

2-[(Cyclohex-3-en-1-ylmethoxy)methyl]-6-phenyl-1,2,4-triazine-3,5(2H,4H)-dione

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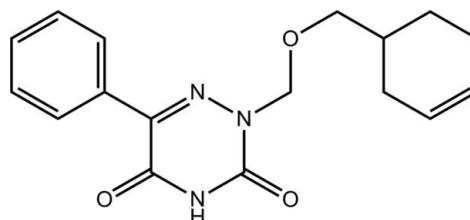
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$; disorder in main residue; R factor = 0.061; wR factor = 0.133; data-to-parameter ratio = 8.3.

In the title 1,2,4-triazine derivative, $C_{17}H_{19}N_3O_3$, the heterocyclic ring is planar (r.m.s. deviation = 0.040 Å) and effectively coplanar with the adjacent phenyl ring [dihedral angle = 4.5 (2)°] but almost perpendicular to the (cyclohex-3-en-1-ylmethoxy)methyl residue [$\text{N}-\text{N}-\text{C}-\text{O}$ torsion angle = 71.6 (5)°], so that the molecule has an ‘L’ shape. Supramolecular chains along [001] are formed in the crystal via $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds where the acceptor O atom is the ether O atom. The adjacent carbonyl O atom forms a complementary $\text{C}-\text{H}\cdots\text{O}$ contact resulting in the formation of a seven-membered $\{\cdots\text{HNCO}\cdots\text{HCO}\}$ heterosynthon; the second carbonyl O atom forms an intramolecular $\text{C}-\text{H}\cdots\text{O}$ contact. Chains are connected into a supramolecular layer in the *ac* plane by $\pi-\pi$ interactions [ring centroid–centroid distance = 3.488 (3) Å]. The central atom in the $-\text{CH}_2\text{CH}_2\text{C}(\text{H})=\text{$ residue of the cyclohexene ring is disordered over two sites, with the major component having a site-occupancy factor of 0.51 (2).

Related literature

For the potential medicinal applications of 1,2,4-triazines, see: Ban *et al.* (2010); Irannejad *et al.* (2010); Sangshetti & Shinde (2010). For the synthesis, see: El-Brollosy (2008).



Experimental

Crystal data

| | |
|-------------------------------|--|
| $C_{17}H_{19}N_3O_3$ | $V = 761.43 (17)\text{ \AA}^3$ |
| $M_r = 313.35$ | $Z = 2$ |
| Monoclinic, Pc | Mo $K\alpha$ radiation |
| $a = 4.7924 (6)\text{ \AA}$ | $\mu = 0.10\text{ mm}^{-1}$ |
| $b = 13.7083 (19)\text{ \AA}$ | $T = 100\text{ K}$ |
| $c = 11.8293 (15)\text{ \AA}$ | $0.35 \times 0.15 \times 0.03\text{ mm}$ |
| $\beta = 101.538 (12)^\circ$ | |

Data collection

| | |
|--|--|
| Agilent SuperNova Dual diffractometer with an Atlas detector | 4230 measured reflections |
| Absorption correction: multi-scan (<i>CrysAlis PRO</i> ; Agilent, 2011) | 1757 independent reflections |
| $T_{\min} = 0.806$, $T_{\max} = 1.000$ | 1158 reflections with $I > 2\sigma(I)$ |
| | $R_{\text{int}} = 0.078$ |

Refinement

| | |
|---------------------------------|---|
| $R[F^2 > 2\sigma(F^2)] = 0.061$ | 4 restraints |
| $wR(F^2) = 0.133$ | H-atom parameters constrained |
| $S = 1.04$ | $\Delta\rho_{\text{max}} = 0.24\text{ e \AA}^{-3}$ |
| 1757 reflections | $\Delta\rho_{\text{min}} = -0.30\text{ e \AA}^{-3}$ |
| 212 parameters | |

Table 1
Hydrogen-bond geometry (Å, °).

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|--|--------------|--------------------|-------------|----------------------|
| $\text{N}1-\text{H}1\cdots\text{O}3^{\text{i}}$ | 0.88 | 2.00 | 2.877 (5) | 174 |
| $\text{C}2-\text{H}2\cdots\text{O}1$ | 0.95 | 2.21 | 2.880 (7) | 127 |
| $\text{C}10-\text{H}10\text{B}\cdots\text{O}2^{\text{ii}}$ | 0.99 | 2.46 | 3.352 (6) | 150 |

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

Data collection: *CrysAlis PRO* (Agilent, 2011); 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: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2425).

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supplementary materials

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2-[(Cyclohex-3-en-1-ylmethoxy)methyl]-6-phenyl-1,2,4-triazine-3,5(2H,4H)-dione

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Comment

Several 1,2,4-triazines have been shown to exhibit herbicidal, anti-viral, anti-microbial, anti-inflammatory, anti-malarial, anti-cancer, anti-proliferative and neuroprotective activities (Ban *et al.*, 2010; Irannejad *et al.*, 2010; Sangshetti & Shinde, 2010). The title compound (I), was originally synthesized as a potential anti-microbial agent (El-Brollosy, 2008). Herein, we describe the results of its crystal structure determination.

