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Synthesis, crystal structure and Hirshfeld surface analysis of *tert*-butyl 4-[4-(difluoromethoxy)phenyl]-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

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The 1,4-dihydropyridine ring of the title compound, $C_{24}H_{29}F_2NO_4$, adopts a distorted boat conformation, while the cyclohexene ring is in an almost twistboat conformation. In the crystal, $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds as well as $C-H\cdots \pi$ interactions connect molecules, forming layers parallel to the (100) plane. These layers are linked by van der Waals forces and $C-H\cdots F$ interactions, which consolidate the crystal structure. Hirshfeld surface analysis shows the major contributions to the crystal packing are from $H\cdots H$ (54.1%), $F\cdots H/H\cdots F$ (16.9%), $O\cdots H/H\cdots O$ (15.4%) and $C\cdots H/H\cdots C$ (12.6%) contacts.

1. Chemical context

Inflammation is the natural and basic response of an organism to signals from tissue damage or pathogenic infections. In this way, the integrity of the organism is preserved. Chronic diseases that cause death and economic losses in the world are constantly increasing. It has been found that chronic diseases occur through inflammation-mediated mechanisms. In recent years, it has been proven that cardiovascular diseases, cancer, diabetes mellitus, chronic kidney disease, non-alcoholic fatty liver disease, autoimmune and neurodegenerative diseases are caused by inflammation. In this context, managing inflammatory mediators and inflammatory processes can be a treatment method for many chronic diseases (Furman *et al.*, 2019; Tu *et al.*, 2022).

Chronic or local inflammation first occurs with the activation of immune system cells such as cytokines, proteases, chemokines, oxygen-independent radicals, which generate signals from damaged cells or pathogens that are dangerous to the tissue. The immune system cells released in the circulatory system increase the pro-inflammatory response and reach the infected tissue area, but if this response is insufficient or excessive, the balance of the immune system is disturbed. This imbalance causes an excessive amount of distress signals and local or systemic tissue damage. This defect in the immune response causes the inflammation to change from acute to chronic, and the disease progresses and results in death. A better understanding of inflammation and its processes enables the discovery of new and effective therapeutic ways to



Figure 1 Structure of nifedipine.

target and regulate inflammation. Drug therapy is widely used for the treatment of inflammation. Therefore, there is a need for new molecules that are more active and have minimal side effects (Tu et al., 2022). The 1,4-DHP ring, which is a partially saturated derivative of the pyridine ring, is involved in the structure of many bioactive compounds. Nifedipine, which has a 1,4-DHP structure, was introduced as an antihypertensive treatment about 50 years ago (Fig. 1). The therapeutic success of nifedipine has led to the preparation of analogue derivatives. In this ongoing process, various compounds such as amlodipine and benidipine, which have a 1,4-DHP structure, are used as antihypertensives. Studies have shown that the 1,4-DHP ring has various activities such as neuroprotective, antiplatelet, anti-ischemic, anti-Alzheimer's, antituberculer, antiulcer and anticancer (Khot et al., 2021; Abdelwahab et al., 2022).

The hexahydroquinoline ring system is obtained by condensing 1,4-DHP with cyclohexane. This ring system also has a variety of pharmacological activities such as calcium channel antagonist, anticancer, antimicrobial, anti-Alzheimer's. In current studies, 1,4-DHP derivatives and condensed analogues were found to be effective inflammation mediators of chronic inflammation in addition to their various biological activities.



In this study, the title compound, *tert*-butyl 4-[4-(difluoromethoxy)phenyl]-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate was obtained by using modified Hantzsch one-pot synthesis (Ghosh *et al.*, 2013). The reaction of 4-difluoromethoxybenzaldehyde with 5,5-dimethylcyclohexane-1,3-dione and *tert*-butyl acetoacetate gives the target compound in methanol in the presence of ammonium acetate as nitrogen source (Çetin *et al.*, 2022). The structure of the compound was elucidated by IR, ¹H-NMR, ¹³C-NMR and HRMS analysis. X-ray analysis was undertaken to determine the crystal structure. Biological activity tests will be conducted in independent studies to determine the inhibition potential of inflammation mediators.

2. Structural commentary

As seen in Fig. 2, the 1,4-dihydropyridine ring (N1/C1/C6–C9) of the title compound adopts a distorted boat conformation [puckering parameters (Cremer & Pople, 1975) are $Q_T = 0.2940$ (18) Å, $\theta = 72.1$ (4)° and $\varphi = 182.9$ (4)°], while the cyclohexene ring (C1–C6) has an almost twist-boat conformation [puckering parameters are $Q_T = 0.4617$ (19) Å, $\theta = 124.5$ (2)° and $\varphi = 313.8$ (3)°]. The 4-[4-(diffuoromethoxy]phenyl ring (C18–C23) makes a dihedral angle of 89.88 (7)° with the mean plane of the quinoline ring system [N1/C1–C9; maximum deviation = 0.358 (2) Å for C4]. The geometrical parameters of the title compound are in agreement with those reported for similar compounds in the *Database survey* section.

