—C8AB | 120.05 (17) |
| C4AA—C8AA—C8A | 119.60 (17) | C7B—C8B—H8B | 120.0 |
| N1A—C8AA—C8A | 121.45 (16) | C8AB—C8B—H8B | 120.0 |
| N1A—C9A—C10A | 115.43 (14) | N1B—C9B—C10B | 113.29 (14) |
| N1A—C9A—H9AA | 108.4 | N1B—C9B—H9B | 108.9 |
| C10A—C9A—H9AA | 108.4 | C10B—C9B—H9B | 108.9 |
| N1A—C9A—H9AB | 108.4 | N1B—C9B—H9D | 108.9 |
| C10A—C9A—H9AB | 108.4 | C10B—C9B—H9D | 108.9 |
| H9AA—C9A—H9AB | 107.5 | H9B—C9B—H9D | 107.7 |
| C11A—C10A—C15A | 118.66 (18) | C11B—C10B—C15B | 118.35 (17) |
| C11A—C10A—C9A | 122.06 (16) | C11B—C10B—C9B | 120.66 (16) |
| C15A—C10A—C9A | 119.08 (17) | C15B—C10B—C9B | 120.98 (17) |
| C10A—C11A—C12A | 120.43 (19) | C10B—C11B—C12B | 120.85 (19) |
| C10A—C11A—H11A | 119.8 | C10B—C11B—H11B | 119.6 |
| C12A—C11A—H11A | 119.8 | C12B—C11B—H11B | 119.6 |
| C13A—C12A—C11A | 119.9 (2) | C13B—C12B—C11B | 120.0 (2) |
| C13A—C12A—H12A | 120.1 | C13B—C12B—H12B | 120.0 |
| C11A—C12A—H12A | 120.1 | C11B—C12B—H12B | 120.0 |
| C14A—C13A—C12A | 120.2 (2) | C14B—C13B—C12B | 119.8 (2) |
| C14A—C13A—H13A | 119.9 | C14B—C13B—H13B | 120.1 |
| C12A—C13A—H13A | 119.9 | C12B—C13B—H13B | 120.1 |
| C13A—C14A—C15A | 120.3 (2) | C13B—C14B—C15B | 120.2 (2) |
| C13A—C14A—H14A | 119.8 | C13B—C14B—H14B | 119.9 |
| C15A—C14A—H14A | 119.8 | C15B—C14B—H14B | 119.9 |
| C14A—C15A—C10A | 120.5 (2) | C14B—C15B—C10B | 120.8 (2) |
| C14A—C15A—H15A | 119.7 | C14B—C15B—H15B | 119.6 |
| C10A—C15A—H15A | 119.7 | C10B—C15B—H15B | 119.6 |
| N3A—C16A—C17A | 111.85 (14) | N3B—C16B—C17B | 113.76 (13) |
| N3A—C16A—H16A | 109.2 | N3B—C16B—H16B | 108.8 |
| C17A—C16A—H16A | 109.2 | C17B—C16B—H16B | 108.8 |
| N3A—C16A—H16C | 109.2 | N3B—C16B—H16D | 108.8 |
| C17A—C16A—H16C | 109.2 | C17B—C16B—H16D | 108.8 |

