Acta Crystallographica Section E
Structure Reports Online
ISSN 1600-5368

Methyl 2-(5-fluoro-1*H*-indol-3-yl)-2-oxoacetate

Shuping Wang, Huajiang Dong, Hong Chen, Kunpeng Zhu and Tieliang Zhu*

The Affiliated Hospital of the Medical College of the Chinese People's Armed Police Forces, Tianjin 300162, People's Republic of China
Correspondence e-mail: shupingwang203@yahoo.com.cn

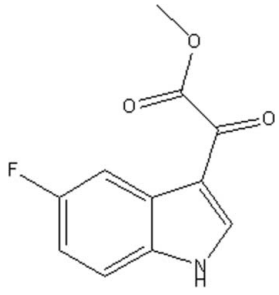
Received 2 December 2009; accepted 9 December 2009

Key indicators: single-crystal X-ray study; T = 113 K; mean σ(C–C) = 0.002 Å; R factor = 0.037; wR factor = 0.097; data-to-parameter ratio = 15.5.

The indolyl portion of the title molecule, $C_{11}H_8FNO_3$, is flat, the five- and six-membered rings making a dihedral angle of 0.815 (6)°. Intermolecular N—H···O hydrogen bonds link adjacent molecules into a linear chain. Slipped π–π stacking interactions between two neighboring indole groups further consolidate the molecules into a three-dimensional supramolecular architecture [centroid–centroid distances = 3.555 (10) and 3.569 (10) Å].

Related literature

For the biological activity of the title compound and its derivatives, see: Kozikowski *et al.* (2006); Albert *et al.* (2002); Jaquith *et al.* (2005). For the preparation, see: Alawar *et al.* (2004).



Experimental

Crystal data

$C_{11}H_8FNO_3$
M_r = 221.18
Monoclinic, $P2_1/c$
a = 7.0584 (14) Å
b = 20.586 (4) Å
c = 7.3286 (15) Å
β = 112.01 (3)°
V = 987.3 (3) Å^3
Z = 4
Mo $K\alpha$ radiation
μ = 0.12 mm^{-1}
T = 113 K
0.30 × 0.24 × 0.20 mm

Data collection

Rigaku Saturn CCD area-detector diffractometer
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2004)
T_{min} = 0.964, T_{max} = 0.976
9472 measured reflections
2328 independent reflections
2097 reflections with $I > 2\sigma(I)$
R_{int} = 0.024

Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.037
$wR(F^2)$ = 0.097
S = 1.06
2328 reflections
150 parameters
H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{max}$ = 0.38 e Å^{-3}
$\Delta\rho_{min}$ = −0.20 e Å^{-3}

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

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O3^{i}	0.932 (17)	2.420 (16)	3.1598 (13)	136.3 (13)
N1—H1···O1^{i}	0.932 (17)	1.932 (17)	2.7861 (13)	151.3 (14)

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

Data collection: *CrystalClear* (Rigaku/MSC, 2004); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXL97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg, 1999); software used to prepare material for publication: *publCIF* (Westrip, 2009).

This work was supported by the National Natural Science Foundation of China (No. 30873363), the Great Program of the Science Foundation of Tianjin (09ZCKFNC01200) and the Program of the Science Foundation of Tianjin (08JCYBJC070000).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: NG2701).

