

Crystal structure of 1*H*-imidazol-3-i um 2-(1,3-dioxoisoindolin-2-yl)acetate

Shaaban K. Mohamed,^{a,b} Mehmet Akkurt,^{c*} Herman Potgieter^d and Muizz Ali^a

^aChemistry and Environmental Division, Manchester Metropolitan University, Manchester M1 5GD, England, ^bChemistry Department, Faculty of Science, Minia University, 61519 El-Minia, Egypt, ^cDepartment of Physics, Faculty of Sciences, Erciyes University, 38039 Kayseri, Turkey, and ^dAnalytical Development Division, Manchester Metropolitan University, Manchester M1 5GD, England. *Correspondence e-mail: akkurt@erciyes.edu.tr

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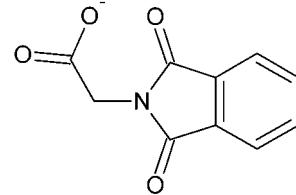
The title salt, $\text{C}_3\text{H}_5\text{N}_2^+\cdot\text{C}_{10}\text{H}_6\text{NO}_4^-$, was obtained during a study of the co-crystallization of *N*'-[bis(1*H*-imidazol-1-yl)methylene]isonicotinohydrazide with (1,3-dioxoisoindolin-2-yl)acetic acid under aqueous conditions. The 1,3-dioxoisoindolinyl ring system of the anion is essentially planar [maximum deviation = 0.023 (2) Å]. In the crystal, cations and anions are linked via classical N—H···O hydrogen bonds and weak C—H···O hydrogen bonds, forming a three-dimensional network. Weak C—H···π interactions and π—π stacking interactions [centroid–centroid distances = 3.4728 (13) and 3.7339 (13) Å] also occur in the crystal.

Keywords: crystal structure; 1*H*-imidazol-3-i um salt; 2-(1,3-dioxoisoindolin-2-yl)acetate salt; hydrogen bonding; π—π stacking interactions; co-crystallization; pharmaceuticals.

CCDC reference: 1017262

1. Related literature

For the use of co-crystals in drug design, see: Babu & Nangia (2011); Sekhon (2013); Frantz (2006); Pan *et al.* (2008); Vermeire *et al.* (2001).



2. Experimental

2.1. Crystal data

$\text{C}_3\text{H}_5\text{N}_2^+\cdot\text{C}_{10}\text{H}_6\text{NO}_4^-$
 $M_r = 273.25$
Monoclinic, $P2_1/c$
 $a = 9.8750$ (7) Å
 $b = 18.0543$ (15) Å
 $c = 7.0942$ (5) Å
 $\beta = 100.955$ (7)°

$V = 1241.75$ (16) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.11$ mm⁻¹
 $T = 150$ K
 $0.09 \times 0.02 \times 0.02$ mm

2.2. Data collection

Agilent SuperNova, Single source at offset, Eos diffractometer
Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2013)
 $T_{\min} = 0.859$, $T_{\max} = 1.000$

4781 measured reflections
2756 independent reflections
1993 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

2.3. Refinement

$R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.118$
 $S = 1.06$
2756 reflections
189 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.26$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻³

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

Cg4 is the centroid of the N2/N3/C11–C13 ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N2—H2N···O4 ⁱ	0.99 (3)	1.69 (3)	2.6846 (19)	178 (4)
N3—H3N···O4 ⁱⁱ	0.97 (2)	1.71 (2)	2.680 (2)	175 (2)
C3—H3···O4 ⁱⁱⁱ	0.95	2.45	3.321 (3)	153
C5—H5···O3 ^{iv}	0.95	2.48	3.266 (3)	141
C9—H9B···O2 ⁱ	0.99	2.41	3.397 (3)	172
C11—H11···O3 ⁱⁱ	0.95	2.40	2.987 (3)	120
C13—H13···O1 ^v	0.95	2.54	3.352 (2)	143
C2—H2···Cg4	0.95	2.87	3.805 (2)	166

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

Data collection: *CrysAlis PRO* (Agilent, 2013); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS2013* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2013* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON* (Spek, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: XU5807).