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|-------------------|--------------|-------------------|--------------|
| H16A—C16A—H16C | 107.9 | H16B—C16B—H16D | 107.7 |
| C18A—C17A—C22A | 118.84 (19) | C22B—C17B—C18B | 118.39 (16) |
| C18A—C17A—C16A | 121.02 (17) | C22B—C17B—C16B | 120.26 (16) |
| C22A—C17A—C16A | 120.13 (19) | C18B—C17B—C16B | 121.31 (16) |
| C19A—C18A—C17A | 120.79 (19) | C19B—C18B—C17B | 121.10 (18) |
| C19A—C18A—H18A | 119.6 | C19B—C18B—H18B | 119.5 |
| C17A—C18A—H18A | 119.6 | C17B—C18B—H18B | 119.5 |
| C20A—C19A—C18A | 119.7 (2) | C20B—C19B—C18B | 119.9 (2) |
| C20A—C19A—H19A | 120.1 | C20B—C19B—H19B | 120.0 |
| C18A—C19A—H19A | 120.1 | C18B—C19B—H19B | 120.0 |
| C21A—C20A—C19A | 120.2 (2) | C19B—C20B—C21B | 119.68 (18) |
| C21A—C20A—H20A | 119.9 | C19B—C20B—H20B | 120.2 |
| C19A—C20A—H20A | 119.9 | C21B—C20B—H20B | 120.2 |
| C20A—C21A—C22A | 120.1 (2) | C20B—C21B—C22B | 120.58 (19) |
| C20A—C21A—H21A | 120.0 | C20B—C21B—H21B | 119.7 |
| C22A—C21A—H21A | 120.0 | C22B—C21B—H21B | 119.7 |
| C21A—C22A—C17A | 120.4 (2) | C17B—C22B—C21B | 120.34 (18) |
| C21A—C22A—H22A | 119.8 | C17B—C22B—H22B | 119.8 |
| C17A—C22A—H22A | 119.8 | C21B—C22B—H22B | 119.8 |
| | | | |
| C8AA—N1A—C2A—O1A | -175.32 (16) | C8AB—N1B—C2B—O1B | 177.61 (15) |
| C9A—N1A—C2A—O1A | 10.7 (3) | C9B—N1B—C2B—O1B | -1.9 (2) |
| C8AA—N1A—C2A—N3A | 5.7 (3) | C8AB—N1B—C2B—N3B | -1.8 (2) |
| C9A—N1A—C2A—N3A | -168.28 (14) | C9B—N1B—C2B—N3B | 178.73 (13) |
| C4A—N3A—C2A—O1A | -179.83 (17) | C4B—N3B—C2B—O1B | 179.24 (16) |
| C16A—N3A—C2A—O1A | -3.4 (3) | C16B—N3B—C2B—O1B | -2.5 (2) |
| C4A—N3A—C2A—N1A | -0.8 (3) | C4B—N3B—C2B—N1B | -1.4 (2) |
| C16A—N3A—C2A—N1A | 175.60 (15) | C16B—N3B—C2B—N1B | 176.87 (13) |
| C2A—N3A—C4A—O2A | 177.49 (17) | C2B—N3B—C4B—O2B | -178.43 (16) |
| C16A—N3A—C4A—O2A | 1.1 (3) | C16B—N3B—C4B—O2B | 3.4 (2) |
| C2A—N3A—C4A—C4AA | -3.4 (3) | C2B—N3B—C4B—C4AB | 2.9 (2) |
| C16A—N3A—C4A—C4AA | -179.81 (14) | C16B—N3B—C4B—C4AB | -175.28 (13) |
| O2A—C4A—C4AA—C8AA | -177.75 (17) | C8AB—C4AB—C4B—O2B | 179.88 (16) |
| N3A—C4A—C4AA—C8AA | 3.2 (2) | C5B—C4AB—C4B—O2B | -0.7 (3) |
| O2A—C4A—C4AA—C5A | 1.8 (3) | C8AB—C4AB—C4B—N3B | -1.5 (2) |
| N3A—C4A—C4AA—C5A | -177.27 (15) | C5B—C4AB—C4B—N3B | 177.94 (15) |
| C8AA—C4AA—C5A—C6A | -0.2 (3) | C8AB—C4AB—C5B—C6B | 1.3 (3) |
| C4A—C4AA—C5A—C6A | -179.71 (17) | C4B—C4AB—C5B—C6B | -178.20 (16) |
| C4AA—C5A—C6A—C7A | -1.2 (3) | C4AB—C5B—C6B—C7B | -0.9 (3) |
| C5A—C6A—C7A—C8A | 1.1 (3) | C5B—C6B—C7B—C8B | -0.2 (3) |
| C6A—C7A—C8A—C8AA | 0.3 (3) | C5B—C4AB—C8AB—C8B | -0.5 (2) |
| C5A—C4AA—C8AA—N1A | -178.40 (15) | C4B—C4AB—C8AB—C8B | 178.98 (15) |
| C4A—C4AA—C8AA—N1A | 1.1 (2) | C5B—C4AB—C8AB—N1B | 179.26 (15) |
| C5A—C4AA—C8AA—C8A | 1.6 (3) | C4B—C4AB—C8AB—N1B | -1.3 (2) |
| C4A—C4AA—C8AA—C8A | -178.84 (16) | C2B—N1B—C8AB—C4AB | 3.1 (2) |
| C2A—N1A—C8AA—C4AA | -5.9 (2) | C9B—N1B—C8AB—C4AB | -177.48 (14) |
| C9A—N1A—C8AA—C4AA | 167.91 (15) | C2B—N1B—C8AB—C8B | -177.20 (15) |
| C2A—N1A—C8AA—C8A | 174.10 (16) | C9B—N1B—C8AB—C8B | 2.3 (2) |

