

N-(4-Cyanophenyl)-2,6-difluoro-benzamide

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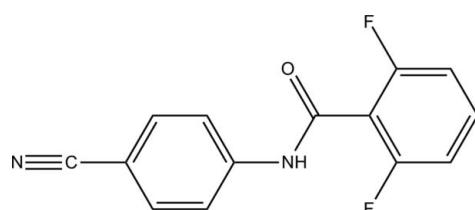
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.043; wR factor = 0.118; data-to-parameter ratio = 20.3.

In the title compound, $\text{C}_{14}\text{H}_8\text{F}_2\text{N}_2\text{O}$, the amide plane is inclined at dihedral angles of $28.12(12)$ and $32.89(12)^\circ$ with respect to the two benzene rings; the dihedral angle between the two rings is $5.58(5)^\circ$. In the crystal, intermolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{F}$ hydrogen bonds link adjacent molecules into a double-chain structure along the b axis.

Related literature

For general background to and applications of the title compound, see: Ashwood *et al.* (1990); Kees *et al.* (1989); Ragavan *et al.* (2010); Carmellino *et al.* (1994); Rauko *et al.* (2001). For a closely related benzamide structure, see: Cronin *et al.* (2000). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).

**Experimental***Crystal data*
 $M_r = 258.22$

Monoclinic, $P2_1/c$
 $a = 9.337(11)\text{ \AA}$
 $b = 5.0793(6)\text{ \AA}$
 $c = 24.500(3)\text{ \AA}$
 $\beta = 100.202(3)^\circ$
 $V = 1143.6(2)\text{ \AA}^3$

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.12\text{ mm}^{-1}$

$T = 100\text{ K}$
 $0.27 \times 0.14 \times 0.14\text{ mm}$

Data collection

Bruker APEXII DUO CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)
 $T_{\min} = 0.968$, $T_{\max} = 0.984$

27219 measured reflections
4135 independent reflections
3176 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.118$
 $S = 1.03$
4135 reflections

204 parameters
All H-atom parameters refined
 $\Delta\rho_{\max} = 0.39\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.23\text{ e \AA}^{-3}$

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

| $D-\text{H}\cdots A$ | $D-\text{H}$ | $\text{H}\cdots A$ | $D\cdots A$ | $D-\text{H}\cdots A$ |
|--|--------------|--------------------|-------------|----------------------|
| $\text{N}1-\text{H}1\text{N}1\cdots\text{O}1^{\text{i}}$ | 0.863 (15) | 2.107 (15) | 2.9029 (12) | 153.3 (14) |
| $\text{C}12-\text{H}12\cdots\text{F}1^{\text{ii}}$ | 0.940 (16) | 2.473 (16) | 3.4066 (14) | 172.4 (13) |

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IS2630).

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§ Thomson Reuters ResearcherID: C-7576-2009.

supporting information

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S1. Comment

A number of benzamide derivatives were reported as anti-hypertensive (Ashwood *et al.*, 1990), anti-diabetic (Kees *et al.*, 1989), anti-bacterial (Ragavan *et al.*, 2010), anti-fungal (Carmellino *et al.*, 1994) and anti-cancer (Rauko *et al.*, 2001) activities. On the basis of these considerations, our particular attention was paid for the synthesis of some benzamide derivatives.

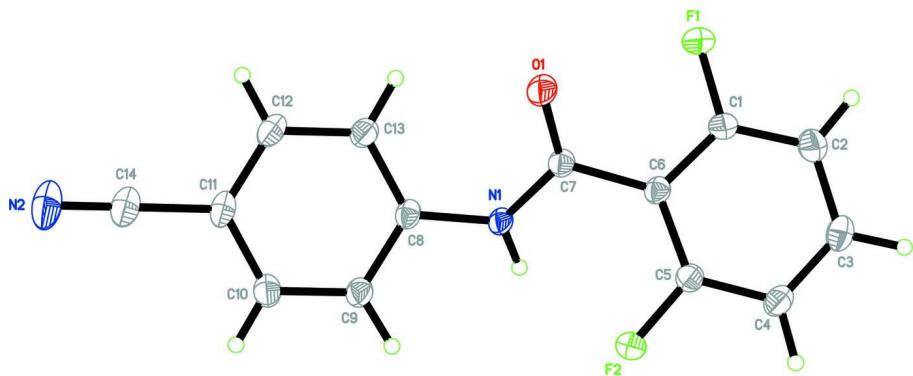
In the title benzamide derivative, the amino moiety (C7/N1/O1) is essentially planar, as indicated by the C7—O1—N1—H1N1 torsion angle of -1.4 (18)°. The mean plane through the amido moiety is inclined at dihedral angles of 32.89 (12) and 28.12 (12)°, respectively, with the C1—C6 and C8—C13 benzene rings. The dihedral angle between the two benzene rings being 5.58 (5)°. All bond lengths and angles are comparable to values observed in a closely related benzamide structure (Cronin *et al.*, 2000). In the crystal packing, adjacent molecules are interconnected into two-molecule-wide infinite chains propagating along the [010] direction (Fig. 2) *via* intermolecular N1—H1N1···O1 and C12—H12···F1 hydrogen bonds (Table 1).

