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5-Acetyl-2-amino-4-(2-fluorophenyl)-6-methyl-4H-pyran-3-carbonitrile dichloromethane hemisolvate

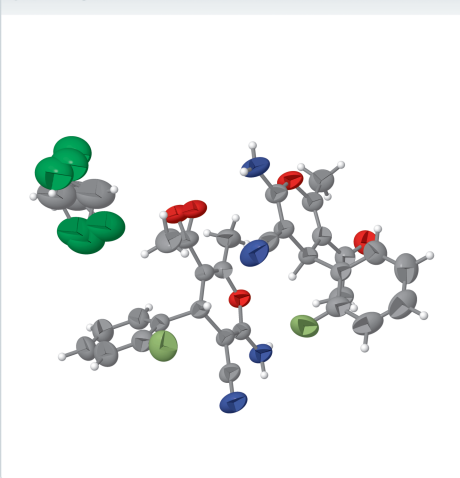
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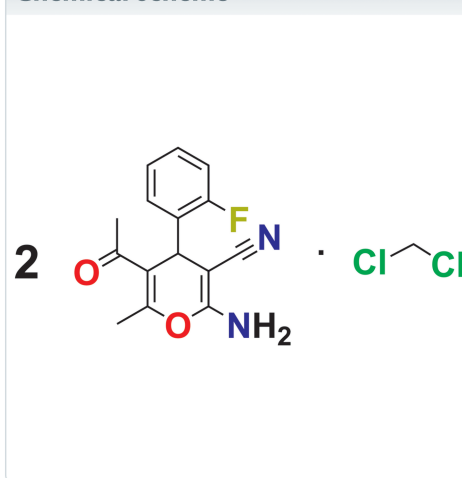
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The title compound, $2C_{15}H_{13}FN_2O_2 \cdot CH_2Cl_2$, crystallizes with two main molecules in the asymmetric unit with a disordered dichloromethane solvent molecule. The dihedral angles between the fluoro phenyl ring and 4H-pyran ring are $74.36(15)$ and $80.69(15)^\circ$ in the two molecules. In the crystal, $N-H \cdots N$ and $N-H \cdots O$ hydrogen bonds link the molecules into a two-dimensional supramolecular network propagating in the (100) plane.

3D view



Chemical scheme



Structure description

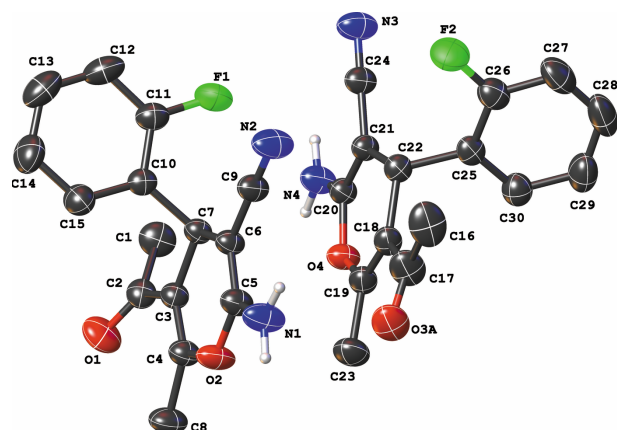
Amino-4H-pyran derivatives are useful building blocks in creating pharmacologically active heterocycles in multicomponent reactions, such as anti-tumor (Fouda, 2016), antibacterial (Kathrotiya & Patel, 2012), antimycobacterial (Alvey *et al.*, 2009) anti-leishmanial (Narender & Gupta, 2004) and antiproliferative agents (Mansouri *et al.*, 2011). Our previous study on aryl-based 2-amino carbonitriles identified different hydrogen-bonding patterns in the crystal structures (Zamisa *et al.*, 2022). A structural analysis using the Cambridge Structural Database (CSD version 5.46, November 2024 update); Groom *et al.*, 2016) showed various types of hydrogen-bonding patterns driven by substituents on the 4H-pyran core. These were dominated by $N-H \cdots N$ and $N-H \cdots O$ interactions involving the amino and carbonyl/cyano groups. This analysis indicated that the binding affinity of these compounds towards calf thymus deoxyribonucleic acid may be associated with hydrogen bonds involving their amino functional groups. This finding is consistent with our exploration of potential anticancer agents (Zamisa *et al.*, 2022). The current study continues our investigation into the structures of 4H-pyran derivatives as potential anticancer agents and reports the synthesis and structure of the title compound.

