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5-(Trifluoromethoxy)isatin

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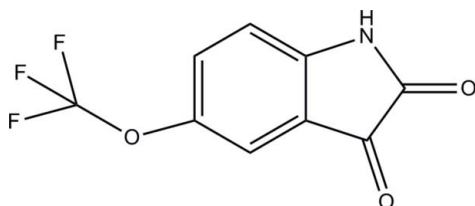
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.036; wR factor = 0.102; data-to-parameter ratio = 14.9.

The title compound [systematic name: 5-(trifluoromethoxy)-1*H*-indole-2,3-dione], $\text{C}_9\text{H}_4\text{F}_3\text{NO}_3$, crystallized with two molecules in the asymmetric unit. Intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds link the molecules to form layers parallel to the *ab* plane. In addition, π - π stacking interactions are observed with a centroid-centroid distance of 3.721 (1) Å. The near planarity of the two isatin ring systems is illustrated by the maximum deviations of 0.023 (1) and 0.025 (1) Å for the N atom in each case.

Related literature

For similar structures and background information on isatin as a biological agent, see Schutte *et al.* (2012); Garden *et al.* (2006); Goldschmidt & Llewellyn (1950); Frolova *et al.* (1988); Wei *et al.* (2004); Palmer *et al.* (1987); Akkurt *et al.* (2006). For reaction kinetic data on similar structures, see: Schutte *et al.* (2011).



Experimental

Crystal data

$\text{C}_9\text{H}_4\text{F}_3\text{NO}_3$
 $M_r = 231.13$
 Monoclinic, $P2_1/n$
 $a = 14.329$ (5) Å
 $b = 6.082$ (5) Å
 $c = 20.401$ (5) Å
 $\beta = 91.608$ (5)°

$V = 1777.2$ (16) Å³
 $Z = 8$
 Mo $K\alpha$ radiation
 $\mu = 0.17$ mm⁻¹
 $T = 100$ K
 $0.51 \times 0.18 \times 0.10$ mm

Data collection

Bruker APEXII CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2008)
 $T_{\min} = 0.152$, $T_{\max} = 0.394$

26198 measured reflections
 4428 independent reflections
 3513 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.029$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.036$
 $wR(F^2) = 0.102$
 $S = 1.04$
 4418 reflections
 297 parameters

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.34$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.33$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{O4}^{\text{i}}$	0.84 (2)	1.99 (2)	2.7615 (18)	152.8 (18)
$\text{N2}-\text{H2}\cdots\text{O1}^{\text{ii}}$	0.89 (2)	2.03 (2)	2.8776 (18)	157.4 (19)
$\text{N2}-\text{H2}\cdots\text{O4}^{\text{iii}}$	0.89 (2)	2.55 (2)	2.9850 (18)	111.2 (16)
$\text{C16}-\text{H16}\cdots\text{F3}^{\text{i}}$	0.93	2.39	3.171 (2)	142
$\text{C18}-\text{H18}\cdots\text{O2}^{\text{iv}}$	0.93	2.47	3.327 (3)	153

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

Data collection: APEX2 (Bruker, 2008); cell refinement: SAINT-Plus (Bruker, 2008); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: DIAMOND (Brandenburg & Putz, 2005); software used to prepare material for publication: WinGX (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FY2072).

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supplementary materials

Acta Cryst. (2012). E68, o3472 [doi:10.1107/S1600536812044297]

