

exo-4-[(1*H*-Benzimidazol-2-yl)methyl]-10-oxa-4-azatricyclo[5.2.1.0^{2,6}]decane-3,5-dione

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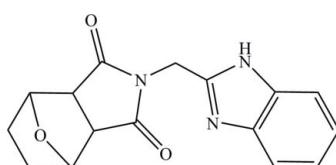
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.035; wR factor = 0.087; data-to-parameter ratio = 10.8.

In the title compound, $\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}_3$, the dihedral angle between the approximately planar benzimidazolyl group (r.m.s. deviation = 0.010 \AA) and the pyrrolidine ring is $78.20(6)^\circ$. The $\text{C}-\text{C}-\text{N}$ bond angle of the bridging CH_2 group is $112.14(16)^\circ$. In the crystal, molecules are linked via $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds, forming infinite chains parallel to [101] and [101].

Related literature

For the bioactivity of norcantharidin (systematic name 4,10-dioxatricyclo[5.2.1.0^{2,6}]decane-3,5-dione), see: Wang (1989). For the use of norcantharidin in synthesis, see: Hill *et al.* (2007). For a related structure, see: Zhu & Lin (2009).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}_3$

$M_r = 297.31$

Orthorhombic, $Fdd2$
 $a = 17.4294(2)\text{ \AA}$
 $b = 48.2746(6)\text{ \AA}$
 $c = 6.7947(1)\text{ \AA}$
 $V = 5717.04(13)\text{ \AA}^3$

$Z = 16$
Mo $K\alpha$ radiation
 $\mu = 0.10\text{ mm}^{-1}$
 $T = 296\text{ K}$
 $0.24 \times 0.17 \times 0.10\text{ mm}$

Data collection

Bruker SMART APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $R_{\text{int}} = 0.038$
 $T_{\text{min}} = 0.981$, $T_{\text{max}} = 0.990$

22741 measured reflections
2159 independent reflections
1690 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.087$
 $S = 1.04$
2159 reflections
199 parameters

1 restraint
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.10\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.17\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|---|--------------|--------------------|-------------|----------------------|
| N1—H1A \cdots N2 ⁱ | 0.86 | 1.97 | 2.827 (2) | 175 |
| Symmetry code: (i) $x + \frac{1}{4}, -y + \frac{1}{4}, z + \frac{1}{4}$. | | | | |

Data collection: *APEX2* (Bruker, 2006); cell refinement: *SAINT* (Bruker, 2006); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: QK2008).

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supporting information

Acta Cryst. (2011). E67, o1974 [doi:10.1107/S1600536811026602]

exo-4-[(1*H*-Benzimidazol-2-yl)methyl]-10-oxa-4-azatricyclo-[5.2.1.0^{2,6}]decane-3,5-dione

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S1. Comment

Norcantharidin is an important compound that has a variety of bioactivities, such as protein kinase inhibition and antitumor activity (Wang, 1989). As a contribution to structure-activity studies in this field, the title compound, (I), a norcantharidin imide (Hill *et al.*, 2007), was synthesized and investigated by X-ray crystallography. A related norcantharidin imide was reported by Zhu & Lin (2009).