3. Supramolecular features and Hirshfeld surface analysis

The molecules in the crystal are connected by $N-H\cdots O$ and $C-H\cdots O$ hydrogen bonds, as well as $C-H\cdots \pi$ interactions, resulting in the formation of layers parallel to the (100) plane (see Table 1; Figs. 3 and 4). These layers are linked by van der





View of the title molecule. Displacement ellipsoids are drawn at the 30% probability level.

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

Cg3 is the centroid of the C18–C23 ring.

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1-H1N\cdotsO1^{i}$	0.91 (2)	1.96 (2)	2.866 (2)	176.6 (18)
$C12 - H12A \cdots O2$	0.98	2.25	2.800(2)	114
$C16-H16A\cdots O2$	0.98	2.36	2.938 (2)	117
C17−H17C···O2	0.98	2.37	2.958 (3)	118
$C20-H20A\cdots F1$	0.95	2.46	2.989 (2)	115
$C24 - H24A \cdots O1^{ii}$	1.00	2.35	3.230 (2)	147
$C2-H2A\cdots Cg3^{iii}$	0.99	2.74	3.6959 (19)	162

Symmetry codes: (i) $x, -y + \frac{3}{2}, z - \frac{1}{2}$; (ii) x, y - 1, z; (iii) $x, -y + \frac{1}{2}, z - \frac{3}{2}$.

Table 2

Summary of short interatomic contacts (Å) in the title compound.

$H11C \cdots H10A$	2.49	$2 - x, -\frac{1}{2} + y, \frac{1}{2} - z$
$F2 \cdot \cdot \cdot H19A$	2.51	$x, \frac{1}{2} - y, \frac{1}{2} + z$
$O1 \cdot \cdot \cdot H24A$	2.35	$x, \tilde{1} + y, \tilde{z}$
$O1 \cdot \cdot \cdot H1N$	1.96	$x, \frac{1}{2} - y, -\frac{1}{2} + z$
$H12A \cdots O2$	2.61	1 - x, 1 - y, -z
$H15A \cdots H12A$	2.40	$1 - x, \frac{1}{2} + y, \frac{1}{2} - z$
$H22A \cdot \cdot \cdot H16B$	2.38	$1 - x, \bar{1} - y, \bar{1} - z$

Waals forces and $C-H\cdots$ F interactions, which consolidate the crystal structure (Fig. 5).

The Hirshfeld surfaces and their corresponding twodimensional fingerprint plots were calculated using the *Crystal Explorer 17.5* (Spackman *et al.*, 2021) software package. The d_{norm} surfaces are mapped over a fixed colour scale from -0.5814 (red) to +1.6362 (blue) a.u. Red spots on the surface correspond to $N \cdots H/H \cdots N$ and $O \cdots H/H \cdots O$ interactions (Tables 1 and 2; Fig. 6*a*,*b*).

Fingerprint plots of the most important non-covalent interactions for the title compound are shown in Fig. 7. The major contributions to the crystal packing are from $H \cdots H$ (54.1%), $F \cdots H/H \cdots F$ (16.9%), $O \cdots H/H \cdots O$ (15.4%) and



Figure 3

A view of the molecular packing of the title compound along the *a* axis by the N-H···O, C-H···O hydrogen bonds and C-H··· π interactions (dashed lines).



View of the molecular packing along [010]. Hydrogen bonds are shown as dashed lines.

C···H/H···C (12.6) contacts. N···H/H···N (0.5%), F···N/ N···F (0.3%) and F···F (0.2%) contacts, which contribute less than 1%, are not shown in Fig. 7.



View of the molecular packing along [001]. Hydrogen bonds are shown as dashed lines.



Figure 6

(a) Front and (b) back views of the three-dimensional Hirshfeld surface for the title compound.



Figure 7

The two-dimensional fingerprint plots for the title compound showing (*a*) all interactions, and delineated into (*b*) $H \cdots H$, (*c*) $F \cdots H/H \cdots F$, (*d*) $O \cdots H/H \cdots O$ and (*e*) $C \cdots H/H \cdots C$ interactions. The d_i and d_e values are the closest internal and external distances (in Å) from given points on the Hirshfeld surface.

4. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.42, update of September 2021; Groom *et al.*, 2016) for similar structures with the 1,4,5,6,7,8-hexahydroquinoline group showed that the nine most closely related to the title compound are WEZJUK (Yıldırım *et al.*, 2023), ECUCUE (Yıldırım *et al.*, 2022), LOQCAX (Steiger *et al.*, 2014), NEQMON (Öztürk Yıldırım *et al.*, 2013), PECPUK (Gündüz *et al.*, 2012), IMEJOA (Linden *et al.*, 2011), PUGCIE (Mookiah *et al.*, 2009), UCOLOO (Linden *et al.*, 2006) and DAYJET (Linden *et al.*, 2005). In all these compounds, molecules are linked by N–H···O hydrogen bonds. Furthermore, C–H···O hydrogen bonds in WEZJUK, ECUCUE, NEQMON, IMEJOA and PUGCIE and C–H··· π interactions in WEZJUK and ECUCUE were also observed.

5. Synthesis and crystallization

The target compound was synthesized by refluxing 5,5-dimethylcyclohexane-1,3-dione (1 mmol), 4-difluoromethoxy-

Table 3	
Experimental details.	
Crystal data	
Chemical formula	$C_{24}H_{29}F_2NO_4$
M _r	433.48
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	100
a, b, c (Å)	17.6062 (11), 9.7588 (7), 13.1509 (9)
β (°)	95.905 (2)
$V(Å^3)$	2247.5 (3)
Ζ	4
Radiation type	Μο Κα
$\mu \ (\mathrm{mm}^{-1})$	0.10
Crystal size (mm)	$0.26 \times 0.20 \times 0.14$
Data collection	
Diffractometer	Bruker D8 Quest with Photon 2 detector
Absorption correction	Multi-scan (SADABS; Bruker, 2018)
T_{\min}, T_{\max}	0.657, 0.746
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	31708, 4599, 3208
R _{int}	0.111
$(\sin \theta / \lambda)_{\rm max} ({\rm \AA}^{-1})$	0.625
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.046, 0.110, 1.02
No. of reflections	4599
No. of parameters	290
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta \rho_{\rm max}, \Delta \rho_{\rm min} \ ({ m e} \ { m \AA}^{-3})$	0.21, -0.26

Computer programs: *APEX2* and *SAINT* (Bruker, 2018), *SHELXT* (Sheldrick, 2015*a*), *SHELXL* (Sheldrick, 2015*b*), *ORTEP-3 for Windows* (Farrugia, 2012) and *PLATON* (Spek, 2020).

benzaldehyde (1 mmol), *tert*-butylacetoacetate (1 mmol) and ammonium acetate (5 mmol) for 8 h in absolute methanol (10 ml). The reaction mixture was monitored by TLC, and after completion of the reaction was cooled to room temperature. The obtained precipitate was filtered and recrystallized from methanol for further purification. The synthetic route is shown in Fig. 8.

Yellow solid, m.p. 487–488 K; yield: 65.32%. IR (ν , cm⁻¹) 3211 (N–H, stretching), 3080 (C–H stretching, aromatic), 2968 (C–H stretching, aliphatic) 1697 (C=O stretching, ester), 1641 (C=O stretching, ketone). ¹H NMR (DMSO- d_6) δ : 0.84 (3H; s; 7-CH₃), 1.00 (3H; s; 7-CH₃), 1.31 [9H, s, C(CH₃)₃], 1.95–1.99 (2H; d; J = 16 Hz; quinoline H8), 2.13–2.16 (H; d; J = 16.1; quinoline H8), 2.25 (3H; s; 2-CH₃), 2.26–2.30 (H; d; J = 16.95 quinoline H6), 2.37–2.41 (H; d; J = 16.95quinoline H6), 4.78 (1H; s; quinoline H4), 6.99–7.01 (2H, d, J =8.5 Hz Ar–H3), 7.14 (1H; t; J = 74.4 Hz; OCHF₂), 7.17–7.18 (2H, d, J = 10 Ar–H2), 8.99 (1H,s; NH). ¹³C NMR (DMSO-



Figure 8 Synthetic scheme.

*d*₆) δ : 18.7 (2-CH₃), 27.0 (7-CH₃), 28.3 [COOC(CH₃)₃], 29.4 (C-7), 32.0 (C-8), 36.2 (C-4), 50.6 (C-6), 79.2 [COOC(CH₃)₃], 105.4 (C-3), 110.0 (C-4a), 114.8 (C₃'), 116.9, 118.4, 118.9 (OCHF₂), 129.4 (C₂'), 144.5 (C₁'), 145.3 (C-2), 149.3 (C-8a), 150.0 (C₄'), 166.7 [COOC(CH₃)₃], 194.6 (C-5). HRMS (ESI/Q-TOF) *m*/*z*: [*M* + H]⁺ Calculated for C₂₄H₂₉F₂NO₄ 433.2065; found 434.2328 (*M* + H).

