

N'-(*Z*)-4-Methylbenzylidene]-4-nitro-benzohydrazide monohydrate

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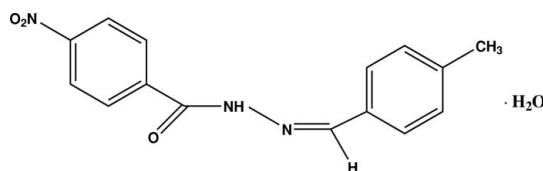
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(C-C) = 0.001$ Å;
R factor = 0.037; wR factor = 0.119; data-to-parameter ratio = 35.0.

In the title compound, $C_{15}H_{13}N_3O_3 \cdot H_2O$, the two benzene rings form a dihedral angle of $2.03(2)^\circ$. In the crystal structure, adjacent hydrazide molecules are linked into dimers by water molecules; these dimers are then stacked along the b axis. Intermolecular O—H···O, O—H···N and C—H···O hydrogen bonds and a π — π stacking interaction between the nitrobenzene and tolyl rings with a centroid–centroid distance of $3.8208(3)$ Å are observed. There is also a short O···N contact [$2.6824(7)$ Å].

Related literature

For related literature on hydrazones, see: Sridhar & Perumal (2003). For the biological applications of hydrazides/hydrazones, see: Bedia *et al.* (2006). For a related structure, see: Fun *et al.* (2008). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

$C_{15}H_{13}N_3O_3 \cdot H_2O$
 $M_r = 301.30$
Triclinic, $P\bar{1}$
 $a = 6.5387(1)$ Å
 $b = 6.9730(1)$ Å
 $c = 15.9064(3)$ Å

$\alpha = 80.524(1)^\circ$
 $\beta = 82.628(1)^\circ$
 $\gamma = 85.036(1)^\circ$
 $V = 707.85(2)$ Å³
 $Z = 2$
Mo $K\alpha$ radiation

$\mu = 0.10$ mm⁻¹
 $T = 100.0(1)$ K

$0.68 \times 0.44 \times 0.23$ mm

Data collection

Bruker SMART APEXII CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
 $T_{\min} = 0.932$, $T_{\max} = 0.976$

31311 measured reflections
7380 independent reflections
6571 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.118$
 $S = 1.05$
7380 reflections
211 parameters
4 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.48$ e Å⁻³
 $\Delta\rho_{\min} = -0.59$ e Å⁻³

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
N2—H1N2···O1W	0.864 (8)	1.978 (9)	2.8191 (7)	164.4 (11)
O1W—H2W1···O1 ⁱ	0.837 (9)	2.013 (9)	2.8327 (7)	166.1 (11)
O1W—H1W1···O1 ⁱⁱ	0.851 (9)	2.258 (11)	2.9430 (6)	137.7 (10)
O1W—H1W1···N1 ⁱⁱ	0.851 (9)	2.357 (9)	3.1287 (7)	151.0 (11)
C1—H1A···O1W ⁱⁱⁱ	0.93	2.50	3.4090 (7)	165
C4—H4A···O2 ^{iv}	0.93	2.58	3.4565 (8)	157
C7—H7A···O1W	0.93	2.55	3.2393 (7)	132

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

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 2005); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2358).

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supporting information

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N'-(Z)-4-Methylbenzylidene]-4-nitrobenzohydrazide monohydrate

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S1. Comment

Hydrazones are versatile intermediates and important building blocks. Aryl hydrazones are important building blocks for the synthesis of a variety of heterocyclic compounds such as pyrazolines and pyrazoles (Sridhar & Perumal, 2003).

Hydrazones of aliphatic and aromatic methyl ketones yield pyrazole-4-carboxaldehyde on diformylation by the treatment with Vilsmeier reagent. A series of hydrazide-hydrazone were reported to possess good antituberculosis activity (Bedia *et al.*, 2006). Prompted by these review and in continuation of our work (Fun *et al.*, 2008), we here in report the crystal structure of the title compound, (I).