The six ring atoms comprising the 2,4-dihydro-1,2,4-triazine-3,5-dione residue in (I), Fig. 1, are co-planar with a r.m.s. = 0.040 Å; the maximum deviations from their least-squares plane being 0.035 (5) Å for the C7 atom and -0.039 (5) Å for the C8 atom. The dihedral angle between this ring and the attached phenyl ring of 4.5 (2)° is consistent with an almost co-planar relationship. The (cyclohex-3-en-1-ylmethoxy)methyl residue lies perpendicular to the rest of the molecule as seen in the value of the N3—N2—C10—O3 torsion angle of 71.6 (5)°, so that to a first approximation the molecule has an *L*-shape.

The crystal structure features supramolecular chains along [001]. These are mediated by, perhaps surprisingly, N—H···O hydrogen bonds where the O atom is the ether O atom, rather than carbonyl O atoms. The adjacent carbonyl-O2 atom forms a complementary C—H···O contact resulting in the formation of a seven-membered {···HNCO···HCO} heterosynthon, Fig. 2 and Table 1. The carbonyl-O1 atom forms an intramolecular C—H..O interaction. The chains are connected into supramolecular layers in the *ac* plane by π — π interactions [ring centroid(N1—N3,C7—C9)···centroid(C1—C6)ⁱ = 3.488 (3) Å and tilt angle = 4.5 (2)°; symmetry code: (i) $x+1, y, z$], Fig. 3. Supramolecular layers stack along the *b* axis with no specific interactions between them, Fig. 4.

Experimental

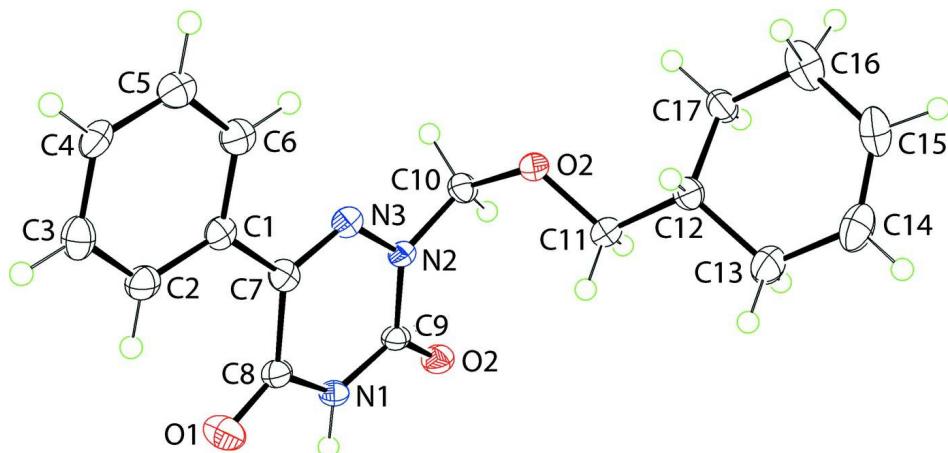
5-Phenyl-6-azauracil (0.189 g, 1 mmol) was stirred in dry acetonitrile (15 ml) under nitrogen and *N,O*-bis-trimethylsilylacetamide (0.87 ml, 3.5 mmol) was added. After a clear solution was obtained (10 min), the mixture was cooled to 223 K and trimethylsilyl trifluoromethanesulphonate (0.18 ml, 1 mmol) was added followed by the drop wise addition of bis(3-cyclohexen-1-ylmethoxy)methane (0.472 g, 2 mmol). The reaction mixture was stirred at room temperature for 5 h. The reaction was quenched by addition of sat. *aq.* NaHCO₃ solution (5 ml). The mixture was evaporated under reduced pressure and the residue was extracted with ether (3 x 50 ml). The combined ether fractions were collected, dried (MgSO₄) and evaporated under reduced pressure. The residue was purified on a silica gel column using 1:5 petroleum ether / chloroform to give the title compound in 64% (0.199 g) yield. Colourless crystals were obtained upon crystallization from its ethanol solution (El-Brollosy, 2008).

Refinement

H-atoms were placed in calculated positions [N—H = 0.88 and C—H = 0.95 to 0.99 Å, $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N}, \text{C})$] and were included in the refinement in the riding model approximation. The amino H-atom was refined freely. In the absence of significant anomalous scattering effects, 799 Friedel pairs were averaged in the final refinement. The C16 atom of the cyclohexene ring was disordered over two positions. From anisotropic refinement, the major component of the disorder had a site occupancy factor = 0.51 (2). The pairs of the respective C15—C16/C16'—C17 bond lengths were restrained to be within 0.01 Å of each other; the anisotropic displacement parameters for the disordered atoms were constrained to be equal.

