

2-(1,3-Dioxoisoindolin-2-yl)propanoic acid

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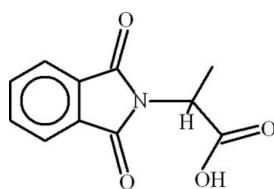
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.030; wR factor = 0.082; data-to-parameter ratio = 9.0.

The crystal structure of the title compound, $\text{C}_{11}\text{H}_9\text{NO}_4$, consists of infinite one-dimensional polymeric chains due to intermolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds between the carboxylate and carbonyl groups. The phthalimide ring system and the $\text{C}-\text{COO}$ group are planar, with r.m.s. deviations of 0.0253 and 0.0067 \AA , respectively, from their mean square planes and the dihedral angle between them is $66.41(7)^\circ$. The molecules are stabilized by $\text{C}=\text{O}\cdots\pi$ interactions and weak intramolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For the medicinal properties of isocoumarin, see: Matsuda *et al.* (1999). For related crystal structures, see: Li & Liang (2006); Raza *et al.* (2009); Wheeler *et al.* (2004). For the graph-set analysis of hydrogen-bond patterns in crystal structures, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{11}\text{H}_9\text{NO}_4$
 $M_r = 219.19$
Monoclinic, $P2_1$
 $a = 9.3056(8)\text{ \AA}$
 $b = 5.9768(4)\text{ \AA}$
 $c = 9.7583(8)\text{ \AA}$
 $\beta = 110.988(3)^\circ$

$V = 506.73(7)\text{ \AA}^3$
 $Z = 2$
Mo $K\alpha$ radiation
 $\mu = 0.11\text{ mm}^{-1}$
 $T = 296\text{ K}$
 $0.30 \times 0.25 \times 0.23\text{ mm}$

Data collection

Bruker Kappa APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
 $T_{\min} = 0.968$, $T_{\max} = 0.974$

5705 measured reflections
1381 independent reflections
1302 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.082$
 $S = 1.06$
1381 reflections
153 parameters
1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.16\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.13\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O3—H3A \cdots O1 ⁱ	0.80 (4)	1.96 (4)	2.750 (2)	172 (4)
C9—H9 \cdots O2	0.96 (3)	2.48 (2)	2.899 (2)	106.6 (17)
C8—O2 \cdots Cg1 ⁱⁱ	1.20 (1)	3.09 (1)	4.0543 (17)	138 (1)

Symmetry codes: (i) $-x + 1, y - \frac{1}{2}, -z + 1$; (ii) $-x, y - \frac{1}{2}, -z$. Cg1 is the centroid of 5-membered heterocyclic ring.

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); 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, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SI2209).

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supporting information

Acta Cryst. (2009). E65, o2678 [https://doi.org/10.1107/S1600536809040434]

2-(1,3-Dioxoisooindolin-2-yl)propanoic acid

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S1. Comment

Isocoumarin are important components among natural products that exhibit a broad range of biological activities including anti-microbial, anti-allergic and immunomodulatory (Matsuda *et al.*, 1999). Isocoumarins are also useful intermediates in synthesis of many important compounds *e.g.*, isoquinoline alkaloids. The titled compound (I, Fig. 1), is an intermediate towards the synthesis of isocoumarins. The title compound has also been prepared for complexation with various metals.

We have recently reported the crystal structure of (II) (*2R*)-2-(1,3-Dioxoisooindolin-2-yl)-4-(methylsulfanyl)butanoic acid (Raza *et al.*, 2009) which contain the same isoindoline. The crystal structures of (III) DL-2-(1,3-Dioxoisooindolin-2-yl)propanoic acid (Wheeler *et al.*, 2004), (IV) (*S*)-2-(1,3-Dioxoisooindolin-2-yl)propanoic acid (Li & Liang, 2006) have also been reported which are the racemate of (I).