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supporting information

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Crystal structure of 1*H*-imidazol-3-ium 2-(1,3-dioxoisooindolin-2-yl)acetate

Shaaban K. Mohamed, Mehmet Akkurt, Herman Potgieter and Muizz Ali

S1. Comment

The use of co-crystals in drug design see and delivery and as functional materials with potential applications as pharmaceuticals has recently attracted a significant amount of interest in the pharmaceutical industry (Babu & Nangia, 2011). Co-crystallization in particular is a reliable technique for the modification of the physical properties of a drug as it enables the control of physical properties of Active Pharmaceutical Ingredient (API) molecules such as dissolution, stability, solubility, bioavailability, hygroscopicity and compressibility without changing the chemical composition of the API (Sekhon, 2013). Multi-API co-crystals are also possible solid forms for the delivery of combination drugs that can be tested to overcome problems related with traditional combination drugs (Frantz, 2006). Another benefit of multi-API co-crystal is the ability to reduce the number of pills being taken by a patient due to the improvement of patients long-term medication compliance in long-term drug therapy, since fewer pills need to be taken (Pan *et al.*, 2008; Vermeire *et al.*, 2001). The title compound was obtained during our study on co-crystallization reaction of *N'*-(di-1*H*-imidazol-1-ylmethylene)isonicotinohydrazide with (1,3-dioxoisooindolin-2-yl)acetic acid under aqueous condition.

Fig. 1 shows one 1*H*-imidazol-3-ium cation and one (1,3-dioxoisooindolin-2-yl)acetate anion in the asymmetric unit of the title compound (I).

The five-membered ring (N2/N3/C11—C13) of the 1*H*-imidazol-3-ium cation is essentially planar [maximum deviation = 0.003 (2) Å for C12]. The nine-membered ring system (N1/C1—C8) of the (1,3-dioxoisooindolin-2-yl)acetate anion is also essentially planar [maximum deviation = -0.023 (2) Å for C8].

In the crystal structure, the anions and cations of (I) are linked *via* N—H···O and C—H···O hydrogen bonds (Table 1, Fig. 2), forming three dimensional network. Further C—H···π interactions (Table 1) and face-to-face π-π stacking interactions [$Cg1 \cdots Cg2 (x, 1/2 - y, -1/2 + z) = 3.4728 (13)$ Å, $Cg2 \cdots Cg2 (x, 1/2 - y, 1/2 + z) = 3.7339 (13)$ Å, where $Cg1$ and $Cg2$ are the centroids of the N1/C1/C6—C8 and C1—C6 rings, respectively] presents in the three-dimensional framework.

S2. Experimental

A mixture of 1 mmol (281 mg) of *N'*-(di-1*H*-imidazol-1-ylmethylene)isonicotinohydrazide and 1 mmol (205 mg) of (1,3-dioxoisooindolin-2-yl)acetic acid was stirred in 30 ml ethanol at room temperature. Few drops of glacial acetic acid as a catalyst was added to the reaction mixture and allowed to reflux at 351 K for 5 h. The reaction progress was monitored by TLC using a mixture of cyclohexane and ethyl acetate (1:1) as an eluent. On completion, the reaction mixture was poured on crushed ice (50 g). The resulting solid was filtered off, washed with cold ethanol dried under vacuum and recrystallized from ethanol to yield colourless blocks of the title compound (74% yield).

S3. Refinement

H atoms attached to carbon were placed in calculated positions (C—H = 0.95 and 0.99 Å) and were included as riding contributions with isotropic displacement parameters 1.2 those of the attached atoms. H-atoms attached to nitrogen were

placed in locations derived from a difference map and they were refined freely.