There are two symmetrically independent $C_{15}H_{13}FN_2O_2$ molecules in the asymmetric unit (Fig. 1). Each molecule comprises a cycloalkanone, 4H-pyran core with attached



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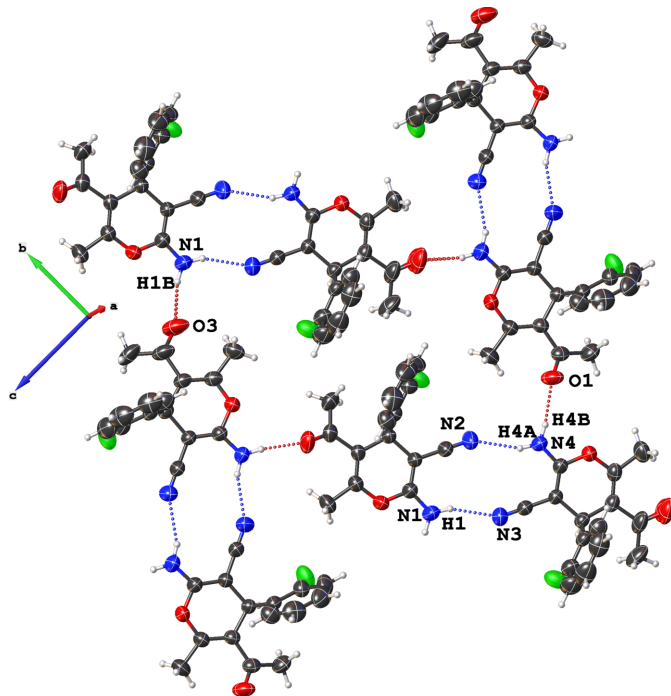
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**Figure 1**

The molecular structure of the title compound with displacement ellipsoids drawn at the 40% probability level. The disordered components, including the aromatic and methine hydrogen atoms, are omitted for clarity.

phenyl moiety, cyano, and amino groups. The dihedral angle between the fluorophenyl and 4*H*-pyran rings are 74.36 (15)° (C1 molecule) and 80.69 (15)° (C16 molecule) and are similar to those of related compounds in the literature (Zamisa *et al.*, 2022, 2023).

In the crystal, the molecules are linked by N—H···N and N—H···O hydrogen bonds (Table 1, Fig. 2) engendered by the amine functional group. One of these hydrogen atoms, H1*B* or H4*B*, interacts with the nitrogen atom N3 or N2 of a neighbouring molecule *via* an N—H···N link with the graph-

**Figure 2**

A projection of the crystal packing along the *a*-axis. Dashed lines denote N—H···N and N—H···O hydrogen bonds. Motifs **I** and **II** (see text) are depicted by blue and red dotted lines, respectively.

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
N1—H1 <i>A</i> ···N3 ⁱ	0.86	2.26	3.066 (4)	156
N1—H1 <i>B</i> ···O3 ⁱⁱ	0.86	2.03	2.844 (8)	159
N1—H1 <i>B</i> ···O3 <i>A</i> ⁱⁱ	0.86	2.02	2.855 (7)	165
N4—H4 <i>A</i> ···N2 ⁱⁱⁱ	0.86	2.20	3.016 (4)	157
N4—H4 <i>B</i> ···O1 ^{iv}	0.86	2.06	2.910 (3)	170

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

Table 2

Experimental details.

Crystal data	
Chemical formula	2C ₁₅ H ₁₃ FN ₂ O ₂ ·CH ₂ Cl ₂
<i>M</i> _r	629.47
Crystal system, space group	Monoclinic, <i>P</i> ₂ /c
Temperature (K)	296
<i>a</i> , <i>b</i> , <i>c</i> (Å)	10.8670 (4), 15.2671 (6), 19.1569 (8)
β (°)	97.463 (2)
<i>V</i> (Å ³)	3151.4 (2)
<i>Z</i>	4
Radiation type	Mo <i>K</i> α
μ (mm ^{−1})	0.26
Crystal size (mm)	0.31 × 0.22 × 0.16
Data collection	
Diffractometer	Bruker SMART APEXII CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Krause <i>et al.</i> , 2015)
<i>T</i> _{min} , <i>T</i> _{max}	0.485, 0.746
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	84575, 6190, 4405
<i>R</i> _{int}	0.063
(sin θ/λ) _{max} (Å ^{−1})	0.617
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.074, 0.239, 1.08
No. of reflections	6190
No. of parameters	419
No. of restraints	59
H-atom treatment	H-atom parameters constrained
Δρ _{max} , Δρ _{min} (e Å ^{−3})	0.71, −0.61

Computer programs: *APEX2* and *SAINT* (Bruker, 2016), *SHELXT2018/2* (Sheldrick, 2015*a*), *SHELXL2018/3* (Sheldrick, 2015*b*) and *OLEX2* 1.5 (Dolomanov *et al.*, 2009).

set descriptor *R*₂²(12) (involving the amino and the cyano group) (Motif **I**). An N—H···O hydrogen bond graph with graph-set descriptor *C*₂²(10) with the carbonyl group oxygen atom O3 or O1 of a neighbouring molecule acting as acceptor to the amine (Motif **II**), Fig. 2 (Nyapola *et al.*, 2024; Zamisa *et al.*, 2022) is also observed. These two motifs combined create a supramolecular structure that propagates in the (100) plane of the crystal.

