

2-Amino-4-methylpyridinium 2-(3-methylphenyl)-acetate

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Received 5 July 2016

Accepted 6 July 2016

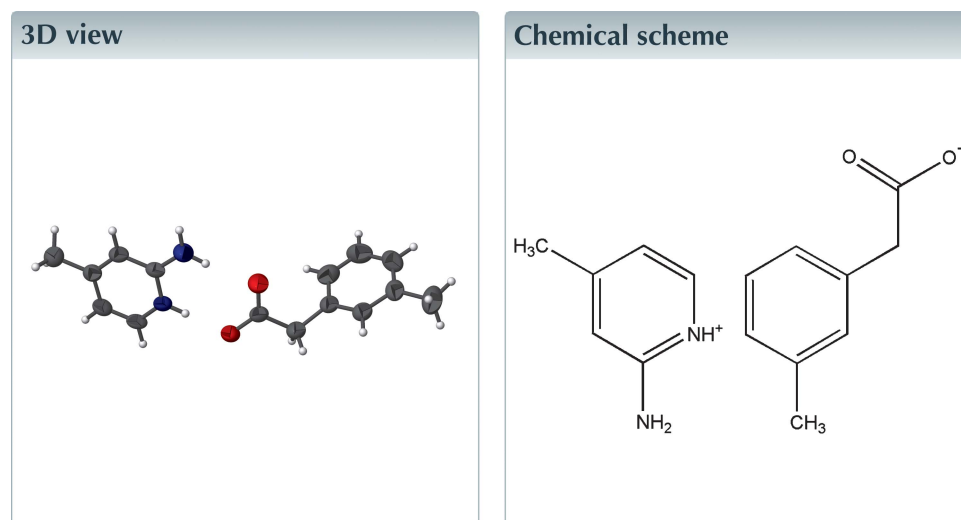
Edited by L. Van Meervelt, Katholieke Universiteit Leuven, Belgium

Keywords: molecular salt; crystal structure; hydrogen bonding.

CCDC reference: 1491153

Structural data: full structural data are available from iucrdata.iucr.org

In the title molecular salt, $C_6H_9N_2^+ \cdot C_9H_9O_2^-$, the cation is protonated at the pyridine N atom and the anion is deprotonated at the hydroxy O atom. The dihedral angle between the benzene and pyridine rings is $66.58(10)^\circ$. In the molecular structure, a pair of $N-H \cdots O$ hydrogen bonds links the anion and cation, generating an $R_2^2(8)$ ring motif. These ring motifs are connected to adjacent anions and cations *via* intermolecular $N-H \cdots O$ hydrogen bonding, generating a bifurcated $R_2^2(8)$ ring motif. $C-H \cdots O$, $C-H \cdots \pi$ and $\pi-\pi$ [centroid-to-centroid distances = $3.7053(11)$ and $3.9547(13)$ Å] interactions lead to the formation of a three-dimensional network.



Structure description

Pyridine derivatives exhibit antifungal, anticancer and anti-inflammatory activities (Luo & Hu, 2006; Liu & Hu, 2002). We herein report the synthesis and the crystal structure of the title compound (Fig. 1). The geometric parameters are comparable with those reported for similar structures (Divya Bharathi *et al.*, 2015; Sivakumar *et al.*, 2016).

The asymmetric unit of (Fig. 1) consists of a 2-amino-4-methylpyridinium cation, which is protonated at atom N1, and a 3-methylphenylacetate anion, which is deprotonated at the hydroxy O atom. The dihedral angle between the benzene ring (C1–C6) and the pyridine ring (N2/C10–C14) is $66.58(10)^\circ$. $N1-H1 \cdots O1$ and $N2-H2A \cdots O2$ hydrogen bonds (Table 1) link the anion and cation, generating an $R_2^2(8)$ ring motif (Fig. 2). These ring motifs are connected with adjacent anions and cations *via* $N2-H2B \cdots O2^i$ hydrogen bonds (Table 1), generating a bifurcated $R_2^2(8)$ ring motif (Fig. 2). Intermolecular $C-H \cdots O$, $C-H \cdots \pi$ (Table 1) and $\pi-\pi$ [$Cg1 \cdots Cg1^i = 3.9547(13)$ Å; $Cg2 \cdots Cg2 = 3.7053(11)$ Å; symmetry codes: (i) $1-x, 2-y, 1-z$; (ii) $-x, 1-y, -z$; Cg1 and Cg2 are

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

Cg1 is the centroid of the C1–C6 ring.

