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3-[(3-Hydroxypropyl)amino]-1-phenylbut-2-en-1-one

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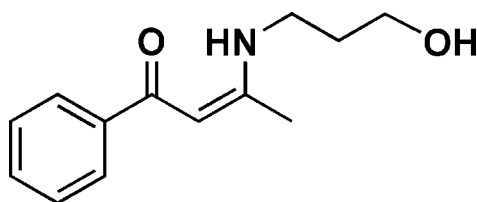
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.028; wR factor = 0.079; data-to-parameter ratio = 8.1.

The title compound, $\text{C}_{13}\text{H}_{17}\text{NO}_2$, has an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond, forming a planar six-membered ring with a mean deviation of 0.015 (5) Å from the plane. This plane makes a dihedral angle of 7.19 (8)° with the adjacent phenyl ring. Through an intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond, the molecules with their 2_1 screw and b -translation equivalents form a helical chain running parallel to the b axis.

Related literature

For general background, see: Morozova *et al.* (2007). For a related structure, see: Shi (2005).



Experimental

Crystal data

 $\text{C}_{13}\text{H}_{17}\text{NO}_2$ $M_r = 219.28$ Orthorhombic, $P2_12_12_1$ $a = 5.9131$ (3) Å $b = 8.0101$ (4) Å $c = 24.9626$ (13) Å $V = 1182.34$ (10) Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.08$ mm⁻¹ $T = 293$ (2) K

0.30 × 0.20 × 0.20 mm

Data collection

Bruker Kappa APEXII CCD diffractometer

Absorption correction: multi-scan (SADABS; Bruker, 1999)

 $T_{\min} = 0.944$, $T_{\max} = 0.984$

11541 measured reflections

1236 independent reflections

1168 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.022$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.079$ $S = 1.05$

1236 reflections

153 parameters

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\text{max}} = 0.12$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.09$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H2A}\cdots\text{O1}^1$	0.82	1.99	2.805 (2)	176
$\text{N1}-\text{H1N}\cdots\text{O1}$	0.85 (2)	1.94 (2)	2.642 (2)	139.1 (18)

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

Data collection: APEX2 (Bruker, 2004); cell refinement: SAINT-Plus (Bruker, 2004); data reduction: SAINT-Plus; program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 (Farrugia, 1997) and Mercury (Macrae *et al.*, 2006); software used to prepare material for publication: SHELXL97.

The authors express their thanks to the Sophisticated Analytical Instruments Facility, Indian Institute of Technology Madras, Chennai, for the collection of X-ray diffraction data. The authors are also grateful to the Department of Science and Technology (DST), India, for financial support (SR/S3/ME/03/2005-SERC) to carry out this work.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2371).

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supporting information

Acta Cryst. (2009). E65, o206 [doi:10.1107/S1600536808043183]

3-[(3-Hydroxypropyl)amino]-1-phenylbut-2-en-1-one

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S1. Comment

The Schiff base 1-phenyl-3-[(3-hydroxypropyl)amino]-1-butanone could be a good chelating ligand and may find use in the field of coordination chemistry of transition metal complexes. The compound could act as a bidentate ligand through the N and O atoms. The replacement of oxygen by nitrogen in the ligand can increase the covalency of the complexes (Morozova *et al.*, 2007).

Figure 1 gives *ORTEP* representation of the molecule with atoms represented as 50% anisotropic ellipsoids. Figure 2 gives packing of the molecules showing hydrogen bonded interactions. The molecules and their 2_1 screw translation equivalents are bound through O2—H2A \cdots O1 hydrogen bonds (Table 1). These H-bonded pairs are further linked with their b-translation equivalents to form an one-dimensional hydrogen bonded network parallel to *b* axis. There is an intramolecular N1—H1 \cdots O1 hydrogen bond between the imino hydrogen and the keto oxygen (Table 1). The packing is further stabilized through van der Waals interactions. The crystal is found to cleave easily through the (001) plane. The closely related compound, C₁₂H₁₅O₂N, (3-[(2-hydroxyethyl)amino]-1-phenylbut-2-en-1-one) crystallizes in monoclinic system with centrosymmetric space group *P*2₁/*n*, forming hydrogen bonded dimers in the structure (Shi, 2005), while the title compound crystallizes in polar space group *P*2₁2₁2₁ and with extended hydrogen bonding in the structure.