Synthesis and crystallization

0.015 mmol of 1,3-cyclohexanedione were mixed with 0.015 mmol of malonitrile and 0.015 mmol of benzaldehyde in a microwave vessel. A catalytic amount of triethylamine was added in a tightly sealed 35 ml microwave reaction vessel, and the mixture was subjected to microwave radiation at 150°C for 10 minutes. An off-white solid precipitate was formed and collected by vacuum filtration. The reaction progress was

monitored using thin-layer chromatography with a solvent ratio of 1:1 for ethyl acetate and hexane. The resulting precipitate was isolated and recrystallized from ethanol solution. Crystals of the title compound were obtained through slow isothermal evaporation from absolute dichloromethane solution.

Refinement

Crystallographic data and structure refinement details are summarized in Table 2. SIMU restraints and EADP constraints in *SHELXL* were used to model the disorder of the solvent molecule. The hydrogen atoms were positioned geometrically with N—H = 0.86 Å and C—H = 0.93–0.96 Å depending on hybridization and allowed to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Acknowledgements

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full crystallographic data

IUCrData (2025). **10**, x250337 [<https://doi.org/10.1107/S2414314625003372>]

5-Acetyl-2-amino-4-(2-fluorophenyl)-6-methyl-4*H*-pyran-3-carbonitrile dichloromethane hemisolvate

Carren Nyapola, Sizwe J. Zamisa, Bernard Omondi and Eric M. Njogu

5-Acetyl-2-amino-4-(2-fluorophenyl)-6-methyl-4*H*-pyran-3-carbonitrile dichloromethane hemisolvate

Crystal data

$2\text{C}_{15}\text{H}_{13}\text{FN}_2\text{O}_2 \cdot \text{CH}_2\text{Cl}_2$

$M_r = 629.47$

Monoclinic, $P2_1/c$

$a = 10.8670$ (4) Å

$b = 15.2671$ (6) Å

$c = 19.1569$ (8) Å

$\beta = 97.463$ (2)°

$V = 3151.4$ (2) Å³

$Z = 4$

$F(000) = 1304$

$D_x = 1.327$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 9748 reflections

$\theta = 2.3\text{--}27.0^\circ$

$\mu = 0.26$ mm⁻¹

$T = 296$ K

Block, colourless

$0.31 \times 0.22 \times 0.16$ mm

Data collection

Bruker SMART APEXII CCD
diffractometer

Radiation source: microfocus sealed X-ray tube,
Incoatec I μ s

Mirror optics monochromator

Detector resolution: 7.9 pixels mm⁻¹

ω and ϕ scans

Absorption correction: multi-scan
(SADABS; Krause *et al.*, 2015)

$T_{\min} = 0.485$, $T_{\max} = 0.746$

84575 measured reflections

6190 independent reflections

4405 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.063$

$\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 1.7^\circ$

$h = -13 \rightarrow 13$

$k = -18 \rightarrow 18$

$l = -23 \rightarrow 23$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.074$

$wR(F^2) = 0.239$

$S = 1.08$

6190 reflections

419 parameters

59 restraints

Primary atom site location: dual

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.1098P)^2 + 1.8679P]$

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

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

$\Delta\rho_{\max} = 0.71$ e Å⁻³

$\Delta\rho_{\min} = -0.60$ e Å⁻³

Special details

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ($\text{\AA}^2$)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
F1	0.6616 (2)	0.83522 (16)	0.26782 (10)	0.0953 (7)	
O1	0.7001 (3)	0.93283 (16)	0.55386 (12)	0.0889 (7)	
O2	0.6515 (2)	0.66060 (13)	0.52455 (10)	0.0754 (6)	
N1	0.6603 (3)	0.53973 (16)	0.46177 (14)	0.0874 (9)	
H1A	0.664961	0.509947	0.424097	0.105*	
H1B	0.655578	0.513218	0.500938	0.105*	
N2	0.6820 (4)	0.60700 (19)	0.28440 (15)	0.0938 (10)	
C1	0.6117 (3)	0.9631 (2)	0.43838 (18)	0.0780 (9)	
H1C	0.672777	0.970770	0.406908	0.117*	
H1D	0.592906	1.018817	0.457700	0.117*	
H1E	0.537620	0.938604	0.413090	0.117*	
C2	0.6612 (3)	0.90264 (18)	0.49666 (15)	0.0590 (7)	
C3	0.6610 (2)	0.80710 (17)	0.48275 (13)	0.0512 (6)	
C4	0.6532 (3)	0.74986 (18)	0.53509 (14)	0.0592 (7)	
C5	0.6604 (3)	0.62692 (18)	0.45944 (14)	0.0616 (7)	
C6	0.6671 (3)	0.67926 (16)	0.40317 (13)	0.0524 (6)	
C7	0.6665 (2)	0.77841 (15)	0.40747 (13)	0.0487 (6)	
H7	0.591720	0.799862	0.378372	0.058*	
C8	0.6404 (4)	0.7660 (2)	0.61059 (16)	0.0832 (10)	
H8A	0.720099	0.760602	0.638425	0.125*	
H8B	0.584349	0.723771	0.626143	0.125*	
H8C	0.608424	0.823891	0.615798	0.125*	
C9	0.6762 (3)	0.63938 (18)	0.33777 (15)	0.0617 (7)	
C10	0.7795 (2)	0.81656 (16)	0.37852 (13)	0.0518 (6)	
C11	0.7726 (3)	0.84350 (19)	0.30958 (16)	0.0651 (7)	
C12	0.8707 (4)	0.8794 (2)	0.2811 (2)	0.0873 (11)	
H12	0.861562	0.896505	0.234132	0.105*	
C13	0.9811 (4)	0.8896 (3)	0.3224 (3)	0.0961 (12)	
H13	1.048277	0.913937	0.303786	0.115*	
C14	0.9936 (3)	0.8638 (3)	0.3919 (2)	0.0920 (11)	
H14	1.068962	0.871236	0.420324	0.110*	
C15	0.8935 (3)	0.8266 (2)	0.41938 (18)	0.0727 (8)	
H15	0.903207	0.808215	0.466051	0.087*	
F2	0.3253 (2)	0.65517 (16)	0.13286 (11)	0.1017 (7)	
O4	0.30673 (19)	0.84069 (12)	0.38524 (9)	0.0618 (5)	
N3	0.3590 (3)	0.89205 (18)	0.14758 (15)	0.0865 (9)	
N4	0.3104 (3)	0.96089 (15)	0.32094 (13)	0.0688 (7)	
H4A	0.316273	0.990322	0.283234	0.083*	
H4B	0.300445	0.987702	0.359285	0.083*	
C16	0.3714 (4)	0.5364 (2)	0.3085 (2)	0.0996 (13)	
H16A	0.297518	0.526520	0.276201	0.149*	
H16B	0.399092	0.482090	0.330364	0.149*	
H16C	0.435043	0.559851	0.283542	0.149*	
C17	0.3444 (4)	0.5994 (2)	0.3631 (2)	0.0860 (10)	
C18	0.3294 (2)	0.69418 (17)	0.34666 (15)	0.0562 (6)	