References

Alawar, R. S., Ray, J. E., Hecker, K. A., Joseph, S., Huang, J. P., Shih, C., Brooks, H. B., Spencer, C. D., Watkins, S. A., Schultz, R. M., Considine, E. L., Faul, M. M., Sullivan, K. A., Kolis, S. P., Carr, M. A. & Zhang, F. M. (2004). *Bioorg. Med. Chem. Lett.* **14**, 3925–3928.
Albert, R., Cooke, N. G., Cottens, S., Ehrhardt, C., Evenou, J. P., Sedrani, R., Von, M. P., Wagner, J. & Zenke, G. (2002). PCT Int. Appl. 60.
Brandenburg, K. (1999). *DIAMOND*. Crystal Impact GbR, Bonn, Germany.
Jaquith, J. B., Gillard, J. W. & Laurent, A. (2005). PCT Int. Appl. 50.
Kozikowski, A. P., Gaisina, I. N., Petukhov, P. A., Sridhar, J., King, L. S. T., Blond, S. Y., Duka, T., Rusnak, M. & Sidhu, A. (2006). *ChemMedChem*, **1**, 256–266.
Rigaku/MSC (2004). *CrystalClear*. Rigaku/MSC Inc., The Woodlands, Texas, USA.
Sheldrick, G. M. (2008). *Acta Cryst.* A**64**, 112–122.
Westrip, S. P. (2009). *publCIF*. In preparation.

supporting information

Acta Cryst. (2010). E**66**, o155 [doi:10.1107/S1600536809053033]

Methyl 2-(5-fluoro-1*H*-indol-3-yl)-2-oxoacetate

Shuping Wang, Huajiang Dong, Hong Chen, Kunpeng Zhu and Tieliang Zhu

S1. Comment

The synthesis of the title compound (I) and its derivatives has received considerable attention on account of their biological activity especially for both enzymic and cell-based study (Kozikowski *et al.*, 2006). They also can be used as neuroprotective (Albert *et al.*, 2002) and anti-proliferative agents (Jaquith *et al.*, 2005).

In this work, the title molecule, $C_{11}H_8FNO_3$, (Fig. 1), has been synthesized. In an asymmetric unit of (I) one molecule can be observed. (I) crystalizes in the Monoclinic P2(1)/n space group, a = 7.0584 (14) Å, b = 20.586 (4) Å, c = 7.3286 (15) Å, β = 112.01 (3)°. The heterocyclic rings (N1, C1, C6, C7, C8) make a small dihedral angle of 0.815 (6)° with benzene rings (C1, C2, C3, C4, C5, C6,). N atoms in the molecule act as hydrogen-bond donors to O atoms in the neighbouring molecule forming intermolecular N1—H1···O1 (symmetry code: *x*, *y*, *z* - 1) and N1—H1···O3 (symmetry code: *x*, *y*, *z* - 1) hydrogen bonds. These N—H···O hydrogen bonds, C3—H3···F1(symmetry code: *x*, -*y* + 1/2, *z* - 1/2), and intra-molecular C5—H5···O1, C8—H8···O2 hydrogen bonds stabilize the crystal structure and extend molecules (I) into a double-tape structure along the c direction. π–π interactions between the indole rings are also present, the centroid-centroid distances [symmetry code: 1 - *x*,-*y*,2 - *z*; 2 - *x*,-*y*,2 - *z*] are 3.555 (10) and 3.569 (10) Å. N—H···π interactions [symmetry code: 1 - *x*,-*y*,2 - *z*] are 3.218 (21) Å. These Parallel slipped π-π stacking interactions between two neighboring indole groups and N—H···π interactions also further consolidate (I) into the three-dimensional supramolecular architecture.

Perspective drawing with the atomic numbering scheme is illustrated in Figure 1. Selected geometric parameters (Å, °) for (I) are listed in table 1. Selected hydrogen-bonding geometric parameters (Å, °) for (I) are listed in table 2. The corresponding N—H···O, C—H···O and C—H···F hydrogen bonds are shown in Figure 2. The π-π stacking interactions and N—H···π interactions are shown in Ffigure 3. The three-dimensional supramolecular packing architecture of (I) is shown in Figure 4.

S2. Experimental

Methyl 2-(5-fluoro-1*H*-indol-3-yl)-2-oxoacetate was prepared according to the previous literature (Alawar *et al.*, 2004). Colorless block Single crystals were obtained by slow evaporation from a methanol and water mixed solution (150 ml) of (I) at room temperature.

