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The title sterically congested piperazine derivative,  $C_{20}H_{27}FN_2O_2$ , was prepared using a modified Bruylants approach. A search of the Cambridge Structural Database identified 51 compounds possessing an *N-tert*-butyl piperazine substructure. Of these only 14 were asymmetrically substituted on the piperazine ring and none with a synthetically useful second nitrogen. Given the novel chemistry generating a pharmacologically useful core, determination of the crystal structure for this compound was necessary. The piperazine ring is present in a chair conformation with di-equatorial substitution. Of the two N atoms, one is  $sp^3$  hybridized while the other is  $sp^2$  hybridized. Intermolecular interactions resulting from the crystal packing patterns were investigated using Hirshfeld surface analysis and fingerprint analysis. Directional weak hydrogenbond-like interactions as the major source of attraction are present in the crystal packing.

#### 1. Chemical context

In the course of designing novel sigma-2 ligands, it was necessary to synthesize 1-(2-methyl-4-phenylbutan-2-yl)piperazines. These could be prepared in several steps from the corresponding alkyne **1** shown in Fig. 1. The challenge of synthesizing quaternary carbons (Wei *et al.*, 2020; Liu *et al.*, 2015; Volla *et al.*, 2014; Fuji, 1993; Martin, 1980), particularly amine-bearing quaternary carbons (Zhu *et al.*, 2019; Yeung *et al.*, 2019; Xu *et al.*, 2019; Velasco-Rubio *et al.*, 2019; Vasu *et al.*,







2019; Trost et al., 2019; Ling & Rivas, 2016; Hager et al., 2016; Clavden et al., 2011: Fu et al., 2008: Riant & Hannedouche, 2007), is well established. The presence of the N-gem-dimethyl group of **1** presented a significant synthetic challenge arising from steric congestion. Nucleophilic attack by an organometallic reagent into a transient 1-N-ethylidenepiperazinium has a literature precedent, but nucleophilic attack into the more sterically congested 1-N-propylidenepiperazinium intermediate by an alkynyl Grignard reagent is presented here for the first time. Four potential synthetic routes were identified including Katritzky benzotriazole trapping of an iminium (Monbaliu et al., 2013; Ingram et al., 2006; Katritzky, 1998; Katritzky et al., 1989, 1991, 2005; Katritzky & Rogovoy, 2003; Katritzky & Saczewski, 1990), a Bruylants (Bruylants, 1924) trapping of an iminium, sequential addition of two methyl groups into an amide, and rearrangement to the gem-dimethyl group. All in-house attempts at the Katritzky benzotriazole (Tang et al., 2013; Pierce et al., 2012; Albaladejo et al., 2012) or triazole (Prashad et al., 2005) reactions failed. A variation on the Bruylants reaction (Liu et al., 2014; Beaufort-Droal et al., 2006; Prashad et al., 2005; Kudzma et al., 1988; Bernardi et al., 2003) described herein was successful. The traditional Bruylants reaction captures a trapped iminium as the corresponding  $\alpha$ -amino nitrile. In a subsequent reaction, the  $\alpha$ amino nitrile transiently forms an iminium that is then trapped with excess Grignard reagent. Conversion of the terminal alkyne 4 to the corresponding magnesiobromide acetylide proceeded under established conditions. Attack of an alkynyl magnesium bromide into the transient iminium is precedented to yield tertiary carbon products. Generation of a quaternary carbon product in an analogous manner has not been described. A single paper details addition of a copper acetylide into a Brulyants adduct (Tang et al., 2013). Given the pharmacological importance of this compound and its tractable synthesis with novel chemistry, careful structural characterization by X-ray crystallographic analysis was necessary. Optimization of this reaction, subsequent structural elaboration, and specific pharmacological relevance will be detailed in later publications.



2. Structural commentary

The title compound, prepared from achiral reagents as a racemic mixture, crystallizes in the chiral monoclinic space group  $P2_1$  with one molecule in the asymmetric unit as shown in the Scheme and Fig. 2. No heavy atoms are present in the

