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(E)-N'-(3,4-Dihydroxybenzylidene)-2,4-dimethylbenzohydrazide monohydrate

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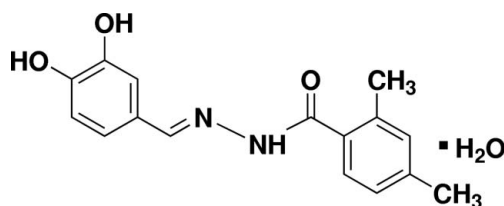
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.040; wR factor = 0.111; data-to-parameter ratio = 13.1.

In the title compound, $\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_3 \cdot \text{H}_2\text{O}$, the dihedral angle between the benzene rings is $30.27(7)^\circ$. In the crystal, the components are linked by $\text{N}-\text{H} \cdots \text{O}$, $\text{O}-\text{H} \cdots \text{O}$ and $\text{C}-\text{H} \cdots \text{O}$ interactions into a three-dimensional network.

Related literature

For the applications and biological activity of Schiff bases, see: Musharraf *et al.* (2012); Khan *et al.* (2012). For the crystal structures of related compounds, see: Taha *et al.* (2012); Baharudin *et al.* (2012).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_3 \cdot \text{H}_2\text{O}$
 $M_r = 302.32$
Monoclinic, $P2_1/n$
 $a = 8.1373(3)$ Å
 $b = 13.9025(5)$ Å

$c = 13.7886(5)$ Å
 $\beta = 92.913(1)^\circ$
 $V = 1557.87(10)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.09$ mm⁻¹
 $T = 298$ K

0.30 × 0.10 × 0.10 mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Bruker, 2000)
 $T_{\min} = 0.973$, $T_{\max} = 0.991$

9012 measured reflections
2897 independent reflections
2535 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.016$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.111$
 $S = 1.05$
2897 reflections
221 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.19$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.28$ e Å⁻³

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

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
$\text{O1W}-\text{H1W1} \cdots \text{N2}^{\text{i}}$	0.87 (2)	2.20 (2)	3.059 (2)	169 (2)
$\text{O1W}-\text{H2W1} \cdots \text{O1}$	0.94 (2)	2.01 (2)	2.935 (2)	173 (2)
$\text{N1}-\text{H1A} \cdots \text{O3}^{\text{ii}}$	0.91 (2)	2.08 (2)	2.962 (2)	163 (2)
$\text{O2}-\text{H2A} \cdots \text{O1}^{\text{i}}$	0.88 (2)	1.94 (2)	2.791 (2)	162 (2)
$\text{O3}-\text{H3A} \cdots \text{O1W}^{\text{iii}}$	0.85 (2)	1.79 (2)	2.629 (2)	172 (2)
$\text{C8}-\text{H8A} \cdots \text{O3}^{\text{ii}}$	0.93	2.58	3.382 (2)	145
$\text{C15}-\text{H15B} \cdots \text{O2}^{\text{i}}$	0.96	2.52	3.351 (2)	144

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

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); 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: SHELXTL, PARST (Nardelli, 1995) and PLATON (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PV2622).

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supplementary materials

Acta Cryst. (2013). E69, o490 [doi:10.1107/S1600536813005692]

(*E*)-*N'*-(3,4-Dihydroxybenzylidene)-2,4-dimethylbenzohydrazide monohydrate

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Comment

Structurally diverse range of benzohydrazides have been extensively studied in order to explore the structural features that may be responsible for different biological activities (Musharraf *et al.*, 2012; Khan *et al.*, 2012). The title compound is yet another benzohydrazide monohydrate, obtained as a part of our ongoing research that has been studied by X-ray crystallographic method and reported in this article.