In (I) the phthalimide ring system A (C1—C8/N1/O1/O2) and the group B (C9/C10/O3/O4) are planar with r.m.s. deviations of 0.0253 and 0.0067 Å respectively, from their mean square planes. The dihedral angle between A/B is 66.41 (7)°, whereas it is 86.7 (3)° as observed in (IV). The title compound is stabilized in the form of infinite one dimensional polymeric chains due to intermolecular H-bondings (Table 1, Fig. 2). There exist a weak intramolecular H-bondings forming S(5) ring motifs (Fig. 1) (Bernstein *et al.*, 1995). The C=O···Cg1 [Cg1 is the centroid of five membered ring (C1/C2/C7/C8/N1)] interaction (Table 1), may also be responsible for stabilizing of the molecules.

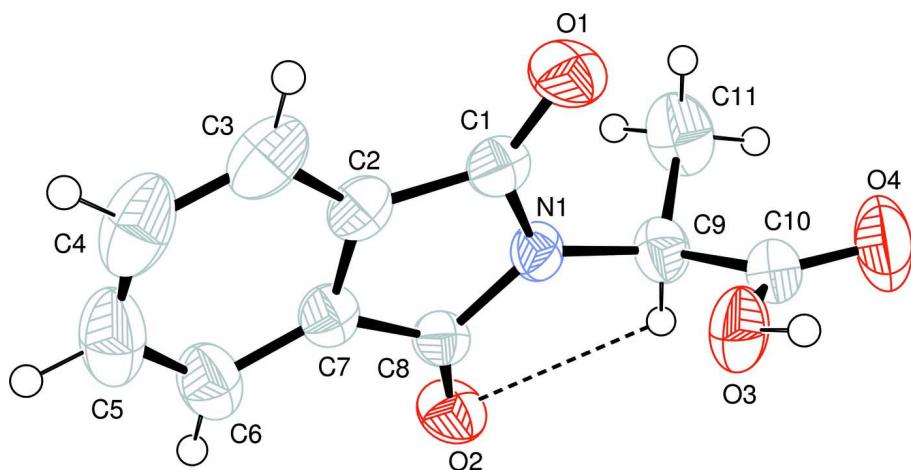
S2. Experimental

The (*S*)-alanine (1.96 g, 22 mmol) and phthalic anhydride (3.6 g, 24.3 mmol) were added to a flask with constant stirring. The temperature of oil bath was kept at 433 K. Three hours later the flask was removed from oil bath, brought to room temperature and the crystals of phthalic anhydride on the walls of the flask were removed manually. The solid crude product was purified by crystallization from ethanol:water (8:2) that yielded (70%) colorless prisms of the title compound (I).