U	1	· /	
F1-C1	1.359 (3)	N2-C16	1.336 (3)
O1-C16	1.211 (3)	C1-C6	1.351 (4)
O2-C16	1.345 (3)	$C7 \cdot \cdot \cdot N1$	3.508 (3)
C12-N1-C14	108.36 (16)	C16-N2-C13	120.9 (2)
C12-N1-C9	113.89 (18)	C15-N2-C13	112.8 (2)
C14-N1-C9	113.48 (16)	N1-C12-C13	110.77 (19)
C16-N2-C15	126.30 (19)	N2-C15-C14	110.1 (2)

structure and data were collected using Mo K $\alpha$  radiation. Thus, the absolute structure of the randomly chosen crystals could not be determined reliably (Parsons et al., 2013; Zhou et al., 2015). In the molecule, the NC(=O)O group of the carbamate exists in resonance. The bond lengths between carbon and other atoms (Table 1) are in the expected ranges with the bond length between O1-C16 [1.211 (3) Å] being the shortest, followed by N2-C16 [1.336 (3) Å], O2-C16 [1.345 (3) Å], and F1-C1 [1.359 (3) Å] owing to the presence of the more electronegative atoms oxygen, nitrogen and fluorine. The bond length between C1–C6 [1.351 (4) Å] is the shortest among all the bond lengths in the phenyl group, possibly due to the inductive effect of fluorine. The spatial distance between the extreme atoms of propargylamine groups ( $C7 \cdot \cdot \cdot N1$ ) was observed to be 3.508 (3) Å, which is slightly longer than for the other reported propargylamines (3.372-3.478 Å; Marvelli et al., 2004; Sidorov et al., 1999, 2000), and possibly due to the open L-shaped structure of the molecule. Also, the piperazine ring is shown in its most stable chair form conformation in Fig. 3, as evidenced by the bond angles (Table 1) between N1-C12-C13  $[110.77 (19)^{\circ}]$  and N2-C15-C14 [110.1 (2) $^{\circ}$ ], which are close to the ideal bond angle of 109.5° for a chair conformation. The sum of the bond angles around N1 (335.73°) indicate  $sp^3$  hybridization, while the sum of the bond angles around N2 (360°) indicates  $sp^2$ hybridization. This is also evidenced by the tetragonal molecular geometry of C12-N1-C9 [113.89 (18)°], C14-N1-C9 [113.48 (16)°], and C12-N1-C14 [108.36 (16)°] and the trigonal planar molecular geometry of C16-N2-C15  $[126.30 (19)^{\circ}]$ , C16-N2-C13  $[120.9 (2)^{\circ}]$ , and C15-N2-C13 [112.8 (2) $^{\circ}$ ]. The delocalization of the lone pair of N2 into the  $\pi$  bond of carbonyl group causes  $sp^2$  hybridization of N2.



Figure 2

30% probability ellipsoid plot for the crystal structure solution of *tert*butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate. Hydrogen atoms are omitted for clarity.





40% probability plot of the molecular crystal structure solution of *tert*butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate showing the L-shaped structure and the chair conformation of the piperazine ring.

#### 3. Supramolecular features

Hirshfeld surface analysis and fingerprint analysis were performed using CrystalExplorer (Spackman & Jayatilaka, 2009, Spackman & McKinnon, 2002, McKinnon *et al.*, 2007). In the absence of acidic hydrogen atoms, there cannot be any conventional hydrogen bonds; however, there are directional interactions present between C2–H2···O1 and C–H··· $\pi$ interactions between C19–H19···C1, as shown in the crystal packing along the *a*-axis (Fig. 4). These interactions are



Figure 4

30% probability plot of crystal packing of *tert*-butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate viewed down the *a* axis showing weak hydrogen-bond-like interactions between C2– H2···O1 and C-H··· $\pi$  interactions between C19–H19···C1 due to dispersion interactions. Hydrogen atoms not involved in intermolecular interactions are omitted for clarity.



Hirshfeld surface for *tert*-butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate mapped over  $d_{norm}$  showing weak hydrogenbond-like interactions between C2-H2···O1 and C-H··· $\pi$  interactions between C19-H19···C1.

represented by the faint red spots between C2–H2···O1 and C19–H19···C1 on the Hirshfeld surface mapped over  $d_{norm}$  in Fig. 5. The directional C2–H2···O1  $[d(H \cdot \cdot O) = 2.595 \text{ Å}]$  present in the crystal packing could be weak C–H···O hydrogen-bond-like interactions (Desiraju & Steiner, 1999) and the C19–H19···C1  $[d(C \cdot \cdot H) = 2.804 \text{ Å}]$  interactions could be  $C-H \cdot \cdot \pi$  interactions with dispersion interactions as the major source of attraction. Fingerprint analysis (Fig. 6) complemented the Hirshfeld analysis by showing a minimal contact surface between O···H (3.1%) and F···H (5.4%), as shown in Fig. 6b and Fig. 6c. These could be the directional C–H···F close contacts attributed to the proximity of the F atom to the C–H··· $\pi$  interactions. Please see Table 2 for the interatomic



#### Figure 6

The two-dimensional fingerprint plots of *tert*-butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate showing contributions from different contacts.

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 Table 2

 Short interatomic contact distances (Å).