Computing details

Data collection: *CrysAlis PRO* (Agilent, 2011); cell refinement: *CrysAlis PRO* (Agilent, 2011); data reduction: *CrysAlis PRO* (Agilent, 2011); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

**Figure 1**

The molecular structure of (I) showing the atom-labelling scheme and displacement ellipsoids at the 50% probability level. Only the major component of the disordered C16 atom is shown.

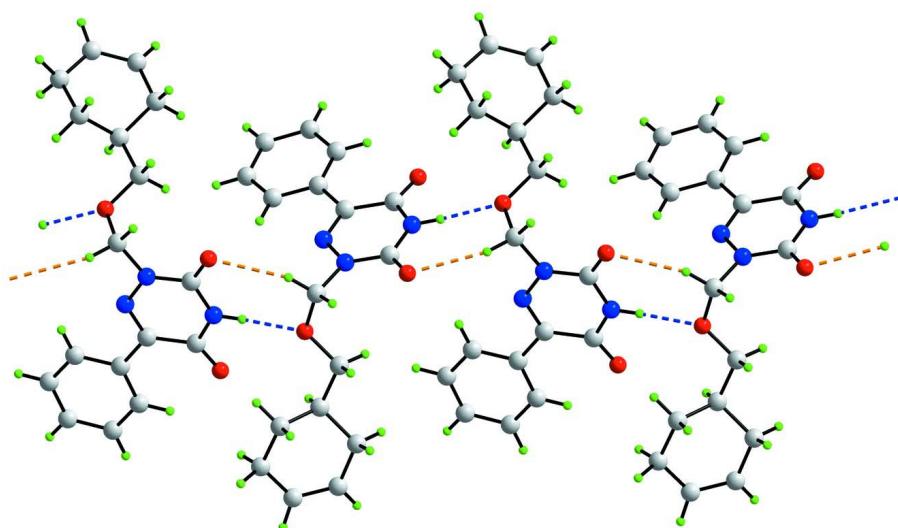
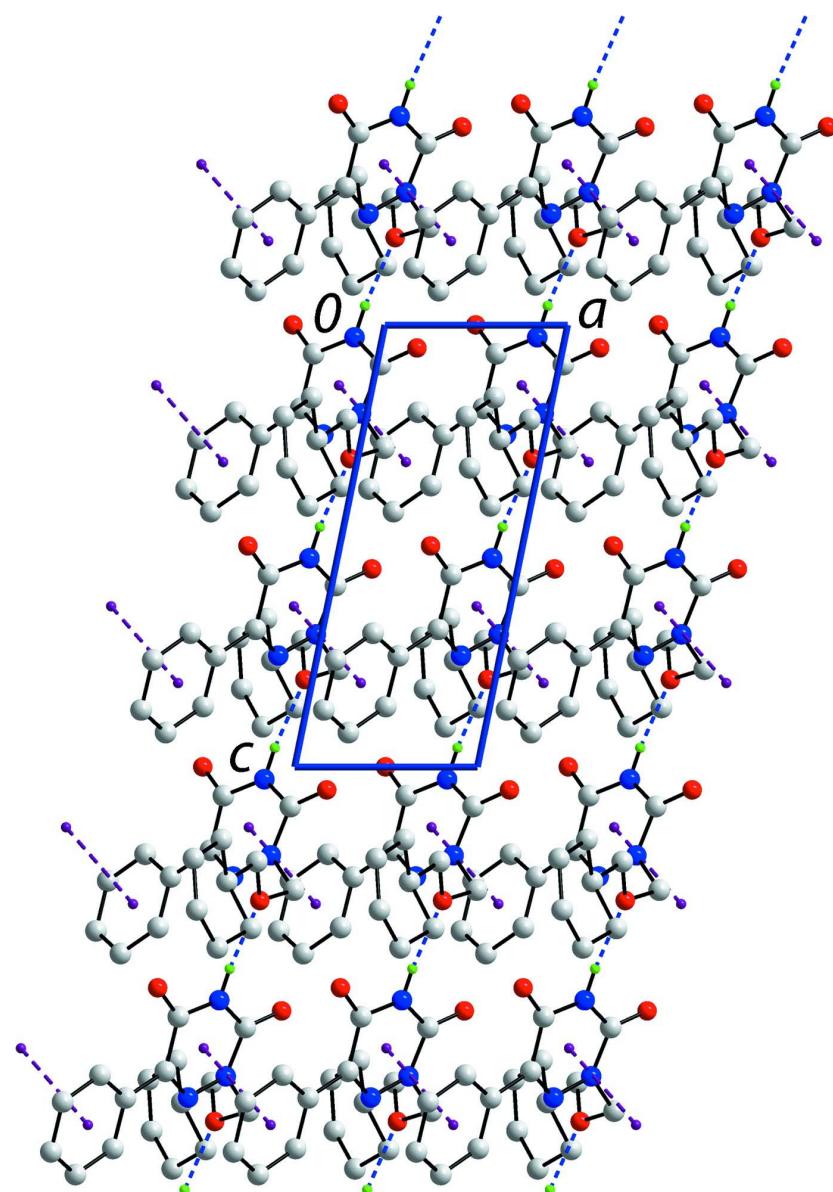
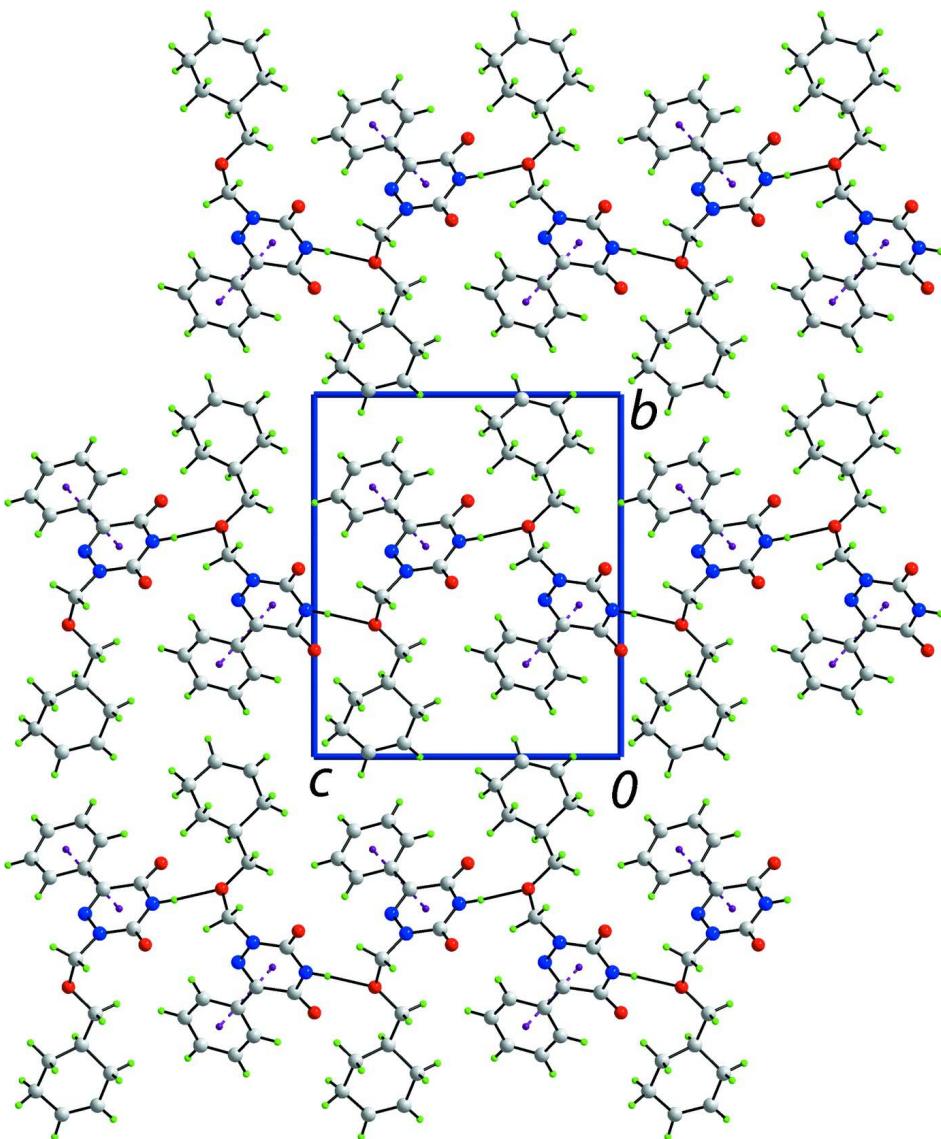