In the title compound (Fig. 1) dimethyl and dihydroxy substituted benzene rings (C1–C6 and C9–C14, respectively) are each planar with a dihedral angle $30.27(7)^\circ$ between their mean-planes. The azomethine double bond, N2=C8 (1.2729(19) Å) adopts an *E* configuration. The bond lengths and angle are similar to the corresponding bond lengths and angles reported in structurally related benzohydrazide derivatives (Taha *et al.*, 2012; Baharudin *et al.*, 2012). The crystal structure is stabilized by N1—H1A···N2, O2—H2A···O1, C8—H8A···O3 and C15—H15B···O2 intermolecular interactions. The interactions further extend the structure to a three dimensional network *via* O1W—H2W1···O1, O1W—H1A···O3 and O3—H3A···O1W interactions involving the water of hydration (Table 2 and Fig. 2).

Experimental

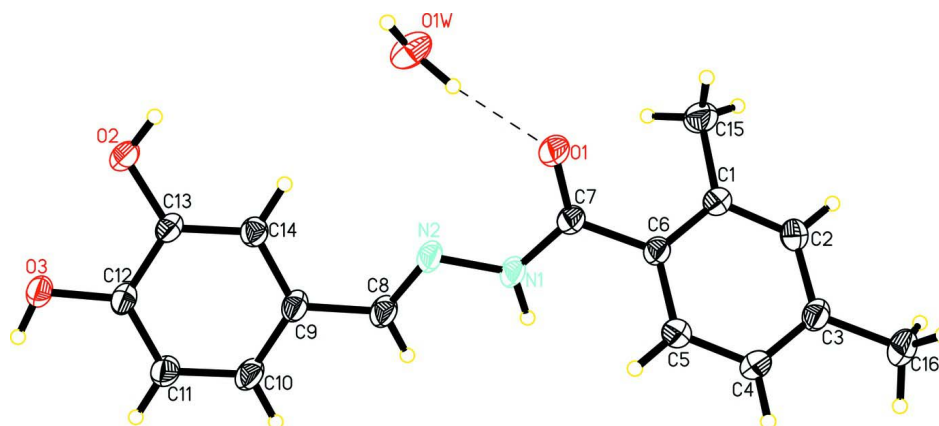
The title compound was synthesized by reacting (0.328 g, 2 mmol) 2,4-dimethylbenzohydrazide and (0.276 g, 2 mmol) 3,4-dihydroxybenzaldehyde as starting material under the same conditions and solvents as described previously for the synthesis of benzohydrazides (Taha *et al.*, 2012). The title compound was recrystallized by dissolving in methanol to obtain colorless needles (0.499 g, 88% yield). All chemicals were purchased by Sigma Aldrich Germany.

Refinement

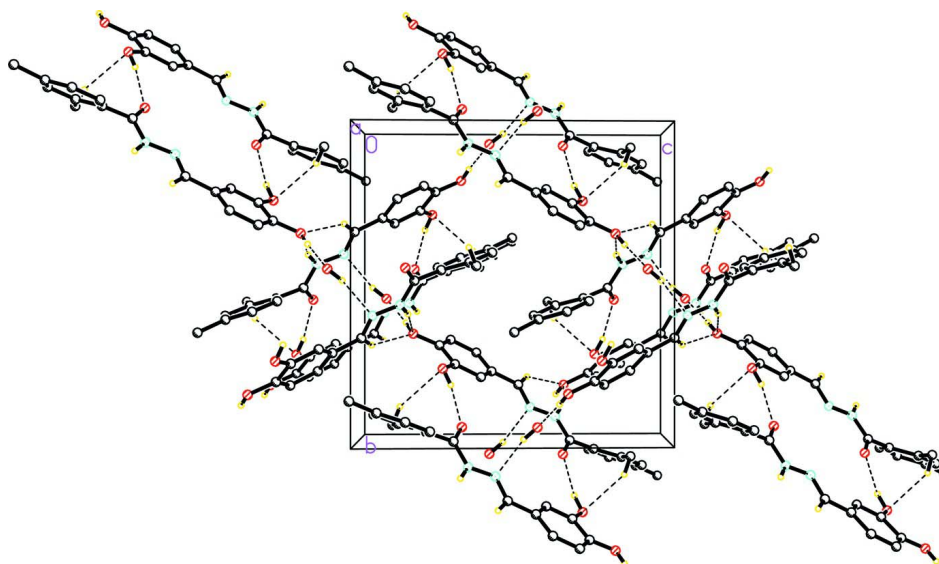
H atoms on methyl and benzene ring were positioned geometrically with C—H = 0.96 and 0.93 Å, respectively and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{benzene})$ or $1.5U_{\text{eq}}(\text{methyl})$. The H atoms on oxygen and nitrogen were located in difference Fourier map and refined isotropically. A rotating group model was applied to the methyl groups.