5-(Trifluoromethoxy)isatin

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Comment

5-(Trifluoromethoxy)isatin crystallized in the $P2_1/n$ spacegroup and with two independent molecules in the asymmetric unit. The use of 5-(trifluoromethoxy)isatin forms part of a current study, where water soluble O,*O'*-donor bidentate ligands are coordinated to the rhenium tricarbonyl core in order to study the reactivity and activation of these potential radiopharmaceutical compounds. Previous studies have shown that the O,*O'*-bidentate ligands activate the Re(I) core to a great extent and we would like to explore this further (Schutte *et al.* (2011)). The isatin molecule also has a variety of biological activities. It can act as an anticonvulsant agent, it has sedative effects but can also cause anxiety. Palmer *et al.*, 1987; Goldschmidt & Llewellyn, 1950; Wei *et al.*, 2004; Frolova *et al.*, 1988; Akkurt *et al.*, 2006). Due to its diverse pharmacological properties, a lot of interest has been shown in the isatin molecule and its derivatives. All bond lengths and angles compare well to the structure of *N*-benzylisatin reported by Schutte *et al.* (2012). Five intermolecular hydrogen interactions are observed in the structure (C—H \cdots O, N—H \cdots O and C—H \cdots F) as well as π - π stacking with a centroid to centroid distance of 3.721 (1) Å. The molecules pack in a head to head fashion in alternating layers with the benzyl ring overlapping with the next molecules' pyrrolidinedione ring and *vice versa*. The π - π stacking and the packing in the unit cell are illustrated in Figure 2. The planarity of the ring systems, N1—C11—C12—C13—C14—C15—C16—C17—C18 and N2—C21—C22—C23—C24—C25—C26—C27—C28, are illustrated by very small deviations of all the atoms from these planes, with the largest deviations 0.023 (1) Å for N1 and 0.025 (1) for N2 respectively.

Experimental

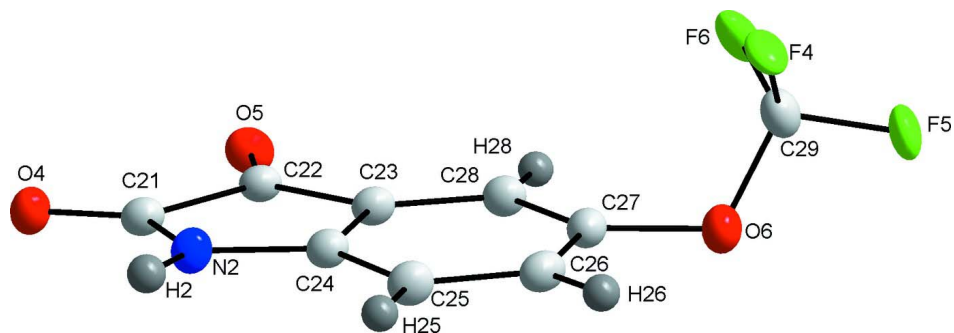
5-(Trifluoromethoxy)isatin was dissolved in water (pH 4). The crystals were grown from a vapour diffusion setup with chloroform at 5 °C in a commercial refrigerator.

Refinement

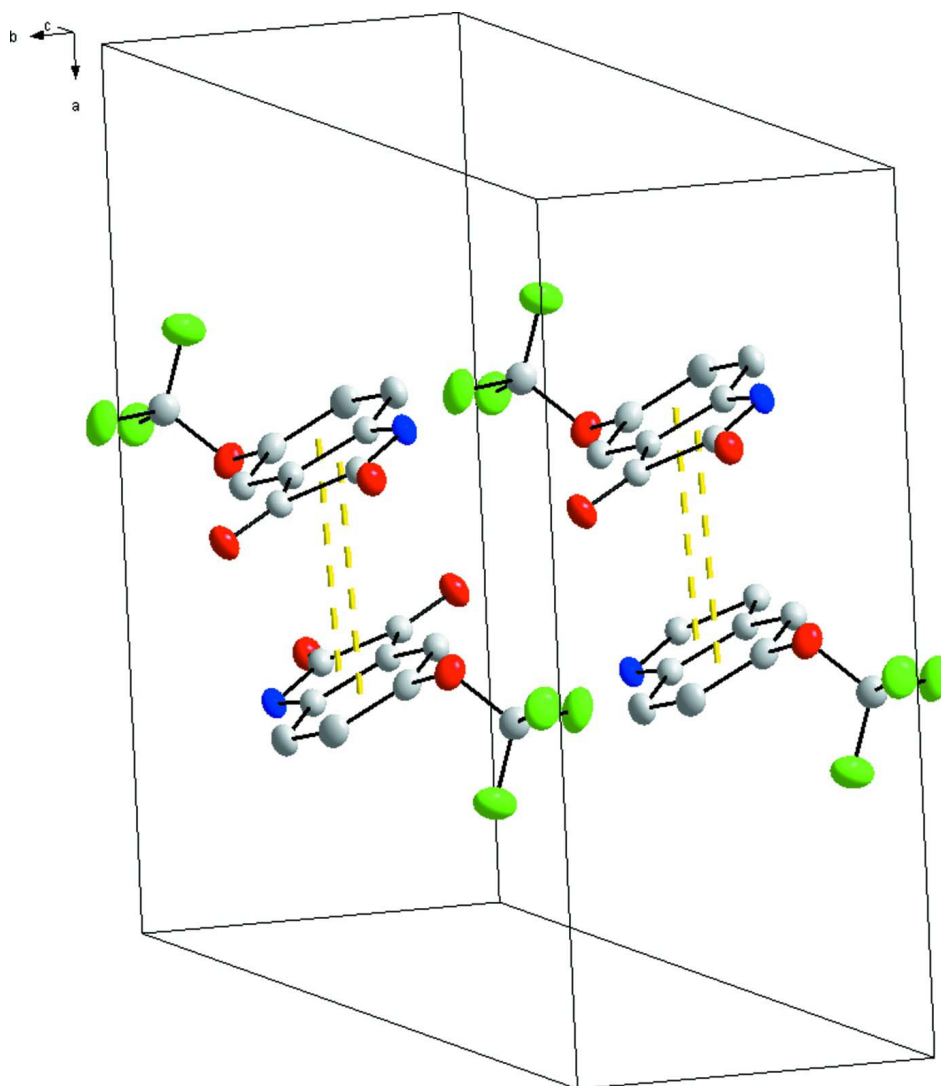
Aromatic H atoms were positioned geometrically and allowed to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{parent})$ and with a C—H distance of 0.93 Å. N-bound hydrogen atoms were placed from the electron density map and refined freely. 10 reflections with large negative intensities [$I < -4\sigma(I)$] were excluded from the refinement.