Figure 2

A view of the supramolecular chain in (I) mediated by N—H···O hydrogen bonding and reinforced by C—H···O interactions, shown as blue and orange dashed lines, respectively.

**Figure 3**

A view of the supramolecular layer parallel to the ac plane in (I). The $\text{N}—\text{H}\cdots\text{O}$ and $\pi—\pi$ interactions are shown as blue and purple dashed lines, respectively. Hydrogen atoms not participating in intermolecular interactions have been omitted.

**Figure 4**

A view in projection down the a axis of the unit-cell contents for (I). The N—H···O and π — π interactions are shown as blue and purple dashed lines, respectively.

2-[(Cyclohex-3-en-1-ylmethoxy)methyl]-6-phenyl-1,2,4-triazine- 3,5(2*H*,4*H*)-dione

Crystal data

$C_{17}H_{19}N_3O_3$
 $M_r = 313.35$
Monoclinic, Pc
Hall symbol: P -2yc
 $a = 4.7924 (6)$ Å
 $b = 13.7083 (19)$ Å
 $c = 11.8293 (15)$ Å
 $\beta = 101.538 (12)^\circ$
 $V = 761.43 (17)$ Å³
 $Z = 2$

$F(000) = 332$
 $D_x = 1.367 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 801 reflections
 $\theta = 2.3\text{--}27.5^\circ$
 $\mu = 0.10 \text{ mm}^{-1}$
 $T = 100$ K
Plate, colourless
 $0.35 \times 0.15 \times 0.03$ mm

Data collection

Agilent SuperNova Dual
diffractometer with an Atlas detector
Radiation source: SuperNova (Mo) X-ray
Source
Mirror monochromator
Detector resolution: 10.4041 pixels mm⁻¹
 ω scan
Absorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2011)

$T_{\min} = 0.806$, $T_{\max} = 1.000$
4230 measured reflections
1757 independent reflections
1158 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.078$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 2.3^\circ$
 $h = -5 \rightarrow 6$
 $k = -17 \rightarrow 17$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.061$
 $wR(F^2) = 0.133$
 $S = 1.04$
1757 reflections
212 parameters
4 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
 $w = 1/[\sigma^2(F_o^2) + (0.0421P)^2]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.002$
 $\Delta\rho_{\max} = 0.24 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.30 \text{ e } \text{\AA}^{-3}$

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> | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|----|-------------|------------|------------|----------------------------------|-----------|
| O1 | 0.4970 (7) | 0.7062 (3) | 0.4998 (3) | 0.0321 (9) | |
| O2 | 1.2011 (7) | 0.4812 (3) | 0.5519 (3) | 0.0245 (8) | |
| O3 | 0.9583 (7) | 0.3645 (2) | 0.8023 (3) | 0.0228 (8) | |
| N1 | 0.8476 (9) | 0.5948 (3) | 0.5278 (4) | 0.0205 (9) | |
| H1 | 0.8696 | 0.6052 | 0.4567 | 0.025* | |
| N2 | 0.9682 (7) | 0.5123 (3) | 0.7000 (3) | 0.0190 (10) | |
| N3 | 0.7856 (7) | 0.5669 (3) | 0.7480 (3) | 0.0194 (9) | |
| C1 | 0.4436 (9) | 0.6924 (4) | 0.7489 (4) | 0.0201 (11) | |
| C2 | 0.2789 (10) | 0.7697 (4) | 0.6974 (5) | 0.0253 (12) | |
| H2 | 0.2844 | 0.7872 | 0.6202 | 0.030* | |
| C3 | 0.1041 (10) | 0.8224 (4) | 0.7579 (5) | 0.0275 (13) | |
| H3 | -0.0072 | 0.8754 | 0.7221 | 0.033* | |
| C4 | 0.0957 (11) | 0.7960 (4) | 0.8709 (5) | 0.0244 (12) | |
| H4 | -0.0211 | 0.8315 | 0.9125 | 0.029* | |
| C5 | 0.2553 (10) | 0.7192 (4) | 0.9223 (5) | 0.0260 (12) | |
| H5 | 0.2458 | 0.7011 | 0.9989 | 0.031* | |