Bond lengths and angles in (I) (Fig. 1) are found to have normal values (Allen *et al.*, 1987). The two benzene rings are essentially planar with the maximum deviation from planarity being -0.004 (1) Å for atom C6 and 0.002 (1) Å for atom C12, respectively. The dihedral angle formed by the benzene (C1—C6) and (C9—C14) rings is 2.03 (2)°.

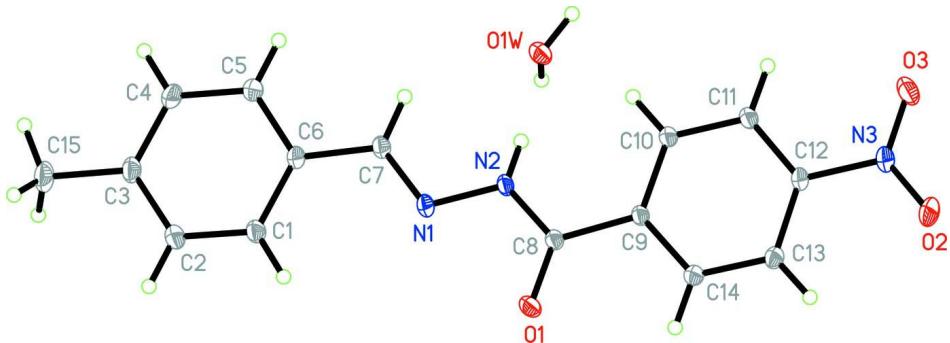
The crystal packing is consolidated by O—H···O, O—H···N, C—H···O and N—H···O inter and intramolecular hydrogen bonding (Table 1). Furthermore, the packing is strengthened by π — π stacking interactions involving the benzene (C1—C6) ($Cg1$) and the symmetry related (C9—C14) ring ($Cg2$) [$Cg1\cdots Cg2^i = 3.8208$ (3) Å; symmetry code: (i) 2-x, 1-y, -z] together with O···N short contacts [2.6824 (7) Å]. In the crystal packing, adjacent molecules are linked into dimers by water molecules and the dimers were then stacked down the [010] direction (Fig. 2).

S2. Experimental

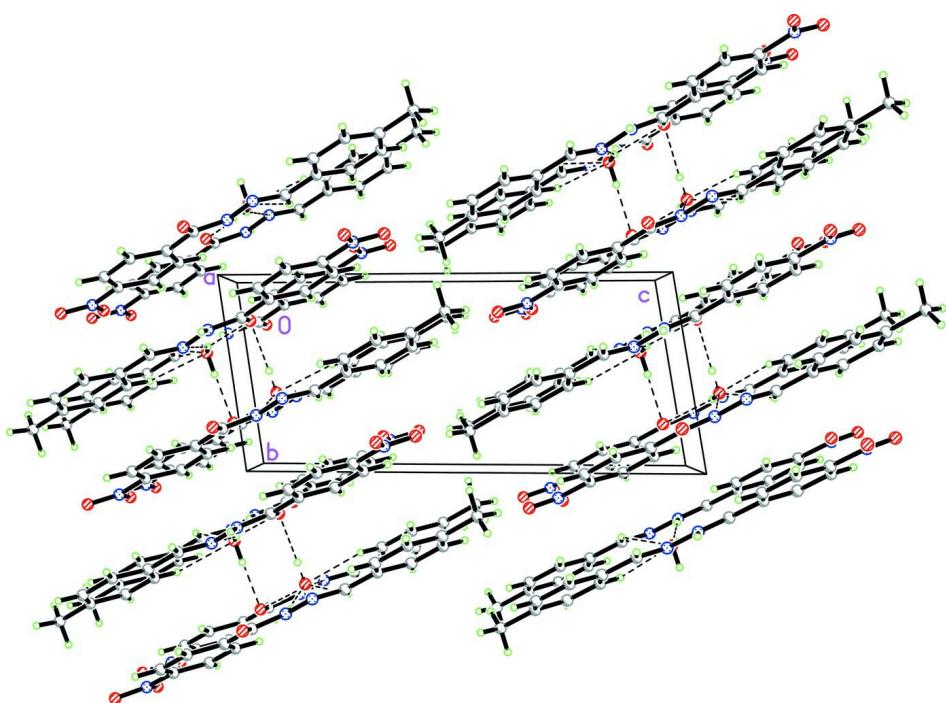
The title compound, $C_{15}H_{15}N_3O_4$, was obtained by refluxing 4-nitrobenzohydrazide (0.01 mol) and 4-methylbenzaldehyde (0.01 mol) in ethanol (30 ml) by adding 3 drops of concentrated sulfuric acid for 3 h. Excess ethanol was removed from the reaction mixture under reduced pressure. The solid product obtained was filtered, washed with water and dried. Crystals suitable for X-ray analysis were obtained from ethanol by slow evaporation.

