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(Z)-3-[(2-Aminobenzyl)amino]-1-phenylbut-2-en-1-oneVedavalli Sairaj,^a Thothadri Srinivasan,^b Muthusamy Kandaswamy^a and Devadasan Velmurugan^{b*}^aDepartment of Inorganic Chemistry, University of Madras, Guindy Campus, Chennai 600 025, India, and ^bCentre of Advanced Study in Crystallography and Biophysics, University of Madras, Guindy Campus, Chennai 600 025, India
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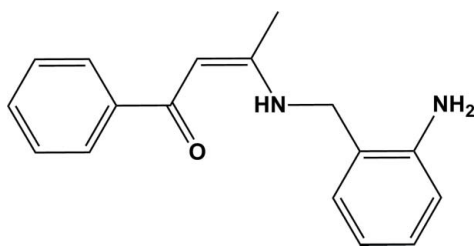
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.050; wR factor = 0.156; data-to-parameter ratio = 21.4.

In the title compound, $\text{C}_{17}\text{H}_{18}\text{N}_2\text{O}$, the aromatic rings are almost normal to one another, making a dihedral angle of $89.00(8)^\circ$. There is an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond in the molecule enclosing an $S(6)$ ring motif. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, forming chains along $[010]$.

Related literature

For the biological activity of chalcones, see: Di Carlo *et al.* (1999); Lin *et al.* (2002). For a related structure, see: Ranjith *et al.* (2010).



Experimental

Crystal data

 $\text{C}_{17}\text{H}_{18}\text{N}_2\text{O}$
 $M_r = 266.33$
Monoclinic, $P2_1/c$
 $a = 11.3197(4)$ Å $b = 9.8341(3)$ Å
 $c = 13.4207(4)$ Å
 $\beta = 106.387(2)^\circ$
 $V = 1433.29(8)$ Å³ $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹ $T = 293$ K
 $0.30 \times 0.25 \times 0.20$ mm

Data collection

Bruker SMART APEXII area-detector diffractometer
Absorption correction: multi-scan (SADABS; Bruker, 2008)
 $T_{\min} = 0.698$, $T_{\max} = 0.746$ 14860 measured reflections
3921 independent reflections
2528 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.156$
 $S = 1.03$
3921 reflections183 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2A}\cdots\text{O1}$	0.86	1.99	2.6619 (17)	134
$\text{N1}-\text{H1A}\cdots\text{O1}^i$	0.86	2.27	3.0009 (19)	143

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

Data collection: APEX2 (Bruker, 2008); cell refinement: SAINT (Bruker, 2008); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 2012); software used to prepare material for publication: SHELXL97 and PLATON (Spek, 2009).

The authors thank the TBI X-ray facility, CAS in Crystallography and Biophysics, University of Madras, India for data collection. TS also thanks the DST for an Inspire Fellowship. The UGC (SAP-CAS) is acknowledged for departmental facilities.

Supporting information for this paper is available from the IUCr electronic archives (Reference: SU2735).

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supporting information

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(Z)-3-[(2-Aminobenzyl)amino]-1-phenylbut-2-en-1-one**Vedavalli Sairaj, Thothadri Srinivasan, Muthusamy Kandaswamy and Devadasan Velmurugan****S1. Comment**

Chalcones are a major class of natural products with widespread distribution in fruits, vegetables, spices, tea and soy based foodstuff and have recently been the subject of great interest for their interesting pharmacological activities (Di Carlo *et al.*, 1999). Chalcones and flavonoids have been reported to be active anti-tuberculosis agents (Lin *et al.*, 2002). Against this background and in order to obtain detailed information on molecular conformation in the solid state, an X-ray study of the title compound was carried out.

In the title compound, Fig. 1, the aminobenzyl ring (C1-C6) and the phenyl ring (C12-C17) are normal to one another with a dihedral angle of 89.00 (8)°. The amine N atom, N1, attached to phenyl ring (C1-C6), deviates by only -0.0020 (16) Å from the ring plane. There is an intramolecular N-H...O hydrogen bonds enclosing an S(6) ring motif.