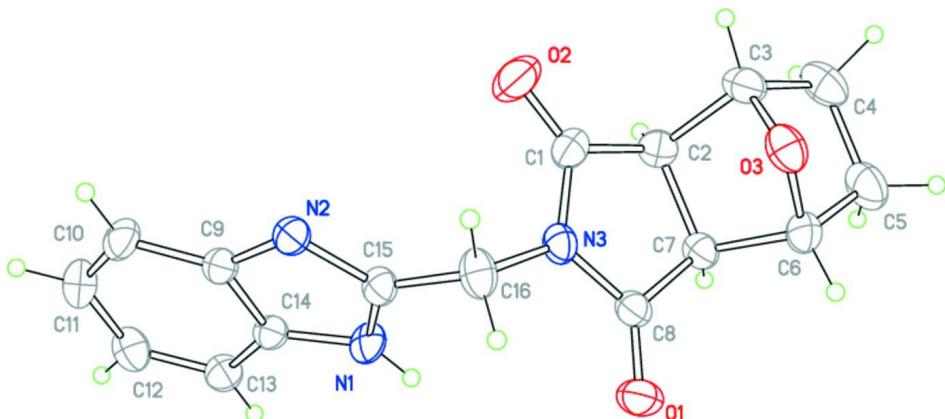
X-ray crystallography confirmed the anticipated molecular structure and the atom connectivity for the title compound, as illustrated in Fig. 1. The norcantharidin imide and the benzimidazole moiety of (I) are rigid and feature usual bond lengths and angles. The dihedral angle between the flat benzimidazolyl group (C9—C15,N1,N2) and the pyrrolidine ring (C1,C2,C7,C8,N3) is 78.20 (6)°. The bond angle of the bridging CH₂ group is C15—C16—N3 = 112.1 (2)°. In the crystal structure the molecules are linked *via* the hydrogen bond N1—H1A···N2ⁱ, N1···N2ⁱ = 2.827 (2) Å ((i): x+1/4, -y+1/4, z+1/4), to form infinite hydrogen bonded chains parallel to [101] and [10[−]1].

S2. Experimental

A mixture of 0.5 mmol norcantharidin, 0.5 mmol 2-(aminomethyl)benzimidazole, 0.5 mmol cadmium acetate as a promoter, and 10 mL distilled water was sealed in a 25 mL Teflon-lined stainless vessel and heated at 433 K for 3 d, then cooled slowly to room temperature. The solution was filtered and gave then block-like transparent crystals.

S3. Refinement

The structure was solved by direct methods and successive Fourier difference synthesis. The H atoms were positioned geometrically and refined using a riding model [aromatic C—H = 0.93 Å, aliphatic C—H = 0.97 Å, and N—H = 0.86 Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$]. In the absence of significant anomalous scattering, the Friedel pairs of the polar structure were merged.

**Figure 1**

A view of the molecule of (I) showing the atom-labelling scheme with displacement ellipsoids drawn at 30% probability.

exo-8-[(1*H*-Benzimidazol-2-yl)methyl]-10-oxa-8-azatricyclo[4.3.0.1^{2.5}]decane-7,9-dione

Crystal data

$C_{16}H_{15}N_3O_3$
 $M_r = 297.31$
Orthorhombic, $Fdd2$
Hall symbol: F 2 -2d
 $a = 17.4294 (2)$ Å
 $b = 48.2746 (6)$ Å
 $c = 6.7947 (1)$ Å
 $V = 5717.04 (13)$ Å³
 $Z = 16$

$F(000) = 2496$
 $D_x = 1.382$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 5637 reflections
 $\theta = 2.5\text{--}29.8^\circ$
 $\mu = 0.10$ mm⁻¹
 $T = 296$ K
Block, colourless
 $0.24 \times 0.17 \times 0.10$ mm

Data collection

Bruker SMART APEXII CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.990$

22741 measured reflections
2159 independent reflections
1690 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$
 $\theta_{\max} = 29.8^\circ$, $\theta_{\min} = 2.5^\circ$
 $h = -24 \rightarrow 24$
 $k = -67 \rightarrow 66$
 $l = -9 \rightarrow 8$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.035$
 $wR(F^2) = 0.087$
 $S = 1.04$
2159 reflections
199 parameters
1 restraint
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0448P)^2 + 1.1097P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.10$ e Å⁻³
 $\Delta\rho_{\min} = -0.17$ e Å⁻³
Absolute structure: 1753 Friedel pairs were
merged