Computing details

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINT* (Bruker, 2000); data reduction: *SAINT* (Bruker, 2000); 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: *SHELXTL* (Sheldrick, 2008), *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2009).


Figure 1

The molecular structure of the title compound with displacement ellipsoids drawn at 30% probability level.


Figure 2

The crystal packing of the title compound. Only hydrogen atoms involved in hydrogen bonding are shown.

(*E*)-*N'*-(3,4-Dihydroxybenzylidene)-2,4-dimethylbenzohydrazide monohydrate

Crystal data

$C_{16}H_{16}N_2O_3 \cdot H_2O$

$M_r = 302.32$

Monoclinic, $P2_1/n$

Hall symbol: $-P 2_1/n$

$a = 8.1373 (3) \text{ \AA}$

$b = 13.9025 (5) \text{ \AA}$

$c = 13.7886 (5) \text{ \AA}$

$\beta = 92.913 (1)^\circ$

$V = 1557.87 (10) \text{ \AA}^3$

$Z = 4$

$F(000) = 640$

$D_x = 1.289 \text{ Mg m}^{-3}$

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

Cell parameters from 4856 reflections

$\theta = 2.8\text{--}28.2^\circ$

$\mu = 0.09 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Block, brown

$0.30 \times 0.10 \times 0.10 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector diffractometer	9012 measured reflections
Radiation source: fine-focus sealed tube	2897 independent reflections
Graphite monochromator	2535 reflections with $I > 2\sigma(I)$
ω scan	$R_{\text{int}} = 0.016$
Absorption correction: multi-scan (SADABS; Bruker, 2000)	$\theta_{\text{max}} = 25.5^\circ$, $\theta_{\text{min}} = 2.1^\circ$
$T_{\text{min}} = 0.973$, $T_{\text{max}} = 0.991$	$h = -9 \rightarrow 9$
	$k = -16 \rightarrow 16$
	$l = -16 \rightarrow 13$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.040$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.111$	$w = 1/[\sigma^2(F_o^2) + (0.0596P)^2 + 0.3352P]$
$S = 1.05$	where $P = (F_o^2 + 2F_c^2)/3$
2897 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
221 parameters	$\Delta\rho_{\text{max}} = 0.19 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.28 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	

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	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.15363 (13)	0.55077 (8)	-0.16328 (8)	0.0569 (3)
O1W	-0.12355 (15)	0.45259 (10)	-0.07769 (9)	0.0612 (3)
O2	-0.04800 (12)	0.26761 (9)	0.22605 (8)	0.0505 (3)
O3	0.16469 (13)	0.15980 (8)	0.32684 (7)	0.0460 (3)
N1	0.34634 (15)	0.43523 (9)	-0.14057 (9)	0.0454 (3)
N2	0.28104 (14)	0.40196 (9)	-0.05580 (8)	0.0435 (3)
C1	0.30375 (16)	0.56844 (10)	-0.36100 (10)	0.0395 (3)
C2	0.40413 (18)	0.60048 (11)	-0.43273 (10)	0.0446 (3)
H2C	0.3561	0.6153	-0.4935	0.054*
C3	0.57324 (17)	0.61142 (11)	-0.41767 (10)	0.0435 (3)
C4	0.64410 (17)	0.58608 (11)	-0.32796 (11)	0.0455 (4)
H4A	0.7574	0.5911	-0.3166	0.055*
C5	0.54846 (16)	0.55358 (10)	-0.25556 (10)	0.0423 (3)
H5A	0.5981	0.5371	-0.1956	0.051*
C6	0.37828 (16)	0.54484 (9)	-0.27016 (10)	0.0366 (3)
C7	0.28131 (16)	0.51150 (10)	-0.18789 (10)	0.0397 (3)