S3. Refinement

The amino and water H atoms were located in a difference map and refined with restraints of N—H = 0.85 (1) Å and O—H = 0.84 (1) Å. The remaining H atoms were positioned geometrically [C—H = 0.93 Å (aromatic) or 0.96 Å (methyl)] and refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}$ (aromatic C) and $1.5U_{eq}$ (methyl C). A rotating group model was used for the methyl group.

**Figure 1**

The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atom numbering scheme.

**Figure 2**

The crystal packing of the title compound, viewed down the a axis, showing stacking of the dimers along the b axis.

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Crystal data

$C_{15}H_{13}N_3O_3 \cdot H_2O$

$M_r = 301.30$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.5387(1) \text{ \AA}$

$b = 6.9730(1) \text{ \AA}$

$c = 15.9064(3) \text{ \AA}$

$\alpha = 80.524(1)^\circ$

$\beta = 82.628(1)^\circ$

$\gamma = 85.036(1)^\circ$

$V = 707.85(2) \text{ \AA}^3$

$Z = 2$

$F(000) = 316$

$D_x = 1.414 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 9969 reflections

$\theta = 2.6\text{--}26.3^\circ$

$\mu = 0.11 \text{ mm}^{-1}$

$T = 100\text{ K}$
Block, colourless

Data collection

Bruker SMART APEXII CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2005)
 $T_{\min} = 0.932$, $T_{\max} = 0.976$

$0.68 \times 0.44 \times 0.23\text{ mm}$

31311 measured reflections
7380 independent reflections
6571 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$
 $\theta_{\max} = 37.5^\circ$, $\theta_{\min} = 2.6^\circ$
 $h = -11 \rightarrow 11$
 $k = -11 \rightarrow 10$
 $l = -27 \rightarrow 27$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.119$
 $S = 1.05$
7380 reflections
211 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 atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0681P)^2 + 0.1283P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.48\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.59\text{ e \AA}^{-3}$

Special details

Experimental. The data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	1.27580 (6)	0.77076 (7)	-0.04714 (3)	0.01642 (8)
O2	0.75581 (8)	1.18850 (8)	-0.38841 (3)	0.02159 (10)
O3	0.46360 (8)	1.16591 (9)	-0.30724 (3)	0.02477 (11)
N1	1.08155 (7)	0.62939 (7)	0.10666 (3)	0.01228 (8)
N2	0.97488 (7)	0.71707 (7)	0.03900 (3)	0.01180 (8)
N3	0.65301 (8)	1.14188 (7)	-0.31919 (3)	0.01517 (9)
C1	1.25937 (9)	0.43357 (8)	0.25978 (3)	0.01394 (9)
H1A	1.3528	0.4556	0.2105	0.017*
C2	1.32689 (9)	0.33958 (9)	0.33630 (4)	0.01558 (9)
H2A	1.4660	0.2992	0.3375	0.019*
C3	1.19044 (10)	0.30421 (9)	0.41173 (4)	0.01618 (10)
C4	0.98236 (10)	0.36711 (9)	0.40850 (4)	0.01732 (10)