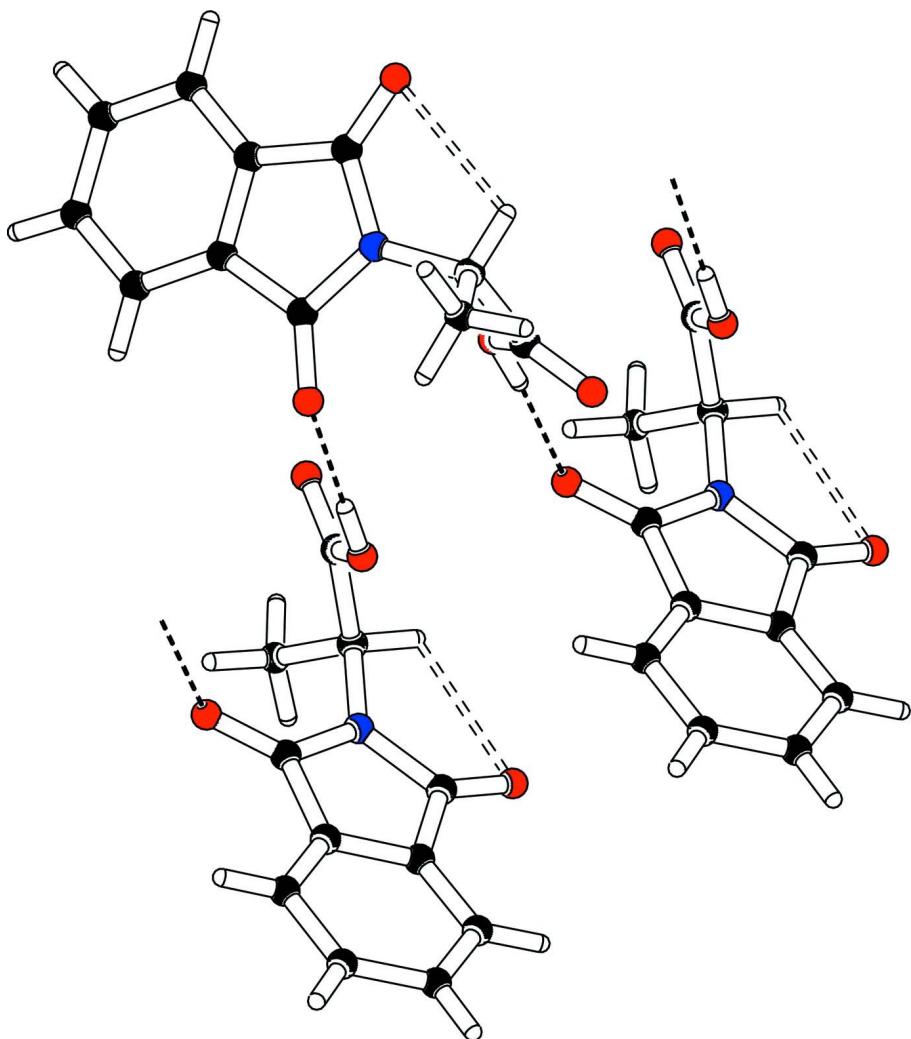
S3. Refinement

In the absence of significant anomalous dispersion effects, Friedel pairs were averaged using MERG 3.

The coordinates of H3A and H9 were refined. The H-atoms were positioned geometrically with C—H = 0.93 and 0.96 Å for aromatic and methyl H atoms respectively and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C}, \text{O})$, where $x = 1.5$ for methyl and 1.2 for all other H atoms.

**Figure 1**

View of (I) with the atom numbering scheme. The thermal ellipsoids are drawn at the 50% probability level. H-atoms are shown by small circles of arbitrary radii. The dotted line represent the intramolecular H-bonding.

**Figure 2**

The partial packing (*PLATON*; Spek, 2009) which shows that molecules form infinite one dimensional polymeric chains.

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$C_{11}H_9NO_4$
 $M_r = 219.19$
Monoclinic, $P2_1$
Hall symbol: P 2yb
 $a = 9.3056 (8)$ Å
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 $c = 9.7583 (8)$ Å
 $\beta = 110.988 (3)^\circ$
 $V = 506.73 (7)$ Å³
 $Z = 2$

$F(000) = 228$
 $D_x = 1.437$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2319 reflections
 $\theta = 2.2\text{--}28.3^\circ$
 $\mu = 0.11$ mm⁻¹
 $T = 296$ K
Prism, colorless
 $0.30 \times 0.25 \times 0.23$ mm

Data collection

Bruker Kappa APEXII CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 7.40 pixels mm⁻¹
 ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2005)
 $T_{\min} = 0.968$, $T_{\max} = 0.974$

5705 measured reflections
1381 independent reflections
1302 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 2.2^\circ$
 $h = -12 \rightarrow 12$
 $k = -7 \rightarrow 7$
 $l = -12 \rightarrow 13$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.030$
 $wR(F^2) = 0.082$
 $S = 1.06$
1381 reflections
153 parameters
1 restraint
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
 $w = 1/[\sigma^2(F_o^2) + (0.0443P)^2 + 0.0584P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.16 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.13 \text{ e } \text{\AA}^{-3}$
Extinction correction: SHELXL97 (Sheldrick,
2008)
Extinction coefficient: 0.132 (13)