S2. Experimental

The title Schiff base ligand was synthesized by the condensation of 3-amino-1-propanol and benzoylacetone. To 0.1 molar solution of 3-amino-1-propanol (dissolved in 5 ml of ethanol) was slowly added to a 0.1 molar solution (in ethanol) of benzoylacetone. The reaction mixture was refluxed for 30 min. The solution was cooled overnight and the precipitate was washed with ethanol. The compound was crystallized in ethanol by slow evaporation (m.p. = 394 K). Anal. Calc. for C₁₃H₁₈O₂N: (Found %): C 70.88 (69.96), H 8.24 (8.13), N 6.35 (6.26). IR (KBr, cm⁻¹): 3170 = ν (O—H); 3340, ν (N—H); 1596, ν [(C—N)—C=C]; 1265, ν (C—O). ¹H NMR (400 MHz, CDCl₃) δ values at 1.8, 2.0, 3.4, 3.6, 5.7 and 7.4 p.p.m. for CH₃, CH₂, NH—CH₂, CH₂—OH, H—C=C and aromatic protons respectively. ¹³C NMR (400 MHz, CDCl₃) δ values at 14.83, 32.53, 40.06, 92.46, 128, 140.39 and 187 for CH₃, CH₂, -HN—CH₂, -CH₂OH, Aromatic, C—CH=C— and C=O carbon respectively. Mass Spectra: *M*⁺, *m/z* = 220.29, (*I* = 19%); *M* [L-(O—CH₂—CH₂—CH₂—N)]⁺, 148.10, (22); *M* [(C₆H₅-C=O)]⁺, 104.52, (100); *M* [(O—CH₂—CH₂—CH₂—N)]⁺, 74.63, (65).

S3. Refinement

All the hydrogen atoms could be located in a difference Fourier map. However, the H atoms except that of NH, were fixed at geometrically meaningful positions and refined using riding model. The riding tertiary CH₃ hydrogen atoms were assigned 1.5 times the equivalent displacement parameters of parent atoms, while 1.2 times was assigned for CH₂ and aromatic H atoms. The aromatic C—H distances were fixed at 0.93 Å while the secondary CH₂ and tertiary CH₃ were assigned 0.97 Å and 0.96 Å respectively. The isotropic displacement parameter of hydroxyl hydrogen was refined. In the

absence of significant anomalous scattering effects, Friedel pairs have been merged.