Contact	Distance
C2-H2···O1	2.595
C19-H19···C1	2.804
C19-H19···F1	3.163

contact distances. These data also suggested the absence of  $\pi$ - $\pi$  stacking as C···C contacts contribute 0% of the Hirshfeld surfaces (Fig. 6*d*).

#### 4. Database survey

A search in the Cambridge Structural Database (Version 5.41 update of March 2020; (Groom et al., 2016)) for compounds possessing an N-tert-butyl piperazine substructure identified 51 compounds. These compounds were several variations of BuckyBall adducts, diketopiperazine derivatives, and ligands. There were only 14 compounds viz. DIYWAK (McDermott et al., 2008), HEHZOL (Legnani et al., 2012), HICYID, HICYOJ (Sinha et al., 2013b), JIFHEO (Zhong et al., 2018), OFUDAW (Korotaev et al., 2012), PUYNUS (Jin & Liebscher, 2002), RIPWUJ (Bobeck et al., 2007), TILJIJ (Sinha et al., 2013a), UPIBIF, UPIBOL (Wiedner & Vedejs, 2010), UYIHOB (Chen & Cao, 2017), WANTAJ (Golubev & Krasavin, 2017), and WINMAH (Brouant & Giorgi, 1995) that were asymmetrically substituted on the piperazine ring, and none with a synthetically useful second nitrogen. All were effectively 'nonintermediate' compounds that could not reasonably serve for additional substitution at the second nitrogen and none had alkyne substitutions. The quaternary carbon piperazines were explored by Sinha et al. (2013a,b) using an Ugi reaction; however, the present structure is the only compound containing an  $\alpha, \alpha$ -dimethyl carbon attached to an alkyne and an amine. This new methodology required the X-ray studies to confirm the generated structure. In summary, to the best of the authors' knowledge, there is no published crystal structure like the title compound, for a molecule containing asymmetrical substitutions on the piperazine ring, having a synthetically useful second nitrogen, and an  $\alpha, \alpha$ -dimethyl carbon attached to an alkyne and an amine.

#### 5. Synthesis and crystallization

*tert*-Butyl 4-(2-cyanopropan-2-yl)piperazine-1-carboxylate (3): Ethereal HCl (40.3 mL of a 2.0 M in Et<sub>2</sub>O, 80.6 mmol, 1.5 eq. titrated against standardized 1 N NaOH to a phenolph-thalein pink end-point) was added dropwise to a stirred solution of *tert*-butyl piperazine-1-carboxylate 2 (12.6 g, 53.7 mmol, 1.0 eq.) in MeOH (60 mL) and CH<sub>2</sub>Cl<sub>2</sub> (60 mL) at 273 K under Argon. The resulting mixture was stirred at 273 K for 1 h, after which the solvent and excess HCl were removed under reduced pressure and the white residual solid was dissolved in water (150 mL). In a well-ventilated fume hood, solid NaCN (2.63 g, 53.7 mmol, 1.0 eq.) and then a solution of acetone (9.4 g, 11.8 mL, 161.2 mmol, 3.0 eq.) in water (20 mL)

Table 3	
Experimental details.	
Crystal data	
Chemical formula	$C_{20}H_{27}FN_2O_2$
M <sub>r</sub>	346.43
Crystal system, space group	Monoclinic, P2 <sub>1</sub>
Temperature (K)	293
<i>a</i> , <i>b</i> , <i>c</i> (Å)	10.2576 (11), 9.5127 (10), 10.5318 (11)
$\beta$ (°)	104.691 (2)
$V(Å^3)$	994.07 (18)
Z	2
Radiation type	Μο Κα
$\mu \text{ (mm}^{-1})$	0.08
Crystal size (mm)	$0.65\times0.50\times0.17$
Data collection	
Diffractometer	Bruker SMART APEXII
Absorption correction	Multi-scan ( <i>SADABS</i> ; Sheldrick, 2002)
$T_{\min}, T_{\max}$	0.704, 0.746
No. of measured, independent and	10640, 5058, 3662
observed $[I > 2\sigma(I)]$ reflections	
R <sub>int</sub>	0.017
$(\sin \theta / \lambda)_{\max} ( \text{\AA}^{-1} )$	0.675
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.040, 0.109, 1.04
No. of reflections	5058
No. of parameters	231
No. of restraints	1
H-atom treatment	H-atom parameters constrained
$\Delta \rho_{\rm max},  \Delta \rho_{\rm min} \ ({ m e} \ { m \AA}^{-3})$	0.12, -0.11

Computer programs: SMART and SAINT (Bruker, 1998), SHELXS97 (Sheldrick, 2008), SHELXL2018/3 (Sheldrick, 2015) and CrystalMaker (Palmer, 2014).