| | | | | | |
|------|-------------|-------------|-------------|-------------|----------|
| C6 | 0.4310 (11) | 0.6673 (4) | 0.8634 (5) | 0.0251 (12) | |
| H6 | 0.5428 | 0.6148 | 0.9004 | 0.030* | |
| C7 | 0.6321 (9) | 0.6348 (4) | 0.6886 (4) | 0.0205 (11) | |
| C8 | 0.6423 (10) | 0.6503 (4) | 0.5652 (4) | 0.0209 (11) | |
| C9 | 1.0207 (10) | 0.5252 (4) | 0.5896 (4) | 0.0185 (11) | |
| C10 | 1.1326 (10) | 0.4421 (4) | 0.7787 (4) | 0.0217 (11) | |
| H10A | 1.2888 | 0.4159 | 0.7441 | 0.026* | |
| H10B | 1.2188 | 0.4752 | 0.8517 | 0.026* | |
| C11 | 0.8986 (11) | 0.2912 (4) | 0.7141 (5) | 0.0252 (12) | |
| H11A | 1.0778 | 0.2701 | 0.6920 | 0.030* | |
| H11B | 0.7715 | 0.3182 | 0.6449 | 0.030* | |
| C12 | 0.7570 (10) | 0.2050 (4) | 0.7590 (4) | 0.0221 (11) | |
| H12 | 0.5764 | 0.2283 | 0.7801 | 0.027* | |
| C13 | 0.6801 (13) | 0.1266 (4) | 0.6644 (5) | 0.0376 (14) | |
| H13A | 0.5451 | 0.1545 | 0.5980 | 0.045* | |
| H13B | 0.8544 | 0.1070 | 0.6373 | 0.045* | |
| C14 | 0.5488 (12) | 0.0383 (4) | 0.7073 (6) | 0.0365 (14) | |
| H14 | 0.4158 | -0.0003 | 0.6557 | 0.044* | |
| C15 | 0.6234 (12) | 0.0133 (4) | 0.8234 (6) | 0.0360 (14) | |
| H15 | 0.5694 | -0.0490 | 0.8468 | 0.043* | |
| C16 | 0.779 (5) | 0.0777 (12) | 0.9093 (9) | 0.032 (5) | 0.51 (2) |
| H16A | 0.9165 | 0.0382 | 0.9644 | 0.039* | 0.51 (2) |
| H16B | 0.6429 | 0.1066 | 0.9529 | 0.039* | 0.51 (2) |
| C16' | 0.837 (5) | 0.0637 (12) | 0.9049 (10) | 0.032 (5) | 0.49 |
| H16C | 1.0033 | 0.0199 | 0.9266 | 0.039* | 0.49 (2) |
| H16D | 0.7602 | 0.0758 | 0.9753 | 0.039* | 0.49 (2) |
| C17 | 0.9405 (10) | 0.1602 (4) | 0.8654 (5) | 0.0253 (12) | |
| H17A | 1.1188 | 0.1347 | 0.8463 | 0.030* | |
| H17B | 0.9912 | 0.2105 | 0.9261 | 0.030* | |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|--------------|-------------|--------------|
| O1 | 0.032 (2) | 0.039 (2) | 0.026 (2) | 0.0098 (18) | 0.0082 (17) | 0.0077 (19) |
| O2 | 0.0274 (18) | 0.027 (2) | 0.0197 (19) | 0.0025 (16) | 0.0073 (16) | -0.0009 (16) |
| O3 | 0.032 (2) | 0.0200 (19) | 0.0175 (18) | -0.0002 (16) | 0.0074 (16) | -0.0005 (15) |
| N1 | 0.028 (2) | 0.021 (2) | 0.014 (2) | 0.0002 (18) | 0.0081 (18) | 0.0006 (17) |
| N2 | 0.024 (2) | 0.018 (2) | 0.015 (2) | 0.0027 (18) | 0.0042 (19) | -0.0017 (19) |
| N3 | 0.016 (2) | 0.022 (2) | 0.021 (2) | -0.0021 (18) | 0.0038 (18) | -0.0020 (19) |
| C1 | 0.017 (2) | 0.020 (3) | 0.023 (3) | -0.006 (2) | 0.004 (2) | -0.002 (2) |
| C2 | 0.029 (3) | 0.024 (3) | 0.023 (3) | -0.007 (2) | 0.005 (2) | -0.002 (2) |
| C3 | 0.022 (3) | 0.023 (3) | 0.037 (3) | 0.002 (2) | 0.005 (3) | -0.002 (3) |
| C4 | 0.027 (3) | 0.022 (3) | 0.026 (3) | -0.003 (2) | 0.011 (2) | -0.008 (2) |
| C5 | 0.025 (3) | 0.029 (3) | 0.025 (3) | -0.006 (2) | 0.008 (2) | -0.002 (2) |
| C6 | 0.024 (3) | 0.026 (3) | 0.026 (3) | -0.005 (2) | 0.006 (2) | -0.001 (3) |
| C7 | 0.017 (3) | 0.021 (3) | 0.023 (3) | -0.004 (2) | 0.004 (2) | 0.001 (2) |
| C8 | 0.025 (3) | 0.021 (3) | 0.017 (3) | -0.002 (2) | 0.005 (2) | -0.001 (2) |
| C9 | 0.022 (3) | 0.020 (3) | 0.014 (3) | -0.005 (2) | 0.004 (2) | 0.000 (2) |
| C10 | 0.024 (3) | 0.021 (3) | 0.021 (3) | -0.005 (2) | 0.005 (2) | 0.002 (2) |
| C11 | 0.035 (3) | 0.022 (3) | 0.019 (2) | -0.001 (2) | 0.005 (2) | -0.003 (2) |

| | | | | | | |
|------|-----------|-----------|-----------|------------|------------|------------|
| C12 | 0.021 (2) | 0.021 (3) | 0.024 (3) | 0.003 (2) | 0.004 (2) | 0.001 (2) |
| C13 | 0.047 (3) | 0.027 (3) | 0.033 (3) | -0.004 (3) | -0.006 (3) | -0.004 (3) |
| C14 | 0.034 (3) | 0.030 (3) | 0.048 (4) | -0.011 (3) | 0.012 (3) | -0.011 (3) |
| C15 | 0.033 (3) | 0.021 (3) | 0.053 (4) | -0.003 (3) | 0.007 (3) | 0.004 (3) |
| C16 | 0.014 (8) | 0.029 (6) | 0.055 (5) | 0.013 (6) | 0.008 (4) | 0.012 (4) |
| C16' | 0.014 (8) | 0.029 (6) | 0.055 (5) | 0.013 (6) | 0.008 (4) | 0.012 (4) |
| C17 | 0.028 (3) | 0.022 (3) | 0.025 (3) | -0.002 (2) | 0.002 (2) | 0.005 (2) |