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> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|------|---------------|-------------|-------------|----------------------------------|
| N2 | -0.06276 (8) | 0.10833 (3) | 0.2792 (3) | 0.0454 (4) |
| N1 | 0.04528 (8) | 0.11596 (3) | 0.4450 (3) | 0.0428 (4) |
| H1A | 0.0893 | 0.1229 | 0.4740 | 0.051* |
| N3 | -0.02809 (8) | 0.16452 (3) | 0.1037 (3) | 0.0431 (4) |
| O2 | -0.07386 (9) | 0.14915 (3) | -0.1930 (3) | 0.0629 (4) |
| O1 | -0.00312 (11) | 0.18963 (4) | 0.3811 (3) | 0.0765 (5) |
| O3 | -0.02560 (7) | 0.22474 (3) | -0.0973 (2) | 0.0556 (4) |
| C1 | -0.07375 (10) | 0.16595 (3) | -0.0624 (3) | 0.0439 (4) |
| C2 | -0.12089 (10) | 0.19215 (3) | -0.0513 (3) | 0.0431 (4) |
| H2B | -0.1761 | 0.1887 | -0.0634 | 0.052* |
| C3 | -0.09207 (12) | 0.21403 (4) | -0.1970 (3) | 0.0527 (5) |
| H3A | -0.0813 | 0.2068 | -0.3288 | 0.063* |
| C4 | -0.14630 (14) | 0.23883 (4) | -0.1956 (5) | 0.0713 (7) |
| H4A | -0.1995 | 0.2330 | -0.2016 | 0.086* |
| H4B | -0.1357 | 0.2513 | -0.3041 | 0.086* |
| C5 | -0.12741 (14) | 0.25219 (4) | 0.0024 (4) | 0.0700 (7) |
| H5A | -0.1088 | 0.2710 | -0.0139 | 0.084* |
| H5B | -0.1716 | 0.2523 | 0.0892 | 0.084* |
| C6 | -0.06469 (12) | 0.23307 (4) | 0.0784 (4) | 0.0552 (5) |
| H6A | -0.0311 | 0.2417 | 0.1763 | 0.066* |
| C7 | -0.09930 (11) | 0.20533 (4) | 0.1463 (3) | 0.0464 (5) |
| H7A | -0.1434 | 0.2077 | 0.2341 | 0.056* |
| C8 | -0.03878 (12) | 0.18644 (4) | 0.2306 (3) | 0.0492 (5) |
| C16 | 0.02831 (10) | 0.14281 (4) | 0.1375 (4) | 0.0519 (5) |
| H16A | 0.0760 | 0.1511 | 0.1814 | 0.062* |
| H16B | 0.0384 | 0.1333 | 0.0145 | 0.062* |
| C15 | 0.00198 (10) | 0.12236 (3) | 0.2871 (3) | 0.0410 (4) |
| C9 | -0.06167 (9) | 0.09153 (3) | 0.4452 (3) | 0.0391 (4) |
| C10 | -0.11496 (11) | 0.07221 (4) | 0.5106 (3) | 0.0488 (5) |
| H10A | -0.1597 | 0.0687 | 0.4402 | 0.059* |
| C11 | -0.09925 (12) | 0.05847 (4) | 0.6828 (4) | 0.0571 (5) |
| H11A | -0.1339 | 0.0453 | 0.7288 | 0.068* |
| C12 | -0.03280 (13) | 0.06372 (5) | 0.7903 (4) | 0.0600 (6) |
| H12A | -0.0247 | 0.0543 | 0.9078 | 0.072* |
| C13 | 0.02144 (11) | 0.08263 (4) | 0.7270 (3) | 0.0526 (5) |