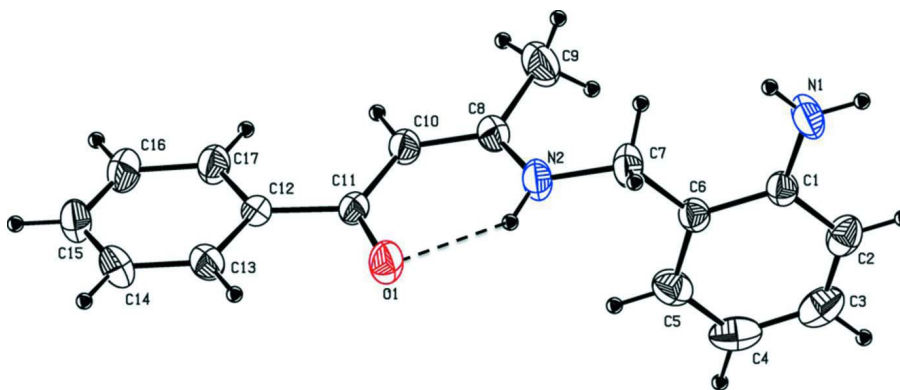
In the crystal, molecules are linked by N-H...O hydrogen bonds forming chains along the b axis direction (Table 1 and Fig. 2).

S2. Experimental

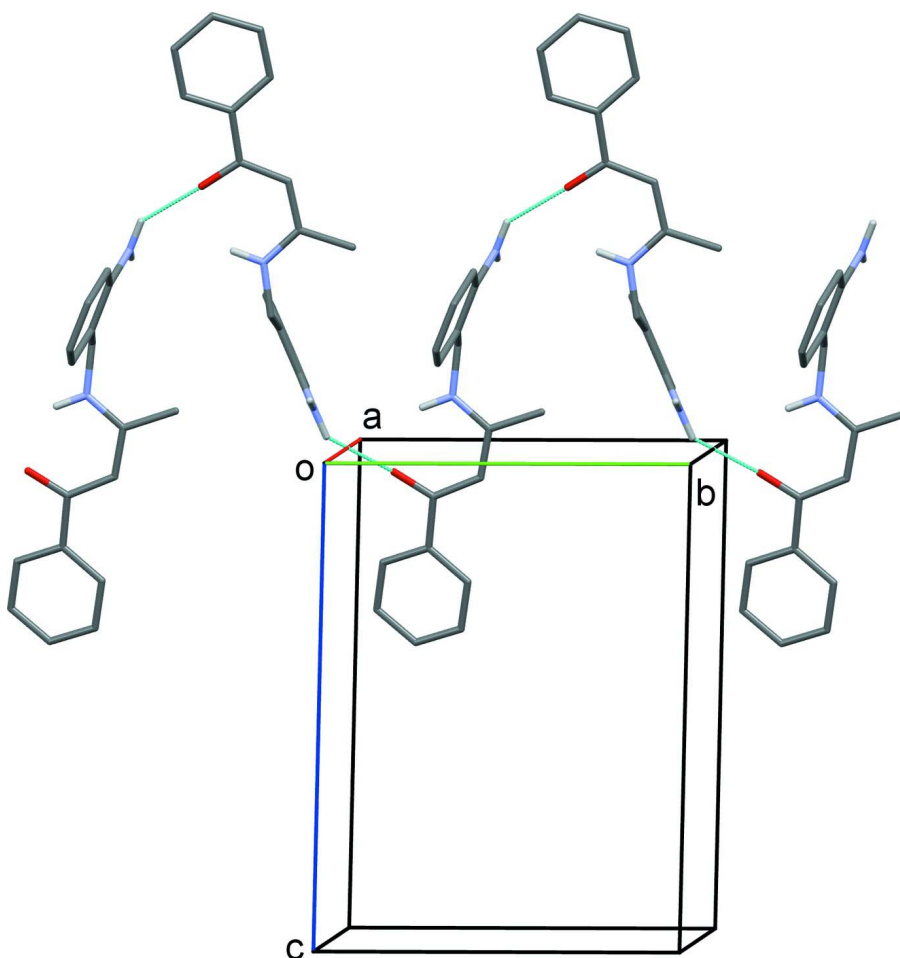
To a warm ethanolic solution (25 ml) of 2-aminobenzylamine (0.25 g, 0.2 mmol), an ethanolic solution of benzylacetone (0.3 g, 0.2 mmol) was added dropwise and the resulting solution was refluxed for 3 h. The solution was then filtered hot and allowed to stand at room temperature. After slow evaporation of the solvent at 298 K, block-like colourless crystals of the title compound were obtained. They were filtered off, washed with cold methanol and dried [Yield 0.45 g, 83%].

S3. Refinement

Hydrogen atoms were placed in calculated positions and refined as riding atoms: N-H = 0.86 Å, C-H = 0.93- 0.97 Å, with $U_{iso}(H) = 1.5U_{eq}(C\text{-methyl})$ and $= 1.2U_{eq}(N,C)$ for other H atoms.