C19	0.3188 (3)	0.75174 (18)	0.39809 (15)	0.0584 (7)	
C20	0.3169 (2)	0.87347 (16)	0.31995 (13)	0.0516 (6)	
C21	0.3321 (2)	0.82051 (16)	0.26544 (13)	0.0497 (6)	
C22	0.3258 (2)	0.72140 (15)	0.27028 (14)	0.0516 (6)	
H22	0.398628	0.696739	0.252122	0.062*	
C23	0.3164 (4)	0.7376 (3)	0.47495 (17)	0.0837 (10)	
H23A	0.238184	0.712484	0.482371	0.100*	
H23B	0.327234	0.792666	0.499163	0.100*	
H23C	0.382327	0.698536	0.492790	0.100*	
C24	0.3476 (3)	0.85970 (17)	0.20039 (15)	0.0597 (7)	
C25	0.2090 (3)	0.68716 (16)	0.22507 (14)	0.0550 (6)	
C26	0.2138 (3)	0.6560 (2)	0.15778 (17)	0.0715 (8)	
C27	0.1119 (5)	0.6249 (3)	0.1150 (2)	0.0957 (12)	
H27	0.119165	0.604588	0.069968	0.115*	
C28	−0.0002 (4)	0.6245 (3)	0.1398 (3)	0.1032 (14)	
H28	−0.069957	0.603421	0.111521	0.124*	
C29	−0.0109 (3)	0.6550 (3)	0.2067 (2)	0.0913 (11)	
H29	−0.087412	0.654609	0.223550	0.110*	
C30	0.0939 (3)	0.6862 (2)	0.24847 (18)	0.0711 (8)	
H30	0.086499	0.707098	0.293346	0.085*	
O3A	0.3692 (6)	0.5749 (4)	0.4238 (4)	0.0913 (13)	0.5
O3	0.2981 (7)	0.5700 (4)	0.4175 (4)	0.107 (2)	0.5
Cl2	−0.0289 (13)	0.5107 (10)	0.3527 (10)	0.183 (2)	0.3333
Cl3	−0.0044 (12)	0.3416 (6)	0.4020 (5)	0.1845 (17)	0.3333
C1A	0.0493 (19)	0.4488 (8)	0.4202 (10)	0.182 (2)	0.3333
H1AA	0.029605	0.468328	0.465648	0.219*	0.3333
H1AB	0.138398	0.452374	0.419922	0.219*	0.3333
Cl3A	0.0105 (8)	0.3565 (4)	0.4722 (4)	0.1845 (17)	0.3333
C1B	−0.045 (2)	0.3910 (15)	0.3879 (10)	0.182 (2)	0.3333
H1BA	0.002048	0.358295	0.356847	0.219*	0.3333
H1BB	−0.129373	0.369594	0.378881	0.219*	0.3333
Cl2A	−0.0494 (14)	0.4896 (11)	0.3592 (10)	0.183 (2)	0.3333
Cl1	0.0398 (7)	0.5174 (6)	0.3831 (4)	0.183 (2)	0.3333
Cl4	0.0405 (12)	0.3494 (5)	0.4256 (5)	0.1845 (17)	0.3333
C1C	−0.0492 (18)	0.4449 (9)	0.4279 (13)	0.182 (2)	0.3333
H1CA	−0.054082	0.464065	0.475786	0.219*	0.3333
H1CB	−0.132263	0.437309	0.403269	0.219*	0.3333