| | | | | |
|------|--------------|-------------|------------|------------|
| H13A | 0.0660 | 0.0861 | 0.7984 | 0.063* |
| C14 | 0.00597 (10) | 0.09617 (3) | 0.5518 (3) | 0.0395 (4) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|--------------|--------------|
| N2 | 0.0332 (7) | 0.0520 (8) | 0.0510 (10) | -0.0018 (6) | -0.0111 (7) | 0.0068 (8) |
| N1 | 0.0308 (7) | 0.0445 (7) | 0.0532 (10) | -0.0010 (6) | -0.0124 (8) | -0.0003 (7) |
| N3 | 0.0394 (8) | 0.0408 (7) | 0.0490 (10) | 0.0022 (6) | -0.0058 (7) | 0.0065 (7) |
| O2 | 0.0768 (10) | 0.0467 (7) | 0.0652 (11) | 0.0086 (7) | -0.0172 (9) | -0.0142 (8) |
| O1 | 0.0876 (11) | 0.0950 (13) | 0.0468 (11) | 0.0081 (9) | -0.0214 (10) | -0.0072 (9) |
| O3 | 0.0472 (7) | 0.0489 (7) | 0.0708 (11) | -0.0026 (6) | 0.0051 (8) | 0.0100 (7) |
| C1 | 0.0451 (9) | 0.0351 (8) | 0.0514 (12) | -0.0021 (7) | -0.0074 (9) | 0.0015 (8) |
| C2 | 0.0380 (9) | 0.0387 (8) | 0.0525 (11) | 0.0014 (7) | -0.0062 (9) | -0.0002 (8) |
| C3 | 0.0624 (12) | 0.0469 (10) | 0.0489 (13) | 0.0087 (9) | -0.0070 (11) | 0.0047 (9) |
| C4 | 0.0740 (15) | 0.0457 (10) | 0.094 (2) | 0.0098 (10) | -0.0194 (15) | 0.0099 (13) |
| C5 | 0.0703 (13) | 0.0407 (9) | 0.099 (2) | 0.0100 (10) | 0.0017 (15) | -0.0043 (12) |
| C6 | 0.0576 (12) | 0.0427 (9) | 0.0654 (15) | -0.0041 (8) | -0.0085 (11) | -0.0086 (10) |
| C7 | 0.0447 (10) | 0.0481 (9) | 0.0466 (12) | 0.0029 (8) | 0.0035 (9) | -0.0034 (9) |
| C8 | 0.0493 (11) | 0.0568 (11) | 0.0414 (12) | -0.0011 (9) | -0.0015 (10) | 0.0022 (9) |
| C16 | 0.0386 (9) | 0.0511 (10) | 0.0660 (15) | 0.0073 (8) | -0.0002 (10) | 0.0147 (10) |
| C15 | 0.0324 (8) | 0.0416 (8) | 0.0490 (12) | 0.0040 (7) | -0.0080 (8) | 0.0041 (8) |
| C9 | 0.0334 (8) | 0.0393 (8) | 0.0444 (11) | 0.0033 (6) | -0.0079 (8) | -0.0022 (8) |
| C10 | 0.0394 (9) | 0.0502 (10) | 0.0570 (14) | -0.0041 (8) | -0.0060 (9) | -0.0017 (9) |
| C11 | 0.0535 (11) | 0.0506 (10) | 0.0671 (15) | -0.0001 (8) | 0.0027 (12) | 0.0114 (11) |
| C12 | 0.0610 (12) | 0.0633 (12) | 0.0558 (14) | 0.0089 (10) | -0.0030 (11) | 0.0175 (11) |
| C13 | 0.0470 (11) | 0.0628 (11) | 0.0480 (12) | 0.0076 (9) | -0.0156 (10) | 0.0025 (10) |
| C14 | 0.0333 (8) | 0.0388 (8) | 0.0463 (12) | 0.0042 (7) | -0.0079 (8) | -0.0021 (8) |

Geometric parameters (\AA , $^\circ$)

| | | | |
|--------|-----------|----------|-----------|
| N2—C15 | 1.317 (2) | C5—C6 | 1.521 (3) |
| N2—C9 | 1.389 (2) | C5—H5A | 0.9700 |
| N1—C15 | 1.348 (3) | C5—H5B | 0.9700 |
| N1—C14 | 1.381 (2) | C6—C7 | 1.540 (3) |
| N1—H1A | 0.8600 | C6—H6A | 0.9800 |
| N3—C8 | 1.378 (3) | C7—C8 | 1.507 (3) |
| N3—C1 | 1.383 (3) | C7—H7A | 0.9800 |
| N3—C16 | 1.455 (2) | C16—C15 | 1.490 (3) |
| O2—C1 | 1.202 (2) | C16—H16A | 0.9700 |
| O1—C8 | 1.206 (3) | C16—H16B | 0.9700 |
| O3—C6 | 1.432 (3) | C9—C10 | 1.389 (3) |
| O3—C3 | 1.438 (3) | C9—C14 | 1.402 (2) |
| C1—C2 | 1.510 (2) | C10—C11 | 1.373 (3) |
| C2—C3 | 1.532 (3) | C10—H10A | 0.9300 |
| C2—C7 | 1.533 (3) | C11—C12 | 1.393 (3) |
| C2—H2B | 0.9800 | C11—H11A | 0.9300 |
| C3—C4 | 1.525 (3) | C12—C13 | 1.383 (3) |