were added sequentially at room temperature (296 K). The resulting mixture was stirred at room temperature under air for an additional 48 h. Water (100 mL) was added and the mixture was extracted with EtOAc ( $3 \times 100$  mL) then NaCl (sat, aq.). The combined organic extracts were dried (MgSO<sub>4</sub>) and the solvent was removed under reduced pressure to give *tert*-butyl 4-(2-cyanopropan-2-yl)piperazine-1-carboxylate **3** as a white crystalline solid, 11 g (64%). MP: 381.2 K (reported 381–383 K) matching the literature (Firth *et al.*, 2016). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>:  $\delta$ 3.50 (*dd*, *J* = 4.8 Hz, 4H), 2.62 (*dd*, *J* = 4.8 Hz, 4H), 1.54 (*s*, 6H), 1.49 (*s*, 9H) matches literature (Firth *et al.*, 2016).

Note: the aqueous extracts (pH > 10) were collected and the residual cyanide was oxidized to cyanate with sodium hypochlorite (Gerritsen & Margerum, 1990) and absence of a cyanide ion was confirmed with an MQuant<sup>(m)</sup> Koening Cyanide test indicator from EM sciences.

#### *tert*-Butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate (1):

A 250 mL flame-dried, round-bottom flask was cooled under argon and then charged with 1-ethynyl-4-fluorobenzene 4 (1.98 g, 16.5 mmol) in 50 mL of anhydrous THF. This solution was cooled with an external ice-bath. A commercial solution of methyl magnesium bromide (5.25 mL, 16.5 mmol) (Acros, ~3.2 *M* in THF, assayed against anhydrous diphenyl acetic acid with 2 mg 1,10-phenanthroline as an indicator) was added with slow dropwise addition over 10 minutes. The internal temperature was maintained between 274–275 K. This mixture

was stirred at ice-bath temperature for an additional 20 minutes, which resulted in a pale-yellow solution. A solution of tert-butyl 4-(2-cyanopropan-2-yl)piperazine-1-carboxylate 3 (Firth et al., 2016) (2.33 g, 9.2 mmol) in 25 mL THF was added dropwise to this mixture over 10 minutes; the internal temperature was maintained between 274-275.3 K. This deepyellow solution was permitted to stir with the external ice-bath slowly melting and rising to room temperature, while progress was monitored by TLC ( $R_f$  of product at 0.6 1:1 H:EA, SiO<sub>2</sub> plates, SWUV and I<sub>2</sub> visualization). Following stirring for 12 h at 296 K, the crude reaction mixture was cooled to ice-bath temperature and the reaction was quenched with the addition of 10 mL of ice-cold water at a rate of addition that maintained the internal temperature below 278 K. After quenching the organo-base, an additional 50 mL of water were added. Small aliquots of brine and ethanol were used, as required, to break the emulsion in the following extraction. This mixture was extracted with  $3 \times 20$  mL of ethyl acetate, washed (3  $\times$ 10 mL H<sub>2</sub>O,  $3 \times 10$  mL brine) dried (Na<sub>2</sub>SO<sub>4</sub>), decanted, and the solvent removed under reduced pressure to afford 30.6 g of a yellow solid. This was separated on 50 g of  $SiO_2$  with hexane/ethyl acetate (1/1) as the eluent to yield tert-butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxvlate 1 as a white powder, 2.74 g (86.3%). This compound was recrystallized from ethyl acetate as colorless plates, having a melting point of 388.1 K. <sup>1</sup>H NMR (400 MHz, chloroform-d)  $\delta$  7.36 (*dd*, *J* = 8.2, 5.6 Hz, 2H), 6.96 (*t*, *J* = 8.5 Hz, 2H), 3.46 (*s*, 5H), 2.63 (s, 4H), 1.45 (s, 16H). HRMS: (C<sub>20</sub>H<sub>27</sub>FN<sub>2</sub>O<sub>2</sub>) calculated for  $[M + H]^+$  347.2129, found 347.2127.

#### 6. Refinement

Crystal data, data collection, and structure refinement details are summarized in Table 3. H atoms were localized in a difference-Fourier map. C-bound H atoms were treated as riding, with C-H = 0.93, 0.96 or 0.97 Å, and with  $U_{iso}(H) =$  $1.2U_{eq}(C)$  for aromatic and  $1.5U_{eq}(C)$  for methyl groups.