Computing details

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINTE-Plus* (Bruker, 2008); data reduction: *SAINTE-Plus* (Bruker, 2008); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *DIAMOND* (Brandenburg & Putz, 2005); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

**Figure 1**

Representation of the title compound, showing the numbering scheme and displacement ellipsoids (50% probability).

**Figure 2**

Observed π - π stacking (indicated by dashed lines) and packing in the crystal structure (hydrogen atoms omitted for clarity).

5-(trifluoromethoxy)-1*H*-indole-2,3-dione

Crystal data

C₉H₄F₃NO₃

$M_r = 231.13$

Monoclinic, $P2_1/n$

$a = 14.329 (5) \text{ \AA}$

$b = 6.082 (5) \text{ \AA}$

$c = 20.401 (5) \text{ \AA}$

$\beta = 91.608 (5)^\circ$

$V = 1777.2 (16) \text{ \AA}^3$

$Z = 8$

$F(000) = 928$

$D_x = 1.728 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71069 \text{ \AA}$

Cell parameters from 9825 reflections

$\theta = 2.8\text{--}28.2^\circ$

$\mu = 0.17 \text{ mm}^{-1}$

$T = 100 \text{ K}$

Needle, orange

$0.51 \times 0.18 \times 0.10 \text{ mm}$

Data collection

Bruker APEXII CCD

diffractometer

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2008)

$T_{\min} = 0.152$, $T_{\max} = 0.394$

26198 measured reflections

4428 independent reflections

3513 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$

$\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 1.8^\circ$

$h = -19 \rightarrow 19$

$k = -7 \rightarrow 8$

$l = -27 \rightarrow 27$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.036$

$wR(F^2) = 0.102$

$S = 1.04$

4418 reflections

297 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

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0458P)^2 + 0.9713P]$

where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.34 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.33 \text{ e \AA}^{-3}$

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$
O2	0.44882 (8)	0.39123 (17)	0.43829 (5)	0.0258 (2)
O4	0.25592 (7)	0.62417 (16)	0.33404 (5)	0.0222 (2)
O3	0.41251 (8)	0.06479 (18)	0.69260 (5)	0.0255 (2)
O6	0.01035 (8)	-0.39384 (17)	0.38480 (5)	0.0258 (2)
N1	0.33820 (9)	-0.1165 (2)	0.43057 (6)	0.0207 (3)