**Figure 1**

The molecular structure of the title compound, with atom labelling. Displacement ellipsoids are drawn at the 30% probability level. The intramolecular N-H...O hydrogen bond is shown as a dashed line (see Table 1 for details).

**Figure 2**

A partial view along the *a* axis of the crystal packing of the title compound. The hydrogen bonds are shown as dashed lines (see Table 1 for details).

(Z)-3-[(2-Aminobenzyl)amino]-1-phenylbut-2-en-1-one*Crystal data*C₁₇H₁₈N₂O $M_r = 266.33$ Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

 $a = 11.3197 (4) \text{ \AA}$ $b = 9.8341 (3) \text{ \AA}$ $c = 13.4207 (4) \text{ \AA}$ $\beta = 106.387 (2)^\circ$ $V = 1433.29 (8) \text{ \AA}^3$ $Z = 4$ $F(000) = 568$ $D_x = 1.234 \text{ Mg m}^{-3}$ Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 3921 reflections

 $\theta = 1.9\text{--}29.3^\circ$ $\mu = 0.08 \text{ mm}^{-1}$ $T = 293 \text{ K}$

Block, colourless

 $0.30 \times 0.25 \times 0.20 \text{ mm}$ *Data collection*Bruker SMART APEXII area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 ω and φ scansAbsorption correction: multi-scan
(*SADABS*; Bruker, 2008) $T_{\min} = 0.698$, $T_{\max} = 0.746$

14860 measured reflections

3921 independent reflections

2528 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.028$ $\theta_{\max} = 29.3^\circ$, $\theta_{\min} = 1.9^\circ$ $h = -15 \rightarrow 15$ $k = -13 \rightarrow 13$ $l = -10 \rightarrow 18$ *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.050$ $wR(F^2) = 0.156$ $S = 1.03$

3921 reflections

183 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0715P)^2 + 0.2241P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.23 \text{ e \AA}^{-3}$ $\Delta\rho_{\min} = -0.16 \text{ e \AA}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.020 (3)

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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	0.37143 (13)	0.36601 (14)	-0.34891 (10)	0.0447 (3)
C2	0.24361 (15)	0.37189 (17)	-0.38045 (13)	0.0588 (4)
H2	0.2035	0.4073	-0.4455	0.071*

C3	0.17631 (16)	0.3268 (2)	-0.31782 (17)	0.0701 (5)
H3	0.0908	0.3311	-0.3407	0.084*
C4	0.23319 (19)	0.27475 (19)	-0.22096 (17)	0.0724 (5)
H4	0.1867	0.2442	-0.1783	0.087*
C5	0.36069 (17)	0.26853 (18)	-0.18788 (13)	0.0601 (4)
H5	0.3995	0.2335	-0.1224	0.072*
C6	0.43142 (13)	0.31337 (15)	-0.25025 (10)	0.0445 (3)
C7	0.56975 (14)	0.3028 (2)	-0.21832 (11)	0.0602 (4)
H7A	0.5930	0.2264	-0.2545	0.072*
H7B	0.6031	0.3844	-0.2407	0.072*
C8	0.66303 (14)	0.38414 (16)	-0.03882 (11)	0.0516 (4)
C9	0.6376 (2)	0.5280 (2)	-0.07480 (15)	0.0830 (6)
H9A	0.6843	0.5498	-0.1222	0.125*
H9B	0.6607	0.5880	-0.0161	0.125*
H9C	0.5513	0.5384	-0.1092	0.125*
C10	0.72605 (14)	0.35770 (15)	0.06411 (11)	0.0492 (4)
H10	0.7488	0.4313	0.1090	0.059*
C11	0.75754 (13)	0.22683 (14)	0.10462 (10)	0.0438 (3)
C12	0.83849 (12)	0.20920 (14)	0.21386 (10)	0.0414 (3)
C13	0.88339 (15)	0.08133 (17)	0.24592 (12)	0.0559 (4)
H13	0.8632	0.0086	0.1999	0.067*
C14	0.95824 (18)	0.0598 (2)	0.34596 (13)	0.0714 (5)
H14	0.9888	-0.0267	0.3663	0.086*
C15	0.98725 (17)	0.1654 (2)	0.41484 (12)	0.0709 (5)
H15	1.0376	0.1509	0.4819	0.085*
C16	0.94187 (17)	0.2926 (2)	0.38466 (12)	0.0691 (5)
H16	0.9604	0.3641	0.4318	0.083*
C17	0.86870 (15)	0.31536 (17)	0.28458 (11)	0.0568 (4)
H17	0.8395	0.4025	0.2645	0.068*
N1	0.43740 (14)	0.41274 (17)	-0.41456 (10)	0.0704 (4)
H1A	0.3993	0.4452	-0.4745	0.084*
H1B	0.5165	0.4093	-0.3954	0.084*
N2	0.62611 (13)	0.28532 (14)	-0.10721 (9)	0.0572 (4)
H2A	0.6362	0.2033	-0.0841	0.069*
O1	0.72160 (13)	0.12117 (11)	0.05299 (8)	0.0710 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0496 (8)	0.0388 (8)	0.0424 (7)	0.0033 (6)	0.0076 (6)	-0.0008 (5)
C2	0.0501 (9)	0.0528 (9)	0.0654 (9)	0.0066 (7)	0.0031 (7)	-0.0081 (7)
C3	0.0493 (9)	0.0607 (11)	0.0986 (14)	-0.0023 (8)	0.0179 (9)	-0.0194 (10)
C4	0.0741 (12)	0.0607 (11)	0.0985 (14)	-0.0168 (9)	0.0506 (11)	-0.0133 (10)
C5	0.0728 (11)	0.0578 (10)	0.0551 (8)	-0.0043 (8)	0.0270 (8)	0.0019 (7)
C6	0.0493 (8)	0.0446 (8)	0.0381 (6)	0.0003 (6)	0.0101 (5)	0.0000 (6)
C7	0.0511 (8)	0.0879 (13)	0.0373 (7)	0.0072 (8)	0.0052 (6)	0.0062 (7)
C8	0.0517 (8)	0.0490 (9)	0.0492 (7)	0.0033 (7)	0.0065 (6)	0.0074 (6)
C9	0.1066 (16)	0.0575 (12)	0.0705 (11)	0.0052 (11)	0.0013 (11)	0.0193 (9)