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# supporting information

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Crystal structure of *tert*-butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate

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## **Computing details**

Data collection: *SMART* and *SAINT* (Bruker, 1998); cell refinement: *SMART* and *SAINT* (Bruker, 1998); data reduction: *SMART* and *SAINT* (Bruker, 1998); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015); molecular graphics: *CrystalMaker* (Palmer, 2014); software used to prepare material for publication: *SHELXL2018/3* (Sheldrick, 2015).

tert-Butyl 4-[4-(4-fluorophenyl)-2-methylbut-3-yn-2-yl]piperazine-1-carboxylate

Crystal data

 $C_{20}H_{27}FN_2O_2$   $M_r = 346.43$ Monoclinic,  $P2_1$  a = 10.2576 (11) Å b = 9.5127 (10) Å c = 10.5318 (11) Å  $\beta = 104.691$  (2)° V = 994.07 (18) Å<sup>3</sup> Z = 2

Data collection

Bruker SMART APEXII diffractometer  $\varphi$  and  $\omega$  Scans scans Absorption correction: multi-scan (SADABS; Sheldrick, 2002)  $T_{\min} = 0.704$ ,  $T_{\max} = 0.746$ 10640 measured reflections

## Refinement

Refinement on  $F^2$ Least-squares matrix: full  $R[F^2 > 2\sigma(F^2)] = 0.040$  $wR(F^2) = 0.109$ S = 1.045058 reflections 231 parameters 1 restraint F(000) = 372  $D_x = 1.157 \text{ Mg m}^{-3}$ Mo K\alpha radiation,  $\lambda = 0.71073 \text{ Å}$ Cell parameters from 3739 reflections  $\theta = 2.9-23.9^{\circ}$   $\mu = 0.08 \text{ mm}^{-1}$  T = 293 KPlate, colorless  $0.65 \times 0.50 \times 0.17 \text{ mm}$ 

5058 independent reflections 3662 reflections with  $I > 2\sigma(I)$   $R_{int} = 0.017$   $\theta_{max} = 28.7^{\circ}, \ \theta_{min} = 2.0^{\circ}$   $h = -13 \rightarrow 13$   $k = -12 \rightarrow 12$  $l = -14 \rightarrow 14$ 

Primary atom site location: structure-invariant direct methods Secondary atom site location: difference Fourier map Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained  $w = 1/[\sigma^2(F_o^2) + (0.0543P)^2 + 0.0286P]$ where  $P = (F_o^2 + 2F_c^2)/3$  $(\Delta/\sigma)_{\text{max}} < 0.001$   $\Delta \rho_{\rm max} = 0.12 \text{ e } \text{\AA}^{-3}$  $\Delta \rho_{\rm min} = -0.11 \text{ e } \text{\AA}^{-3}$ 

## 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	1.18988 (19)	0.1239 (2)	-0.11543 (15)	0.1037 (6)
01	0.36605 (17)	0.51912 (19)	0.18415 (19)	0.0806 (5)
O2	0.51029 (17)	0.70053 (17)	0.25327 (19)	0.0744 (5)
N1	0.75069 (16)	0.3139 (2)	0.49328 (17)	0.0532 (4)
N2	0.54352 (19)	0.4993 (2)	0.3602 (3)	0.0786 (7)
C1	1.1321 (3)	0.1417 (3)	-0.0138 (2)	0.0675 (6)
C2	1.1908 (2)	0.2311 (3)	0.0845 (2)	0.0657 (6)
H2	1.267513	0.281757	0.081485	0.079*
C3	1.1337 (2)	0.2446 (3)	0.1888 (2)	0.0607 (5)
Н3	1.173183	0.304622	0.257504	0.073*
C4	1.0186 (2)	0.1709 (2)	0.1937 (2)	0.0551 (5)
C5	0.9617 (3)	0.0830 (3)	0.0901 (3)	0.0760 (7)
Н5	0.884011	0.032892	0.090838	0.091*
C6	1.0188 (3)	0.0687 (4)	-0.0148 (3)	0.0830 (8)
H6	0.980055	0.010008	-0.084752	0.100*
C7	0.9619 (2)	0.1869 (3)	0.3042 (2)	0.0623 (5)
C8	0.9197 (2)	0.1996 (3)	0.3994 (2)	0.0614 (5)
C9	0.8686 (2)	0.2172 (3)	0.5184 (2)	0.0594 (5)
C10	0.8289 (3)	0.0738 (3)	0.5623 (3)	0.0761 (7)
H10A	0.907916	0.016550	0.591129	0.114*
H10B	0.766913	0.028670	0.490045	0.114*
H10C	0.786779	0.086116	0.633205	0.114*
C11	0.9824 (3)	0.2766 (3)	0.6290 (2)	0.0766 (7)
H11A	1.053887	0.208928	0.652455	0.115*
H11B	0.948419	0.296461	0.704074	0.115*
H11C	1.016014	0.361555	0.599756	0.115*
C12	0.6330 (2)	0.2618 (2)	0.3960 (3)	0.0623 (5)
H12A	0.612475	0.167143	0.419119	0.075*
H12B	0.652638	0.258352	0.310685	0.075*
C13	0.5133 (2)	0.3551 (2)	0.3890 (3)	0.0758 (7)
H13A	0.437271	0.321130	0.321152	0.091*
H13B	0.488758	0.352041	0.472069	0.091*
C14	0.7810(2)	0.4556 (2)	0.4548 (2)	0.0629 (6)
H14A	0.799613	0.452268	0.369024	0.075*
H14B	0.860821	0.490898	0.517127	0.075*
C15	0.6649 (2)	0.5530 (3)	0.4505 (3)	0.0753 (7)