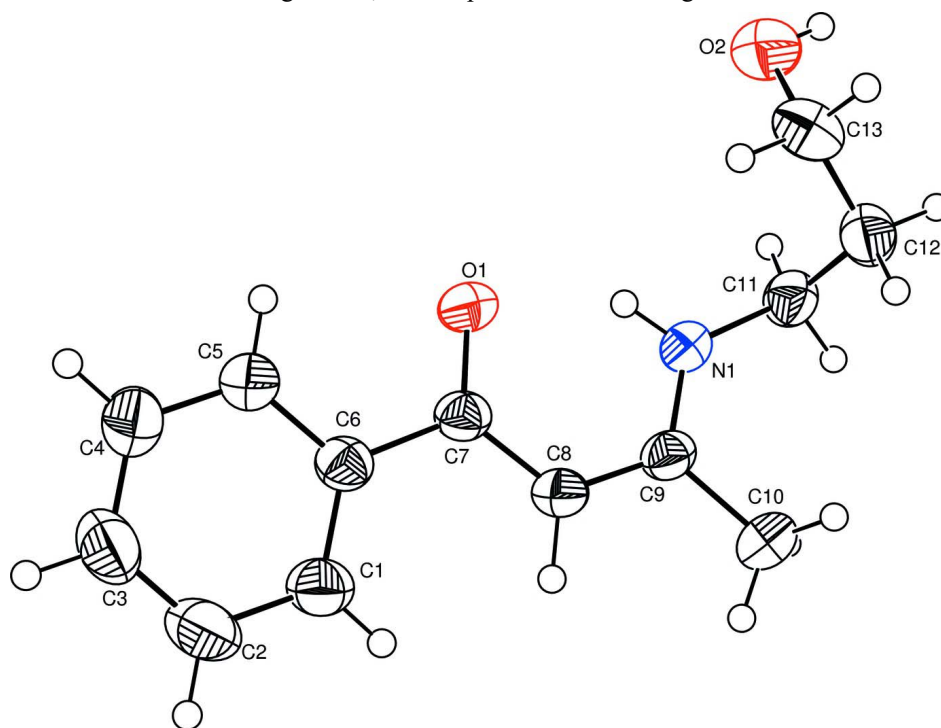
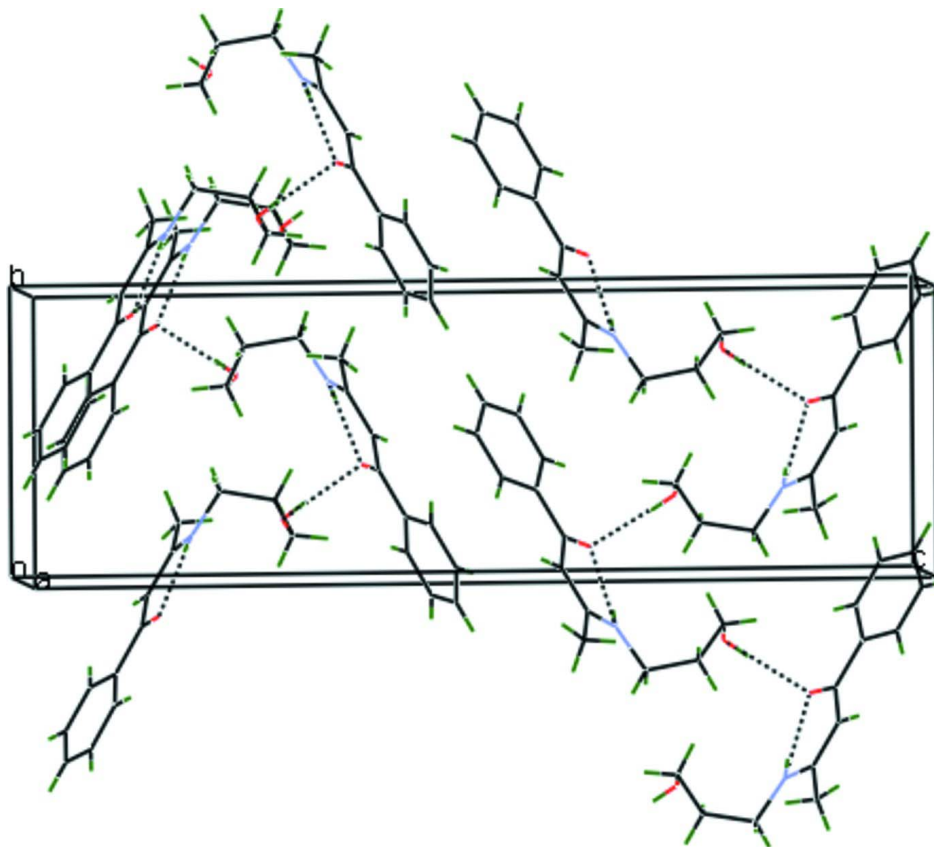


Figure 1

The *ORTEP* representation of the molecule with atoms represented as 50% probability ellipsoid.

**Figure 2**

Packing of molecules in the unit cell. Intra and intermolecular interactions are shown with dotted lines.

3-[(3-Hydroxypropyl)amino]-1-phenylbut-2-en-1-one

Crystal data

$C_{13}H_{17}NO_2$

$M_r = 219.28$

Orthorhombic, $P2_12_12_1$

Hall symbol: P 2ac 2ab

$a = 5.9131$ (3) Å

$b = 8.0101$ (4) Å

$c = 24.9626$ (13) Å

$V = 1182.34$ (10) Å³

$Z = 4$

$F(000) = 472$

$D_x = 1.232$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 7505 reflections

$\theta = 2.5\text{--}31.0^\circ$

$\mu = 0.08$ mm⁻¹

$T = 293$ K

Needle, colourless

0.30 × 0.20 × 0.20 mm

Data collection

Bruker Kappa APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and ϕ scan

Absorption correction: multi-scan
(*SADABS*; Bruker, 1999)