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 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.34281 (15)	1.0490 (3)	0.31137 (15)	0.0551 (4)
O2	-0.00322 (14)	0.5031 (3)	0.09136 (15)	0.0520 (4)
O3	0.41497 (15)	0.5761 (3)	0.42520 (15)	0.0550 (5)
O4	0.57532 (16)	0.5164 (5)	0.30695 (19)	0.0817 (7)
N1	0.19830 (14)	0.7462 (3)	0.19714 (14)	0.0358 (4)
C1	0.22202 (19)	0.9473 (3)	0.27082 (17)	0.0382 (4)
C2	0.07493 (19)	1.0033 (3)	0.28948 (17)	0.0402 (5)
C3	0.0385 (3)	1.1806 (4)	0.3606 (2)	0.0562 (6)
C4	-0.1092 (3)	1.1832 (5)	0.3656 (2)	0.0690 (8)
C5	-0.2125 (3)	1.0146 (6)	0.3031 (2)	0.0684 (9)
C6	-0.1755 (2)	0.8353 (5)	0.2313 (2)	0.0539 (6)
C7	-0.02955 (19)	0.8354 (3)	0.22485 (16)	0.0392 (4)
C8	0.04667 (17)	0.6700 (3)	0.16030 (16)	0.0361 (4)
C9	0.31769 (19)	0.6226 (4)	0.1658 (2)	0.0432 (5)
C10	0.45173 (19)	0.5679 (4)	0.3068 (2)	0.0470 (5)

C11	0.3710 (3)	0.7438 (6)	0.0556 (2)	0.0642 (8)
H3	0.10903	1.29359	0.40332	0.0674*
H3A	0.490 (4)	0.564 (6)	0.497 (4)	0.0660*
H4	-0.13859	1.30102	0.41199	0.0828*
H5	-0.30990	1.02072	0.30905	0.0820*
H6	-0.24537	0.72111	0.18964	0.0647*
H9	0.273 (2)	0.482 (5)	0.125 (2)	0.0518*
H11A	0.28411	0.77266	-0.03225	0.0962*
H11B	0.44407	0.65239	0.03244	0.0962*
H11C	0.41850	0.88289	0.09692	0.0962*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0464 (7)	0.0507 (8)	0.0545 (7)	-0.0167 (7)	0.0015 (5)	-0.0070 (7)
O2	0.0474 (7)	0.0484 (8)	0.0563 (7)	-0.0153 (6)	0.0139 (6)	-0.0153 (7)
O3	0.0409 (6)	0.0693 (10)	0.0459 (7)	0.0152 (7)	0.0046 (5)	0.0073 (7)
O4	0.0412 (7)	0.1193 (17)	0.0869 (11)	0.0252 (10)	0.0257 (8)	0.0306 (13)
N1	0.0288 (6)	0.0363 (7)	0.0363 (6)	0.0000 (5)	0.0042 (5)	-0.0024 (6)
C1	0.0385 (8)	0.0368 (8)	0.0310 (7)	-0.0025 (7)	0.0024 (6)	-0.0004 (6)
C2	0.0470 (9)	0.0382 (9)	0.0311 (7)	0.0044 (7)	0.0088 (6)	0.0021 (7)
C3	0.0773 (13)	0.0452 (11)	0.0407 (8)	0.0125 (10)	0.0147 (9)	-0.0034 (8)
C4	0.0893 (16)	0.0758 (17)	0.0461 (10)	0.0356 (15)	0.0293 (10)	0.0019 (11)
C5	0.0612 (12)	0.098 (2)	0.0532 (11)	0.0276 (14)	0.0292 (10)	0.0102 (13)
C6	0.0444 (9)	0.0727 (14)	0.0478 (9)	0.0048 (10)	0.0203 (8)	0.0056 (10)
C7	0.0382 (7)	0.0451 (9)	0.0314 (7)	0.0023 (7)	0.0089 (6)	0.0036 (7)
C8	0.0327 (7)	0.0384 (8)	0.0325 (7)	-0.0032 (7)	0.0060 (5)	0.0005 (6)
C9	0.0346 (8)	0.0473 (10)	0.0444 (8)	0.0009 (8)	0.0102 (6)	-0.0067 (8)
C10	0.0338 (7)	0.0475 (10)	0.0566 (10)	0.0031 (8)	0.0126 (7)	0.0069 (9)
C11	0.0556 (11)	0.0928 (18)	0.0468 (10)	0.0056 (13)	0.0217 (8)	0.0027 (12)