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

| | | | |
|------------|-----------|---------------|------------|
| O1—C8 | 1.207 (6) | C10—H10A | 0.9900 |
| O2—C9 | 1.210 (5) | C10—H10B | 0.9900 |
| O3—C10 | 1.415 (6) | C11—C12 | 1.510 (7) |
| O3—C11 | 1.435 (6) | C11—H11A | 0.9900 |
| N1—C9 | 1.375 (6) | C11—H11B | 0.9900 |
| N1—C8 | 1.385 (6) | C12—C17 | 1.514 (7) |
| N1—H1 | 0.8800 | C12—C13 | 1.543 (8) |
| N2—N3 | 1.359 (5) | C12—H12 | 1.0000 |
| N2—C9 | 1.390 (5) | C13—C14 | 1.499 (8) |
| N2—C10 | 1.456 (6) | C13—H13A | 0.9900 |
| N3—C7 | 1.301 (6) | C13—H13B | 0.9900 |
| C1—C2 | 1.387 (7) | C14—C15 | 1.390 (9) |
| C1—C6 | 1.410 (7) | C14—H14 | 0.9500 |
| C1—C7 | 1.486 (6) | C15—C16' | 1.435 (11) |
| C2—C3 | 1.406 (7) | C15—C16 | 1.437 (11) |
| C2—H2 | 0.9500 | C15—H15 | 0.9500 |
| C3—C4 | 1.392 (7) | C16—C17 | 1.521 (9) |
| C3—H3 | 0.9500 | C16—H16A | 0.9900 |
| C4—C5 | 1.371 (7) | C16—H16B | 0.9900 |
| C4—H4 | 0.9500 | C16'—C17 | 1.519 (10) |
| C5—C6 | 1.391 (7) | C16'—H16C | 0.9900 |
| C5—H5 | 0.9500 | C16'—H16D | 0.9900 |
| C6—H6 | 0.9500 | C17—H17A | 0.9900 |
| C7—C8 | 1.486 (6) | C17—H17B | 0.9900 |
| | | | |
| C10—O3—C11 | 115.1 (3) | O3—C11—H11B | 109.8 |
| C9—N1—C8 | 127.1 (4) | C12—C11—H11B | 109.8 |
| C9—N1—H1 | 116.4 | H11A—C11—H11B | 108.3 |
| C8—N1—H1 | 116.4 | C11—C12—C17 | 112.5 (4) |
| N3—N2—C9 | 125.1 (4) | C11—C12—C13 | 110.4 (4) |
| N3—N2—C10 | 114.1 (4) | C17—C12—C13 | 109.5 (4) |
| C9—N2—C10 | 120.6 (4) | C11—C12—H12 | 108.1 |
| C7—N3—N2 | 120.7 (4) | C17—C12—H12 | 108.1 |
| C2—C1—C6 | 118.6 (5) | C13—C12—H12 | 108.1 |
| C2—C1—C7 | 122.8 (4) | C14—C13—C12 | 111.9 (5) |
| C6—C1—C7 | 118.6 (4) | C14—C13—H13A | 109.2 |
| C1—C2—C3 | 120.9 (5) | C12—C13—H13A | 109.2 |
| C1—C2—H2 | 119.6 | C14—C13—H13B | 109.2 |
| C3—C2—H2 | 119.6 | C12—C13—H13B | 109.2 |
| C4—C3—C2 | 119.3 (5) | H13A—C13—H13B | 107.9 |