S3. Refinement

H atoms on N atoms were located in a difference Fourier map and refined isotropically, with restrains of N—H = 0.90±0.01 Å. Other H atoms were positioned geometrically with C—H = 0.93 and 0.96 Å, for indole H atoms, respectively, and refined using a riding model, with U_{iso}(H) =1.2Ueq(C)(1.5 for methyl groups).

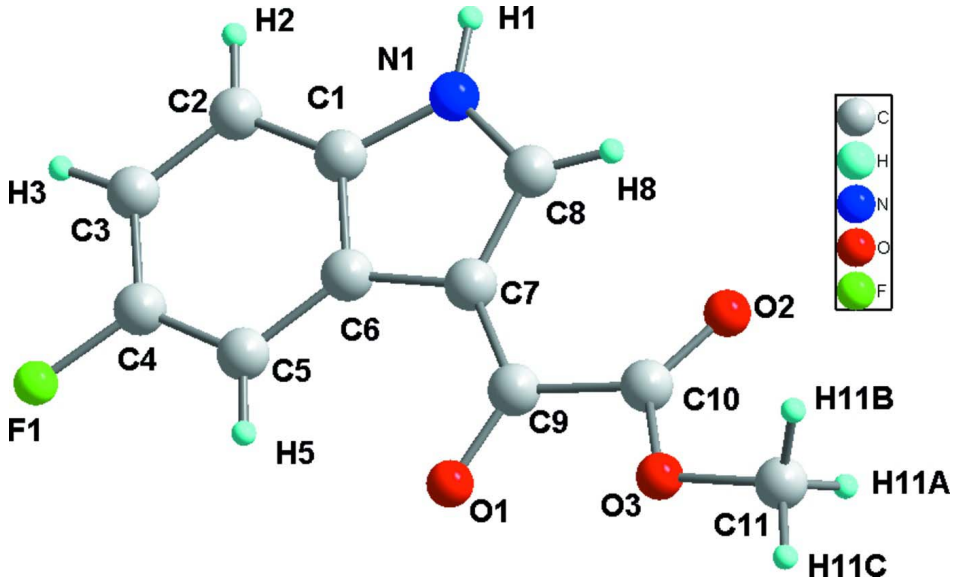


Figure 1

The molecular structure and atom-labeling scheme of (I).

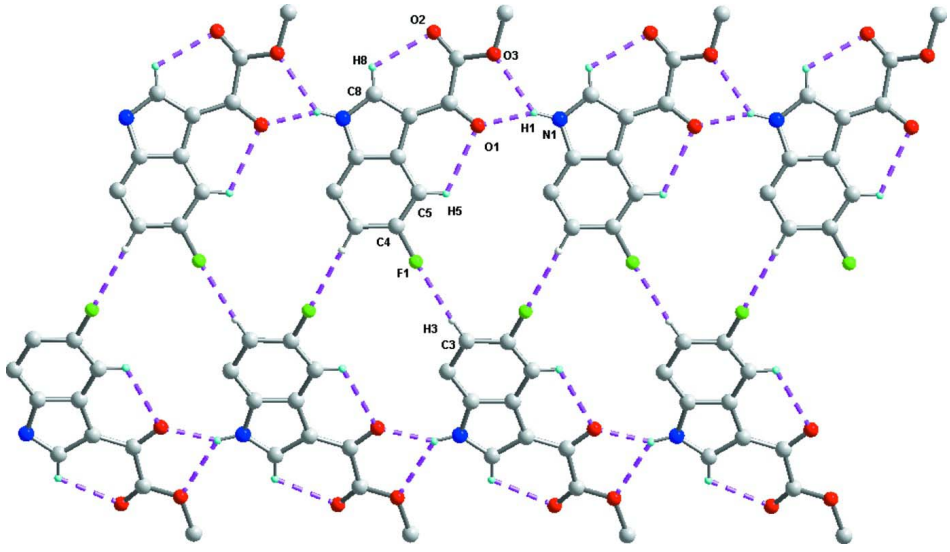


Figure 2

The corresponding N—H···O, C—H···O and C—H···F hydrogen bonds stabling the packing structure of (I). Hydrogen bonds are shown as dashed lines.