<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>
N1–H1···O1	0.88 (2)	1.75 (2)	2.627 (2)	172 (2)
N2–H2A···O2	0.86	2.01	2.845 (2)	163
N2–H2B···O2 ⁱ	0.86	2.10	2.829 (2)	142
C14–H14···O1 ⁱⁱ	0.93	2.43	3.347 (2)	168
C11–H11···Cg1 ⁱ	0.93	2.81	3.706 (2)	161

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

the centroids of the rings (C1–C6) and (N2/C10–C14), respectively] interactions lead to the formation of a three dimensional network (Fig. 3).

Synthesis and crystallization

The title compound was synthesized using the raw materials *m*-tolylacetic acid (1.001 g) and 2-amino-4-methylpyridine (0.72 g) in an equimolar ratio. These reactants were dissolved in 15 ml acetone and kept for slow evaporation at room temperature. Crystals suitable for X-ray diffraction were harvested after two weeks.

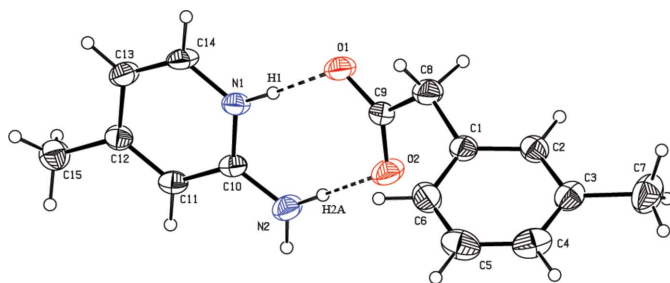


Figure 1
The molecular structure of the title molecular salt, showing the atom labelling and 30% probability displacement ellipsoids. Hydrogen bonds are shown as dashed lines.

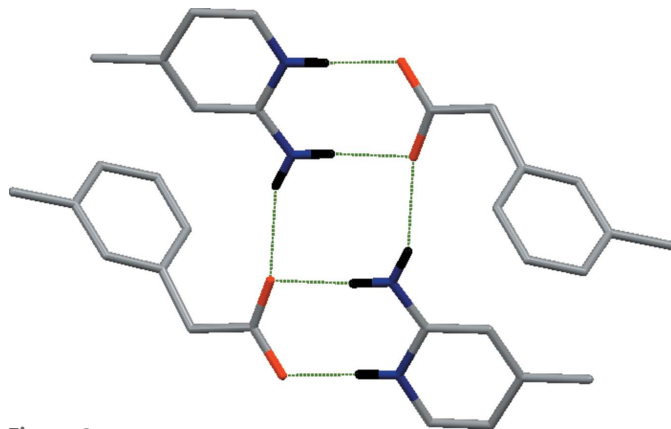


Figure 2
A partial view of the crystal packing of the title molecular salt, showing the ring graph-set motifs. Hydrogen bonds are shown as dashed lines and C-bound H atoms which are not involved in interactions have been omitted for clarity.