F3	0.35956 (8)	0.40475 (15)	0.70118 (5)	0.0375 (3)
O5	0.21790 (7)	0.35661 (17)	0.44843 (5)	0.0230 (2)
F2	0.38339 (8)	0.20705 (17)	0.78654 (5)	0.0366 (2)
F5	-0.09269 (8)	-0.57440 (18)	0.43734 (6)	0.0482 (3)
O1	0.37037 (8)	0.09137 (17)	0.33911 (5)	0.0241 (2)
N2	0.16708 (9)	0.3571 (2)	0.28133 (6)	0.0202 (3)
C13	0.39620 (10)	0.1071 (2)	0.51369 (7)	0.0188 (3)
C14	0.34985 (10)	-0.0899 (2)	0.49875 (7)	0.0196 (3)
C11	0.37229 (10)	0.0574 (2)	0.39792 (7)	0.0202 (3)
C23	0.13214 (10)	0.1310 (2)	0.36813 (7)	0.0184 (3)
F6	-0.05978 (9)	-0.2481 (2)	0.46963 (6)	0.0564 (4)
C18	0.41720 (10)	0.1665 (2)	0.57812 (7)	0.0202 (3)
H18	0.449	0.2958	0.5885	0.024*
F1	0.26737 (7)	0.1395 (2)	0.72116 (5)	0.0436 (3)
F4	-0.13875 (8)	-0.2999 (2)	0.38080 (7)	0.0564 (4)
C15	0.32098 (11)	-0.2309 (2)	0.54708 (8)	0.0244 (3)
H15	0.2897	-0.361	0.5369	0.029*
C17	0.38837 (10)	0.0232 (2)	0.62607 (7)	0.0217 (3)
C26	0.03648 (10)	-0.1692 (2)	0.29098 (7)	0.0221 (3)
H26	0.0038	-0.2727	0.2659	0.027*
C24	0.12267 (10)	0.1618 (2)	0.30027 (7)	0.0190 (3)
C21	0.20969 (10)	0.4565 (2)	0.33289 (7)	0.0195 (3)
C28	0.09416 (10)	-0.0507 (2)	0.39831 (7)	0.0198 (3)
H28	0.101	-0.0734	0.4433	0.024*
C27	0.04544 (10)	-0.1968 (2)	0.35822 (7)	0.0209 (3)
C12	0.41347 (10)	0.2177 (2)	0.45094 (7)	0.0196 (3)
C29	-0.06921 (12)	-0.3765 (3)	0.41759 (9)	0.0324 (4)
C22	0.18838 (10)	0.3137 (2)	0.39389 (7)	0.0182 (3)
C25	0.07594 (10)	0.0119 (2)	0.26051 (7)	0.0217 (3)
H25	0.071	0.0311	0.2153	0.026*
C19	0.35582 (11)	0.2012 (3)	0.72439 (7)	0.0251 (3)
C16	0.34070 (11)	-0.1702 (2)	0.61182 (8)	0.0250 (3)
H16	0.3217	-0.2602	0.6458	0.03*
H1	0.3123 (13)	-0.224 (3)	0.4120 (9)	0.028 (5)*
H2	0.1668 (14)	0.403 (3)	0.2399 (11)	0.041 (6)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O2	0.0326 (6)	0.0228 (5)	0.0219 (5)	-0.0080 (4)	-0.0022 (4)	0.0025 (4)
O4	0.0266 (5)	0.0198 (5)	0.0204 (5)	-0.0040 (4)	0.0004 (4)	-0.0013 (4)
O3	0.0346 (6)	0.0267 (5)	0.0150 (5)	0.0069 (4)	-0.0031 (4)	0.0020 (4)
O6	0.0279 (6)	0.0206 (5)	0.0291 (6)	-0.0038 (4)	0.0033 (4)	0.0012 (4)
N1	0.0249 (6)	0.0178 (6)	0.0193 (6)	-0.0031 (5)	-0.0038 (5)	-0.0016 (5)
F3	0.0583 (7)	0.0238 (5)	0.0308 (5)	0.0088 (4)	0.0079 (5)	0.0058 (4)
O5	0.0240 (5)	0.0285 (5)	0.0165 (5)	-0.0019 (4)	-0.0021 (4)	-0.0034 (4)
F2	0.0518 (6)	0.0408 (6)	0.0171 (5)	0.0045 (5)	0.0000 (4)	0.0003 (4)
F5	0.0507 (7)	0.0394 (6)	0.0552 (7)	-0.0198 (5)	0.0141 (6)	0.0074 (5)
O1	0.0310 (6)	0.0246 (5)	0.0166 (5)	-0.0014 (4)	-0.0024 (4)	0.0000 (4)
N2	0.0259 (6)	0.0198 (6)	0.0147 (6)	-0.0021 (5)	-0.0011 (5)	-0.0005 (5)