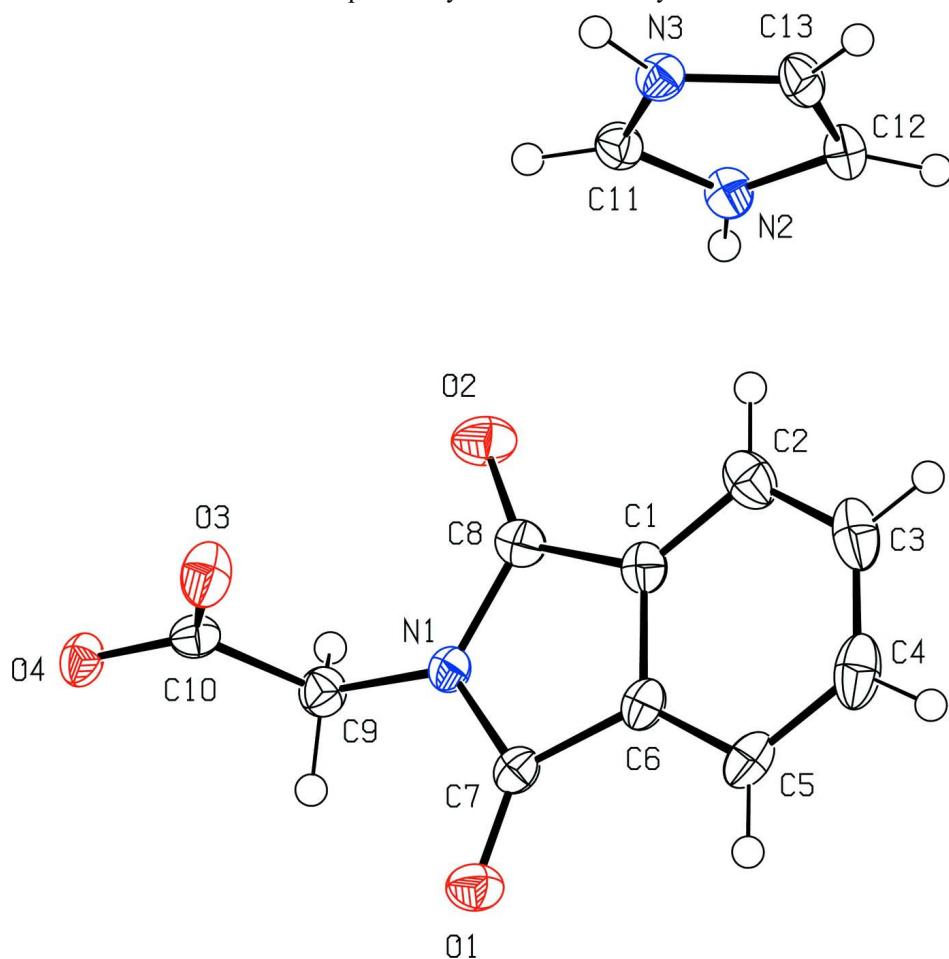


Figure 1

Perspective view of the title compound (**I**). Displacement ellipsoids are drawn at the 50% probability level.

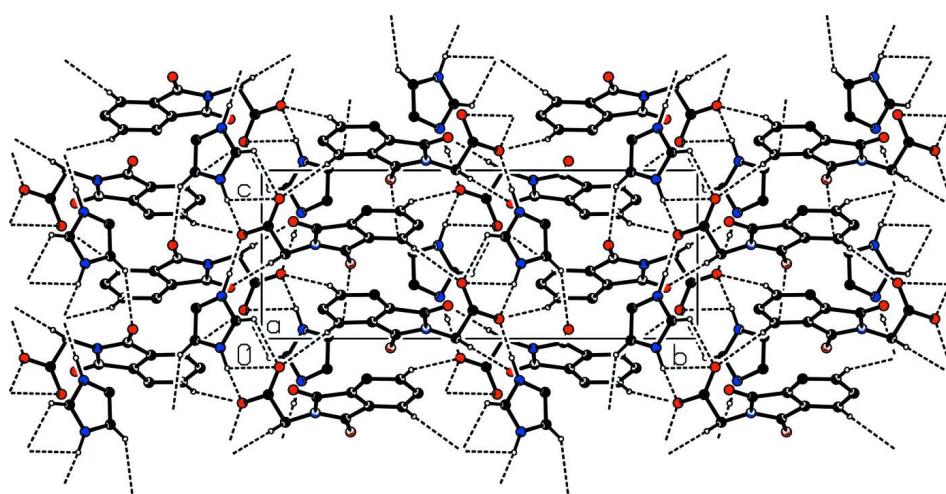


Figure 2

Packing viewed down the a axis showing the intermolecular interactions as dotted lines.

1*H*-imidazol-3-ium (1,3-dioxoisooindolin-2-yl)acetate*Crystal data*
 $M_r = 273.25$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

 $a = 9.8750 (7)$ Å

 $b = 18.0543 (15)$ Å

 $c = 7.0942 (5)$ Å

 $\beta = 100.955 (7)^\circ$
 $V = 1241.75 (16)$ Å³
 $Z = 4$
 $F(000) = 568$
 $D_x = 1.462$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.7107$ Å