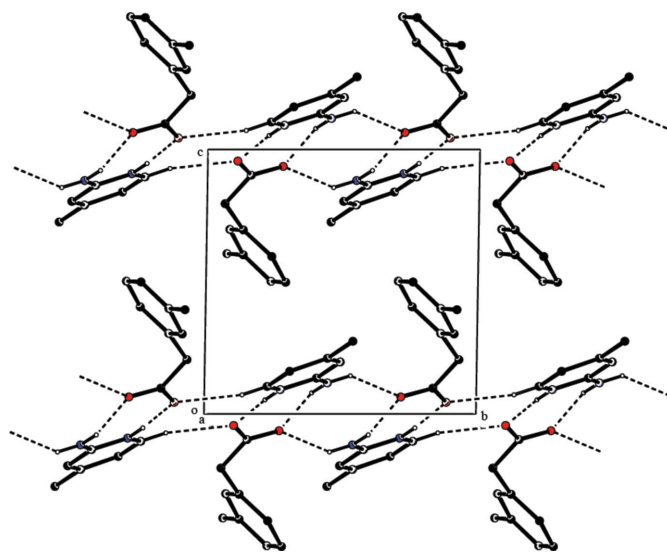


Figure 3
The crystal packing of the title compound, viewed along the *a* axis. Hydrogen bonds are shown as dashed lines and C-bound H atoms which are not involved in interactions have been omitted for clarity.

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2.

Table 2
Experimental details.

Crystal data	
Chemical formula	$C_6H_9N_2^+ \cdot C_9H_9O_2^-$
M_r	258.31
Crystal system, space group	Triclinic, $P\bar{1}$
Temperature (K)	293
<i>a</i> , <i>b</i> , <i>c</i> (Å)	8.3253 (4), 9.5065 (5), 9.5909 (5)
α , β , γ (°)	85.526 (3), 70.885 (3), 78.863 (3)
<i>V</i> (Å ³)	703.62 (6)
<i>Z</i>	2
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.26 × 0.24 × 0.20
Data collection	
Diffractometer	Bruker Kappa APEXII CCD
Absorption correction	Multi-scan (SADABS; Bruker, 2004)
T_{min} , T_{max}	0.979, 0.984
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	15018, 3059, 1885
R_{int}	0.035
($\sin \theta/\lambda$) _{max} (Å ⁻¹)	0.641
Refinement	
$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, <i>S</i>	0.050, 0.150, 1.02
No. of reflections	3059
No. of parameters	178
No. of restraints	2
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$, $\Delta\rho_{min}$ (e Å ⁻³)	0.22, -0.21

Computer programs: APEX2 and SAINT (Bruker, 2004), SHELXS97 and SHELXL97 (Sheldrick, 2008) and PLATON (Spek, 2009).

Acknowledgements

The authors acknowledge the SAIF, IIT, Madras, for the data collection.

References

Bruker (2004). *APEX2*, *SAINTE* and *SADABS*. Bruker AXS Inc., Madison, Wisconsin, USA.

- Divya Bharathi, M., Ahila, G., Mohana, J., Chakkaravarthi, G. & Anbalagan, G. (2015). *Acta Cryst. E* **71**, o261–o262.
- Liu, T. & Hu, Y. (2002). *Bioorg. Med. Chem. Lett.* **12**, 2411–2413.
- Luo, Y. & Hu, Y. (2006). *Arch. Pharm. Chem. Life Sci.* **339**, 262–266.
- Sheldrick, G. M. (2008). *Acta Cryst. A* **64**, 112–122.
- Sivakumar, P., Sudhakar, S., Gunasekaran, B., Israel, S. & Chakkaravarthi, G. (2016). *IUCrData*, **1**, x160747.
- Spek, A. L. (2009). *Acta Cryst. D* **65**, 148–155.

full crystallographic data

IUCrData (2016). **1**, x161098 [https://doi.org/10.1107/S2414314616010981]

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Crystal data

$C_6H_9N_2^+ \cdot C_9H_9O_2^-$
 $M_r = 258.31$
 Triclinic, $P\bar{1}$
 Hall symbol: $-P\ 1$
 $a = 8.3253$ (4) Å
 $b = 9.5065$ (5) Å
 $c = 9.5909$ (5) Å
 $\alpha = 85.526$ (3)°
 $\beta = 70.885$ (3)°
 $\gamma = 78.863$ (3)°
 $V = 703.62$ (6) Å³