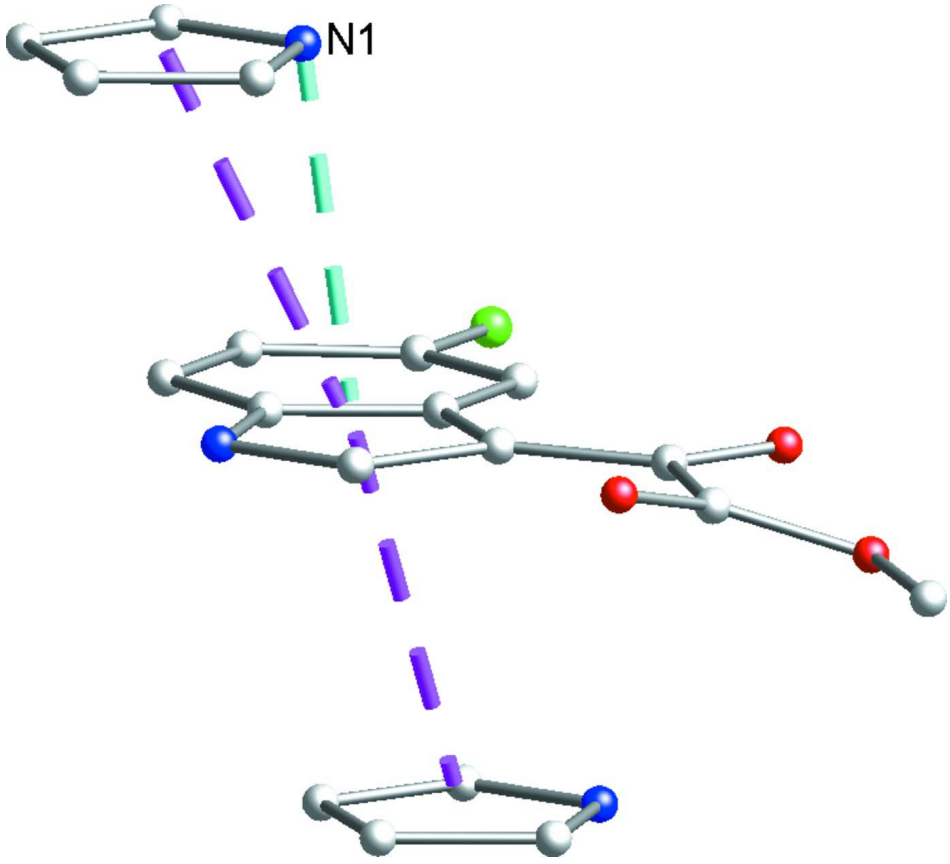


Figure 3

The π-π stacking interactions and N—H···π interactions in (I). Hydrogen bonds are shown as dashed lines.

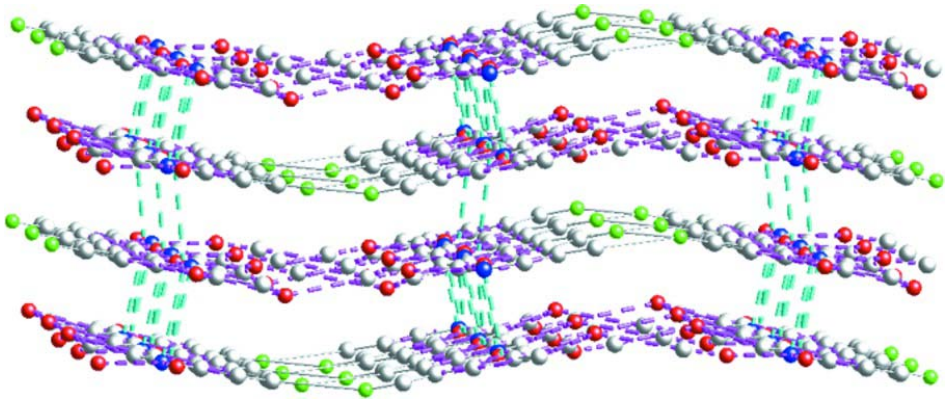


Figure 4

The three-dimensional supramolecular packing architecture of (I). cyan represent π-π stacking interactions and N—H···π interactions while purple represent N—H···O, C—H···F and C—H···O interactions.

