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The molecular conformation of the title compound, $C_{20}H_{15}Cl_2N_3O_2$, is stabilized by an intramolecular $O-H\cdots O$ hydrogen bond, forming an S(6) ring motif. The central pyridine ring is almost planar [maximum deviation = 0.074 (3) Å]. It subtends dihedral angles of 86.10 (15) and 87.17 (14)°, respectively, with the phenyl and dichlorophenyl rings, which are at an angle of 21.28 (15)° to each other. The ==C(-OH)CH₃ group is coplanar. In the crystal, molecules are linked by intermolecular N-H···N and C-H···N hydrogen bonds, and N-H··· π and C-H··· π interactions, forming a three-dimensional network. The most important contributions to the crystal packing are from H···H (33.1%), C···H/H···C (22.5%), Cl···H/H···Cl (14.1%), O···H/H···O (11.9%) and N···H/H···N (9.7%) interactions.

1. Chemical context

The development of effective methods for the construction of small-sized molecules bearing a nitrogen heterocycle is a very important proposition in organic synthesis and catalysis (Abdel-Hafiz *et al.*, 2012; Gurbanov *et al.*, 2018; Zubkov *et al.*, 2018). As members of this family, pyridine derivatives play a key role in flavor chemistry, crystal engineering, and the development of biologically active compounds (Adams & De Kimpe, 2006; Mahmoudi *et al.*, 2019; Mamedov *et al.*, 2020). The pyridine core is a key bioactive fragment of diverse natural products (niacin, pyridoxine, nicotine, NADP⁺) and series of derivatives constitute promising drugs in medicinal chemistry (Mohsin & Ahmad, 2018).







research communications

Table 1	l			
Hydrog	en-bond	geome	etry (Å,	°).

Cg2 is the centroid of the C7–C12 phenyl ring.

D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
0.86 (4)	1.72 (4)	2.514 (3)	153 (4)
0.86 (4)	2.22 (4)	3.032 (4)	159 (3)
0.95	2.62	3.308 (4)	129
0.88(4)	2.88 (4)	3.581 (3)	138 (3)
0.95	2.70	3.564 (4)	151
	0.86 (4) 0.86 (4) 0.95 0.88 (4) 0.95	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$

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

Table 2	
Interatomic contacts of the title compound (Å).	

Contact	Distance	Symmetry operation
Cl1···Cl1	3.6744 (14)	2 - x, 1 - y, 1 - z
$H4 \cdot \cdot \cdot C20$	2.77	1-x, 1-y, 1-z
O1···H9	2.54	$x, \frac{1}{2} - y, -\frac{1}{2} + z$
$N2 \cdot \cdot \cdot H13C$	2.81	x, y, 1 + z
$H3A \cdot \cdot \cdot N2$	2.22 (4)	1 - x, 1 - y, 2 - z
$H16 \cdot \cdot \cdot N2$	2.62	2-x, 1-y, 2-z
H11···H13B	2.54	-1 + x, y, z
H17···H3B	2.54	1 + x, y, z
$H12 \cdot \cdot \cdot C18$	2.93	-1 + x, y, -1 + z

In this study, in the framework of our ongoing structural studies (Naghiyev *et al.*, 2020, 2021*a,b*), we report the crystal structure and Hirshfeld surface analysis of the title compound, (Z)-2-amino-4-(2,6-dichlorophenyl)-5-(1-hydroxyethylidene)-6-oxo-1-phenyl-1,4,5,6-tetrahydropyridine-3-carbonitrile, previously mistakenly reported in the *E* isomeric form (Maharramov *et al.*, 2018). This compound was also previously mentioned as transient intermediate but neither isolated nor characterized (Naghiyeva *et al.*, 2019).

2. Structural commentary

The title compound crystallizes in the monoclinic space group $P2_1/c$ with Z = 4, in which the asymmetric unit comprises one molecule. In the molecule (Fig. 1), the central pyridine ring (N1/C2–C6) is almost planar with a maximum deviation of 0.074 (3) Å for C4. The phenyl (C7–C12) and dichlorophenyl (C14–C19) rings are at an angle of 21.28 (15)°. They form dihedral angles of 86.10 (15) and 87.17 (14)°, respectively, with the central pyridine ring. The =C(–OH)CH₃ group is nearly coplanar with the pyridine ring with C2–C3–C1–O2 and C4–C3–C1–C13 torsion angles of only 5.5 (5) and 3.3 (5)°, respectively. A strong intramolecular O2–H2···O1 hydrogen bond (Fig. 1, Table 1) stabilizes the molecular conformation of the title molecule, creating an *S*(6) ring motif (Bernstein *et al.*, 1995).