$Z = 2$
 $F(000) = 276$
 $D_x = 1.219$ Mg m⁻³
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 4380 reflections
 $\theta = 2.2$ – 27.0 °
 $\mu = 0.08$ mm⁻¹
 $T = 293$ K
 Block, colourless
 $0.26 \times 0.24 \times 0.20$ 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, 2004)
 $T_{\min} = 0.979$, $T_{\max} = 0.984$

15018 measured reflections
 3059 independent reflections
 1885 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.035$
 $\theta_{\max} = 27.1$ °, $\theta_{\min} = 2.2$ °
 $h = -10 \rightarrow 10$
 $k = -12 \rightarrow 12$
 $l = -12 \rightarrow 12$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.050$
 $wR(F^2) = 0.150$
 $S = 1.02$
 3059 reflections
 178 parameters
 2 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
 $w = 1/[\sigma^2(F_o^2) + (0.0613P)^2 + 0.2177P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.22$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.4822 (2)	0.8660 (2)	0.3042 (2)	0.0475 (5)
C2	0.6275 (3)	0.9154 (2)	0.3021 (2)	0.0490 (5)
H2	0.6663	0.9857	0.2329	0.059*
C3	0.7182 (3)	0.8647 (2)	0.3985 (2)	0.0558 (6)
C4	0.6600 (3)	0.7610 (3)	0.4989 (3)	0.0699 (6)
H4	0.7180	0.7261	0.5659	0.084*
C5	0.5160 (4)	0.7072 (3)	0.5021 (3)	0.0730 (7)
H5	0.4787	0.6356	0.5701	0.088*
C6	0.4279 (3)	0.7596 (2)	0.4047 (3)	0.0619 (6)
H6	0.3312	0.7230	0.4069	0.074*
C7	0.8749 (3)	0.9245 (3)	0.3939 (3)	0.0870 (9)
H7A	0.8417	1.0247	0.4162	0.131*
H7B	0.9586	0.9121	0.2973	0.131*
H7C	0.9243	0.8747	0.4654	0.131*
C8	0.3800 (3)	0.9344 (2)	0.2050 (3)	0.0600 (6)
H8A	0.2665	0.9784	0.2672	0.072*
H8B	0.4360	1.0111	0.1492	0.072*
C9	0.3557 (2)	0.83992 (19)	0.0971 (2)	0.0458 (5)
C10	0.2100 (2)	0.58859 (19)	-0.1350 (2)	0.0409 (4)
C11	0.1335 (2)	0.5018 (2)	-0.1964 (2)	0.0443 (5)
H11	0.1923	0.4106	-0.2283	0.053*
C12	-0.0260 (2)	0.5491 (2)	-0.2101 (2)	0.0447 (5)
C13	-0.1123 (3)	0.6872 (2)	-0.1610 (2)	0.0526 (5)
H13	-0.2218	0.7217	-0.1680	0.063*
C14	-0.0352 (3)	0.7693 (2)	-0.1038 (2)	0.0524 (5)
H14	-0.0921	0.8610	-0.0720	0.063*
C15	-0.1096 (3)	0.4565 (2)	-0.2745 (3)	0.0619 (6)
H15A	-0.0876	0.4791	-0.3776	0.093*
H15B	-0.2320	0.4735	-0.2246	0.093*
H15C	-0.0630	0.3575	-0.2624	0.093*
N1	0.1233 (2)	0.72080 (16)	-0.09179 (18)	0.0446 (4)
N2	0.3643 (2)	0.54781 (18)	-0.1174 (2)	0.0547 (5)
H2A	0.4059	0.6054	-0.0794	0.066*
H2B	0.4226	0.4638	-0.1441	0.066*
O1	0.23722 (19)	0.88996 (14)	0.04377 (17)	0.0591 (4)