H4A	0.8891	0.3455	0.4579	0.021*
C5	0.91322 (9)	0.46166 (9)	0.33225 (4)	0.01511 (9)
H5A	0.7744	0.5032	0.3313	0.018*
C6	1.05015 (8)	0.49509 (8)	0.25695 (3)	0.01205 (8)
C7	0.96681 (8)	0.59124 (8)	0.17866 (3)	0.01265 (9)
H7A	0.8259	0.6261	0.1809	0.015*
C8	1.08418 (8)	0.78209 (7)	-0.03665 (3)	0.01124 (8)
C9	0.96283 (8)	0.87393 (7)	-0.10826 (3)	0.01078 (8)
C10	0.74981 (8)	0.92155 (8)	-0.09685 (3)	0.01259 (9)
H10A	0.6760	0.8932	-0.0428	0.015*
C11	0.64812 (8)	1.01101 (8)	-0.16601 (3)	0.01320 (9)
H11A	0.5067	1.0435	-0.1589	0.016*
C12	0.76204 (8)	1.05088 (8)	-0.24589 (3)	0.01220 (8)
C13	0.97361 (9)	1.00639 (8)	-0.25947 (3)	0.01340 (9)
H13A	1.0465	1.0355	-0.3137	0.016*
C14	1.07314 (8)	0.91711 (8)	-0.18969 (3)	0.01269 (9)
H14A	1.2148	0.8857	-0.1972	0.015*
C15	1.26554 (12)	0.20102 (11)	0.49388 (4)	0.02473 (13)
H15A	1.1517	0.1899	0.5386	0.037*
H15B	1.3687	0.2740	0.5096	0.037*
H15C	1.3242	0.0733	0.4859	0.037*
O1W	0.56488 (7)	0.60570 (7)	0.07749 (3)	0.01669 (8)
H1N2	0.8430 (12)	0.7058 (17)	0.0471 (8)	0.027 (3)*
H2W1	0.5907 (17)	0.4907 (12)	0.0684 (8)	0.031 (3)*
H1W1	0.4410 (13)	0.6364 (17)	0.0668 (8)	0.035 (3)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.00972 (15)	0.0221 (2)	0.01583 (17)	0.00035 (13)	-0.00227 (13)	0.00155 (14)
O2	0.0250 (2)	0.0269 (2)	0.01076 (17)	-0.00148 (17)	-0.00275 (15)	0.00358 (15)
O3	0.01615 (19)	0.0361 (3)	0.0199 (2)	0.00133 (18)	-0.00737 (16)	0.00461 (19)
N1	0.01330 (18)	0.01250 (17)	0.01075 (17)	-0.00018 (13)	-0.00421 (13)	0.00081 (13)
N2	0.01085 (17)	0.01426 (18)	0.00977 (16)	-0.00046 (13)	-0.00339 (13)	0.00113 (13)
N3	0.0174 (2)	0.01585 (19)	0.01215 (18)	-0.00094 (15)	-0.00535 (15)	0.00083 (14)
C1	0.0137 (2)	0.0152 (2)	0.01227 (19)	-0.00081 (16)	-0.00285 (15)	0.00066 (15)
C2	0.0155 (2)	0.0166 (2)	0.0143 (2)	0.00085 (17)	-0.00481 (16)	-0.00018 (16)
C3	0.0208 (2)	0.0158 (2)	0.01158 (19)	0.00194 (17)	-0.00496 (17)	-0.00040 (16)
C4	0.0199 (2)	0.0196 (2)	0.01066 (19)	0.00142 (18)	-0.00093 (17)	0.00081 (16)
C5	0.0148 (2)	0.0174 (2)	0.01210 (19)	0.00041 (16)	-0.00163 (16)	-0.00002 (16)
C6	0.01369 (19)	0.01192 (19)	0.01037 (18)	-0.00115 (15)	-0.00323 (14)	0.00030 (14)
C7	0.01325 (19)	0.0133 (2)	0.01122 (18)	-0.00088 (15)	-0.00333 (15)	0.00004 (15)
C8	0.01101 (18)	0.01159 (18)	0.01084 (18)	0.00006 (14)	-0.00243 (14)	-0.00044 (14)
C9	0.01090 (18)	0.01125 (18)	0.00998 (17)	-0.00020 (14)	-0.00238 (14)	-0.00045 (14)
C10	0.01106 (18)	0.0154 (2)	0.01048 (18)	0.00026 (15)	-0.00182 (14)	0.00036 (15)
C11	0.01170 (19)	0.0154 (2)	0.01187 (19)	0.00029 (15)	-0.00297 (15)	0.00029 (15)
C12	0.01382 (19)	0.01256 (19)	0.01019 (18)	-0.00090 (15)	-0.00400 (15)	0.00040 (14)
C13	0.0141 (2)	0.0151 (2)	0.01041 (18)	-0.00143 (16)	-0.00148 (15)	-0.00005 (15)