Figure 1

The molecular structure of the title compound showing the atomnumbering scheme and displacement ellipsoids at the 50% probability level.

3. Supramolecular features and Hirshfeld surface analysis

Intermolecular N3-H3A···N2 hydrogen bonds, which form an $R_2^2(12)$ ring motif between pairs of molecules along the *b*axis direction and an $R_2^2(16)$ ring motif between pairs of molecules along the *a*-axis direction, together with N3-H3B···Cg2 and C9-H9···Cg2 interactions (Fig. 2, Tables 1 and 2; Cg2 is the centroid of the C7-C12 phenyl ring) create a three-dimensional network in the crystal (Figs. 2 and 3).



Figure 2

A general view of the intra- and intermolecular O-H···O, N-H···N hydrogen bonding and N-H··· π and C-H··· π interactions in the title compound. Symmetry codes: (a) 1 - x, 1 - y, 2 - z; (b) 2 - x, 1 - y, 2 - z; (c) x, $\frac{1}{2} - y$, $\frac{1}{2} + z$.



Figure 3

A view down the a axis of the crystal packing of the title compound based on the intermolecular interactions shown in Fig. 2.



Figure 4 Hirshfeld surface of the title compound mapped with d_{norm} .

The Hirshfeld surfaces were calculated and the twodimensional fingerprint plots generated using Crystal Explorer 17.5 (Turner et al., 2017). The use of various hues and intensities to represent short and long contacts, as well as the relative intensity of the connections, allows Hirshfeld surfaces to depict intermolecular interactions. Fig. 4 shows the threedimensional Hirshfeld surfaces of the title compound plotted over d_{norm} (normalized contact distance) in the range of -0.4290 to 1.5192 a.u. The red patches that appear around N2 are caused by the intermolecular N3-H3A···N2 and C16-H16 \cdots N2 interactions, which are important in the packing of the title molecule. Bright red dots near N2 and amine hydrogen atoms H3A and H3B highlight their functions as hydrogen-bonding acceptors and donors, respectively; these also appear as blue and red areas on the Hirshfeld surface mapped over electrostatic potential (Spackman et al., 2008) in Fig. 5, corresponding to positive and negative potentials. Positive electrostatic potential (hydrogen-bond donors) is



Figure 5

View of the three-dimensional Hirshfeld surface of the title compound plotted over electrostatic potential energy in the range -0.0500 to 0.0500 a.u. using the STO-3 G basis set at the Hartree–Fock level of theory. Hydrogen-bond donors and acceptors are shown as blue and red regions, respectively, around the atoms, corresponding to positive and negative potentials.

shown in blue, whereas negative electrostatic potential is indicated in red (hydrogen-bond acceptors).

In Fig. 6, the overall two-dimensional fingerprint plot for the title compound and those delineated into $H \cdots H$, $C \cdots H/H \cdots C$, $Cl \cdots H/H \cdots Cl$, $O \cdots H/H \cdots O$ and $N \cdots H/H \cdots N$ contacts, as well as their relative contributions to the Hirshfeld surface, are presented, while details of the various contacts are given in Table 2. The percentage contributions to the Hirshfeld surfaces from the various interatomic contacts are as follows: $H \cdots H$ (33.1%; Fig. 6b), $C \cdots H/H \cdots C$ (22.5%; Fig. 6c), $Cl \cdots H/H \cdots Cl$ (14.1%; Fig. 6d), $O \cdots H/H \cdots O$ (11.9%; Fig. 6e) and $N \cdots H/H \cdots N$ (9.7%; Fig. 6f). Other $Cl \cdots C/C \cdots Cl$, $C \cdots C$, $Cl \cdots O/O \cdots Cl$, $Cl \cdots N/N \cdots Cl$, $N \cdots C/C \cdots N$, $O \cdots N/N \cdots O$,







The two-dimensional fingerprint plots of the title compound, showing (a) all interactions, and delineated into (b) $H \cdots H$, (c) $C \cdots H/H \cdots C$, (d) $CI \cdots H/H \cdots CI$, (e) $O \cdots H/H \cdots O$ and (f) $N \cdots H/H \cdots N$ interactions [de and d_i represent the distances from a point on the Hirshfeld surface to the nearest atoms outside (external) and inside (internal) the surface, respectively].

d

Table 3	
Percentage contributions of interatomic contact	cts to the Hirshfeld surface
for the title compound.	