S2. Experimental

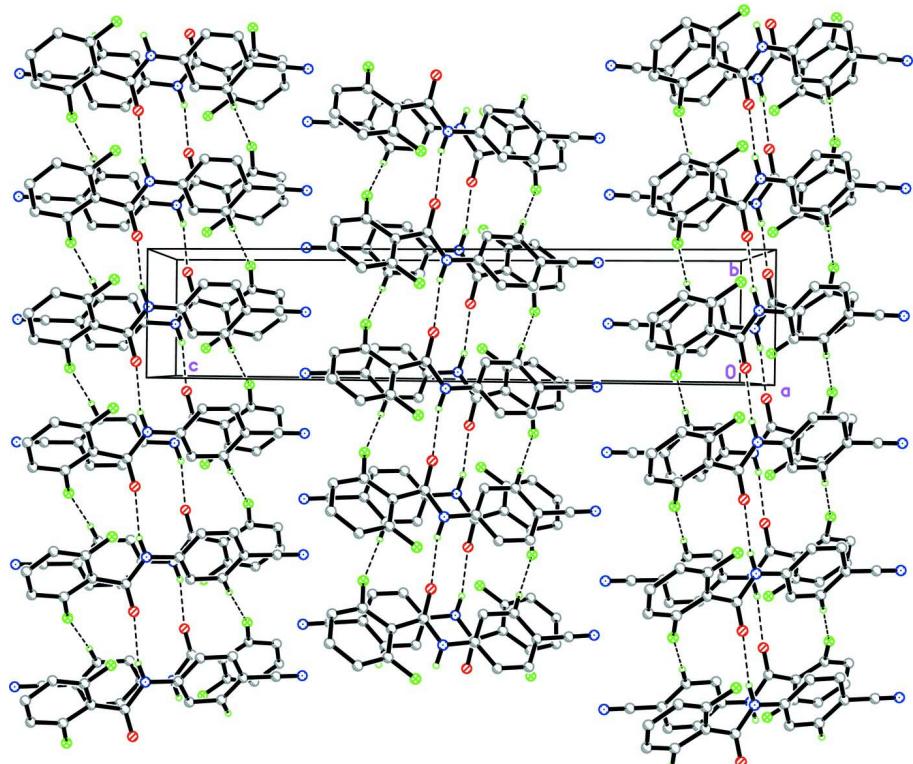
A mixture of 4-amino benzonitrile (4.2 mmol), 2,6-difluorobenzoic acid (4.6 mmol) and triethyl amine (21 mmol) was dissolved in methylene dichloride (5 ml). The resulting solution was cooled to 273 K followed by the drop wise addition of 50 % phosphoric acid cyclic anhydride solution in ethyl acetate (4 ml, 6.3 mmol) and stirred for 12 h. The completion of reaction was checked by TLC. Evaporation of solvent gave 2,6-difluoro-N-(*p*-cyanophenyl)benzamide as solid mass, which was then stirred with saturated NaHCO₃ solution to remove excess of acid. Single crystals suitable for X-ray analysis were obtained by crystallization from acetone under slow evaporation. *M.p.* 418 K.

S3. Refinement

All H atoms were located from a difference Fourier map and allowed to refine freely [refined distances: N—H = 0.860 (16) Å and C—H = 0.933 (15)–0.984 (15) Å].

**Figure 1**

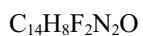
The molecular structure of the title compound, showing 50 % probability displacement ellipsoids for non-H atoms and the atom-numbering scheme.

**Figure 2**

The crystal structure of the title compound, viewed along the a axis, showing two-molecule-wide infinite chains along the [010] direction. H atoms not involved in intermolecular hydrogen bonds (dashed lines) have been omitted for clarity.