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 3. The N-bound H atom was located in a difference Fourier map and refined freely [N1-H1N =0.91 (2) Å]. All C-bound H atoms were positioned geometrically [C-H = 0.95-1.00 Å] and refined using a riding model with $U_{iso}(H) = 1.2$ or 1.5 $U_{eq}(C)$.

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Authors' contributions are as follows. Conceptualization, *RS* and SÖY; methodology, *RS* and EP; investigation, *RS* and SÖY; writing (original draft), EP and MA; writing (review and editing of the manuscript), *RS* and SÖY; crystal data production and validation, RJB and SÖY; visualization, MA; funding acquisition, RJB; resources, AB, RJB and *RS*.

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Synthesis, crystal structure and Hirshfeld surface analysis of *tert*-butyl 4-[4-(di-fluoromethoxy)phenyl]-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydro-quinoline-3-carboxylate

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Computing details

Data collection: *APEX2* (Bruker, 2018); cell refinement: *SAINT* (Bruker, 2018); data reduction: *SAINT* (Bruker, 2018); program(s) used to solve structure: *SHELXT* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL* (Sheldrick, 2015b); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *PLATON* (Spek, 2020).

tert-Butyl 4-[4-(difluoromethoxy)phenyl]-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

Crystal data

 $C_{24}H_{29}F_{2}NO_{4}$ $M_{r} = 433.48$ Monoclinic, $P2_{1}/c$ a = 17.6062 (11) Å b = 9.7588 (7) Å c = 13.1509 (9) Å $\beta = 95.905 (2)^{\circ}$ $V = 2247.5 (3) Å^{3}$ Z = 4

Data collection

Bruker D8 Quest with Photon 2 detector diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Bruker, 2018) $T_{\min} = 0.657, T_{\max} = 0.746$ 31708 measured reflections

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.110$ S = 1.024599 reflections 290 parameters F(000) = 920 $D_x = 1.281 \text{ Mg m}^{-3}$ Mo K\alpha radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 4059 reflections $\theta = 2.3-30.3^{\circ}$ $\mu = 0.10 \text{ mm}^{-1}$ T = 100 KPrism, colorless $0.26 \times 0.20 \times 0.14 \text{ mm}$

4599 independent reflections 3208 reflections with $I > 2\sigma(I)$ $R_{int} = 0.111$ $\theta_{max} = 26.4^\circ, \ \theta_{min} = 2.4^\circ$ $h = -22 \rightarrow 22$ $k = -12 \rightarrow 12$ $l = -16 \rightarrow 16$

0 restraints Primary atom site location: dual Secondary atom site location: difference Fourier map Hydrogen site location: mixed H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0445P)^2 + 0.5785P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta \rho_{\rm max} = 0.21 \text{ e } \text{\AA}^{-3}$ $\Delta \rho_{\rm min} = -0.26 \text{ e } \text{\AA}^{-3}$

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 $(Å^2)$

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$
F1	0.92250 (6)	0.20646 (12)	0.49817 (9)	0.0351 (3)
F2	0.86051 (6)	0.08089 (11)	0.59614 (9)	0.0311 (3)
01	0.78594 (7)	0.91033 (13)	0.37607 (9)	0.0202 (3)
02	0.51155 (7)	0.57880 (16)	0.12973 (10)	0.0338 (4)
03	0.54282 (7)	0.65743 (13)	0.28906 (9)	0.0189 (3)
O4	0.80976 (7)	0.27398 (13)	0.54548 (9)	0.0242 (3)
N1	0.74130 (8)	0.64973 (15)	0.07479 (12)	0.0155 (3)
H1N	0.7556 (11)	0.634 (2)	0.0116 (16)	0.027 (6)*
C1	0.78375 (9)	0.73899 (17)	0.13677 (13)	0.0141 (4)
C2	0.85366 (9)	0.79398 (18)	0.09469 (13)	0.0160 (4)
H2A	0.838042	0.864711	0.042740	0.019*
H2B	0.878600	0.718755	0.060078	0.019*
C3	0.91163 (10)	0.85665 (18)	0.17650 (13)	0.0164 (4)
C4	0.86773 (10)	0.94844 (18)	0.24516 (14)	0.0184 (4)
H4A	0.903915	0.984065	0.301593	0.022*
H4B	0.846852	1.027890	0.204631	0.022*
C5	0.80277 (10)	0.87635 (17)	0.29047 (13)	0.0155 (4)
C6	0.76121 (9)	0.77274 (17)	0.22958 (13)	0.0140 (4)
C7	0.69507 (9)	0.69774 (18)	0.27054 (13)	0.0146 (4)
H7A	0.666791	0.764154	0.310637	0.018*
C8	0.64044 (10)	0.64486 (17)	0.18117 (13)	0.0148 (4)
C9	0.66713 (9)	0.61265 (18)	0.09115 (13)	0.0153 (4)
C10	0.96912 (11)	0.9421 (2)	0.12400 (15)	0.0245 (4)
H10A	1.008339	0.977719	0.175509	0.037*
H10B	0.942644	1.018719	0.087535	0.037*
H10C	0.993191	0.884569	0.075323	0.037*
C11	0.95397 (11)	0.7439 (2)	0.24056 (15)	0.0257 (5)
H11A	0.990465	0.785721	0.292795	0.039*
H11B	0.981403	0.685259	0.196126	0.039*
H11C	0.917130	0.688629	0.273739	0.039*
C12	0.62714 (10)	0.5386 (2)	0.00161 (14)	0.0209 (4)
H12A	0.588198	0.477760	0.025035	0.031*
H12B	0.664249	0.484180	-0.031788	0.031*
H12C	0.602762	0.605142	-0.047102	0.031*
C13	0.55921 (10)	0.62201 (18)	0.19487 (14)	0.0174 (4)
C14	0.46412 (10)	0.64012 (19)	0.31875 (15)	0.0212 (4)