Methyl 2-(5-fluoro-1*H*-indol-3-yl)-2-oxoacetate

Crystal data

$C_{11}H_8FNO_3$
$M_r = 221.18$
Monoclinic, $P2_1/c$
Hall symbol: -P 2ybc
$a = 7.0584$ (14) Å
$b = 20.586$ (4) Å
$c = 7.3286$ (15) Å
$\beta = 112.01$ (3)°
$V = 987.3$ (3) Å^3
$Z = 4$
$F(000) = 456$
$D_x = 1.488$ Mg m^{-3}
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 2958 reflections
$\theta = 3.1$–$27.9°$
$\mu = 0.12$ mm^{-1}
$T = 113$ K
Block, colorless
0.30 × 0.24 × 0.20 mm

Data collection

Rigaku Saturn CCD area-detector diffractometer
Radiation source: rotating anode
Multilayer monochromator
Detector resolution: 7.31 pixels mm^{-1}
ω and φ scans
Absorption correction: multi-scan (*CrystalClear*; Rigaku/MSC, 2004)
$T_{min} = 0.964$, $T_{max} = 0.976$
9472 measured reflections
2328 independent reflections
2097 reflections with $I > 2\sigma(I)$
$R_{int} = 0.024$
$\theta_{max} = 27.9°$, $\theta_{min} = 3.1°$
$h = -9 \rightarrow 8$
$k = -27 \rightarrow 26$
$l = -9 \rightarrow 9$

Refinement

Refinement on F^2
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.037$
$wR(F^2) = 0.097$
$S = 1.06$
2328 reflections
150 parameters
0 restraints
H atoms treated by a mixture of independent and constrained refinement
$w = 1/[\sigma^2(F_o^2) + (0.0503P)^2 + 0.338P]$ where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{max} = 0.001$
$\Delta\rho_{max} = 0.38$ e Å^{-3}
$\Delta\rho_{min} = -0.20$ e Å^{-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 (Å^2)

	x	*y*	*z*	U_{iso}*/U_{eq}
F1	0.12543 (14)	0.29293 (3)	0.65340 (12)	0.0361 (2)
O1	0.26544 (13)	0.51327 (4)	0.99608 (11)	0.02011 (19)
O2	0.38836 (16)	0.66111 (4)	0.83866 (13)	0.0297 (2)
N1	0.26610 (14)	0.52509 (5)	0.37504 (13)	0.0173 (2)
O3	0.28315 (14)	0.63607 (4)	1.08373 (12)	0.0244 (2)
C1	0.23060 (16)	0.46201 (5)	0.42115 (15)	0.0162 (2)