N-(4-Cyanophenyl)-2,6-difluorobenzamide

Crystal data



$M_r = 258.22$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 9.3377 (11) \text{ \AA}$

$b = 5.0793 (6) \text{ \AA}$

$c = 24.500 (3) \text{ \AA}$

$\beta = 100.202 (3)^\circ$

$V = 1143.6 (2) \text{ \AA}^3$
 $Z = 4$
 $F(000) = 528$
 $D_x = 1.500 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 5020 reflections

$\theta = 2.2\text{--}31.3^\circ$
 $\mu = 0.12 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
Block, yellow
 $0.27 \times 0.14 \times 0.14 \text{ mm}$

Data collection

Bruker APEXII DUO CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(SADABS; Bruker, 2009)
 $T_{\min} = 0.968$, $T_{\max} = 0.984$

27219 measured reflections
4135 independent reflections
3176 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.054$
 $\theta_{\max} = 32.6^\circ$, $\theta_{\min} = 2.2^\circ$
 $h = -14 \rightarrow 14$
 $k = -7 \rightarrow 7$
 $l = -36 \rightarrow 37$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.118$
 $S = 1.03$
4135 reflections
204 parameters
0 restraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
All H-atom parameters refined
 $w = 1/[\sigma^2(F_o^2) + (0.0568P)^2 + 0.2429P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.39 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.23 \text{ e \AA}^{-3}$

Special details

Experimental. The crystal was placed in the cold stream of an Oxford Cryosystems Cobra open-flow nitrogen cryostat (Cosier & Glazer, 1986) operating at 100.0 (1)K.

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 | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|----|--------------|--------------|--------------|----------------------------------|
| F1 | 0.81277 (8) | 0.06726 (14) | 0.14252 (3) | 0.03127 (18) |
| F2 | 1.02229 (7) | 0.75952 (13) | 0.05423 (3) | 0.02670 (16) |
| O1 | 0.76102 (9) | 0.12482 (15) | 0.03141 (3) | 0.02539 (18) |
| N1 | 0.75319 (9) | 0.56187 (17) | 0.01066 (3) | 0.01774 (17) |
| N2 | 0.33842 (13) | 0.5436 (2) | -0.24586 (5) | 0.0384 (3) |
| C1 | 0.91364 (12) | 0.2568 (2) | 0.14216 (4) | 0.0207 (2) |
| C2 | 1.01589 (13) | 0.2889 (2) | 0.18972 (5) | 0.0252 (2) |
| C3 | 1.11850 (12) | 0.4873 (2) | 0.19138 (5) | 0.0246 (2) |
| C4 | 1.11804 (11) | 0.6494 (2) | 0.14589 (5) | 0.0228 (2) |