Atomic displacement parameters ($\text{\AA}^2$)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
F1	0.1047 (15)	0.1224 (18)	0.0571 (11)	−0.0045 (12)	0.0040 (10)	0.0216 (10)
O1	0.123 (2)	0.0682 (14)	0.0725 (15)	0.0101 (13)	0.0004 (13)	−0.0274 (11)
O2	0.1291 (19)	0.0538 (11)	0.0446 (10)	−0.0029 (11)	0.0166 (11)	0.0041 (8)
N1	0.156 (3)	0.0467 (14)	0.0599 (15)	−0.0021 (15)	0.0174 (16)	0.0057 (11)
N2	0.159 (3)	0.0625 (16)	0.0641 (17)	−0.0180 (18)	0.0316 (18)	−0.0145 (14)
C1	0.098 (2)	0.0503 (16)	0.086 (2)	0.0146 (15)	0.0117 (18)	−0.0010 (15)
C2	0.0612 (15)	0.0557 (15)	0.0615 (17)	0.0043 (12)	0.0134 (13)	−0.0107 (13)

C3	0.0566 (14)	0.0501 (14)	0.0474 (13)	0.0034 (11)	0.0088 (11)	−0.0045 (11)
C4	0.0758 (18)	0.0559 (15)	0.0466 (14)	0.0005 (13)	0.0101 (12)	−0.0029 (11)
C5	0.087 (2)	0.0474 (14)	0.0506 (15)	0.0006 (13)	0.0084 (13)	0.0000 (11)
C6	0.0701 (16)	0.0441 (13)	0.0430 (13)	−0.0012 (11)	0.0075 (11)	−0.0013 (10)
C7	0.0588 (14)	0.0431 (13)	0.0435 (13)	0.0024 (10)	0.0046 (10)	0.0008 (10)
C8	0.121 (3)	0.083 (2)	0.0489 (17)	−0.007 (2)	0.0238 (17)	−0.0078 (15)
C9	0.087 (2)	0.0458 (14)	0.0530 (16)	−0.0081 (13)	0.0119 (13)	−0.0026 (12)
C10	0.0630 (15)	0.0409 (12)	0.0526 (14)	0.0041 (11)	0.0120 (11)	−0.0025 (10)
C11	0.0790 (19)	0.0589 (16)	0.0594 (17)	0.0044 (14)	0.0160 (14)	0.0053 (13)
C12	0.104 (3)	0.081 (2)	0.085 (2)	0.010 (2)	0.045 (2)	0.0221 (19)
C13	0.091 (3)	0.082 (2)	0.126 (4)	−0.004 (2)	0.055 (3)	0.006 (2)
C14	0.0598 (19)	0.107 (3)	0.112 (3)	−0.0048 (18)	0.0196 (19)	−0.012 (2)
C15	0.0657 (18)	0.084 (2)	0.0692 (19)	0.0023 (15)	0.0119 (14)	−0.0028 (16)
F2	0.1180 (17)	0.1154 (18)	0.0760 (13)	−0.0059 (13)	0.0281 (12)	−0.0268 (12)
O4	0.0872 (13)	0.0507 (10)	0.0483 (10)	0.0004 (9)	0.0120 (9)	0.0018 (8)
N3	0.138 (3)	0.0617 (16)	0.0629 (17)	−0.0072 (16)	0.0236 (16)	0.0076 (13)
N4	0.109 (2)	0.0436 (12)	0.0536 (13)	−0.0007 (12)	0.0116 (13)	−0.0040 (10)
C16	0.113 (3)	0.0431 (17)	0.135 (3)	0.0125 (17)	−0.012 (2)	0.0041 (19)
C17	0.110 (2)	0.0546 (16)	0.088 (2)	−0.0015 (16)	−0.0097 (19)	0.0247 (15)
C18	0.0607 (15)	0.0455 (13)	0.0601 (16)	−0.0011 (11)	−0.0005 (12)	0.0102 (12)
C19	0.0614 (15)	0.0557 (15)	0.0562 (16)	−0.0037 (12)	0.0001 (12)	0.0113 (12)
C20	0.0607 (15)	0.0443 (13)	0.0492 (14)	−0.0031 (11)	0.0049 (11)	0.0016 (10)
C21	0.0600 (14)	0.0405 (12)	0.0487 (13)	−0.0032 (10)	0.0073 (11)	0.0009 (10)
C22	0.0581 (14)	0.0394 (12)	0.0570 (15)	0.0029 (10)	0.0063 (11)	0.0013 (10)
C23	0.105 (3)	0.087 (2)	0.0570 (18)	−0.0057 (19)	0.0022 (17)	0.0175 (16)
C24	0.0834 (19)	0.0422 (13)	0.0542 (16)	−0.0032 (12)	0.0119 (13)	−0.0025 (12)
C25	0.0666 (16)	0.0366 (12)	0.0602 (16)	0.0000 (11)	0.0021 (12)	0.0018 (11)
C26	0.090 (2)	0.0567 (17)	0.0664 (19)	−0.0031 (15)	0.0030 (16)	−0.0045 (14)
C27	0.125 (3)	0.083 (2)	0.072 (2)	−0.013 (2)	−0.016 (2)	−0.0143 (18)
C28	0.106 (3)	0.087 (3)	0.103 (3)	−0.025 (2)	−0.040 (3)	0.007 (2)
C29	0.067 (2)	0.094 (3)	0.108 (3)	−0.0108 (18)	−0.0091 (19)	0.015 (2)
C30	0.0692 (19)	0.0662 (18)	0.076 (2)	−0.0005 (15)	0.0018 (15)	0.0026 (15)
O3A	0.116 (3)	0.060 (2)	0.092 (2)	−0.001 (2)	−0.008 (2)	0.0293 (18)
O3	0.132 (5)	0.073 (3)	0.108 (4)	−0.012 (4)	−0.008 (4)	0.044 (3)
Cl2	0.183 (5)	0.198 (5)	0.172 (4)	0.039 (4)	0.029 (3)	0.042 (3)
Cl3	0.249 (6)	0.148 (2)	0.152 (5)	0.028 (3)	0.007 (4)	−0.019 (3)
C1A	0.181 (5)	0.196 (5)	0.171 (4)	0.040 (4)	0.026 (4)	0.044 (4)
Cl3A	0.249 (6)	0.148 (2)	0.152 (5)	0.028 (3)	0.007 (4)	−0.019 (3)
C1B	0.181 (5)	0.196 (5)	0.171 (4)	0.040 (4)	0.026 (4)	0.044 (4)
Cl2A	0.183 (5)	0.198 (5)	0.172 (4)	0.039 (4)	0.029 (3)	0.042 (3)
Cl1	0.183 (5)	0.198 (5)	0.172 (4)	0.039 (4)	0.029 (3)	0.042 (3)
Cl4	0.249 (6)	0.148 (2)	0.152 (5)	0.028 (3)	0.007 (4)	−0.019 (3)
C1C	0.181 (5)	0.196 (5)	0.171 (4)	0.040 (4)	0.026 (4)	0.044 (4)