Contact	Percentage contribution
$H \cdots H$	33.1
$C \cdots H/H \cdots C$	22.5
$Cl \cdot \cdot \cdot H/H \cdot \cdot \cdot Cl$	14.1
$O \cdots H/H \cdots O$	11.9
$N \cdots H/H \cdots N$	9.7
$Cl \cdot \cdot \cdot C/C \cdot \cdot \cdot Cl$	2.1
$\mathbf{C}\!\cdot\!\cdot\!\cdot\!\mathbf{C}$	1.4
$Cl \cdot \cdot \cdot O/O \cdot \cdot \cdot Cl$	1.2
$Cl \cdot \cdot \cdot N/N \cdot \cdot \cdot Cl$	1.1
$N \cdots C/C \cdots N$	1.0
$O \cdots N/N \cdots O$	0.6
Cl···Cl	0.6
$O \cdots C/C \cdots O$	0.5
$N \cdots N$	0.1

Cl···Cl, O···C/C···O and N···N contacts contribute less than 2.1% to Hirshfeld surface mapping and have little directional influence on molecular packing (Table 3).

4. Database survey

A search of the Cambridge Structural Database (CSD, Version 5.39, update of August 2018; Groom *et al.*, 2016) using *Conquest* (Bruno *et al.*, 2002) for the tetrahydropyridine unit revealed 1339 hits. Some interesting structures related to the title compound based on their tetrahydropyridine moieties include: ethyl 4-hydroxy-2,6-diphenyl-5-(phenylsulfanyl)pyridine-3-carboxylate (refcode SETWOE: Suresh *et al.*, 2007), ethyl 2,6-bis(4-fluorophenyl)-4-hydroxy-5-(4-methylphenyl-sulfanyl)pyridine-3-carboxylate (SETWUK: Suresh *et al.*, 2007), 2,6-diamino-4-chloropyrimidin-1-ium 2-carboxy-3-nitrobenzoate (JEBRAM: Mohana *et al.*, 2017) and 2,6-diamino-4-chloropyrimidin-1-sulfonate monohydrate (JEBREQ: Mohana *et al.*, 2017).

The polysubstituted pyridines, SETWOE (space group: $P2_1/c$) and SETWUK (space group: $P2_1/n$), adopt nearly planar structures. The crystal structure of SETWOE is stabilized by intermolecular C-H···O and C-H··· π interactions. The C-H···O hydrogen bonds generate rings with $R_2^2(14)$ and $R_2^2(20)$ motifs. The crystal structure of SETWUK is stabilized by intermolecular C-H···F and C-H··· π interactions. The C-H···F bond generates a linear chain with a C(14) motif. In addition, in SETWOE and SETWUK, intramolecular O-H···O interactions are found, which generate an S(6) graph-set motif. No significant aryl-aryl or π - π interactions exist in these structures. All this bears some resemblance to the title compound.

In both the related salts, JEBRAM (space group: $P\overline{1}$) and JEBREQ (space group: $P\overline{1}$), the N atom in the 1-position of the pyrimidine ring is protonated. In JEBRAM, the protonated N atom and the amino group of the pyrimidinium cation interact with the carboxylate group of the anion through N-H···O hydrogen bonds, forming a heterosynthon with an $R_2^2(8)$ ring motif. In the hydrated salt JEBREQ, the presence of the water molecule prevents the formation of the familiar

Table 4Experimental details.	
Crystal data	
Chemical formula	$C_{20}H_{15}Cl_2N_3O_2$
M _r	400.25
Crystal system, space group	Monoclinic, $P2_1/c$
Temperature (K)	100
a, b, c (Å)	9.662 (1), 27.010 (3), 7.4782 (8)
β (°)	111.571 (2)
$V(Å^3)$	1814.9 (3)
Ζ	4
Radiation type	Μο Κα
$\mu (\text{mm}^{-1})$	0.38
Crystal size (mm)	$0.24 \times 0.21 \times 0.02$
Data collection	
Diffractometer	Bruker D8 QUEST PHOTON-III CCD
Absorption correction	Multi-scan (SADABS; Krause et al., 2015)
T_{\min}, T_{\max}	0.864, 0.986
No. of measured, independent and observed $[I > 2\sigma(I)]$ reflections	27440, 4180, 2631
R _{int}	0.099
$(\sin \theta/\lambda)_{\rm max} ({\rm \AA}^{-1})$	0.650
Refinement	
$R[F^2 > 2\sigma(F^2)], wR(F^2), S$	0.053, 0.126, 1.01
No. of reflections	4180
No. of parameters	255
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta ho_{ m max}, \Delta ho_{ m min} ({ m e} \; { m \AA}^{-3})$	0.50, -0.37

Computer programs: APEX3 (Bruker, 2018), SAINT (Bruker, 2013), SHELXT2014/5 (Sheldrick, 2015a), SHELXL2018/3 (Sheldrick, 2015b), ORTEP-3 for Windows (Farrugia, 2012) and PLATON (Spek, 2020).