Cell parameters from 1373 reflections

 $\theta = 4.0\text{--}27.4^\circ$
 $\mu = 0.11$ mm⁻¹
 $T = 150$ K

Block, colourless

0.09 × 0.02 × 0.02 mm

Data collection
Agilent SuperNova, Single source at offset, Eos
diffractometer

 $T_{\min} = 0.859$, $T_{\max} = 1.000$

4781 measured reflections

Radiation source: SuperNova (Mo) X-ray
Source

2756 independent reflections

1993 reflections with $I > 2\sigma(I)$

Mirror monochromator

 $R_{\text{int}} = 0.028$

Detector resolution: 8.0714 pixels mm⁻¹
 ω scans

 $\theta_{\max} = 29.1^\circ$, $\theta_{\min} = 3.1^\circ$

Absorption correction: multi-scan
(*CrysAlis PRO*; Agilent, 2013)

 $h = -12 \rightarrow 12$
 $k = -9 \rightarrow 23$
 $l = -9 \rightarrow 5$
Refinement
Refinement on F^2

Hydrogen site location: inferred from
neighbouring sites

Least-squares matrix: full

H atoms treated by a mixture of independent
and constrained refinement

 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $w = 1/[\sigma^2(F_o^2) + (0.0327P)^2 + 0.2765P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.118$
 $(\Delta/\sigma)_{\max} < 0.001$
 $S = 1.06$
 $\Delta\rho_{\max} = 0.26$ e Å⁻³

2756 reflections

 $\Delta\rho_{\min} = -0.26$ e Å⁻³

189 parameters

0 restraints

Special details

Geometry. Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors.

Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating R -factor-obs 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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.16235 (14)	0.29586 (8)	-0.05484 (19)	0.0277 (5)
O2	0.54966 (15)	0.42729 (9)	0.1952 (2)	0.0316 (5)
O3	0.24304 (16)	0.45939 (9)	0.33073 (19)	0.0310 (5)
O4	0.14750 (14)	0.54438 (8)	0.11691 (18)	0.0236 (5)
N1	0.33840 (16)	0.37710 (10)	0.0589 (2)	0.0202 (5)

C1	0.5087 (2)	0.29370 (12)	0.1790 (3)	0.0215 (6)
C2	0.6297 (2)	0.25882 (14)	0.2609 (3)	0.0292 (7)
C3	0.6299 (2)	0.18176 (14)	0.2585 (3)	0.0344 (8)
C4	0.5146 (3)	0.14112 (14)	0.1752 (3)	0.0343 (8)
C5	0.3917 (2)	0.17722 (12)	0.0932 (3)	0.0265 (7)
C6	0.3923 (2)	0.25360 (12)	0.0988 (3)	0.0207 (6)
C7	0.2809 (2)	0.30730 (12)	0.0234 (3)	0.0199 (6)
C8	0.4768 (2)	0.37384 (13)	0.1526 (3)	0.0221 (6)
C9	0.2655 (2)	0.44563 (12)	0.0027 (3)	0.0231 (7)
C10	0.2159 (2)	0.48474 (12)	0.1675 (3)	0.0212 (6)
N2	0.90678 (17)	0.40848 (10)	0.2486 (2)	0.0228 (6)
N3	0.91310 (17)	0.40494 (10)	0.5539 (2)	0.0221 (6)
C11	0.8627 (2)	0.44091 (13)	0.3939 (3)	0.0227 (6)
C12	0.9894 (2)	0.34962 (12)	0.3193 (3)	0.0244 (7)
C13	0.9924 (2)	0.34743 (12)	0.5102 (3)	0.0247 (7)
H2	0.70950	0.28620	0.31660	0.0350*
H3	0.71140	0.15610	0.31560	0.0410*
H4	0.51900	0.08860	0.17370	0.0410*
H5	0.31170	0.15030	0.03640	0.0320*
H9A	0.18490	0.43490	-0.09980	0.0280*
H9B	0.32720	0.47940	-0.05150	0.0280*
H2N	0.885 (3)	0.4263 (15)	0.114 (4)	0.061 (8)*
H3N	0.894 (2)	0.4211 (13)	0.677 (3)	0.040 (7)*
H11	0.80430	0.48310	0.38420	0.0270*
H12	1.03540	0.31680	0.24770	0.0290*
H13	1.04070	0.31250	0.59820	0.0300*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0257 (8)	0.0262 (9)	0.0296 (8)	-0.0052 (7)	0.0014 (6)	-0.0001 (7)
O2	0.0293 (8)	0.0287 (10)	0.0353 (9)	-0.0110 (7)	0.0026 (6)	0.0010 (7)
O3	0.0450 (10)	0.0266 (10)	0.0218 (8)	0.0109 (8)	0.0074 (7)	0.0055 (7)
O4	0.0313 (8)	0.0196 (9)	0.0195 (7)	0.0062 (7)	0.0040 (6)	0.0007 (6)
N1	0.0209 (8)	0.0159 (10)	0.0235 (9)	0.0004 (7)	0.0038 (7)	-0.0011 (7)
C1	0.0241 (10)	0.0227 (12)	0.0191 (10)	0.0015 (9)	0.0080 (8)	0.0019 (9)
C2	0.0248 (11)	0.0383 (15)	0.0259 (11)	0.0047 (11)	0.0083 (9)	0.0046 (10)
C3	0.0383 (14)	0.0365 (16)	0.0308 (13)	0.0171 (12)	0.0130 (10)	0.0083 (11)
C4	0.0523 (15)	0.0262 (14)	0.0280 (12)	0.0147 (12)	0.0168 (11)	0.0067 (11)
C5	0.0395 (13)	0.0205 (13)	0.0209 (11)	-0.0008 (10)	0.0090 (9)	-0.0030 (9)
C6	0.0270 (11)	0.0205 (12)	0.0164 (10)	0.0007 (9)	0.0091 (8)	-0.0002 (9)
C7	0.0237 (10)	0.0199 (12)	0.0174 (10)	-0.0018 (9)	0.0069 (8)	-0.0009 (9)
C8	0.0211 (10)	0.0267 (13)	0.0191 (10)	-0.0018 (9)	0.0057 (8)	0.0015 (9)
C9	0.0257 (11)	0.0199 (12)	0.0233 (11)	0.0017 (9)	0.0039 (8)	0.0016 (9)
C10	0.0217 (10)	0.0206 (12)	0.0206 (10)	-0.0034 (9)	0.0025 (8)	0.0003 (9)
N2	0.0241 (9)	0.0260 (11)	0.0190 (9)	0.0016 (8)	0.0061 (7)	0.0021 (8)
N3	0.0245 (9)	0.0233 (11)	0.0184 (9)	0.0000 (8)	0.0038 (7)	0.0003 (8)
C11	0.0223 (10)	0.0240 (12)	0.0212 (11)	0.0017 (9)	0.0025 (8)	0.0007 (9)