| | | | | |
|------|--------------|--------------|--------------|--------------|
| C5 | 1.01538 (11) | 0.6060 (2) | 0.09898 (4) | 0.01910 (19) |
| C6 | 0.90844 (10) | 0.41080 (19) | 0.09449 (4) | 0.01688 (18) |
| C7 | 0.80070 (11) | 0.35107 (19) | 0.04285 (4) | 0.01768 (19) |
| C8 | 0.66236 (11) | 0.55108 (19) | -0.04175 (4) | 0.01768 (19) |
| C9 | 0.67895 (12) | 0.7489 (2) | -0.07969 (4) | 0.0207 (2) |
| C10 | 0.59456 (12) | 0.7492 (2) | -0.13217 (5) | 0.0224 (2) |
| C11 | 0.49135 (11) | 0.5511 (2) | -0.14682 (4) | 0.0220 (2) |
| C12 | 0.47172 (12) | 0.3569 (2) | -0.10873 (5) | 0.0231 (2) |
| C13 | 0.55706 (11) | 0.3555 (2) | -0.05623 (4) | 0.0209 (2) |
| C14 | 0.40564 (13) | 0.5474 (2) | -0.20187 (5) | 0.0275 (2) |
| H1N1 | 0.7848 (16) | 0.716 (3) | 0.0216 (6) | 0.031 (4)* |
| H2 | 1.0120 (16) | 0.173 (3) | 0.2194 (6) | 0.031 (4)* |
| H3 | 1.1931 (16) | 0.507 (3) | 0.2249 (6) | 0.031 (4)* |
| H4 | 1.1877 (17) | 0.787 (3) | 0.1456 (6) | 0.028 (4)* |
| H9 | 0.7526 (15) | 0.877 (3) | -0.0683 (6) | 0.022 (3)* |
| H10 | 0.6115 (15) | 0.879 (3) | -0.1573 (6) | 0.026 (4)* |
| H12 | 0.3999 (17) | 0.227 (3) | -0.1180 (6) | 0.031 (4)* |
| H13 | 0.5408 (15) | 0.221 (3) | -0.0309 (6) | 0.026 (4)* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|------------|------------|------------|-------------|-------------|-------------|
| F1 | 0.0336 (4) | 0.0283 (4) | 0.0293 (4) | -0.0134 (3) | -0.0015 (3) | 0.0072 (3) |
| F2 | 0.0242 (3) | 0.0266 (3) | 0.0274 (3) | -0.0063 (3) | -0.0006 (3) | 0.0092 (3) |
| O1 | 0.0327 (4) | 0.0133 (3) | 0.0260 (4) | -0.0007 (3) | -0.0061 (3) | -0.0027 (3) |
| N1 | 0.0212 (4) | 0.0123 (4) | 0.0175 (4) | -0.0004 (3) | -0.0026 (3) | -0.0018 (3) |
| N2 | 0.0415 (6) | 0.0392 (6) | 0.0283 (5) | 0.0086 (5) | -0.0108 (5) | -0.0071 (5) |
| C1 | 0.0227 (5) | 0.0172 (5) | 0.0215 (5) | -0.0029 (4) | 0.0021 (4) | 0.0004 (3) |
| C2 | 0.0283 (5) | 0.0265 (5) | 0.0195 (5) | -0.0012 (4) | 0.0011 (4) | 0.0017 (4) |
| C3 | 0.0235 (5) | 0.0285 (6) | 0.0198 (5) | -0.0004 (4) | -0.0018 (4) | -0.0042 (4) |
| C4 | 0.0190 (5) | 0.0222 (5) | 0.0257 (5) | -0.0025 (4) | -0.0003 (4) | -0.0029 (4) |
| C5 | 0.0187 (4) | 0.0174 (4) | 0.0206 (4) | 0.0002 (3) | 0.0020 (3) | 0.0004 (3) |
| C6 | 0.0173 (4) | 0.0145 (4) | 0.0179 (4) | 0.0004 (3) | 0.0007 (3) | -0.0023 (3) |
| C7 | 0.0188 (4) | 0.0144 (4) | 0.0187 (4) | 0.0009 (3) | 0.0002 (3) | -0.0014 (3) |
| C8 | 0.0186 (4) | 0.0156 (4) | 0.0176 (4) | 0.0020 (3) | -0.0003 (3) | -0.0030 (3) |
| C9 | 0.0240 (5) | 0.0170 (4) | 0.0194 (5) | -0.0008 (4) | -0.0007 (4) | -0.0015 (3) |
| C10 | 0.0264 (5) | 0.0202 (5) | 0.0189 (5) | 0.0021 (4) | -0.0002 (4) | -0.0006 (4) |
| C11 | 0.0214 (5) | 0.0226 (5) | 0.0195 (5) | 0.0049 (4) | -0.0037 (4) | -0.0055 (4) |
| C12 | 0.0196 (5) | 0.0206 (5) | 0.0266 (5) | -0.0003 (4) | -0.0030 (4) | -0.0051 (4) |
| C13 | 0.0206 (5) | 0.0181 (5) | 0.0224 (5) | -0.0004 (4) | -0.0008 (4) | -0.0012 (4) |
| C14 | 0.0279 (5) | 0.0259 (5) | 0.0254 (5) | 0.0056 (4) | -0.0043 (4) | -0.0057 (4) |

Geometric parameters (\AA , ^\circ)

| | | | |
|-------|-------------|-------|-------------|
| F1—C1 | 1.3480 (12) | C4—H4 | 0.956 (15) |
| F2—C5 | 1.3563 (12) | C5—C6 | 1.3975 (14) |
| O1—C7 | 1.2246 (12) | C6—C7 | 1.5014 (13) |
| N1—C7 | 1.3569 (13) | C8—C9 | 1.3961 (14) |