| | | | |
|---------------------|--------------|---------------------|--------------|
| C9A—N1A—C8AA—C8A | -12.1 (2) | C6B—C7B—C8B—C8AB | 0.9 (3) |
| C7A—C8A—C8AA—C4AA | -1.7 (3) | C4AB—C8AB—C8B—C7B | -0.6 (2) |
| C7A—C8A—C8AA—N1A | 178.33 (16) | N1B—C8AB—C8B—C7B | 179.66 (15) |
| C2A—N1A—C9A—C10A | -99.51 (18) | C2B—N1B—C9B—C10B | 102.55 (18) |
| C8AA—N1A—C9A—C10A | 86.33 (19) | C8AB—N1B—C9B—C10B | -76.95 (19) |
| N1A—C9A—C10A—C11A | 38.0 (2) | N1B—C9B—C10B—C11B | -60.1 (2) |
| N1A—C9A—C10A—C15A | -147.14 (17) | N1B—C9B—C10B—C15B | 120.61 (18) |
| C15A—C10A—C11A—C12A | 0.1 (3) | C15B—C10B—C11B—C12B | -0.2 (3) |
| C9A—C10A—C11A—C12A | 174.99 (16) | C9B—C10B—C11B—C12B | -179.58 (17) |
| C10A—C11A—C12A—C13A | 0.0 (3) | C10B—C11B—C12B—C13B | -1.5 (3) |
| C11A—C12A—C13A—C14A | -0.5 (3) | C11B—C12B—C13B—C14B | 1.8 (3) |
| C12A—C13A—C14A—C15A | 0.9 (3) | C12B—C13B—C14B—C15B | -0.4 (3) |
| C13A—C14A—C15A—C10A | -0.8 (3) | C13B—C14B—C15B—C10B | -1.4 (3) |
| C11A—C10A—C15A—C14A | 0.3 (3) | C11B—C10B—C15B—C14B | 1.7 (3) |
| C9A—C10A—C15A—C14A | -174.73 (18) | C9B—C10B—C15B—C14B | -178.98 (17) |
| C4A—N3A—C16A—C17A | 88.4 (2) | C4B—N3B—C16B—C17B | -97.31 (18) |
| C2A—N3A—C16A—C17A | -88.3 (2) | C2B—N3B—C16B—C17B | 84.33 (18) |
| N3A—C16A—C17A—C18A | 81.3 (2) | N3B—C16B—C17B—C22B | 76.8 (2) |
| N3A—C16A—C17A—C22A | -97.6 (2) | N3B—C16B—C17B—C18B | -105.44 (19) |
| C22A—C17A—C18A—C19A | 2.4 (3) | C22B—C17B—C18B—C19B | 0.2 (3) |
| C16A—C17A—C18A—C19A | -176.51 (17) | C16B—C17B—C18B—C19B | -177.63 (17) |
| C17A—C18A—C19A—C20A | -2.0 (3) | C17B—C18B—C19B—C20B | -0.8 (3) |
| C18A—C19A—C20A—C21A | 0.1 (3) | C18B—C19B—C20B—C21B | 0.7 (3) |
| C19A—C20A—C21A—C22A | 1.4 (4) | C19B—C20B—C21B—C22B | 0.0 (3) |
| C20A—C21A—C22A—C17A | -0.9 (3) | C18B—C17B—C22B—C21B | 0.5 (3) |
| C18A—C17A—C22A—C21A | -1.0 (3) | C16B—C17B—C22B—C21B | 178.34 (16) |
| C16A—C17A—C22A—C21A | 177.99 (18) | C20B—C21B—C22B—C17B | -0.6 (3) |

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

Cg4 and Cg8 are the centroids of the C17A–C22A and C10B–C15B rings, respectively.

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--|-------|-------------|-------------|---------------|
| C19A—H19A \cdots O2b ⁱ | 0.93 | 2.70 | 3.419 (3) | 134 |
| C6B—H6B \cdots C11a ⁱⁱ | 0.93 | 2.89 | 3.604 (3) | 134 |
| C21B—H21B \cdots C11a ⁱⁱⁱ | 0.93 | 2.80 | 3.600 (3) | 145 |
| C19A—H19A \cdots O2b ⁱ | 0.93 | 2.70 | 3.419 (3) | 134 |
| C7B—H7B \cdots Cg4 | 0.93 | 2.78 | 3.586 (2) | 146 |
| C5A—H5A \cdots Cg8 ⁱⁱ | 0.93 | 2.90 | 3.641 (2) | 138 |

Symmetry codes: (i) $x+1/2, y, -z+3/2$; (ii) $-x+3/2, y-1/2, z$; (iii) $x-1/2, y, -z+3/2$.