| | | | |
|---------------|------------|------------------|------------|
| C4—C3—H3 | 120.3 | C15—C14—C13 | 119.4 (5) |
| C2—C3—H3 | 120.3 | C15—C14—H14 | 120.3 |
| C5—C4—C3 | 120.3 (5) | C13—C14—H14 | 120.3 |
| C5—C4—H4 | 119.9 | C14—C15—C16' | 123.6 (6) |
| C3—C4—H4 | 119.9 | C14—C15—C16 | 122.8 (6) |
| C4—C5—C6 | 120.7 (5) | C14—C15—H15 | 118.6 |
| C4—C5—H5 | 119.6 | C16'—C15—H15 | 115.8 |
| C6—C5—H5 | 119.6 | C16—C15—H15 | 118.6 |
| C5—C6—C1 | 120.2 (5) | C15—C16—C17 | 116.2 (8) |
| C5—C6—H6 | 119.9 | C15—C16—H16A | 108.2 |
| C1—C6—H6 | 119.9 | C17—C16—H16A | 108.2 |
| N3—C7—C1 | 117.0 (4) | C15—C16—H16B | 108.2 |
| N3—C7—C8 | 120.4 (4) | C17—C16—H16B | 108.2 |
| C1—C7—C8 | 122.6 (4) | H16A—C16—H16B | 107.4 |
| O1—C8—N1 | 120.0 (4) | C15—C16'—C17 | 116.5 (8) |
| O1—C8—C7 | 126.4 (5) | C15—C16'—H16C | 108.2 |
| N1—C8—C7 | 113.6 (4) | C17—C16'—H16C | 108.2 |
| O2—C9—N1 | 123.3 (4) | C15—C16'—H16D | 108.2 |
| O2—C9—N2 | 124.0 (4) | C17—C16'—H16D | 108.2 |
| N1—C9—N2 | 112.6 (4) | H16C—C16'—H16D | 107.3 |
| O3—C10—N2 | 111.0 (4) | C16'—C17—C12 | 116.0 (7) |
| O3—C10—H10A | 109.4 | C16—C17—C12 | 109.6 (8) |
| N2—C10—H10A | 109.4 | C16'—C17—H17A | 96.5 |
| O3—C10—H10B | 109.4 | C16—C17—H17A | 109.8 |
| N2—C10—H10B | 109.4 | C12—C17—H17A | 109.7 |
| H10A—C10—H10B | 108.0 | C16'—C17—H17B | 115.5 |
| O3—C11—C12 | 109.3 (4) | C16—C17—H17B | 109.7 |
| O3—C11—H11A | 109.8 | C12—C17—H17B | 109.7 |
| C12—C11—H11A | 109.8 | H17A—C17—H17B | 108.2 |
| | | | |
| C9—N2—N3—C7 | 3.3 (6) | N3—N2—C9—N1 | -5.5 (6) |
| C10—N2—N3—C7 | 177.7 (4) | C10—N2—C9—N1 | -179.5 (4) |
| C6—C1—C2—C3 | 0.4 (7) | C11—O3—C10—N2 | 79.3 (5) |
| C7—C1—C2—C3 | -179.8 (5) | N3—N2—C10—O3 | 71.6 (5) |
| C1—C2—C3—C4 | -0.4 (7) | C9—N2—C10—O3 | -113.8 (4) |
| C2—C3—C4—C5 | -0.3 (8) | C10—O3—C11—C12 | 170.0 (4) |
| C3—C4—C5—C6 | 1.0 (8) | O3—C11—C12—C17 | -59.4 (5) |
| C4—C5—C6—C1 | -1.0 (7) | O3—C11—C12—C13 | 177.9 (4) |
| C2—C1—C6—C5 | 0.3 (7) | C11—C12—C13—C14 | 177.9 (5) |
| C7—C1—C6—C5 | -179.5 (4) | C17—C12—C13—C14 | 53.4 (6) |
| N2—N3—C7—C1 | -178.5 (4) | C12—C13—C14—C15 | -29.4 (7) |
| N2—N3—C7—C8 | 3.3 (7) | C13—C14—C15—C16' | -4.5 (16) |
| C2—C1—C7—N3 | 177.0 (5) | C13—C14—C15—C16 | 12.3 (16) |
| C6—C1—C7—N3 | -3.2 (6) | C14—C15—C16—C17 | -19 (3) |
| C2—C1—C7—C8 | -4.9 (7) | C16'—C15—C16—C17 | 79 (3) |
| C6—C1—C7—C8 | 174.9 (4) | C14—C15—C16'—C17 | 12 (3) |
| C9—N1—C8—O1 | -177.0 (5) | C16—C15—C16'—C17 | -79 (3) |
| C9—N1—C8—C7 | 4.6 (7) | C15—C16'—C17—C16 | 79 (3) |
| N3—C7—C8—O1 | 174.8 (5) | C15—C16'—C17—C12 | 15 (2) |

| | | | |
|--------------|------------|------------------|-------------|
| C1—C7—C8—O1 | −3.2 (8) | C15—C16—C17—C16' | −78 (3) |
| N3—C7—C8—N1 | −6.9 (6) | C15—C16—C17—C12 | 43 (2) |
| C1—C7—C8—N1 | 175.1 (4) | C11—C12—C17—C16' | −170.4 (12) |
| C8—N1—C9—O2 | −178.0 (5) | C13—C12—C17—C16' | −47.1 (12) |
| C8—N1—C9—N2 | 1.1 (7) | C11—C12—C17—C16 | 177.0 (10) |
| N3—N2—C9—O2 | 173.6 (4) | C13—C12—C17—C16 | −59.8 (11) |
| C10—N2—C9—O2 | −0.4 (7) | | |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|-----------------------------|------|-------|-----------|---------|
| N1—H1···O3 ⁱ | 0.88 | 2.00 | 2.877 (5) | 174 |
| C2—H2···O1 | 0.95 | 2.21 | 2.880 (7) | 127 |
| C10—H10B···O2 ⁱⁱ | 0.99 | 2.46 | 3.352 (6) | 150 |

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