Geometric parameters (Å, °)

F1—C11	1.364 (4)	C16—H16A	0.9600
O1—C2	1.213 (3)	C16—H16B	0.9600

O2—C4	1.377 (3)	C16—H16C	0.9600
O2—C5	1.364 (3)	C16—C17	1.477 (6)
N1—H1A	0.8600	C17—C18	1.486 (4)
N1—H1B	0.8600	C17—O3A	1.217 (7)
N1—C5	1.332 (4)	C17—O3	1.293 (8)
N2—C9	1.145 (4)	C18—C19	1.336 (4)
C1—H1C	0.9600	C18—C22	1.517 (4)
C1—H1D	0.9600	C19—C23	1.492 (4)
C1—H1E	0.9600	C20—C21	1.348 (4)
C1—C2	1.494 (4)	C21—C22	1.518 (3)
C2—C3	1.483 (4)	C21—C24	1.412 (4)
C3—C4	1.341 (4)	C22—H22	0.9800
C3—C7	1.516 (3)	C22—C25	1.533 (4)
C4—C8	1.491 (4)	C23—H23A	0.9600
C5—C6	1.351 (4)	C23—H23B	0.9600
C6—C7	1.516 (3)	C23—H23C	0.9600
C6—C9	1.408 (4)	C25—C26	1.381 (4)
C7—H7	0.9800	C25—C30	1.382 (4)
C7—C10	1.527 (4)	C26—C27	1.374 (5)
C8—H8A	0.9600	C27—H27	0.9300
C8—H8B	0.9600	C27—C28	1.363 (6)
C8—H8C	0.9600	C28—H28	0.9300
C10—C11	1.376 (4)	C28—C29	1.383 (6)
C10—C15	1.385 (4)	C29—H29	0.9300
C11—C12	1.372 (5)	C29—C30	1.389 (5)
C12—H12	0.9300	C30—H30	0.9300
C12—C13	1.358 (6)	Cl2—C1A	1.734 (8)
C13—H13	0.9300	Cl3—C1A	1.757 (8)
C13—C14	1.377 (6)	C1A—H1AA	0.9700
C14—H14	0.9300	C1A—H1AB	0.9700
C14—C15	1.390 (5)	Cl3A—C1B	1.730 (16)
C15—H15	0.9300	C1B—H1BA	0.9700
F2—C26	1.359 (4)	C1B—H1BB	0.9700
O4—C19	1.383 (3)	C1B—Cl2A	1.601 (18)
O4—C20	1.365 (3)	Cl1—C1C	1.765 (9)
N3—C24	1.147 (4)	Cl4—C1C	1.758 (9)
N4—H4A	0.8600	C1C—H1CA	0.9700
N4—H4B	0.8600	C1C—H1CB	0.9700
N4—C20	1.337 (3)		
C5—O2—C4	120.4 (2)	C16—C17—C18	120.6 (3)
H1A—N1—H1B	120.0	O3A—C17—C16	115.9 (5)
C5—N1—H1A	120.0	O3A—C17—C18	120.4 (5)
C5—N1—H1B	120.0	O3—C17—C16	118.6 (4)
H1C—C1—H1D	109.5	O3—C17—C18	117.7 (5)
H1C—C1—H1E	109.5	C17—C18—C22	117.4 (3)
H1D—C1—H1E	109.5	C19—C18—C17	120.1 (3)
C2—C1—H1C	109.5	C19—C18—C22	122.5 (2)