C2	0.19687 (18)	0.40616 (6)	0.30676 (16)	0.0215 (2)
H2	0.1971	0.4072	0.1772	0.026*
C3	0.16302 (19)	0.34900 (6)	0.38896 (18)	0.0247 (3)
H3	0.1396	0.3095	0.3167	0.030*
C4	0.16368 (19)	0.35010 (5)	0.57927 (18)	0.0232 (3)
C5	0.20002 (17)	0.40431 (5)	0.69689 (16)	0.0187 (2)
H5	0.2012	0.4025	0.8269	0.022*
C6	0.23508 (16)	0.46224 (5)	0.61507 (15)	0.0149 (2)
C7	0.27633 (15)	0.52844 (5)	0.68412 (15)	0.0146 (2)
C8	0.29297 (16)	0.56415 (5)	0.52914 (15)	0.0161 (2)
H8	0.3194	0.6095	0.5323	0.019*
C9	0.28704 (16)	0.55005 (5)	0.87314 (15)	0.0151 (2)
C10	0.32621 (17)	0.62255 (5)	0.92560 (15)	0.0183 (2)
C11	0.3146 (2)	0.70317 (6)	1.14936 (19)	0.0327 (3)
H11A	0.4534	0.7166	1.1666	0.049*
H11B	0.2962	0.7072	1.2749	0.049*
H11C	0.2154	0.7310	1.0508	0.049*
H1	0.271 (2)	0.5363 (8)	0.253 (2)	0.036 (4)*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
F1	0.0544 (5)	0.0165 (3)	0.0358 (4)	−0.0065 (3)	0.0150 (4)	0.0025 (3)
O1	0.0294 (4)	0.0200 (4)	0.0134 (4)	−0.0006 (3)	0.0109 (3)	0.0013 (3)
O2	0.0479 (6)	0.0199 (4)	0.0264 (5)	−0.0071 (4)	0.0198 (4)	−0.0019 (3)
N1	0.0195 (5)	0.0220 (5)	0.0121 (4)	0.0002 (3)	0.0079 (4)	0.0007 (3)
O3	0.0397 (5)	0.0183 (4)	0.0178 (4)	0.0013 (3)	0.0141 (4)	−0.0039 (3)
C1	0.0150 (5)	0.0207 (5)	0.0134 (5)	0.0013 (4)	0.0058 (4)	−0.0006 (4)
C2	0.0218 (6)	0.0264 (6)	0.0171 (5)	0.0010 (4)	0.0082 (4)	−0.0057 (4)
C3	0.0263 (6)	0.0214 (5)	0.0247 (6)	0.0004 (4)	0.0077 (5)	−0.0078 (4)
C4	0.0266 (6)	0.0162 (5)	0.0254 (6)	−0.0003 (4)	0.0080 (5)	0.0022 (4)
C5	0.0209 (5)	0.0189 (5)	0.0155 (5)	0.0007 (4)	0.0061 (4)	0.0017 (4)
C6	0.0136 (5)	0.0184 (5)	0.0126 (5)	0.0012 (4)	0.0047 (4)	−0.0003 (4)
C7	0.0149 (5)	0.0174 (5)	0.0120 (5)	0.0011 (4)	0.0056 (4)	0.0009 (4)
C8	0.0167 (5)	0.0191 (5)	0.0130 (5)	0.0002 (4)	0.0061 (4)	0.0010 (4)
C9	0.0161 (5)	0.0170 (5)	0.0125 (5)	0.0010 (4)	0.0056 (4)	0.0000 (4)
C10	0.0219 (5)	0.0190 (5)	0.0130 (5)	0.0017 (4)	0.0054 (4)	−0.0007 (4)
C11	0.0547 (9)	0.0191 (6)	0.0251 (6)	0.0038 (5)	0.0158 (6)	−0.0059 (5)

Geometric parameters (Å, º)

F1—C4	1.3650 (13)	C3—H3	0.9500
O1—C9	1.2295 (13)	C4—C5	1.3742 (16)
O2—C10	1.1992 (14)	C5—C6	1.3978 (14)
N1—C8	1.3406 (13)	C5—H5	0.9500
N1—C1	1.3881 (14)	C6—C7	1.4450 (14)
N1—H1	0.932 (17)	C7—C8	1.3943 (14)
O3—C10	1.3330 (13)	C7—C9	1.4297 (14)

O3—C11	1.4520 (14)	C8—H8	0.9500
C1—C2	1.3898 (15)	C9—C10	1.5400 (15)
C1—C6	1.4097 (14)	C11—H11A	0.9800
C2—C3	1.3829 (17)	C11—H11B	0.9800
C2—H2	0.9500	C11—H11C	0.9800
C3—C4	1.3932 (17)		