C15	0.47276 (12)	0.6890 (2)	0.42836 (16)	0.0338 (5)
H15A	0.487156	0.786090	0.430522	0.051*
H15B	0.512507	0.635304	0.467906	0.051*
H15C	0.424214	0.677487	0.457741	0.051*
C16	0.44228 (11)	0.4893 (2)	0.31253 (16)	0.0272 (5)
H16A	0.438276	0.459295	0.241063	0.041*
H16B	0.393032	0.476330	0.339800	0.041*
H16C	0.481471	0.435070	0.352732	0.041*
C17	0.40767 (11)	0.7291 (2)	0.25328 (19)	0.0358 (6)
H17A	0.424720	0.824723	0.257522	0.054*
H17B	0.357177	0.721817	0.278004	0.054*
H17C	0.404679	0.698213	0.182077	0.054*
C18	0.72353 (10)	0.57950 (18)	0.34146 (13)	0.0151 (4)
C19	0.77883 (10)	0.48904 (19)	0.31318 (14)	0.0207 (4)
H19A	0.796943	0.499218	0.248000	0.025*
C20	0.80838 (11)	0.38451 (19)	0.37706 (14)	0.0218 (4)
H20A	0.846835	0.325229	0.356664	0.026*
C21	0.78073 (10)	0.36846 (18)	0.47084 (14)	0.0185 (4)
C22	0.72373 (10)	0.45291 (19)	0.49978 (14)	0.0192 (4)
H22A	0.703948	0.439405	0.563564	0.023*
C23	0.69559 (10)	0.55749 (19)	0.43512 (13)	0.0179 (4)
H23A	0.656394	0.615290	0.455307	0.021*
C24	0.85104 (11)	0.16566 (19)	0.51508 (15)	0.0223 (4)
H24A	0.824329	0.118997	0.453797	0.027*

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
F1	0.0253 (6)	0.0380 (7)	0.0430 (8)	-0.0014 (5)	0.0079 (5)	0.0154 (6)
F2	0.0389 (7)	0.0244 (6)	0.0305 (7)	0.0009 (5)	0.0062 (5)	0.0127 (5)
01	0.0250 (7)	0.0224 (7)	0.0140 (7)	-0.0035 (6)	0.0055 (5)	-0.0038 (5)
O2	0.0210 (7)	0.0590 (10)	0.0214 (8)	-0.0140 (7)	0.0026 (6)	-0.0078 (7)
O3	0.0145 (6)	0.0228 (7)	0.0204 (7)	-0.0024 (5)	0.0062 (5)	-0.0030 (6)
O4	0.0342 (8)	0.0217 (7)	0.0169 (7)	0.0053 (6)	0.0039 (6)	0.0042 (6)
N1	0.0168 (8)	0.0190 (8)	0.0113 (8)	-0.0013 (6)	0.0045 (6)	-0.0025 (6)
C1	0.0139 (9)	0.0133 (9)	0.0149 (9)	0.0020 (7)	0.0001 (7)	0.0013 (7)
C2	0.0175 (9)	0.0158 (9)	0.0156 (9)	0.0005 (7)	0.0051 (7)	0.0010 (7)
C3	0.0165 (9)	0.0166 (9)	0.0166 (9)	-0.0011 (7)	0.0035 (7)	-0.0002 (8)
C4	0.0200 (10)	0.0175 (9)	0.0179 (10)	-0.0043 (8)	0.0036 (8)	-0.0018 (8)
C5	0.0173 (9)	0.0146 (9)	0.0147 (9)	0.0031 (7)	0.0014 (7)	0.0022 (7)
C6	0.0129 (9)	0.0156 (9)	0.0136 (9)	0.0011 (7)	0.0017 (7)	0.0017 (7)
C7	0.0150 (9)	0.0158 (9)	0.0135 (9)	-0.0009(7)	0.0040 (7)	-0.0001 (7)
C8	0.0164 (9)	0.0138 (9)	0.0141 (9)	-0.0002 (7)	0.0008 (7)	0.0015 (7)
C9	0.0150 (9)	0.0150 (9)	0.0158 (9)	-0.0006 (7)	0.0010(7)	0.0025 (7)
C10	0.0211 (10)	0.0255 (11)	0.0287 (11)	-0.0057 (8)	0.0107 (8)	-0.0022 (9)
C11	0.0203 (10)	0.0299 (11)	0.0265 (11)	0.0033 (9)	-0.0001 (8)	0.0027 (9)
C12	0.0220 (10)	0.0240 (10)	0.0167 (10)	-0.0049 (8)	0.0018 (8)	-0.0031 (8)
C13	0.0185 (10)	0.0183 (10)	0.0156 (9)	-0.0012 (7)	0.0024 (8)	0.0009 (8)