C10	0.0554 (8)	0.0400 (8)	0.0446 (7)	-0.0008 (6)	0.0016 (6)	0.0002 (6)
C11	0.0445 (7)	0.0414 (8)	0.0402 (6)	-0.0044 (6)	0.0032 (5)	-0.0026 (5)
C12	0.0389 (7)	0.0425 (8)	0.0401 (6)	-0.0038 (5)	0.0069 (5)	0.0023 (5)
C13	0.0635 (10)	0.0474 (9)	0.0527 (8)	0.0000 (7)	0.0094 (7)	0.0081 (7)
C14	0.0772 (12)	0.0672 (12)	0.0623 (10)	0.0094 (9)	0.0074 (9)	0.0260 (9)
C15	0.0655 (11)	0.0922 (14)	0.0453 (8)	-0.0024 (10)	0.0001 (7)	0.0165 (9)
C16	0.0724 (11)	0.0783 (13)	0.0455 (8)	-0.0077 (10)	-0.0015 (8)	-0.0074 (8)
C17	0.0621 (9)	0.0502 (9)	0.0482 (8)	-0.0006 (7)	-0.0008 (7)	-0.0032 (7)
N1	0.0644 (9)	0.0951 (12)	0.0499 (7)	0.0127 (8)	0.0134 (6)	0.0302 (7)
N2	0.0638 (8)	0.0585 (8)	0.0395 (6)	0.0052 (6)	-0.0012 (6)	0.0044 (5)
O1	0.0959 (9)	0.0427 (7)	0.0538 (6)	-0.0044 (6)	-0.0122 (6)	-0.0065 (5)

Geometric parameters (Å, °)