C8	0.35766 (18)	0.33077 (11)	-0.01702 (11)	0.0464 (4)
H8A	0.4495	0.3069	-0.0465	0.056*
C9	0.30656 (17)	0.28538 (10)	0.07190 (10)	0.0421 (3)
C10	0.41516 (19)	0.22570 (11)	0.12370 (12)	0.0514 (4)
H10A	0.5192	0.2145	0.1011	0.062*
C11	0.37047 (19)	0.18256 (11)	0.20869 (12)	0.0499 (4)
H11A	0.4448	0.1431	0.2432	0.060*
C12	0.21552 (16)	0.19792 (9)	0.24262 (10)	0.0390 (3)
C13	0.10379 (16)	0.25680 (10)	0.18973 (10)	0.0376 (3)
C14	0.14912 (16)	0.29972 (10)	0.10553 (10)	0.0392 (3)
H14A	0.0745	0.3387	0.0705	0.047*
C15	0.12199 (18)	0.55713 (14)	-0.38483 (12)	0.0571 (4)
H15A	0.1029	0.5506	-0.4538	0.086*
H15B	0.0648	0.6128	-0.3628	0.086*
H15C	0.0825	0.5008	-0.3531	0.086*
C16	0.6765 (2)	0.65068 (14)	-0.49624 (13)	0.0615 (5)
H16A	0.6324	0.6293	-0.5584	0.092*
H16B	0.7875	0.6280	-0.4863	0.092*
H16C	0.6755	0.7197	-0.4941	0.092*
H2A	-0.099 (3)	0.3180 (16)	0.1996 (15)	0.077 (6)*
H1A	0.436 (2)	0.4053 (13)	-0.1634 (13)	0.061 (5)*
H1W1	-0.165 (3)	0.4883 (17)	-0.0335 (16)	0.080 (7)*
H3A	0.239 (2)	0.1248 (15)	0.3533 (15)	0.071 (6)*
H2W1	-0.029 (3)	0.4798 (17)	-0.1022 (16)	0.091 (7)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0495 (6)	0.0633 (7)	0.0601 (7)	0.0209 (5)	0.0261 (5)	0.0182 (5)
O1W	0.0486 (6)	0.0766 (9)	0.0595 (7)	-0.0124 (6)	0.0122 (5)	-0.0274 (6)
O2	0.0399 (5)	0.0578 (7)	0.0557 (7)	0.0085 (5)	0.0200 (5)	0.0140 (5)
O3	0.0453 (6)	0.0499 (6)	0.0442 (6)	0.0056 (5)	0.0153 (5)	0.0141 (5)
N1	0.0462 (6)	0.0506 (7)	0.0414 (7)	0.0134 (6)	0.0227 (5)	0.0112 (5)
N2	0.0453 (6)	0.0477 (7)	0.0393 (6)	0.0066 (5)	0.0194 (5)	0.0077 (5)
C1	0.0360 (7)	0.0429 (7)	0.0400 (7)	0.0004 (5)	0.0051 (6)	0.0011 (6)
C2	0.0478 (8)	0.0505 (8)	0.0358 (7)	0.0008 (6)	0.0041 (6)	0.0062 (6)
C3	0.0437 (7)	0.0430 (8)	0.0448 (8)	-0.0010 (6)	0.0136 (6)	0.0037 (6)
C4	0.0335 (7)	0.0506 (8)	0.0530 (9)	-0.0033 (6)	0.0068 (6)	0.0038 (7)
C5	0.0383 (7)	0.0488 (8)	0.0396 (7)	0.0023 (6)	0.0014 (6)	0.0048 (6)
C6	0.0360 (7)	0.0379 (7)	0.0367 (7)	0.0023 (5)	0.0085 (5)	0.0023 (5)
C7	0.0378 (7)	0.0434 (7)	0.0387 (7)	0.0050 (6)	0.0105 (6)	0.0030 (6)
C8	0.0484 (8)	0.0466 (8)	0.0463 (8)	0.0108 (7)	0.0218 (6)	0.0064 (7)
C9	0.0468 (8)	0.0393 (7)	0.0418 (7)	0.0055 (6)	0.0176 (6)	0.0046 (6)
C10	0.0468 (8)	0.0526 (9)	0.0572 (9)	0.0149 (7)	0.0263 (7)	0.0122 (7)
C11	0.0469 (8)	0.0496 (9)	0.0547 (9)	0.0156 (7)	0.0171 (7)	0.0154 (7)
C12	0.0441 (7)	0.0352 (7)	0.0388 (7)	0.0008 (6)	0.0145 (6)	0.0039 (5)
C13	0.0370 (7)	0.0365 (7)	0.0405 (7)	0.0007 (5)	0.0128 (5)	-0.0002 (5)
C14	0.0420 (7)	0.0361 (7)	0.0401 (7)	0.0036 (6)	0.0082 (6)	0.0034 (6)
C15	0.0401 (8)	0.0788 (12)	0.0521 (9)	-0.0036 (7)	-0.0010 (7)	0.0037 (8)
C16	0.0567 (9)	0.0699 (11)	0.0598 (10)	-0.0030 (8)	0.0225 (8)	0.0146 (9)