C14	0.0141 (9)	0.0236 (10)	0.0280 (11)	-0.0036 (8)	0.0119 (8)	-0.0044 (8)
C15	0.0263 (11)	0.0417 (13)	0.0362 (13)	-0.0094 (10)	0.0172 (10)	-0.0157 (11)
C16	0.0265 (11)	0.0233 (11)	0.0347 (12)	-0.0060 (9)	0.0169 (9)	-0.0047 (9)
C17	0.0184 (10)	0.0345 (12)	0.0560 (16)	0.0035 (9)	0.0117 (10)	0.0062 (11)
C18	0.0152 (9)	0.0173 (9)	0.0128 (9)	-0.0051 (7)	0.0016 (7)	-0.0010 (7)
C19	0.0262 (10)	0.0224 (10)	0.0146 (10)	0.0007 (8)	0.0072 (8)	0.0007 (8)
C20	0.0259 (10)	0.0212 (10)	0.0191 (10)	0.0033 (8)	0.0062 (8)	-0.0008 (8)
C21	0.0223 (10)	0.0163 (10)	0.0165 (9)	-0.0032 (8)	-0.0003 (8)	0.0010 (8)
C22	0.0204 (10)	0.0252 (10)	0.0126 (9)	-0.0049 (8)	0.0051 (7)	0.0015 (8)
C23	0.0149 (9)	0.0230 (10)	0.0161 (9)	-0.0027 (7)	0.0035 (7)	-0.0019 (8)
C24	0.0257 (11)	0.0186 (10)	0.0228 (10)	-0.0026 (8)	0.0032 (8)	0.0046 (8)

Geometric parameters (Å, °)

F1—C24	1.360 (2)	C10—H10B	0.9800	
F2—C24	1.346 (2)	C10—H10C	0.9800	
01—C5	1.238 (2)	C11—H11A	0.9800	
O2—C13	1.212 (2)	C11—H11B	0.9800	
O3—C13	1.346 (2)	C11—H11C	0.9800	
O3—C14	1.487 (2)	C12—H12A	0.9800	
O4—C24	1.366 (2)	C12—H12B	0.9800	
O4—C21	1.404 (2)	C12—H12C	0.9800	
N1—C1	1.363 (2)	C14—C15	1.511 (3)	
N1—C9	1.393 (2)	C14—C17	1.519 (3)	
N1—H1N	0.91 (2)	C14—C16	1.521 (3)	
C1—C6	1.362 (2)	C15—H15A	0.9800	
C1—C2	1.500 (2)	C15—H15B	0.9800	
С2—С3	1.532 (2)	C15—H15C	0.9800	
C2—H2A	0.9900	C16—H16A	0.9800	
C2—H2B	0.9900	C16—H16B	0.9800	
C3—C10	1.530 (2)	C16—H16C	0.9800	
C3—C11	1.532 (2)	C17—H17A	0.9800	
C3—C4	1.536 (2)	C17—H17B	0.9800	
C4—C5	1.516 (2)	C17—H17C	0.9800	
C4—H4A	0.9900	C18—C23	1.389 (2)	
C4—H4B	0.9900	C18—C19	1.393 (2)	
C5—C6	1.442 (2)	C19—C20	1.388 (3)	
С6—С7	1.520 (2)	C19—H19A	0.9500	
С7—С8	1.530 (2)	C20—C21	1.381 (3)	
C7—C18	1.535 (2)	C20—H20A	0.9500	
C7—H7A	1.0000	C21—C22	1.382 (3)	
С8—С9	1.355 (2)	C22—C23	1.387 (3)	
C8—C13	1.477 (2)	C22—H22A	0.9500	
C9—C12	1.495 (2)	C23—H23A	0.9500	
C10—H10A	0.9800	C24—H24A	1.0000	
C13—O3—C14	120.42 (13)	C9—C12—H12B	109.5	
C24—O4—C21	118.04 (14)	H12A—C12—H12B	109.5	