C13	0.0205 (7)	0.0176 (6)	0.0183 (7)	0.0003 (5)	-0.0016 (5)	0.0016 (5)
C14	0.0205 (7)	0.0189 (6)	0.0192 (7)	0.0006 (5)	-0.0022 (5)	0.0001 (5)
C11	0.0207 (7)	0.0195 (6)	0.0203 (7)	0.0005 (5)	-0.0020 (5)	-0.0012 (5)
C23	0.0194 (7)	0.0192 (6)	0.0165 (7)	0.0011 (5)	-0.0008 (5)	-0.0029 (5)
F6	0.0623 (8)	0.0617 (8)	0.0470 (7)	-0.0306 (6)	0.0310 (6)	-0.0241 (6)
C18	0.0219 (7)	0.0194 (6)	0.0192 (7)	0.0003 (5)	-0.0023 (5)	0.0002 (5)
F1	0.0322 (6)	0.0599 (7)	0.0391 (6)	-0.0086 (5)	0.0089 (5)	-0.0075 (5)
F4	0.0260 (6)	0.0664 (8)	0.0766 (9)	0.0006 (5)	0.0004 (6)	0.0168 (7)
C15	0.0283 (8)	0.0179 (7)	0.0269 (8)	-0.0025 (6)	-0.0004 (6)	0.0020 (6)
C17	0.0264 (7)	0.0229 (7)	0.0157 (7)	0.0047 (6)	-0.0018 (5)	0.0015 (5)
C26	0.0215 (7)	0.0206 (7)	0.0241 (7)	0.0003 (5)	-0.0013 (6)	-0.0057 (6)
C24	0.0199 (7)	0.0187 (6)	0.0183 (7)	0.0017 (5)	0.0000 (5)	-0.0006 (5)
C21	0.0207 (7)	0.0195 (6)	0.0182 (7)	0.0018 (5)	0.0007 (5)	-0.0020 (5)
C28	0.0211 (7)	0.0210 (7)	0.0175 (7)	0.0009 (5)	0.0002 (5)	-0.0010 (5)
C27	0.0203 (7)	0.0178 (6)	0.0247 (7)	-0.0001 (5)	0.0028 (6)	-0.0006 (5)
C12	0.0201 (7)	0.0204 (7)	0.0181 (7)	-0.0004 (5)	-0.0027 (5)	-0.0001 (5)
C29	0.0317 (9)	0.0316 (8)	0.0340 (9)	-0.0095 (7)	0.0050 (7)	-0.0019 (7)
C22	0.0183 (6)	0.0189 (6)	0.0173 (7)	0.0019 (5)	0.0008 (5)	-0.0019 (5)
C25	0.0240 (7)	0.0232 (7)	0.0178 (7)	0.0000 (6)	-0.0023 (5)	-0.0027 (5)
C19	0.0300 (8)	0.0260 (7)	0.0193 (7)	0.0010 (6)	0.0015 (6)	0.0043 (6)
C16	0.0316 (8)	0.0203 (7)	0.0233 (8)	0.0015 (6)	0.0028 (6)	0.0069 (6)

Geometric parameters (Å, °)