$T_{\min} = 0.944$, $T_{\max} = 0.984$

11541 measured reflections

1236 independent reflections

1168 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.7^\circ$

$h = -7 \rightarrow 5$

$k = -9 \rightarrow 9$

$l = -29 \rightarrow 28$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.028$
 $wR(F^2) = 0.079$
 $S = 1.05$
 1236 reflections
 153 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 atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0436P)^2 + 0.1578P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.12 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.09 \text{ e } \text{\AA}^{-3}$
 Extinction correction: *SHELXL*,
 $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.025 (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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.5484 (4)	-0.3200 (2)	0.04625 (8)	0.0547 (5)
H1	0.6664	-0.2439	0.0430	0.066*
C2	0.5575 (4)	-0.4692 (3)	0.01828 (8)	0.0652 (6)
H2	0.6805	-0.4922	-0.0037	0.078*
C3	0.3864 (4)	-0.5822 (3)	0.02298 (8)	0.0647 (6)
H3	0.3928	-0.6825	0.0043	0.078*
C4	0.2058 (4)	-0.5480 (3)	0.05513 (8)	0.0664 (6)
H4	0.0900	-0.6258	0.0586	0.080*
C5	0.1939 (4)	-0.3984 (3)	0.08253 (7)	0.0554 (5)
H5	0.0680	-0.3753	0.1036	0.066*
C6	0.3664 (3)	-0.2826 (2)	0.07901 (6)	0.0410 (4)
C7	0.3439 (3)	-0.1211 (2)	0.10934 (6)	0.0400 (4)
C8	0.5240 (3)	-0.0080 (2)	0.11068 (6)	0.0428 (4)
H8	0.6541	-0.0342	0.0915	0.048 (5)*
C9	0.5197 (3)	0.1415 (2)	0.13909 (6)	0.0409 (4)
C10	0.7269 (3)	0.2485 (3)	0.14091 (9)	0.0608 (5)
H9A	0.6929	0.3566	0.1264	0.091*
H9B	0.8447	0.1973	0.1201	0.091*
H9C	0.7762	0.2601	0.1774	0.091*
C11	0.3128 (3)	0.3395 (2)	0.19816 (7)	0.0489 (5)
H10A	0.1609	0.3829	0.1936	0.059*
H10B	0.4180	0.4238	0.1857	0.059*
C12	0.3543 (3)	0.3077 (3)	0.25711 (7)	0.0560 (5)

H11A	0.5089	0.2699	0.2618	0.067*
H11B	0.3377	0.4120	0.2765	0.067*
C13	0.1971 (4)	0.1805 (3)	0.28110 (7)	0.0575 (5)
H12A	0.2223	0.0735	0.2639	0.069*
H12B	0.2318	0.1679	0.3189	0.069*
N1	0.3385 (3)	0.19014 (19)	0.16559 (6)	0.0438 (4)
O1	0.1589 (2)	-0.09436 (16)	0.13305 (5)	0.0543 (4)
O2	-0.0328 (2)	0.2249 (2)	0.27548 (6)	0.0679 (4)
H2A	-0.0723	0.2808	0.3014	0.099 (10)*
H1N	0.227 (3)	0.124 (3)	0.1625 (8)	0.049 (5)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0546 (11)	0.0488 (10)	0.0607 (10)	-0.0008 (10)	0.0154 (9)	0.0000 (9)
C2	0.0728 (14)	0.0592 (12)	0.0638 (12)	0.0068 (12)	0.0237 (12)	-0.0057 (10)
C3	0.0883 (16)	0.0500 (11)	0.0559 (10)	-0.0010 (12)	0.0091 (12)	-0.0107 (9)
C4	0.0767 (16)	0.0610 (12)	0.0615 (11)	-0.0212 (12)	0.0108 (12)	-0.0123 (10)
C5	0.0540 (11)	0.0626 (11)	0.0495 (9)	-0.0137 (11)	0.0107 (9)	-0.0108 (9)
C6	0.0435 (9)	0.0453 (9)	0.0341 (7)	-0.0013 (8)	0.0018 (7)	0.0036 (7)
C7	0.0389 (9)	0.0474 (9)	0.0338 (7)	-0.0013 (8)	0.0041 (7)	0.0029 (7)
C8	0.0386 (9)	0.0511 (10)	0.0388 (8)	-0.0040 (8)	0.0076 (8)	-0.0018 (7)
C9	0.0359 (9)	0.0509 (9)	0.0359 (7)	-0.0064 (8)	0.0011 (7)	0.0042 (7)
C10	0.0451 (11)	0.0699 (13)	0.0674 (11)	-0.0171 (10)	0.0066 (9)	-0.0082 (11)
C11	0.0467 (10)	0.0418 (9)	0.0583 (9)	-0.0049 (9)	0.0044 (9)	-0.0059 (8)
C12	0.0481 (10)	0.0662 (12)	0.0538 (9)	0.0008 (11)	-0.0027 (9)	-0.0156 (9)
C13	0.0636 (13)	0.0598 (12)	0.0492 (9)	0.0125 (12)	0.0086 (9)	-0.0018 (9)
N1	0.0384 (8)	0.0449 (8)	0.0481 (7)	-0.0081 (8)	0.0042 (7)	-0.0052 (7)
O1	0.0418 (7)	0.0553 (7)	0.0659 (7)	-0.0085 (7)	0.0168 (6)	-0.0116 (6)
O2	0.0542 (9)	0.0851 (11)	0.0643 (8)	-0.0029 (9)	0.0093 (7)	-0.0059 (9)