C1 N1 $C0$	122(1(15))	C0 C12 U12C	100 5
C1 - N1 - U1N	122.01(15)	C9-C12-H12C	109.5
CI-NI-HIN	117.9 (13)	H12A - C12 - H12C	109.5
C9—NI—HIN	116.9 (12)	H12B—C12—H12C	109.5
C6-C1-N1	119.88 (16)	02 - C13 - 03	122.80 (16)
C6-C1-C2	124.75 (16)	02	125.12 (17)
N1—C1—C2	115.38 (15)	O3—C13—C8	112.06 (15)
C1—C2—C3	113.36 (14)	O3—C14—C15	102.09 (14)
C1—C2—H2A	108.9	O3—C14—C17	111.05 (15)
C3—C2—H2A	108.9	C15—C14—C17	110.84 (17)
C1—C2—H2B	108.9	O3—C14—C16	109.44 (14)
C3—C2—H2B	108.9	C15—C14—C16	110.92 (17)
H2A—C2—H2B	107.7	C17—C14—C16	112.07 (16)
C10—C3—C11	109.42 (15)	C14—C15—H15A	109.5
C10—C3—C2	108.94 (14)	C14—C15—H15B	109.5
C11 - C3 - C2	110 54 (15)	H15A—C15—H15B	109 5
C10-C3-C4	110.11 (15)	C14—C15—H15C	109.5
$C_{11} = C_{3} = C_{4}$	109.96 (15)	$H_{15} - C_{15} - H_{15} C$	109.5
$C_2 C_3 C_4$	107.90(13) 107.85(14)	H15R C15 H15C	109.5
$C_2 = C_3 = C_4$	107.05(14) 112.06(14)	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	109.5
C_{5}	115.90 (14)	C14 - C10 - H10A	109.5
C5—C4—H4A	108.8	CI4—CI6—HI6B	109.5
C3—C4—H4A	108.8	H16A—C16—H16B	109.5
C5—C4—H4B	108.8	C14—C16—H16C	109.5
C3—C4—H4B	108.8	H16A—C16—H16C	109.5
H4A—C4—H4B	107.7	H16B—C16—H16C	109.5
O1—C5—C6	122.52 (16)	C14—C17—H17A	109.5
O1—C5—C4	119.59 (15)	C14—C17—H17B	109.5
C6—C5—C4	117.87 (15)	H17A—C17—H17B	109.5
C1—C6—C5	119.29 (15)	C14—C17—H17C	109.5
C1—C6—C7	120.37 (15)	H17A—C17—H17C	109.5
C5—C6—C7	120.29 (15)	H17B—C17—H17C	109.5
C6—C7—C8	109.52 (14)	C23—C18—C19	117.37 (16)
C6-C7-C18	111 30 (13)	C_{23} C_{18} C_{7}	122 12 (16)
$C_{8} - C_{7} - C_{18}$	110.69 (14)	$C_{19} - C_{18} - C_{7}$	122.12(10) 120.51(16)
C6 $C7$ $H7A$	108 /	$\begin{array}{c} C_{10} \\ C_{20} \\ C_{10} \\ C_{10} \\ C_{18} \\ C_{18$	120.31(10) 122.20(17)
C_{0} C_{7} H_{7}	108.4	$C_{20} = C_{19} = C_{18}$	122.20 (17)
$C_0 - C_1 - \Pi/A$	100.4	C_{20} C_{19} H_{10A}	110.9
$C_{18} - C_{7} - H_{7}$	108.4	C18—C19—H19A	118.9
0 0 0 0	119.93 (16)	$C_{21} = C_{20} = C_{19}$	118.65 (17)
C9—C8—C7	120.16 (15)	С21—С20—Н20А	120.7
C13—C8—C7	119.83 (15)	C19—C20—H20A	120.7
C8—C9—N1	119.31 (16)	C20—C21—C22	120.74 (17)
C8—C9—C12	128.53 (16)	C20—C21—O4	124.23 (16)
N1—C9—C12	112.15 (15)	C22—C21—O4	114.93 (16)
C3-C10-H10A	109.5	C21—C22—C23	119.59 (17)
C3—C10—H10B	109.5	C21—C22—H22A	120.2
H10A—C10—H10B	109.5	C23—C22—H22A	120.2
C3—C10—H10C	109.5	C22—C23—C18	121.37 (17)
H10A—C10—H10C	109.5	C22—C23—H23A	119.3
H10B—C10—H10C	109.5	C18—C23—H23A	119.3