# supporting information

H15A	0.650242	0.561673	0.537584	0.090*	
H15B	0.685611	0.645571	0.422237	0.090*	
C16	0.4649 (2)	0.5686 (2)	0.2590 (2)	0.0613 (5)	
C17	0.4383 (3)	0.8005 (3)	0.1535 (2)	0.0719 (6)	
C18	0.4377 (4)	0.7509 (5)	0.0174 (3)	0.1233 (14)	
H18A	0.411731	0.827031	-0.043585	0.185*	
H18B	0.374709	0.674936	-0.007262	0.185*	
H18C	0.526238	0.719043	0.016399	0.185*	
C19	0.2976 (3)	0.8232 (4)	0.1690 (3)	0.0933 (9)	
H19A	0.255882	0.899276	0.113429	0.140*	
H19B	0.301797	0.845773	0.258749	0.140*	
H19C	0.245633	0.739092	0.144516	0.140*	
C20	0.5234 (5)	0.9306 (4)	0.1912 (4)	0.1246 (14)	
H20A	0.487111	1.005437	0.131485	0.187*	
H20B	0.614143	0.911229	0.187209	0.187*	
H20C	0.523167	0.958052	0.278851	0.187*	

Atomic displacement parameters  $(\mathring{A}^2)$ 

	$U^{11}$	U <sup>22</sup>	U <sup>33</sup>	$U^{12}$	$U^{13}$	<i>U</i> <sup>23</sup>
F1	0.1100 (12)	0.1444 (17)	0.0629 (8)	0.0197 (12)	0.0335 (8)	-0.0031 (9)
01	0.0619 (9)	0.0604 (10)	0.1066 (13)	-0.0065 (8)	-0.0025 (9)	0.0007 (9)
02	0.0717 (10)	0.0445 (8)	0.0986 (12)	-0.0040 (8)	0.0064 (9)	0.0062 (8)
N1	0.0502 (9)	0.0478 (9)	0.0611 (9)	0.0025 (7)	0.0132 (7)	0.0059 (8)
N2	0.0513 (10)	0.0470 (11)	0.1223 (17)	-0.0065 (8)	-0.0061 (11)	0.0149 (11)
C1	0.0731 (15)	0.0785 (16)	0.0515 (12)	0.0157 (13)	0.0170 (11)	0.0065 (11)
C2	0.0561 (12)	0.0724 (16)	0.0695 (14)	0.0003 (11)	0.0173 (11)	-0.0018 (12)
C3	0.0586 (12)	0.0612 (13)	0.0615 (13)	0.0023 (10)	0.0136 (10)	-0.0077 (10)
C4	0.0534 (11)	0.0551 (12)	0.0558 (11)	0.0101 (9)	0.0119 (9)	0.0081 (9)
C5	0.0692 (14)	0.0772 (17)	0.0781 (16)	-0.0132 (13)	0.0125 (12)	-0.0043 (13)
C6	0.0916 (19)	0.0891 (19)	0.0612 (14)	-0.0063 (16)	0.0063 (13)	-0.0167 (13)
C7	0.0591 (12)	0.0619 (13)	0.0654 (13)	0.0095 (10)	0.0151 (10)	0.0103 (10)
C8	0.0592 (12)	0.0600 (13)	0.0664 (13)	0.0088 (10)	0.0181 (10)	0.0108 (11)
С9	0.0569 (11)	0.0610(13)	0.0618 (12)	0.0105 (10)	0.0176 (9)	0.0125 (10)
C10	0.0811 (16)	0.0624 (15)	0.0913 (18)	0.0187 (13)	0.0337 (14)	0.0265 (13)
C11	0.0660 (14)	0.095 (2)	0.0641 (14)	0.0166 (13)	0.0075 (11)	0.0112 (13)
C12	0.0601 (12)	0.0428 (10)	0.0788 (14)	-0.0063 (9)	0.0080 (10)	0.0050 (10)
C13	0.0522 (12)	0.0513 (14)	0.115 (2)	-0.0074 (10)	0.0041 (13)	0.0170 (13)
C14	0.0513 (11)	0.0507 (12)	0.0793 (14)	-0.0079 (9)	0.0028 (10)	0.0037 (11)
C15	0.0606 (13)	0.0452 (12)	0.1076 (19)	-0.0031 (10)	-0.0019 (13)	-0.0023 (12)
C16	0.0480 (11)	0.0445 (11)	0.0910 (16)	0.0004 (9)	0.0165 (11)	-0.0030 (11)
C17	0.0970 (17)	0.0551 (13)	0.0653 (13)	0.0022 (13)	0.0237 (12)	0.0101 (11)
C18	0.165 (4)	0.137 (3)	0.084 (2)	-0.021 (3)	0.063 (2)	-0.012 (2)
C19	0.103 (2)	0.0772 (18)	0.098 (2)	0.0310 (18)	0.0233 (17)	0.0133 (16)
C20	0.168 (4)	0.0650 (19)	0.130 (3)	-0.031 (2)	0.018 (3)	0.0250 (19)