O2—C12	1.2017 (19)	C11—C12	1.560 (2)
O4—C21	1.2157 (19)	C23—C28	1.384 (2)
O3—C19	1.3410 (19)	C23—C24	1.400 (2)
O3—C17	1.4140 (18)	C23—C22	1.462 (2)
O6—C29	1.342 (2)	F6—C29	1.322 (2)
O6—C27	1.4134 (19)	C18—C17	1.382 (2)
N1—C11	1.349 (2)	C18—H18	0.93
N1—C14	1.4055 (19)	F1—C19	1.322 (2)
N1—H1	0.84 (2)	F4—C29	1.316 (2)
F3—C19	1.327 (2)	C15—C16	1.393 (2)
O5—C22	1.2078 (17)	C15—H15	0.93
F2—C19	1.3180 (18)	C17—C16	1.387 (2)
F5—C29	1.316 (2)	C26—C27	1.384 (2)
O1—C11	1.2170 (18)	C26—C25	1.393 (2)
N2—C21	1.3450 (18)	C26—H26	0.93
N2—C24	1.407 (2)	C24—C25	1.381 (2)
N2—H2	0.89 (2)	C21—C22	1.555 (2)
C13—C18	1.388 (2)	C28—C27	1.383 (2)
C13—C14	1.399 (2)	C28—H28	0.93
C13—C12	1.473 (2)	C25—H25	0.93
C14—C15	1.379 (2)	C16—H16	0.93
C19—O3—C17	116.08 (12)	C23—C24—N2	110.67 (12)
C29—O6—C27	116.17 (12)	O4—C21—N2	128.85 (14)
C11—N1—C14	111.35 (12)	O4—C21—C22	124.91 (13)
C11—N1—H1	123.4 (13)	N2—C21—C22	106.24 (12)

C14—N1—H1	125.2 (13)	C27—C28—C23	116.62 (13)
C21—N2—C24	111.32 (12)	C27—C28—H28	121.7
C21—N2—H2	126.2 (14)	C23—C28—H28	121.7
C24—N2—H2	122.4 (14)	C28—C27—C26	122.61 (14)
C18—C13—C14	121.22 (13)	C28—C27—O6	119.79 (13)
C18—C13—C12	131.81 (13)	C26—C27—O6	117.33 (13)
C14—C13—C12	106.98 (12)	O2—C12—C13	132.05 (13)
C15—C14—C13	121.78 (14)	O2—C12—C11	123.61 (13)
C15—C14—N1	127.34 (14)	C13—C12—C11	104.33 (12)
C13—C14—N1	110.88 (12)	F5—C29—F4	107.62 (14)
O1—C11—N1	128.44 (13)	F5—C29—F6	108.41 (15)
O1—C11—C12	125.11 (13)	F4—C29—F6	107.90 (16)
N1—C11—C12	106.43 (12)	F5—C29—O6	108.00 (14)
C28—C23—C24	121.25 (13)	F4—C29—O6	112.56 (15)
C28—C23—C22	131.81 (13)	F6—C29—O6	112.18 (14)
C24—C23—C22	106.91 (12)	O5—C22—C23	132.15 (13)
C17—C18—C13	116.42 (13)	O5—C22—C21	123.01 (13)
C17—C18—H18	121.8	C23—C22—C21	104.81 (12)
C13—C18—H18	121.8	C24—C25—C26	117.14 (14)
C14—C15—C16	117.14 (14)	C24—C25—H25	121.4
C14—C15—H15	121.4	C26—C25—H25	121.4
C16—C15—H15	121.4	F2—C19—F1	108.46 (13)
C18—C17—C16	122.79 (14)	F2—C19—F3	107.71 (13)
C18—C17—O3	119.79 (13)	F1—C19—F3	107.18 (13)
C16—C17—O3	117.31 (13)	F2—C19—O3	108.22 (13)
C27—C26—C25	120.73 (13)	F1—C19—O3	113.15 (14)
C27—C26—H26	119.6	F3—C19—O3	111.94 (13)
C25—C26—H26	119.6	C17—C16—C15	120.63 (14)
C25—C24—C23	121.63 (13)	C17—C16—H16	119.7
C25—C24—N2	127.70 (13)	C15—C16—H16	119.7

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1—H1 \cdots O4 ⁱ	0.84 (2)	1.99 (2)	2.7615 (18)	152.8 (18)
N2—H2 \cdots O1 ⁱⁱ	0.89 (2)	2.03 (2)	2.8776 (18)	157.4 (19)
N2—H2 \cdots O4 ⁱⁱⁱ	0.89 (2)	2.55 (2)	2.9850 (18)	111.2 (16)
C16—H16 \cdots F3 ⁱ	0.93	2.39	3.171 (2)	142
C18—H18 \cdots O2 ^{iv}	0.93	2.47	3.327 (3)	153

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