C3—C11—H11A	109.5	F2—C24—F1	105.56 (14)
C3—C11—H11B	109.5	F2—C24—O4	105.67 (15)
H11A—C11—H11B	109.5	F1—C24—O4	110.52 (15)
C3—C11—H11C	109.5	F2—C24—H24A	111.6
H11A—C11—H11C	109.5	F1—C24—H24A	111.6
H11B—C11—H11C	109.5	O4—C24—H24A	111.6
C9—C12—H12A	109.5		
C9—N1—C1—C6	14.1 (2)	C7—C8—C9—C12	168.75 (17)
C9—N1—C1—C2	-165.54 (15)	C1—N1—C9—C8	-12.6 (3)
C6—C1—C2—C3	18.1 (2)	C1—N1—C9—C12	167.69 (16)
N1—C1—C2—C3	-162.30 (14)	C14—O3—C13—O2	2.1 (3)
C1—C2—C3—C10	-165.09 (15)	C14—O3—C13—C8	-179.04 (14)
C1—C2—C3—C11	74.64 (19)	C9—C8—C13—O2	-2.2 (3)
C1—C2—C3—C4	-45.59 (19)	C7—C8—C13—O2	-178.97 (18)
C10—C3—C4—C5	172.96 (15)	C9—C8—C13—O3	179.00 (16)
C11—C3—C4—C5	-66.40 (19)	C7—C8—C13—O3	2.2 (2)
C2—C3—C4—C5	54.20 (19)	C13—O3—C14—C15	178.94 (16)
C3—C4—C5—O1	147.75 (16)	C13—O3—C14—C17	-62.9 (2)
C3—C4—C5—C6	-34.2 (2)	C13—O3—C14—C16	61.4 (2)
N1—C1—C6—C5	-174.53 (15)	C6-C7-C18-C23	134.13 (17)
C2-C1-C6-C5	5.1 (3)	C8—C7—C18—C23	-103.82 (18)
N1—C1—C6—C7	8.1 (2)	C6-C7-C18-C19	-46.2 (2)
C2-C1-C6-C7	-172.29 (15)	C8—C7—C18—C19	75.88 (19)
O1-C5-C6-C1	-178.91 (16)	C23—C18—C19—C20	-3.1 (3)
C4—C5—C6—C1	3.1 (2)	C7-C18-C19-C20	177.17 (16)
O1—C5—C6—C7	-1.5 (3)	C18—C19—C20—C21	1.3 (3)
C4—C5—C6—C7	-179.50 (15)	C19—C20—C21—C22	1.3 (3)
C1—C6—C7—C8	-27.6 (2)	C19—C20—C21—O4	-174.90 (16)
C5—C6—C7—C8	155.05 (15)	C24—O4—C21—C20	-20.1 (3)
C1—C6—C7—C18	95.14 (19)	C24—O4—C21—C22	163.51 (16)
C5-C6-C7-C18	-82.24 (19)	C20—C21—C22—C23	-1.9 (3)
C6—C7—C8—C9	29.0 (2)	O4—C21—C22—C23	174.61 (15)
C18—C7—C8—C9	-94.12 (19)	C21—C22—C23—C18	0.0 (3)
C6—C7—C8—C13	-154.26 (15)	C19—C18—C23—C22	2.5 (3)
C18—C7—C8—C13	82.65 (19)	C7—C18—C23—C22	-177.84 (15)
C13—C8—C9—N1	172.33 (15)	C21—O4—C24—F2	-169.91 (14)
C7—C8—C9—N1	-10.9 (2)	C21—O4—C24—F1	76.36 (19)
C13—C8—C9—C12	-8.0 (3)		

Hydrogen-bond geometry (Å, °)

Cg3 is the centroid of the C18–C23 ring.

<i>D</i> —Н	H···A	$D \cdots A$	<i>D</i> —H··· <i>A</i>
0.91 (2)	1.96 (2)	2.866 (2)	176.6 (18)
0.98	2.25	2.800 (2)	114
0.98	2.36	2.938 (2)	117
0.98	2.37	2.958 (3)	118
	<i>D</i> —H 0.91 (2) 0.98 0.98 0.98	D—H H···A 0.91 (2) 1.96 (2) 0.98 2.25 0.98 2.36 0.98 2.37	D—HH···AD···A0.91 (2)1.96 (2)2.866 (2)0.982.252.800 (2)0.982.362.938 (2)0.982.372.958 (3)

C20—H20A…F1	0.95	2.46	2.989 (2)	115
C24—H24A···O1 ⁱⁱ	1.00	2.35	3.230 (2)	147
C2—H2 <i>A</i> ··· <i>Cg</i> 3 ⁱⁱⁱ	0.99	2.74	3.6959 (19)	162

Symmetry codes: (i) x, -y+3/2, z-1/2; (ii) x, y-1, z; (iii) x, -y+1/2, z-3/2.