C2—C1—H1D	109.5	O4—C19—C23	107.7 (3)
C2—C1—H1E	109.5	C18—C19—O4	122.0 (2)
O1—C2—C1	119.3 (3)	C18—C19—C23	130.3 (3)
O1—C2—C3	121.9 (3)	N4—C20—O4	110.0 (2)
C3—C2—C1	118.8 (2)	N4—C20—C21	128.5 (2)
C2—C3—C7	117.1 (2)	C21—C20—O4	121.5 (2)
C4—C3—C2	120.4 (2)	C20—C21—C22	122.7 (2)
C4—C3—C7	122.5 (2)	C20—C21—C24	118.0 (2)
O2—C4—C8	107.7 (2)	C24—C21—C22	119.2 (2)
C3—C4—O2	122.4 (2)	C18—C22—C21	109.6 (2)
C3—C4—C8	129.8 (3)	C18—C22—H22	108.4
N1—C5—O2	110.2 (2)	C18—C22—C25	111.6 (2)
N1—C5—C6	128.2 (3)	C21—C22—H22	108.4
C6—C5—O2	121.6 (2)	C21—C22—C25	110.2 (2)
C5—C6—C7	123.1 (2)	C25—C22—H22	108.4
C5—C6—C9	118.1 (2)	C19—C23—H23A	109.5
C9—C6—C7	118.8 (2)	C19—C23—H23B	109.5
C3—C7—C6	110.0 (2)	C19—C23—H23C	109.5
C3—C7—H7	108.2	H23A—C23—H23B	109.5
C3—C7—C10	111.6 (2)	H23A—C23—H23C	109.5
C6—C7—H7	108.2	H23B—C23—H23C	109.5
C6—C7—C10	110.6 (2)	N3—C24—C21	179.3 (4)
C10—C7—H7	108.2	C26—C25—C22	121.0 (3)
C4—C8—H8A	109.5	C26—C25—C30	116.4 (3)
C4—C8—H8B	109.5	C30—C25—C22	122.5 (3)
C4—C8—H8C	109.5	F2—C26—C25	118.3 (3)
H8A—C8—H8B	109.5	F2—C26—C27	118.3 (3)
H8A—C8—H8C	109.5	C27—C26—C25	123.4 (4)
H8B—C8—H8C	109.5	C26—C27—H27	120.6
N2—C9—C6	179.1 (4)	C28—C27—C26	118.8 (4)
C11—C10—C7	121.1 (2)	C28—C27—H27	120.6
C11—C10—C15	116.2 (3)	C27—C28—H28	119.7
C15—C10—C7	122.6 (2)	C27—C28—C29	120.5 (4)
F1—C11—C10	117.9 (3)	C29—C28—H28	119.7
F1—C11—C12	118.5 (3)	C28—C29—H29	120.4
C12—C11—C10	123.6 (3)	C28—C29—C30	119.2 (4)
C11—C12—H12	120.4	C30—C29—H29	120.4
C13—C12—C11	119.2 (4)	C25—C30—C29	121.7 (3)
C13—C12—H12	120.4	C25—C30—H30	119.1
C12—C13—H13	120.1	C29—C30—H30	119.1
C12—C13—C14	119.9 (3)	Cl2—C1A—Cl3	104.0 (8)
C14—C13—H13	120.1	Cl2—C1A—H1AA	111.0
C13—C14—H14	120.0	Cl2—C1A—H1AB	111.0
C13—C14—C15	120.0 (4)	Cl3—C1A—H1AA	111.0
C15—C14—H14	120.0	Cl3—C1A—H1AB	111.0
C10—C15—C14	121.2 (3)	H1AA—C1A—H1AB	109.0
C10—C15—H15	119.4	Cl3A—C1B—H1BA	105.6
C14—C15—H15	119.4	Cl3A—C1B—H1BB	105.6