C14	0.01159 (19)	0.0146 (2)	0.01130 (18)	-0.00053 (15)	-0.00128 (14)	-0.00060 (15)
C15	0.0321 (3)	0.0271 (3)	0.0134 (2)	0.0068 (2)	-0.0081 (2)	0.0011 (2)
O1W	0.01130 (16)	0.0212 (2)	0.01725 (18)	-0.00029 (13)	-0.00349 (13)	-0.00109 (14)

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

O1—C8	1.2400 (6)	C6—C7	1.4605 (7)
O2—N3	1.2246 (7)	C7—H7A	0.9300
O3—N3	1.2298 (7)	C8—C9	1.4970 (7)
N1—C7	1.2881 (7)	C9—C14	1.3994 (7)
N1—N2	1.3859 (6)	C9—C10	1.3994 (7)
N2—C8	1.3491 (7)	C10—C11	1.3898 (7)
N2—H1N2	0.864 (8)	C10—H10A	0.9300
N3—C12	1.4703 (7)	C11—C12	1.3867 (7)
C1—C2	1.3896 (8)	C11—H11A	0.9300
C1—C6	1.4019 (8)	C12—C13	1.3884 (8)
C1—H1A	0.9300	C13—C14	1.3895 (7)
C2—C3	1.4008 (8)	C13—H13A	0.9300
C2—H2A	0.9300	C14—H14A	0.9300
C3—C4	1.3977 (9)	C15—H15A	0.9600
C3—C15	1.5035 (8)	C15—H15B	0.9600
C4—C5	1.3913 (8)	C15—H15C	0.9600
C4—H4A	0.9300	O1W—H2W1	0.837 (8)
C5—C6	1.3999 (8)	O1W—H1W1	0.851 (8)
C5—H5A	0.9300		
C7—N1—N2	114.21 (4)	O1—C8—N2	122.52 (5)
C8—N2—N1	118.48 (4)	O1—C8—C9	120.74 (5)
C8—N2—H1N2	125.4 (8)	N2—C8—C9	116.74 (4)
N1—N2—H1N2	115.2 (8)	C14—C9—C10	119.65 (5)
O2—N3—O3	123.96 (5)	C14—C9—C8	116.93 (4)
O2—N3—C12	118.18 (5)	C10—C9—C8	123.40 (5)
O3—N3—C12	117.85 (5)	C11—C10—C9	120.23 (5)
C2—C1—C6	119.91 (5)	C11—C10—H10A	119.9
C2—C1—H1A	120.0	C9—C10—H10A	119.9
C6—C1—H1A	120.0	C12—C11—C10	118.60 (5)
C1—C2—C3	121.56 (5)	C12—C11—H11A	120.7
C1—C2—H2A	119.2	C10—C11—H11A	120.7
C3—C2—H2A	119.2	C11—C12—C13	122.73 (5)
C4—C3—C2	118.18 (5)	C11—C12—N3	118.38 (5)
C4—C3—C15	120.88 (6)	C13—C12—N3	118.89 (5)
C2—C3—C15	120.95 (6)	C12—C13—C14	118.01 (5)
C5—C4—C3	120.72 (5)	C12—C13—H13A	121.0
C5—C4—H4A	119.6	C14—C13—H13A	121.0
C3—C4—H4A	119.6	C13—C14—C9	120.79 (5)
C4—C5—C6	120.80 (5)	C13—C14—H14A	119.6
C4—C5—H5A	119.6	C9—C14—H14A	119.6
C6—C5—H5A	119.6	C3—C15—H15A	109.5