| | | | |
|--------------|-------------|---------------|-------------|
| C3—H3A | 0.9800 | C12—H12A | 0.9300 |
| C4—C5 | 1.528 (4) | C13—C14 | 1.385 (3) |
| C4—H4A | 0.9700 | C13—H13A | 0.9300 |
| C4—H4B | 0.9700 | | |
| | | | |
| C15—N2—C9 | 104.80 (15) | C5—C6—H6A | 113.7 |
| C15—N1—C14 | 107.39 (14) | C7—C6—H6A | 113.7 |
| C15—N1—H1A | 126.3 | C8—C7—C2 | 104.70 (15) |
| C14—N1—H1A | 126.3 | C8—C7—C6 | 111.46 (16) |
| C8—N3—C1 | 113.23 (15) | C2—C7—C6 | 101.23 (17) |
| C8—N3—C16 | 123.08 (17) | C8—C7—H7A | 112.9 |
| C1—N3—C16 | 123.63 (17) | C2—C7—H7A | 112.9 |
| C6—O3—C3 | 96.35 (15) | C6—C7—H7A | 112.9 |
| O2—C1—N3 | 124.71 (16) | O1—C8—N3 | 124.0 (2) |
| O2—C1—C2 | 126.96 (19) | O1—C8—C7 | 127.3 (2) |
| N3—C1—C2 | 108.33 (16) | N3—C8—C7 | 108.75 (17) |
| C1—C2—C3 | 111.50 (17) | N3—C16—C15 | 112.14 (16) |
| C1—C2—C7 | 104.94 (16) | N3—C16—H16A | 109.2 |
| C3—C2—C7 | 101.50 (14) | C15—C16—H16A | 109.2 |
| C1—C2—H2B | 112.7 | N3—C16—H16B | 109.2 |
| C3—C2—H2B | 112.7 | C15—C16—H16B | 109.2 |
| C7—C2—H2B | 112.7 | H16A—C16—H16B | 107.9 |
| O3—C3—C4 | 102.38 (16) | N2—C15—N1 | 113.22 (17) |
| O3—C3—C2 | 101.96 (16) | N2—C15—C16 | 125.23 (18) |
| C4—C3—C2 | 109.49 (19) | N1—C15—C16 | 121.52 (16) |
| O3—C3—H3A | 113.9 | C10—C9—N2 | 129.97 (16) |
| C4—C3—H3A | 113.9 | C10—C9—C14 | 120.28 (17) |
| C2—C3—H3A | 113.9 | N2—C9—C14 | 109.74 (15) |
| C3—C4—C5 | 101.72 (19) | C11—C10—C9 | 117.63 (18) |
| C3—C4—H4A | 111.4 | C11—C10—H10A | 121.2 |
| C5—C4—H4A | 111.4 | C9—C10—H10A | 121.2 |
| C3—C4—H4B | 111.4 | C10—C11—C12 | 121.7 (2) |
| C5—C4—H4B | 111.4 | C10—C11—H11A | 119.2 |
| H4A—C4—H4B | 109.3 | C12—C11—H11A | 119.2 |
| C6—C5—C4 | 101.42 (18) | C13—C12—C11 | 121.7 (2) |
| C6—C5—H5A | 111.5 | C13—C12—H12A | 119.1 |
| C4—C5—H5A | 111.5 | C11—C12—H12A | 119.1 |
| C6—C5—H5B | 111.5 | C12—C13—C14 | 116.48 (19) |
| C4—C5—H5B | 111.5 | C12—C13—H13A | 121.8 |
| H5A—C5—H5B | 109.3 | C14—C13—H13A | 121.8 |
| O3—C6—C5 | 103.2 (2) | N1—C14—C13 | 132.96 (17) |
| O3—C6—C7 | 101.06 (15) | N1—C14—C9 | 104.84 (16) |
| C5—C6—C7 | 110.35 (16) | C13—C14—C9 | 122.19 (18) |
| O3—C6—H6A | 113.7 | | |
| | | | |
| C8—N3—C1—O2 | 179.8 (2) | C16—N3—C8—O1 | -0.5 (3) |
| C16—N3—C1—O2 | 2.6 (3) | C1—N3—C8—C7 | 0.7 (2) |
| C8—N3—C1—C2 | 0.9 (2) | C16—N3—C8—C7 | 177.93 (16) |