O2	0.45236 (18)	0.72287 (15)	0.06398 (18)	0.0653 (5)
H1	0.170 (2)	0.7762 (19)	-0.053 (2)	0.056 (6)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0480 (11)	0.0395 (10)	0.0534 (12)	-0.0016 (9)	-0.0159 (9)	-0.0084 (9)
C2	0.0526 (11)	0.0435 (11)	0.0482 (12)	-0.0073 (9)	-0.0121 (9)	-0.0053 (9)
C3	0.0525 (12)	0.0572 (13)	0.0575 (14)	0.0028 (10)	-0.0204 (10)	-0.0190 (11)
C4	0.0866 (15)	0.0690 (16)	0.0532 (14)	0.0105 (12)	-0.0327 (13)	-0.0107 (12)
C5	0.0947 (17)	0.0595 (14)	0.0528 (14)	-0.0116 (12)	-0.0110 (13)	0.0098 (12)
C6	0.0562 (13)	0.0530 (13)	0.0717 (16)	-0.0133 (10)	-0.0109 (11)	-0.0044 (12)
C7	0.0654 (15)	0.096 (2)	0.110 (2)	-0.0045 (14)	-0.0388 (15)	-0.0381 (17)
C8	0.0659 (13)	0.0416 (11)	0.0808 (16)	-0.0003 (10)	-0.0380 (12)	-0.0078 (11)
C9	0.0432 (10)	0.0340 (10)	0.0603 (13)	-0.0046 (8)	-0.0180 (10)	-0.0009 (9)
C10	0.0424 (10)	0.0351 (10)	0.0391 (10)	0.0012 (8)	-0.0099 (8)	0.0011 (8)
C11	0.0497 (11)	0.0347 (10)	0.0428 (11)	0.0011 (8)	-0.0116 (9)	-0.0033 (8)
C12	0.0494 (11)	0.0418 (11)	0.0411 (11)	-0.0067 (9)	-0.0134 (9)	0.0026 (8)
C13	0.0489 (11)	0.0459 (11)	0.0617 (14)	0.0041 (9)	-0.0229 (10)	-0.0005 (10)
C14	0.0546 (12)	0.0343 (10)	0.0639 (14)	0.0115 (9)	-0.0232 (10)	-0.0042 (9)
C15	0.0673 (14)	0.0570 (13)	0.0660 (15)	-0.0108 (11)	-0.0271 (12)	-0.0035 (11)
N1	0.0511 (9)	0.0319 (8)	0.0509 (10)	-0.0001 (7)	-0.0205 (8)	-0.0015 (7)
N2	0.0480 (9)	0.0461 (10)	0.0690 (12)	0.0067 (7)	-0.0235 (9)	-0.0115 (8)
O1	0.0656 (9)	0.0387 (8)	0.0813 (11)	0.0062 (7)	-0.0416 (8)	-0.0081 (7)
O2	0.0612 (9)	0.0488 (9)	0.0902 (12)	0.0131 (7)	-0.0384 (8)	-0.0210 (8)

Geometric parameters (Å, °)

C1—C2	1.374 (3)	C9—O1	1.254 (2)
C1—C6	1.377 (3)	C10—N2	1.330 (2)
C1—C8	1.507 (3)	C10—N1	1.344 (2)
C2—C3	1.380 (3)	C10—C11	1.400 (3)
C2—H2	0.9300	C11—C12	1.363 (2)
C3—C4	1.366 (3)	C11—H11	0.9300
C3—C7	1.507 (3)	C12—C13	1.405 (3)
C4—C5	1.381 (4)	C12—C15	1.498 (3)
C4—H4	0.9300	C13—C14	1.346 (3)
C5—C6	1.376 (3)	C13—H13	0.9300
C5—H5	0.9300	C14—N1	1.350 (2)
C6—H6	0.9300	C14—H14	0.9300
C7—H7A	0.9600	C15—H15A	0.9600
C7—H7B	0.9600	C15—H15B	0.9600
C7—H7C	0.9600	C15—H15C	0.9600
C8—C9	1.507 (3)	N1—H1	0.877 (9)
C8—H8A	0.9700	N2—H2A	0.8600
C8—H8B	0.9700	N2—H2B	0.8600
C9—O2	1.236 (2)		