C20—O4—C19	120.4 (2)	H1BA—C1B—H1BB	106.1
H4A—N4—H4B	120.0	Cl2A—C1B—Cl3A	126.8 (17)
C20—N4—H4A	120.0	Cl2A—C1B—H1BA	105.6
C20—N4—H4B	120.0	Cl2A—C1B—H1BB	105.6
H16A—C16—H16B	109.5	Cl1—C1C—H1CA	111.8
H16A—C16—H16C	109.5	Cl1—C1C—H1CB	111.8
H16B—C16—H16C	109.5	Cl4—C1C—Cl1	99.6 (7)
C17—C16—H16A	109.5	Cl4—C1C—H1CA	111.8
C17—C16—H16B	109.5	Cl4—C1C—H1CB	111.8
C17—C16—H16C	109.5	H1CA—C1C—H1CB	109.6
F1—C11—C12—C13	178.8 (3)	O4—C20—C21—C22	6.5 (4)
O1—C2—C3—C4	28.1 (4)	O4—C20—C21—C24	−177.1 (2)
O1—C2—C3—C7	−153.5 (3)	N4—C20—C21—C22	−174.2 (3)
O2—C5—C6—C7	0.5 (5)	N4—C20—C21—C24	2.2 (4)
O2—C5—C6—C9	179.9 (3)	C16—C17—C18—C19	−172.5 (3)
N1—C5—C6—C7	−180.0 (3)	C16—C17—C18—C22	7.9 (5)
N1—C5—C6—C9	−0.5 (5)	C17—C18—C19—O4	179.0 (3)
C1—C2—C3—C4	−151.8 (3)	C17—C18—C19—C23	−1.5 (5)
C1—C2—C3—C7	26.7 (4)	C17—C18—C22—C21	−170.5 (3)
C2—C3—C4—O2	179.4 (3)	C17—C18—C22—C25	67.1 (3)
C2—C3—C4—C8	1.9 (5)	C18—C22—C25—C26	−140.8 (3)
C2—C3—C7—C6	179.3 (2)	C18—C22—C25—C30	39.8 (3)
C2—C3—C7—C10	56.2 (3)	C19—O4—C20—N4	−175.8 (2)
C3—C7—C10—C11	−142.6 (2)	C19—O4—C20—C21	3.6 (4)
C3—C7—C10—C15	37.0 (3)	C19—C18—C22—C21	10.0 (4)
C4—O2—C5—N1	178.3 (3)	C19—C18—C22—C25	−112.4 (3)
C4—O2—C5—C6	−2.1 (4)	C20—O4—C19—C18	−6.2 (4)
C4—C3—C7—C6	−2.3 (4)	C20—O4—C19—C23	174.2 (3)
C4—C3—C7—C10	−125.4 (3)	C20—C21—C22—C18	−12.5 (3)
C5—O2—C4—C3	1.3 (4)	C20—C21—C22—C25	110.7 (3)
C5—O2—C4—C8	179.3 (3)	C21—C22—C25—C26	97.1 (3)
C5—C6—C7—C3	1.6 (4)	C21—C22—C25—C30	−82.3 (3)
C5—C6—C7—C10	125.3 (3)	C22—C18—C19—O4	−1.6 (4)
C6—C7—C10—C11	94.6 (3)	C22—C18—C19—C23	178.0 (3)
C6—C7—C10—C15	−85.8 (3)	C22—C25—C26—F2	1.0 (4)
C7—C3—C4—O2	1.0 (4)	C22—C25—C26—C27	−179.8 (3)
C7—C3—C4—C8	−176.4 (3)	C22—C25—C30—C29	180.0 (3)
C7—C10—C11—F1	0.1 (4)	C24—C21—C22—C18	171.2 (2)
C7—C10—C11—C12	179.0 (3)	C24—C21—C22—C25	−65.6 (3)
C7—C10—C15—C14	−178.3 (3)	C25—C26—C27—C28	0.0 (6)
C9—C6—C7—C3	−177.9 (2)	C26—C25—C30—C29	0.5 (4)
C9—C6—C7—C10	−54.2 (3)	C26—C27—C28—C29	0.2 (6)
C10—C11—C12—C13	−0.2 (5)	C27—C28—C29—C30	0.0 (6)
C11—C10—C15—C14	1.4 (5)	C28—C29—C30—C25	−0.4 (5)
C11—C12—C13—C14	0.2 (6)	C30—C25—C26—F2	−179.5 (3)
C12—C13—C14—C15	0.6 (6)	C30—C25—C26—C27	−0.3 (5)
C13—C14—C15—C10	−1.4 (6)	O3A—C17—C18—C19	−13.1 (6)

C15—C10—C11—F1	−179.5 (3)	O3A—C17—C18—C22	167.4 (5)
C15—C10—C11—C12	−0.6 (4)	O3—C17—C18—C19	27.6 (6)
F2—C26—C27—C28	179.2 (3)	O3—C17—C18—C22	−151.9 (4)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H $\cdots$ <i>A</i>	<i>D</i> —H	H $\cdots$ <i>A</i>	<i>D</i> $\cdots$ <i>A</i>	<i>D</i> —H $\cdots$ <i>A</i>
N1—H1 <i>A</i> $\cdots$ N3 ⁱ	0.86	2.26	3.066 (4)	156
N1—H1 <i>B</i> $\cdots$ O3 ⁱⁱ	0.86	2.03	2.844 (8)	159
N1—H1 <i>B</i> $\cdots$ O3 <i>A</i> ⁱⁱ	0.86	2.02	2.855 (7)	165
N4—H4 <i>A</i> $\cdots$ N2 ⁱⁱⁱ	0.86	2.20	3.016 (4)	157
N4—H4 <i>B</i> $\cdots$ O1 ^{iv}	0.86	2.06	2.910 (3)	170

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