Geometric parameters (Å, °)

C1—C6	1.385 (3)	C9—C10	1.496 (2)
C1—C2	1.385 (3)	C10—H9A	0.9600
C1—H1	0.9300	C10—H9B	0.9600
C2—C3	1.363 (3)	C10—H9C	0.9600
C2—H2	0.9300	C11—N1	1.454 (2)
C3—C4	1.364 (3)	C11—C12	1.513 (2)
C3—H3	0.9300	C11—H10A	0.9700
C4—C5	1.382 (3)	C11—H10B	0.9700
C4—H4	0.9300	C12—C13	1.504 (3)
C5—C6	1.381 (3)	C12—H11A	0.9700
C5—H5	0.9300	C12—H11B	0.9700
C6—C7	1.505 (2)	C13—O2	1.412 (3)
C7—O1	1.262 (2)	C13—H12A	0.9700
C7—C8	1.399 (2)	C13—H12B	0.9700
C8—C9	1.392 (2)	N1—H1N	0.85 (2)

C8—H8	0.9300	O2—H2A	0.8200
C9—N1	1.318 (2)		
C6—C1—C2	120.95 (19)	C9—C10—H9B	109.5
C6—C1—H1	119.5	H9A—C10—H9B	109.5
C2—C1—H1	119.5	C9—C10—H9C	109.5
C3—C2—C1	120.07 (19)	H9A—C10—H9C	109.5
C3—C2—H2	120.0	H9B—C10—H9C	109.5
C1—C2—H2	120.0	N1—C11—C12	112.86 (16)
C2—C3—C4	119.90 (19)	N1—C11—H10A	109.0
C2—C3—H3	120.0	C12—C11—H10A	109.0
C4—C3—H3	120.0	N1—C11—H10B	109.0
C3—C4—C5	120.4 (2)	C12—C11—H10B	109.0
C3—C4—H4	119.8	H10A—C11—H10B	107.8
C5—C4—H4	119.8	C13—C12—C11	113.63 (16)
C6—C5—C4	120.85 (18)	C13—C12—H11A	108.8
C6—C5—H5	119.6	C11—C12—H11A	108.8
C4—C5—H5	119.6	C13—C12—H11B	108.8
C5—C6—C1	117.82 (16)	C11—C12—H11B	108.8
C5—C6—C7	118.66 (15)	H11A—C12—H11B	107.7
C1—C6—C7	123.48 (16)	O2—C13—C12	112.62 (18)
O1—C7—C8	122.59 (15)	O2—C13—H12A	109.1
O1—C7—C6	117.30 (15)	C12—C13—H12A	109.1
C8—C7—C6	120.12 (15)	O2—C13—H12B	109.1
C9—C8—C7	123.76 (15)	C12—C13—H12B	109.1
C9—C8—H8	118.1	H12A—C13—H12B	107.8
C7—C8—H8	118.1	C9—N1—C11	127.51 (16)
N1—C9—C8	121.66 (16)	C9—N1—H1N	113.8 (13)
N1—C9—C10	118.77 (15)	C11—N1—H1N	118.7 (13)
C8—C9—C10	119.55 (15)	C13—O2—H2A	109.5
C9—C10—H9A	109.5		
C6—C1—C2—C3	-0.4 (3)	C1—C6—C7—C8	-7.9 (2)
C1—C2—C3—C4	0.2 (4)	O1—C7—C8—C9	1.8 (3)
C2—C3—C4—C5	0.8 (4)	C6—C7—C8—C9	-177.97 (15)
C3—C4—C5—C6	-1.6 (3)	C7—C8—C9—N1	-2.6 (2)
C4—C5—C6—C1	1.4 (3)	C7—C8—C9—C10	175.92 (16)
C4—C5—C6—C7	179.54 (18)	N1—C11—C12—C13	-59.7 (2)
C2—C1—C6—C5	-0.4 (3)	C11—C12—C13—O2	-57.9 (2)
C2—C1—C6—C7	-178.45 (18)	C8—C9—N1—C11	177.83 (15)
C5—C6—C7—O1	-5.7 (2)	C10—C9—N1—C11	-0.7 (3)
C1—C6—C7—O1	172.33 (17)	C12—C11—N1—C9	-95.5 (2)
C5—C6—C7—C8	174.08 (16)		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O2—H2A \cdots O1 ⁱ	0.82	1.99	2.805 (2)	176

N1—H1N···O1	0.85 (2)	1.94 (2)	2.642 (2)	139.1 (18)
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Symmetry code: (i) $-x, y+1/2, -z+1/2$.