C8—N1—C1	109.72 (9)	C5—C6—C7	134.36 (9)
C8—N1—H1	127.7 (10)	C1—C6—C7	106.36 (9)
C1—N1—H1	122.6 (10)	C8—C7—C9	129.44 (10)
C10—O3—C11	115.49 (9)	C8—C7—C6	106.22 (9)
N1—C1—C2	129.26 (10)	C9—C7—C6	124.29 (9)
N1—C1—C6	107.72 (9)	N1—C8—C7	109.97 (9)
C2—C1—C6	123.02 (10)	N1—C8—H8	125.0
C3—C2—C1	117.37 (10)	C7—C8—H8	125.0
C3—C2—H2	121.3	O1—C9—C7	122.84 (10)
C1—C2—H2	121.3	O1—C9—C10	118.29 (9)
C2—C3—C4	119.07 (10)	C7—C9—C10	118.87 (9)
C2—C3—H3	120.5	O2—C10—O3	124.87 (10)
C4—C3—H3	120.5	O2—C10—C9	125.17 (10)
F1—C4—C5	117.96 (11)	O3—C10—C9	109.96 (9)
F1—C4—C3	117.22 (10)	O3—C11—H11A	109.5
C5—C4—C3	124.82 (11)	O3—C11—H11B	109.5
C4—C5—C6	116.43 (10)	H11A—C11—H11B	109.5
C4—C5—H5	121.8	O3—C11—H11C	109.5
C6—C5—H5	121.8	H11A—C11—H11C	109.5
C5—C6—C1	119.27 (9)	H11B—C11—H11C	109.5

C8—N1—C1—C2	−179.36 (11)	C1—C6—C7—C8	0.36 (11)
C8—N1—C1—C6	0.23 (12)	C5—C6—C7—C9	−1.23 (19)
N1—C1—C2—C3	−179.31 (11)	C1—C6—C7—C9	177.96 (10)
C6—C1—C2—C3	1.15 (17)	C1—N1—C8—C7	0.00 (12)
C1—C2—C3—C4	0.19 (17)	C9—C7—C8—N1	−177.67 (10)
C2—C3—C4—F1	178.67 (11)	C6—C7—C8—N1	−0.23 (12)
C2—C3—C4—C5	−1.38 (19)	C8—C7—C9—O1	178.43 (11)
F1—C4—C5—C6	−178.92 (10)	C6—C7—C9—O1	1.41 (17)
C3—C4—C5—C6	1.13 (18)	C8—C7—C9—C10	−1.25 (17)
C4—C5—C6—C1	0.25 (15)	C6—C7—C9—C10	−178.28 (10)
C4—C5—C6—C7	179.36 (12)	C11—O3—C10—O2	0.50 (17)
N1—C1—C6—C5	178.98 (10)	C11—O3—C10—C9	179.76 (10)
C2—C1—C6—C5	−1.40 (16)	O1—C9—C10—O2	164.82 (12)
N1—C1—C6—C7	−0.36 (11)	C7—C9—C10—O2	−15.49 (17)
C2—C1—C6—C7	179.26 (10)	O1—C9—C10—O3	−14.44 (14)
C5—C6—C7—C8	−178.83 (12)	C7—C9—C10—O3	165.25 (9)

Hydrogen-bond geometry (Å, º)

D—H···*A*	*D*—H	H···*A*	*D*···*A*	*D*—H···*A*
N1—H1···O3^{i}	0.932 (17)	2.420 (16)	3.1598 (13)	136.3 (13)
N1—H1···O1^{i}	0.932 (17)	1.932 (17)	2.7861 (13)	151.3 (14)
C3—H3···F1^{ii}	0.95	2.41	3.3532 (15)	173
C5—H5···O1	0.95	2.55	3.0484 (14)	113
C8—H8···O2	0.95	2.36	2.9049 (14)	116

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