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

O1—C1	1.213 (2)	C5—C6	1.390 (4)
O2—C8	1.200 (2)	C6—C7	1.382 (3)
O3—C10	1.318 (2)	C7—C8	1.482 (2)
O4—C10	1.190 (3)	C9—C10	1.525 (3)
O3—H3A	0.80 (4)	C9—C11	1.520 (3)
N1—C8	1.402 (2)	C3—H3	0.9300
N1—C9	1.455 (3)	C4—H4	0.9300
N1—C1	1.377 (2)	C5—H5	0.9300
C1—C2	1.482 (3)	C6—H6	0.9300
C2—C3	1.374 (3)	C9—H9	0.96 (3)
C2—C7	1.382 (2)	C11—H11A	0.9600
C3—C4	1.393 (4)	C11—H11B	0.9600
C4—C5	1.375 (4)	C11—H11C	0.9600
O1···O3		C5···C11 ^{viii}	
		3.023 (2)	
		3.551 (3)	

O1···C10	3.055 (3)	C6···O4 ^{ix}	3.283 (3)
O1···C10 ⁱ	3.268 (3)	C7···O2 ^{viii}	3.361 (2)
O1···C11	3.177 (3)	C8···O2 ^{viii}	3.074 (2)
O1···O3 ⁱⁱ	2.750 (2)	C8···C3 ^{iv}	3.534 (3)
O2···C7 ⁱⁱⁱ	3.361 (2)	C10···O1	3.055 (3)
O2···N1 ⁱⁱⁱ	3.149 (2)	C10···O1 ^{iv}	3.268 (3)
O2···C3 ^{iv}	3.168 (3)	C11···C5 ⁱⁱⁱ	3.551 (3)
O2···C8 ⁱⁱⁱ	3.074 (2)	C11···O1	3.177 (3)
O2···C1 ⁱⁱⁱ	3.405 (2)	C1···H3A ⁱⁱ	2.91 (4)
O3···N1	2.615 (2)	C1···H11C	2.9300
O3···O1	3.023 (2)	C2···H4 ^x	3.0100
O3···C1	2.912 (2)	C3···H4 ^x	3.0800
O3···O1 ^v	2.750 (2)	C5···H3 ^x	2.9800
O4···C6 ^{vi}	3.283 (3)	C5···H11A ^{viii}	2.9200
O1···H3A ⁱⁱ	1.96 (4)	H3···C5 ^{xi}	2.9800
O1···H11C	2.6300	H3A···O1 ^v	1.96 (4)
O2···H11A ⁱⁱⁱ	2.8300	H3A···C1 ^v	2.91 (4)
O2···H9	2.48 (2)	H4···O4 ^{xii}	2.8000
O4···H4 ^{vii}	2.8000	H4···C2 ^{xi}	3.0100
O4···H6 ^{vi}	2.6400	H4···C3 ^{xi}	3.0800
O4···H11B	2.6400	H6···O4 ^{ix}	2.6400
N1···O3	2.615 (2)	H9···O2	2.48 (2)
N1···O2 ^{viii}	3.149 (2)	H11A···O2 ^{viii}	2.8300
C1···O3	2.912 (2)	H11A···C5 ⁱⁱⁱ	2.9200
C1···O2 ^{viii}	3.405 (2)	H11B···O4	2.6400
C3···C8 ⁱ	3.534 (3)	H11C···O1	2.6300
C3···O2 ⁱ	3.168 (3)	H11C···C1	2.9300
C10—O3—H3A	110 (3)	N1—C9—C10	111.01 (15)
C1—N1—C9	124.14 (15)	O3—C10—C9	113.43 (16)
C8—N1—C9	123.75 (16)	O4—C10—C9	122.29 (18)
C1—N1—C8	112.08 (15)	O3—C10—O4	124.24 (19)
O1—C1—C2	129.45 (17)	C2—C3—H3	122.00
N1—C1—C2	106.19 (15)	C4—C3—H3	122.00
O1—C1—N1	124.35 (17)	C3—C4—H4	119.00
C1—C2—C3	129.93 (19)	C5—C4—H4	119.00
C3—C2—C7	122.00 (19)	C4—C5—H5	119.00
C1—C2—C7	108.03 (15)	C6—C5—H5	119.00
C2—C3—C4	116.8 (2)	C5—C6—H6	122.00
C3—C4—C5	121.2 (3)	C7—C6—H6	122.00
C4—C5—C6	121.9 (3)	N1—C9—H9	106.4 (13)
C5—C6—C7	116.6 (2)	C10—C9—H9	106.2 (13)
C2—C7—C6	121.44 (18)	C11—C9—H9	109.1 (12)
C2—C7—C8	108.22 (16)	C9—C11—H11A	109.00
C6—C7—C8	130.30 (18)	C9—C11—H11B	109.00
O2—C8—N1	124.46 (16)	C9—C11—H11C	109.00
O2—C8—C7	130.15 (16)	H11A—C11—H11B	109.00
N1—C8—C7	105.39 (14)	H11A—C11—H11C	109.00