Geometric parameters (Å, °)

F1—C1	1.359 (3)	С10—Н10С	0.9600
O1—C16	1.211 (3)	C11—H11A	0.9600
O2—C16	1.345 (3)	C11—H11B	0.9600
O2—C17	1.470 (3)	C11—H11C	0.9600
N1—C12	1.457 (3)	C12—C13	1.502 (3)
N1—C14	1.463 (3)	C12—H12A	0.9700
N1—C9	1.489 (3)	C12—H12B	0.9700
N2—C16	1.336 (3)	C13—H13A	0.9700
N2—C15	1.454 (3)	C13—H13B	0.9700
N2—C13	1.456 (3)	C14—C15	1.501 (4)
C1—C6	1.351 (4)	C14—H14A	0.9700
C1—C2	1.357 (4)	C14—H14B	0.9700
C2—C3	1.376 (3)	C15—H15A	0.9700
С2—Н2	0.9300	C15—H15B	0.9700
C3—C4	1.385 (3)	C17—C18	1.508 (4)
С3—Н3	0.9300	C17—C19	1.508 (4)
C4—C5	1.381 (3)	C17—C20	1.509 (4)
C4—C7	1.434 (3)	C18—H18A	0.9600
C5—C6	1.382 (4)	C18—H18B	0.9600
С5—Н5	0.9300	C18—H18C	0.9600
С6—Н6	0.9300	C19—H19A	0.9600
С7—С8	1.195 (3)	C19—H19B	0.9600
C8—C9	1.486 (3)	C19—H19C	0.9600
C9—C10	1.528 (4)	C20—H20A	0.9600
C9—C11	1.532 (3)	C20—H20B	0.9600
C10—H10A	0.9600	C20—H20C	0.9600
C10—H10B	0.9600		
C16-02-C17	121 26 (19)	N1H12B	109.5
$C_{12} = N_1 = C_{14}$	108 36 (16)	$C_{13}$ $C_{12}$ $H_{12B}$	109.5
C12 $N1$ $C9$	113 89 (18)	H12A $C12$ $H12B$	108.1
C12 = N1 = C9	113.48 (16)	$N_{2}$ C13 C12	110.6(2)
C16 - N2 - C15	126 30 (19)	N2	109 5
C16 - N2 - C13	120.90(1))	C12— $C13$ — $H13A$	109.5
C15 = N2 = C13	112 8 (2)	N2-C13-H13B	109.5
C6-C1-C2	122.8 (2)	C12— $C13$ — $H13B$	109.5
C6-C1-F1	122.0(2) 1184(2)	$H_{13A}$ $-C_{13}$ $-H_{13B}$	108.1
$C^2$ — $C^1$ — $F^1$	118.1(2) 118.8(2)	N1-C14-C15	110 74 (18)
C1 - C2 - C3	118.1(2)	N1-C14-H14A	109 5
C1-C2-H2	120.9	C15-C14-H14A	109.5
C3—C2—H2	120.9	N1—C14—H14B	109.5
$C_2 - C_3 - C_4$	121.5 (2)	C15—C14—H14B	109.5
C2—C3—H3	119.3	H14A—C14—H14B	108.1
C4—C3—H3	119.3	N2-C15-C14	110.1 (2)
C5—C4—C3	118.0 (2)	N2—C15—H15A	109.6
C5—C4—C7	121.9 (2)	C14—C15—H15A	109.6