| | | | |
|--------------|--------------|-----------------|--------------|
| C16—N3—C1—C2 | -176.28 (16) | C2—C7—C8—O1 | 176.4 (2) |
| O2—C1—C2—C3 | -71.9 (2) | C6—C7—C8—O1 | 67.8 (3) |
| N3—C1—C2—C3 | 106.99 (19) | C2—C7—C8—N3 | -2.0 (2) |
| O2—C1—C2—C7 | 179.06 (19) | C6—C7—C8—N3 | -110.60 (19) |
| N3—C1—C2—C7 | -2.10 (19) | C8—N3—C16—C15 | 77.7 (2) |
| C6—O3—C3—C4 | 55.9 (2) | C1—N3—C16—C15 | -105.4 (2) |
| C6—O3—C3—C2 | -57.41 (16) | C9—N2—C15—N1 | 0.1 (2) |
| C1—C2—C3—O3 | -77.96 (19) | C9—N2—C15—C16 | 178.43 (18) |
| C7—C2—C3—O3 | 33.32 (17) | C14—N1—C15—N2 | -0.4 (2) |
| C1—C2—C3—C4 | 174.13 (18) | C14—N1—C15—C16 | -178.74 (17) |
| C7—C2—C3—C4 | -74.6 (2) | N3—C16—C15—N2 | 55.2 (3) |
| O3—C3—C4—C5 | -35.0 (2) | N3—C16—C15—N1 | -126.70 (19) |
| C2—C3—C4—C5 | 72.7 (2) | C15—N2—C9—C10 | -179.28 (19) |
| C3—C4—C5—C6 | 1.0 (2) | C15—N2—C9—C14 | 0.14 (19) |
| C3—O3—C6—C5 | -55.61 (16) | N2—C9—C10—C11 | -179.60 (19) |
| C3—O3—C6—C7 | 58.61 (15) | C14—C9—C10—C11 | 1.0 (3) |
| C4—C5—C6—O3 | 33.5 (2) | C9—C10—C11—C12 | 0.6 (3) |
| C4—C5—C6—C7 | -73.8 (2) | C10—C11—C12—C13 | -1.3 (4) |
| C1—C2—C7—C8 | 2.40 (19) | C11—C12—C13—C14 | 0.4 (3) |
| C3—C2—C7—C8 | -113.79 (16) | C15—N1—C14—C13 | -178.2 (2) |
| C1—C2—C7—C6 | 118.36 (16) | C15—N1—C14—C9 | 0.44 (19) |
| C3—C2—C7—C6 | 2.16 (17) | C12—C13—C14—N1 | 179.8 (2) |
| O3—C6—C7—C8 | 73.7 (2) | C12—C13—C14—C9 | 1.3 (3) |
| C5—C6—C7—C8 | -177.5 (2) | C10—C9—C14—N1 | 179.13 (16) |
| O3—C6—C7—C2 | -37.14 (17) | N2—C9—C14—N1 | -0.36 (19) |
| C5—C6—C7—C2 | 71.6 (2) | C10—C9—C14—C13 | -2.0 (3) |
| C1—N3—C8—O1 | -177.7 (2) | N2—C9—C14—C13 | 178.48 (17) |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|--------------------------|------|-------|-----------|---------|
| N1—H1A···N2 ⁱ | 0.86 | 1.97 | 2.827 (2) | 175 |

Symmetry code: (i) $x+1/4, -y+1/4, z+1/4$.