C5—C6—C1	118.83 (5)	C3—C15—H15B	109.5
C5—C6—C7	118.05 (5)	H15A—C15—H15B	109.5
C1—C6—C7	123.11 (5)	C3—C15—H15C	109.5
N1—C7—C6	122.33 (5)	H15A—C15—H15C	109.5
N1—C7—H7A	118.8	H15B—C15—H15C	109.5
C6—C7—H7A	118.8	H2W1—O1W—H1W1	106.0 (10)
C7—N1—N2—C8	172.77 (5)	N2—C8—C9—C14	-171.45 (5)
C6—C1—C2—C3	-0.09 (9)	O1—C8—C9—C10	-168.59 (5)
C1—C2—C3—C4	-0.32 (9)	N2—C8—C9—C10	10.18 (8)
C1—C2—C3—C15	179.55 (6)	C14—C9—C10—C11	0.09 (8)
C2—C3—C4—C5	0.20 (9)	C8—C9—C10—C11	178.42 (5)
C15—C3—C4—C5	-179.67 (6)	C9—C10—C11—C12	0.20 (8)
C3—C4—C5—C6	0.33 (9)	C10—C11—C12—C13	-0.47 (8)
C4—C5—C6—C1	-0.73 (9)	C10—C11—C12—N3	178.76 (5)
C4—C5—C6—C7	178.66 (5)	O2—N3—C12—C11	175.31 (5)
C2—C1—C6—C5	0.61 (8)	O3—N3—C12—C11	-4.94 (8)
C2—C1—C6—C7	-178.75 (5)	O2—N3—C12—C13	-5.43 (8)
N2—N1—C7—C6	178.63 (5)	O3—N3—C12—C13	174.32 (6)
C5—C6—C7—N1	179.37 (5)	C11—C12—C13—C14	0.42 (8)
C1—C6—C7—N1	-1.26 (9)	N3—C12—C13—C14	-178.81 (5)
N1—N2—C8—O1	-1.88 (8)	C12—C13—C14—C9	-0.10 (8)
N1—N2—C8—C9	179.38 (4)	C10—C9—C14—C13	-0.14 (8)
O1—C8—C9—C14	9.78 (8)	C8—C9—C14—C13	-178.58 (5)

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

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N2—H1N2…O1W	0.86 (1)	1.98 (1)	2.8191 (7)	164 (1)
O1W—H2W1…O1 ⁱ	0.84 (1)	2.01 (1)	2.8327 (7)	166 (1)
O1W—H1W1…O1 ⁱⁱ	0.85 (1)	2.26 (1)	2.9430 (6)	138 (1)
O1W—H1W1…N1 ⁱⁱ	0.85 (1)	2.36 (1)	3.1287 (7)	151 (1)
C1—H1A…O1W ⁱⁱⁱ	0.93	2.50	3.4090 (7)	165
C4—H4A…O2 ^{iv}	0.93	2.58	3.4565 (8)	157
C7—H7A…O1W	0.93	2.55	3.2393 (7)	132

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