| | | | |
|-------------|--------------|-----------------|--------------|
| N1—C8 | 1.4091 (12) | C8—C13 | 1.3985 (14) |
| N1—H1N1 | 0.860 (16) | C9—C10 | 1.3839 (15) |
| N2—C14 | 1.1474 (15) | C9—H9 | 0.953 (14) |
| C1—C2 | 1.3791 (15) | C10—C11 | 1.3955 (15) |
| C1—C6 | 1.3992 (14) | C10—H10 | 0.933 (15) |
| C2—C3 | 1.3861 (16) | C11—C12 | 1.3922 (16) |
| C2—H2 | 0.943 (15) | C11—C14 | 1.4414 (15) |
| C3—C4 | 1.3849 (16) | C12—C13 | 1.3880 (15) |
| C3—H3 | 0.984 (15) | C12—H12 | 0.939 (15) |
| C4—C5 | 1.3777 (14) | C13—H13 | 0.952 (15) |
| | | | |
| C7—N1—C8 | 125.52 (8) | O1—C7—C6 | 120.86 (9) |
| C7—N1—H1N1 | 118.5 (10) | N1—C7—C6 | 115.57 (8) |
| C8—N1—H1N1 | 115.9 (10) | C9—C8—C13 | 119.89 (9) |
| F1—C1—C2 | 117.31 (9) | C9—C8—N1 | 117.32 (9) |
| F1—C1—C6 | 118.94 (9) | C13—C8—N1 | 122.78 (9) |
| C2—C1—C6 | 123.75 (10) | C10—C9—C8 | 120.48 (10) |
| C1—C2—C3 | 118.83 (10) | C10—C9—H9 | 122.3 (8) |
| C1—C2—H2 | 117.6 (9) | C8—C9—H9 | 117.2 (8) |
| C3—C2—H2 | 123.6 (9) | C9—C10—C11 | 119.47 (10) |
| C4—C3—C2 | 120.33 (10) | C9—C10—H10 | 118.6 (9) |
| C4—C3—H3 | 120.8 (9) | C11—C10—H10 | 121.9 (9) |
| C2—C3—H3 | 118.8 (9) | C12—C11—C10 | 120.36 (9) |
| C5—C4—C3 | 118.60 (10) | C12—C11—C14 | 120.06 (10) |
| C5—C4—H4 | 119.0 (9) | C10—C11—C14 | 119.58 (10) |
| C3—C4—H4 | 122.3 (9) | C13—C12—C11 | 120.18 (10) |
| F2—C5—C4 | 117.17 (9) | C13—C12—H12 | 119.3 (9) |
| F2—C5—C6 | 118.68 (9) | C11—C12—H12 | 120.5 (9) |
| C4—C5—C6 | 124.13 (10) | C12—C13—C8 | 119.60 (10) |
| C5—C6—C1 | 114.34 (9) | C12—C13—H13 | 118.5 (9) |
| C5—C6—C7 | 124.92 (9) | C8—C13—H13 | 121.9 (9) |
| C1—C6—C7 | 120.62 (9) | N2—C14—C11 | 179.41 (13) |
| O1—C7—N1 | 123.57 (9) | | |
| | | | |
| F1—C1—C2—C3 | -178.04 (10) | C1—C6—C7—O1 | -31.41 (15) |
| C6—C1—C2—C3 | 1.20 (18) | C5—C6—C7—N1 | -35.04 (14) |
| C1—C2—C3—C4 | -0.17 (18) | C1—C6—C7—N1 | 149.05 (10) |
| C2—C3—C4—C5 | -1.07 (17) | C7—N1—C8—C9 | -149.79 (11) |
| C3—C4—C5—F2 | -176.83 (10) | C7—N1—C8—C13 | 31.27 (16) |
| C3—C4—C5—C6 | 1.43 (17) | C13—C8—C9—C10 | -1.75 (16) |
| F2—C5—C6—C1 | 177.77 (9) | N1—C8—C9—C10 | 179.28 (10) |
| C4—C5—C6—C1 | -0.47 (15) | C8—C9—C10—C11 | 0.62 (16) |
| F2—C5—C6—C7 | 1.63 (15) | C9—C10—C11—C12 | 0.98 (16) |
| C4—C5—C6—C7 | -176.61 (10) | C9—C10—C11—C14 | -178.58 (10) |
| F1—C1—C6—C5 | 178.35 (9) | C10—C11—C12—C13 | -1.44 (16) |
| C2—C1—C6—C5 | -0.88 (16) | C14—C11—C12—C13 | 178.11 (10) |
| F1—C1—C6—C7 | -5.33 (15) | C11—C12—C13—C8 | 0.31 (16) |
| C2—C1—C6—C7 | 175.44 (10) | C9—C8—C13—C12 | 1.28 (16) |

| | | | |
|-------------|-------------|---------------|--------------|
| C8—N1—C7—O1 | −5.34 (17) | N1—C8—C13—C12 | −179.81 (10) |
| C8—N1—C7—C6 | 174.18 (9) | C7—O1—N1—H1N1 | −1.4 (18) |
| C5—C6—C7—O1 | 144.50 (11) | | |

Hydrogen-bond geometry (Å, °)

| D—H···A | D—H | H···A | D···A | D—H···A |
|----------------------------|------------|------------|-------------|------------|
| N1—H1N1···O1 ⁱ | 0.863 (15) | 2.107 (15) | 2.9029 (12) | 153.3 (14) |
| C12—H12···F1 ⁱⁱ | 0.940 (16) | 2.473 (16) | 3.4066 (14) | 172.4 (13) |

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