C3—C4—C7	120.1 (2)	N2—C15—H15B	109.6
C4—C5—C6	120.7 (2)	C14—C15—H15B	109.6
С4—С5—Н5	119.6	H15A—C15—H15B	108.1
С6—С5—Н5	119.6	O1—C16—N2	124.3 (2)
C1—C6—C5	118.8 (2)	O1—C16—O2	125.2 (2)
С1—С6—Н6	120.6	N2—C16—O2	110.50 (19)
С5—С6—Н6	120.6	O2—C17—C18	110.9 (3)
C8—C7—C4	177.4 (2)	O2—C17—C19	109.7 (2)
C7—C8—C9	179.2 (3)	C18—C17—C19	112.0 (3)
C8—C9—N1	111.34 (17)	O2—C17—C20	101.0 (2)
C8—C9—C10	109.5 (2)	C18—C17—C20	111.7 (3)
N1—C9—C10	109.80 (17)	C19—C17—C20	111.1 (3)
C8—C9—C11	108.59 (18)	C17—C18—H18A	109.5
N1—C9—C11	109.55 (19)	C17—C18—H18B	109.5
C10—C9—C11	108.0 (2)	H18A—C18—H18B	109.5
С9—С10—Н10А	109.5	C17—C18—H18C	109.5
С9—С10—Н10В	109.5	H18A—C18—H18C	109.5
H10A—C10—H10B	109.5	H18B—C18—H18C	109.5
С9—С10—Н10С	109.5	С17—С19—Н19А	109.5
H10A—C10—H10C	109.5	C17—C19—H19B	109.5
H10B—C10—H10C	109.5	H19A—C19—H19B	109.5
С9—С11—Н11А	109.5	С17—С19—Н19С	109.5
C9—C11—H11B	109.5	H19A—C19—H19C	109.5
H11A—C11—H11B	109.5	H19B—C19—H19C	109.5
C9—C11—H11C	109.5	C17—C20—H20A	109.5
H11A—C11—H11C	109.5	C17—C20—H20B	109.5
H11B—C11—H11C	109.5	H20A—C20—H20B	109.5
N1—C12—C13	110.77 (19)	С17—С20—Н20С	109.5
N1—C12—H12A	109.5	H20A—C20—H20C	109.5
C13—C12—H12A	109.5	H20B-C20-H20C	109.5
C6—C1—C2—C3	1.7 (4)	C16—N2—C13—C12	125.9 (3)
F1-C1-C2-C3	-177.9 (2)	C15—N2—C13—C12	-53.4 (3)
C1—C2—C3—C4	-0.7 (3)	N1—C12—C13—N2	56.6 (3)
C2—C3—C4—C5	-0.3 (3)	C12—N1—C14—C15	60.9 (2)
C2—C3—C4—C7	179.7 (2)	C9—N1—C14—C15	-171.60 (19)
C3—C4—C5—C6	0.5 (4)	C16—N2—C15—C14	-125.5 (3)
C7—C4—C5—C6	-179.6 (2)	C13—N2—C15—C14	53.7 (3)
C2-C1-C6-C5	-1.5 (4)	N1-C14-C15-N2	-57.5 (3)
F1—C1—C6—C5	178.1 (2)	C15—N2—C16—O1	178.3 (3)
C4—C5—C6—C1	0.4 (4)	C13—N2—C16—O1	-0.9 (4)
C12—N1—C9—C8	64.1 (2)	C15—N2—C16—O2	-2.3 (4)
C14—N1—C9—C8	-60.5 (2)	C13—N2—C16—O2	178.6 (2)
C12—N1—C9—C10	-57.3 (2)	C17—O2—C16—O1	1.8 (4)
C14—N1—C9—C10	178.12 (19)	C17—O2—C16—N2	-177.7 (2)
C12—N1—C9—C11	-175.76 (19)	C16—O2—C17—C18	-63.8 (3)
C14—N1—C9—C11	59.6 (2)	C16—O2—C17—C19	60.4 (3)
C14—N1—C12—C13	-60.2 (2)	C16—O2—C17—C20	177.7 (3)

C9—N1—C12—C13	172.46 (18)		
Selected bond lengths $(A)$ and	bond angles (°)		
F1—C1		1.359 (3)	
O1—C16		1.211 (3)	
O2—C16		1.345 (3)	
N2		1.336 (3)	
C1—C6		1.351 (4)	
C7—N2		3.508	
N1-C12-C13		110.77 (19)	
N2-C15-C14		110.1 (2)	
C12—N1—C9		113.89 (18)	
C14—N1—C9		113.48 (16)	
C12—N1—C14		108.36 (16)	
C16—N2—C15		126.30 (19)	
C16—N2—C13		120.9 (2)	
C15—N2—C13		112.8 (2)	