C2—C1—C6	118.3 (2)	O2—C9—O1	124.30 (18)
C2—C1—C8	120.02 (19)	O2—C9—C8	119.97 (17)
C6—C1—C8	121.54 (19)	O1—C9—C8	115.70 (16)
C1—C2—C3	122.6 (2)	N2—C10—N1	118.07 (17)
C1—C2—H2	118.7	N2—C10—C11	123.63 (17)
C3—C2—H2	118.7	N1—C10—C11	118.30 (16)
C4—C3—C2	118.0 (2)	C12—C11—C10	120.93 (17)
C4—C3—C7	121.4 (2)	C12—C11—H11	119.5
C2—C3—C7	120.6 (2)	C10—C11—H11	119.5
C3—C4—C5	120.9 (2)	C11—C12—C13	118.45 (18)
C3—C4—H4	119.6	C11—C12—C15	121.33 (18)
C5—C4—H4	119.6	C13—C12—C15	120.22 (18)
C6—C5—C4	120.0 (2)	C14—C13—C12	119.53 (18)
C6—C5—H5	120.0	C14—C13—H13	120.2
C4—C5—H5	120.0	C12—C13—H13	120.2
C5—C6—C1	120.2 (2)	C13—C14—N1	121.13 (18)
C5—C6—H6	119.9	C13—C14—H14	119.4
C1—C6—H6	119.9	N1—C14—H14	119.4
C3—C7—H7A	109.5	C12—C15—H15A	109.5
C3—C7—H7B	109.5	C12—C15—H15B	109.5
H7A—C7—H7B	109.5	H15A—C15—H15B	109.5
C3—C7—H7C	109.5	C12—C15—H15C	109.5
H7A—C7—H7C	109.5	H15A—C15—H15C	109.5
H7B—C7—H7C	109.5	H15B—C15—H15C	109.5
C9—C8—C1	117.78 (17)	C10—N1—C14	121.64 (17)
C9—C8—H8A	107.9	C10—N1—H1	119.1 (14)
C1—C8—H8A	107.9	C14—N1—H1	119.2 (13)
C9—C8—H8B	107.9	C10—N2—H2A	120.0
C1—C8—H8B	107.9	C10—N2—H2B	120.0
H8A—C8—H8B	107.2	H2A—N2—H2B	120.0
C6—C1—C2—C3	-1.4 (3)	C1—C8—C9—O2	-18.8 (3)
C8—C1—C2—C3	174.70 (19)	C1—C8—C9—O1	162.85 (19)
C1—C2—C3—C4	0.3 (3)	N2—C10—C11—C12	179.30 (18)
C1—C2—C3—C7	-178.7 (2)	N1—C10—C11—C12	-1.0 (3)
C2—C3—C4—C5	0.9 (3)	C10—C11—C12—C13	0.0 (3)
C7—C3—C4—C5	179.9 (2)	C10—C11—C12—C15	-179.46 (18)
C3—C4—C5—C6	-0.9 (4)	C11—C12—C13—C14	0.8 (3)
C4—C5—C6—C1	-0.2 (4)	C15—C12—C13—C14	-179.8 (2)
C2—C1—C6—C5	1.4 (3)	C12—C13—C14—N1	-0.5 (3)
C8—C1—C6—C5	-174.7 (2)	N2—C10—N1—C14	-178.95 (18)
C2—C1—C8—C9	121.8 (2)	C11—C10—N1—C14	1.3 (3)
C6—C1—C8—C9	-62.2 (3)	C13—C14—N1—C10	-0.6 (3)

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C1–C6 ring.

<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>
N1—H1···O1	0.88 (2)	1.75 (2)	2.627 (2)	172 (2)
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