N1—C9—C11	112.0 (2)	H11B—C11—H11C	109.00
C10—C9—C11	111.83 (17)		
C8—N1—C1—O1	-178.37 (16)	C1—C2—C7—C6	176.60 (16)
C8—N1—C1—C2	2.54 (18)	C1—C2—C7—C8	-1.21 (18)
C9—N1—C1—O1	3.8 (3)	C3—C2—C7—C6	-1.4 (3)
C9—N1—C1—C2	-175.31 (15)	C3—C2—C7—C8	-179.17 (16)
C1—N1—C8—O2	177.22 (16)	C2—C3—C4—C5	0.5 (3)
C1—N1—C8—C7	-3.24 (18)	C3—C4—C5—C6	-0.5 (4)
C9—N1—C8—O2	-4.9 (3)	C4—C5—C6—C7	-0.4 (3)
C9—N1—C8—C7	174.62 (15)	C5—C6—C7—C2	1.3 (3)
C1—N1—C9—C10	57.8 (2)	C5—C6—C7—C8	178.61 (18)
C1—N1—C9—C11	-67.9 (2)	C2—C7—C8—O2	-177.83 (18)
C8—N1—C9—C10	-119.78 (18)	C2—C7—C8—N1	2.66 (17)
C8—N1—C9—C11	114.5 (2)	C6—C7—C8—O2	4.6 (3)
O1—C1—C2—C3	-2.0 (3)	C6—C7—C8—N1	-174.88 (18)
O1—C1—C2—C7	-179.76 (18)	N1—C9—C10—O3	20.4 (3)
N1—C1—C2—C3	177.02 (18)	N1—C9—C10—O4	-161.9 (3)
N1—C1—C2—C7	-0.73 (18)	C11—C9—C10—O3	146.2 (2)
C1—C2—C3—C4	-177.09 (18)	C11—C9—C10—O4	-36.0 (4)
C7—C2—C3—C4	0.4 (3)		

Symmetry codes: (i) $x, y+1, z$; (ii) $-x+1, y+1/2, -z+1$; (iii) $-x, y-1/2, -z$; (iv) $x, y-1, z$; (v) $-x+1, y-1/2, -z+1$; (vi) $x+1, y, z$; (vii) $x+1, y-1, z$; (viii) $-x, y+1/2, -z$; (ix) $x-1, y, z$; (x) $-x, y-1/2, -z+1$; (xi) $-x, y+1/2, -z+1$; (xii) $x-1, y+1, z$.

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

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
O3—H3A \cdots O1 ^v	0.80 (4)	1.96 (4)	2.750 (2)	172 (4)
C9—H9 \cdots O2	0.96 (3)	2.48 (2)	2.899 (2)	106.6 (17)
C8—O2 \cdots Cg1 ⁱⁱⁱ	1.20 (1)	3.09 (1)	4.0543 (17)	138 (1)

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