Geometric parameters (Å, °)

O1—C7	1.2364 (16)	C5—C6	1.3945 (19)
O1W—H1W1	0.87 (2)	C5—H5A	0.9300
O1W—H2W1	0.94 (2)	C6—C7	1.4884 (18)
O2—C13	1.3645 (15)	C8—C9	1.4581 (19)
O2—H2A	0.88 (2)	C8—H8A	0.9300
O3—C12	1.3599 (16)	C9—C10	1.384 (2)
O3—H3A	0.85 (2)	C9—C14	1.3988 (19)
N1—C7	1.3398 (18)	C10—C11	1.382 (2)
N1—N2	1.3877 (15)	C10—H10A	0.9300
N1—H1A	0.913 (19)	C11—C12	1.3835 (19)
N2—C8	1.2729 (19)	C11—H11A	0.9300
C1—C2	1.3878 (19)	C12—C13	1.4004 (19)
C1—C6	1.4026 (19)	C13—C14	1.3725 (19)
C1—C15	1.5069 (19)	C14—H14A	0.9300
C2—C3	1.390 (2)	C15—H15A	0.9600
C2—H2C	0.9300	C15—H15B	0.9600
C3—C4	1.384 (2)	C15—H15C	0.9600
C3—C16	1.507 (2)	C16—H16A	0.9600
C4—C5	1.373 (2)	C16—H16B	0.9600
C4—H4A	0.9300	C16—H16C	0.9600
H1W1—O1W—H2W1	112 (2)	C10—C9—C14	119.07 (13)
C13—O2—H2A	110.5 (13)	C10—C9—C8	119.44 (12)
C12—O3—H3A	110.4 (13)	C14—C9—C8	121.48 (13)
C7—N1—N2	121.02 (11)	C11—C10—C9	120.62 (13)
C7—N1—H1A	119.8 (11)	C11—C10—H10A	119.7
N2—N1—H1A	119.2 (11)	C9—C10—H10A	119.7
C8—N2—N1	114.37 (11)	C10—C11—C12	120.24 (14)
C2—C1—C6	117.91 (12)	C10—C11—H11A	119.9
C2—C1—C15	119.00 (13)	C12—C11—H11A	119.9
C6—C1—C15	123.05 (12)	O3—C12—C11	123.41 (13)
C1—C2—C3	122.88 (13)	O3—C12—C13	117.06 (11)
C1—C2—H2C	118.6	C11—C12—C13	119.52 (12)
C3—C2—H2C	118.6	O2—C13—C14	123.35 (12)
C4—C3—C2	118.05 (13)	O2—C13—C12	116.69 (12)
C4—C3—C16	120.85 (13)	C14—C13—C12	119.96 (12)
C2—C3—C16	121.10 (13)	C13—C14—C9	120.57 (13)
C5—C4—C3	120.53 (13)	C13—C14—H14A	119.7
C5—C4—H4A	119.7	C9—C14—H14A	119.7
C3—C4—H4A	119.7	C1—C15—H15A	109.5
C4—C5—C6	121.27 (13)	C1—C15—H15B	109.5
C4—C5—H5A	119.4	H15A—C15—H15B	109.5
C6—C5—H5A	119.4	C1—C15—H15C	109.5
C5—C6—C1	119.32 (12)	H15A—C15—H15C	109.5
C5—C6—C7	118.57 (12)	H15B—C15—H15C	109.5
C1—C6—C7	122.11 (12)	C3—C16—H16A	109.5
O1—C7—N1	122.17 (12)	C3—C16—H16B	109.5
O1—C7—C6	123.86 (12)	H16A—C16—H16B	109.5