C12	0.0225 (10)	0.0234 (13)	0.0285 (12)	0.0052 (9)	0.0079 (8)	0.0010 (10)
C13	0.0236 (11)	0.0208 (13)	0.0289 (12)	0.0059 (9)	0.0033 (9)	0.0057 (9)

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

O1—C7	1.214 (2)	C2—C3	1.391 (4)
O2—C8	1.207 (3)	C3—C4	1.389 (3)
O3—C10	1.227 (3)	C4—C5	1.403 (3)
O4—C10	1.285 (3)	C5—C6	1.380 (3)
N1—C7	1.386 (3)	C6—C7	1.488 (3)
N1—C8	1.403 (3)	C9—C10	1.524 (3)
N1—C9	1.449 (3)	C2—H2	0.9500
N2—C11	1.329 (3)	C3—H3	0.9500
N2—C12	1.375 (3)	C4—H4	0.9500
N3—C11	1.320 (3)	C5—H5	0.9500
N3—C13	1.371 (3)	C9—H9A	0.9900
N2—H2N	0.99 (3)	C9—H9B	0.9900
N3—H3N	0.97 (2)	C12—C13	1.350 (3)
C1—C8	1.485 (3)	C11—H11	0.9500
C1—C2	1.378 (3)	C12—H12	0.9500
C1—C6	1.386 (3)	C13—H13	0.9500
C7—N1—C8	112.13 (17)	O3—C10—O4	125.81 (19)
C7—N1—C9	124.17 (16)	O3—C10—C9	120.46 (19)
C8—N1—C9	123.69 (18)	O4—C10—C9	113.72 (17)
C11—N2—C12	108.47 (16)	C3—C2—H2	121.00
C11—N3—C13	108.44 (16)	C1—C2—H2	121.00
C12—N2—H2N	127.4 (16)	C2—C3—H3	119.00
C11—N2—H2N	124.1 (16)	C4—C3—H3	119.00
C13—N3—H3N	130.3 (13)	C3—C4—H4	120.00
C11—N3—H3N	121.2 (13)	C5—C4—H4	120.00
C2—C1—C8	130.2 (2)	C4—C5—H5	122.00
C6—C1—C8	108.52 (18)	C6—C5—H5	122.00
C2—C1—C6	121.3 (2)	C10—C9—H9A	109.00
C1—C2—C3	117.1 (2)	N1—C9—H9B	109.00
C2—C3—C4	122.1 (2)	H9A—C9—H9B	108.00
C3—C4—C5	120.4 (2)	N1—C9—H9A	109.00
C4—C5—C6	117.0 (2)	C10—C9—H9B	109.00
C1—C6—C5	122.19 (19)	N2—C11—N3	108.95 (19)
C1—C6—C7	107.83 (18)	N2—C12—C13	106.68 (18)
C5—C6—C7	129.97 (19)	N3—C13—C12	107.47 (18)
N1—C7—C6	106.14 (17)	N2—C11—H11	126.00
O1—C7—N1	124.34 (19)	N3—C11—H11	126.00
O1—C7—C6	129.5 (2)	N2—C12—H12	127.00
N1—C8—C1	105.36 (18)	C13—C12—H12	127.00
O2—C8—C1	130.18 (19)	N3—C13—H13	126.00
O2—C8—N1	124.5 (2)	C12—C13—H13	126.00
N1—C9—C10	113.57 (17)		