C1—N1	1.384 (2)	C9—H9C	0.9600
C1—C2	1.389 (2)	C10—C11	1.403 (2)
C1—C6	1.4053 (18)	C10—H10	0.9300
C2—C3	1.358 (3)	C11—O1	1.2516 (16)
C2—H2	0.9300	C11—C12	1.5036 (17)
C3—C4	1.376 (3)	C12—C13	1.379 (2)
C3—H3	0.9300	C12—C17	1.387 (2)
C4—C5	1.386 (3)	C13—C14	1.387 (2)
C4—H4	0.9300	C13—H13	0.9300
C5—C6	1.384 (2)	C14—C15	1.367 (3)
C5—H5	0.9300	C14—H14	0.9300
C6—C7	1.506 (2)	C15—C16	1.369 (3)
C7—N2	1.4571 (17)	C15—H15	0.9300
C7—H7A	0.9700	C16—C17	1.383 (2)
C7—H7B	0.9700	C16—H16	0.9300
C8—N2	1.3210 (19)	C17—H17	0.9300
C8—C10	1.3888 (19)	N1—H1A	0.8600
C8—C9	1.496 (2)	N1—H1B	0.8600
C9—H9A	0.9600	N2—H2A	0.8600
C9—H9B	0.9600		
N1—C1—C2	119.65 (13)	H9B—C9—H9C	109.5
N1—C1—C6	121.19 (13)	C8—C10—C11	124.07 (13)
C2—C1—C6	119.16 (14)	C8—C10—H10	118.0
C3—C2—C1	121.04 (16)	C11—C10—H10	118.0
C3—C2—H2	119.5	O1—C11—C10	122.65 (12)
C1—C2—H2	119.5	O1—C11—C12	117.23 (12)
C2—C3—C4	120.75 (16)	C10—C11—C12	120.12 (12)
C2—C3—H3	119.6	C13—C12—C17	118.34 (13)
C4—C3—H3	119.6	C13—C12—C11	118.62 (13)
C3—C4—C5	119.09 (16)	C17—C12—C11	123.03 (13)
C3—C4—H4	120.5	C12—C13—C14	120.79 (16)
C5—C4—H4	120.5	C12—C13—H13	119.6
C6—C5—C4	121.33 (16)	C14—C13—H13	119.6

C6—C5—H5	119.3	C15—C14—C13	120.23 (17)
C4—C5—H5	119.3	C15—C14—H14	119.9
C5—C6—C1	118.63 (14)	C13—C14—H14	119.9
C5—C6—C7	122.58 (13)	C14—C15—C16	119.68 (15)
C1—C6—C7	118.75 (12)	C14—C15—H15	120.2
N2—C7—C6	114.79 (13)	C16—C15—H15	120.2
N2—C7—H7A	108.6	C15—C16—C17	120.48 (16)
C6—C7—H7A	108.6	C15—C16—H16	119.8
N2—C7—H7B	108.6	C17—C16—H16	119.8
C6—C7—H7B	108.6	C16—C17—C12	120.47 (16)
H7A—C7—H7B	107.5	C16—C17—H17	119.8
N2—C8—C10	121.76 (14)	C12—C17—H17	119.8
N2—C8—C9	118.48 (13)	C1—N1—H1A	120.0
C10—C8—C9	119.75 (15)	C1—N1—H1B	120.0
C8—C9—H9A	109.5	H1A—N1—H1B	120.0
C8—C9—H9B	109.5	C8—N2—C7	125.86 (14)
H9A—C9—H9B	109.5	C8—N2—H2A	117.1
C8—C9—H9C	109.5	C7—N2—H2A	117.1
H9A—C9—H9C	109.5		
N1—C1—C2—C3	179.97 (16)	C8—C10—C11—C12	172.88 (14)
C6—C1—C2—C3	-0.5 (2)	O1—C11—C12—C13	9.5 (2)
C1—C2—C3—C4	0.4 (3)	C10—C11—C12—C13	-170.01 (14)
C2—C3—C4—C5	-0.2 (3)	O1—C11—C12—C17	-169.53 (15)
C3—C4—C5—C6	0.0 (3)	C10—C11—C12—C17	11.0 (2)
C4—C5—C6—C1	0.0 (2)	C17—C12—C13—C14	-0.9 (2)
C4—C5—C6—C7	-177.47 (16)	C11—C12—C13—C14	-179.89 (15)
N1—C1—C6—C5	179.81 (15)	C12—C13—C14—C15	0.9 (3)
C2—C1—C6—C5	0.3 (2)	C13—C14—C15—C16	0.1 (3)
N1—C1—C6—C7	-2.6 (2)	C14—C15—C16—C17	-1.1 (3)
C2—C1—C6—C7	177.83 (14)	C15—C16—C17—C12	1.1 (3)
C5—C6—C7—N2	-19.4 (2)	C13—C12—C17—C16	-0.1 (2)
C1—C6—C7—N2	163.10 (14)	C11—C12—C17—C16	178.86 (15)
N2—C8—C10—C11	0.8 (3)	C10—C8—N2—C7	-174.01 (15)
C9—C8—C10—C11	-178.05 (17)	C9—C8—N2—C7	4.9 (3)
C8—C10—C11—O1	-6.6 (3)	C6—C7—N2—C8	-90.1 (2)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N2—H2A...O1	0.86	1.99	2.6619 (17)	134
N1—H1A...O1 ⁱ	0.86	2.27	3.0009 (19)	143

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