N1—C7—C6	113.96 (11)	C3—C16—H16C	109.5
N2—C8—C9	122.36 (12)	H16A—C16—H16C	109.5
N2—C8—H8A	118.8	H16B—C16—H16C	109.5
C9—C8—H8A	118.8		
C7—N1—N2—C8	-178.10 (14)	C5—C6—C7—N1	-45.73 (18)
C6—C1—C2—C3	-1.1 (2)	C1—C6—C7—N1	134.90 (14)
C15—C1—C2—C3	-178.87 (15)	N1—N2—C8—C9	-179.45 (13)
C1—C2—C3—C4	2.2 (2)	N2—C8—C9—C10	-163.72 (16)
C1—C2—C3—C16	-177.21 (15)	N2—C8—C9—C14	17.1 (2)
C2—C3—C4—C5	-1.7 (2)	C14—C9—C10—C11	-1.5 (2)
C16—C3—C4—C5	177.66 (15)	C8—C9—C10—C11	179.34 (15)
C3—C4—C5—C6	0.2 (2)	C9—C10—C11—C12	0.6 (3)
C4—C5—C6—C1	0.9 (2)	C10—C11—C12—O3	-178.29 (14)
C4—C5—C6—C7	-178.49 (13)	C10—C11—C12—C13	0.6 (2)
C2—C1—C6—C5	-0.5 (2)	O3—C12—C13—O2	-2.27 (19)
C15—C1—C6—C5	177.23 (14)	C11—C12—C13—O2	178.79 (14)
C2—C1—C6—C7	178.88 (13)	O3—C12—C13—C14	178.14 (12)
C15—C1—C6—C7	-3.4 (2)	C11—C12—C13—C14	-0.8 (2)
N2—N1—C7—O1	-5.9 (2)	O2—C13—C14—C9	-179.69 (13)
N2—N1—C7—C6	172.84 (12)	C12—C13—C14—C9	-0.1 (2)
C5—C6—C7—O1	133.03 (16)	C10—C9—C14—C13	1.3 (2)
C1—C6—C7—O1	-46.3 (2)	C8—C9—C14—C13	-179.59 (14)

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
O1 <i>W</i> —H1 <i>W</i> 1 \cdots N2 ⁱ	0.87 (2)	2.20 (2)	3.059 (2)	169 (2)
N1—H1 <i>A</i> \cdots O3 ⁱⁱ	0.91 (2)	2.08 (2)	2.962 (2)	163 (2)
O1 <i>W</i> —H2 <i>W</i> 1 \cdots O1	0.94 (2)	2.01 (2)	2.935 (2)	173 (2)
O2—H2 <i>A</i> \cdots O1 ⁱ	0.88 (2)	1.94 (2)	2.791 (2)	162 (2)
O3—H3 <i>A</i> \cdots O1 <i>W</i> ⁱⁱⁱ	0.85 (2)	1.79 (2)	2.629 (2)	172 (2)
C8—H8 <i>A</i> \cdots O3 ⁱⁱ	0.93	2.58	3.382 (2)	145
C15—H15 <i>B</i> \cdots O2 ⁱ	0.96	2.52	3.351 (2)	144

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