C9—N1—C7—C6	178.15 (17)	C2—C1—C6—C5	-1.2 (3)
C7—N1—C8—O2	178.4 (2)	C6—C1—C2—C3	0.3 (3)
C9—N1—C8—O2	-0.4 (3)	C8—C1—C6—C7	-1.6 (2)
C7—N1—C8—C1	-0.3 (2)	C8—C1—C6—C5	177.06 (19)
C8—N1—C7—O1	179.48 (19)	C6—C1—C8—O2	-177.4 (2)
C9—N1—C7—O1	-1.7 (3)	C6—C1—C8—N1	1.2 (2)
C8—N1—C7—C6	-0.7 (2)	C1—C2—C3—C4	0.9 (3)
C8—N1—C9—C10	-79.5 (2)	C2—C3—C4—C5	-1.3 (3)
C9—N1—C8—C1	-179.10 (17)	C3—C4—C5—C6	0.4 (3)
C7—N1—C9—C10	101.9 (2)	C4—C5—C6—C7	179.2 (2)
C12—N2—C11—N3	-0.4 (2)	C4—C5—C6—C1	0.8 (3)
C11—N2—C12—C13	0.5 (2)	C1—C6—C7—O1	-178.8 (2)
C13—N3—C11—N2	0.0 (2)	C5—C6—C7—N1	-177.1 (2)
C11—N3—C13—C12	0.3 (2)	C1—C6—C7—N1	1.4 (2)
C2—C1—C8—O2	0.7 (4)	C5—C6—C7—O1	2.7 (4)
C2—C1—C8—N1	179.3 (2)	N1—C9—C10—O4	-177.97 (17)
C2—C1—C6—C7	-179.89 (19)	N1—C9—C10—O3	2.7 (3)
C8—C1—C2—C3	-177.6 (2)	N2—C12—C13—N3	-0.5 (2)

Hydrogen-bond geometry (Å, °)

Cg4 is the centroid of the N2/N3/C11–C13 ring.

D—H···A	D—H	H···A	D···A	D—H···A
N2—H2N···O4 ⁱ	0.99 (3)	1.69 (3)	2.6846 (19)	178 (4)
N3—H3N···O3 ⁱⁱ	0.97 (2)	2.54 (2)	3.087 (2)	115.4 (16)
N3—H3N···O4 ⁱⁱ	0.97 (2)	1.71 (2)	2.680 (2)	175 (2)
C3—H3···O4 ⁱⁱⁱ	0.95	2.45	3.321 (3)	153
C5—H5···O3 ^{iv}	0.95	2.48	3.266 (3)	141
C9—H9A···O1	0.99	2.55	2.891 (3)	100
C9—H9B···O2 ⁱ	0.99	2.41	3.397 (3)	172
C11—H11···O3 ⁱⁱ	0.95	2.40	2.987 (3)	120
C13—H13···O1 ^v	0.95	2.54	3.352 (2)	143
C2—H2···Cg4	0.95	2.87	3.805 (2)	166

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