 $R_2^2(8)$ ring motif. Instead, an expanded ring [*i.e.* $R_2^3(8)$] is formed involving the sulfonate group, the pyrimidinium cation and the water molecule. Both salts form a supramolecular homosynthon [$R_2^2(8)$ ring motif] through N-H···N hydrogen bonds. The molecular structures are further stabilized by π - π stacking, and C=O··· π , C-H···O and C-H···Cl interactions. None of these are found in the crystal packing of the title compound. It appears that the protonation state of the pyrimidine ring influences the intermolecular interactions within the crystal lattices to a substantial extent.

5. Synthesis and crystallization

The title compound was synthesized using our previously reported procedure (Maharramov *et al.*, 2018), and colorless prisms were obtained upon recrystallization from its methanol solution.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 4. The positional parameters of the H atoms of the hydroxy and amine groups were determined from difference electron-density maps and were refined freely [O2-H2 = 0.86 (4) Å, N3-H3A = 0.86 (4) Å and N3-H3B =0.88 (4) Å]. Their isotropic displacement parameters were refined using a riding model with $U_{iso}(H)$ set to either $1.2U_{eq}(N)$ for the NH₂ group or $1.5U_{eq}(O)$ for the OH group. The C-bound H atoms were positioned geometrically (C-H = 0.95–1.00 Å) and allowed to ride on their parent atoms, with $U_{iso}(H) = 1.5U_{eq}(C)$ for the methyl group and $U_{iso}(H) = 1.2U_{eq}(C)$ for aromatic and methine H atoms.

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Authors' contributions are as follows. Conceptualization, FNN and IGM; methodology, FNN and IGM; investigation, FNN, AVP and AAA; writing (original draft), MA and ANK; writing (review and editing of the manuscript), MA and ANK; visualization, MA, FNN and IGM; funding acquisition, VNK and FNN; resources, AAA, VNK and FNN; supervision, IGM and MA.

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Crystal structure and Hirshfeld surface analysis of (*Z*)-2-amino-4-(2,6-dichloro-phenyl)-5-(1-hydroxyethylidene)-6-oxo-1-phenyl-1,4,5,6-tetrahydropyridine-3-carbonitrile

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Computing details

Data collection: *APEX3* (Bruker, 2018); cell refinement: *SAINT* (Bruker, 2013); data reduction: *SAINT* (Bruker, 2013); program(s) used to solve structure: *SHELXT2014/5* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *PLATON* (Spek, 2020).

(Z) - 2 - Amino - 4 - (2, 6 - dichlorophenyl) - 5 - (1 - hydroxyethylidene) - 6 - oxo - 1 - phenyl - 1, 4, 5, 6 - tetrahydropyridine - 3 - carbonitrile

Crystal data

 $C_{20}H_{15}Cl_2N_3O_2$ $M_r = 400.25$ Monoclinic, $P2_1/c$ a = 9.662 (1) Å b = 27.010 (3) Å c = 7.4782 (8) Å $\beta = 111.571$ (2)° V = 1814.9 (3) Å³ Z = 4

Data collection

Bruker D8 QUEST PHOTON-III CCD diffractometer φ and ω scans Absorption correction: multi-scan (SADABS; Krause *et al.*, 2015) $T_{\min} = 0.864$, $T_{\max} = 0.986$ 27440 measured reflections

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.053$ $wR(F^2) = 0.126$ S = 1.014180 reflections F(000) = 824 $D_x = 1.465 \text{ Mg m}^{-3}$ Mo K\alpha radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 2887 reflections $\theta = 2.3-25.6^{\circ}$ $\mu = 0.38 \text{ mm}^{-1}$ T = 100 KPlate, colourless $0.24 \times 0.21 \times 0.02 \text{ mm}$

4180 independent reflections 2631 reflections with $I > 2\sigma(I)$ $R_{int} = 0.099$ $\theta_{max} = 27.5^\circ, \ \theta_{min} = 2.3^\circ$ $h = -12 \rightarrow 12$ $k = -35 \rightarrow 35$ $l = -9 \rightarrow 9$

255 parameters 0 restraints Primary atom site location: dual Secondary atom site location: dual Hydrogen site location: mixed H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0387P)^2 + 2.0157P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$
$$\begin{split} &\Delta\rho_{\rm max} = 0.50 \ {\rm e} \ {\rm \AA}^{-3} \\ &\Delta\rho_{\rm min} = -0.37 \ {\rm e} \ {\rm \AA}^{-3} \\ & {\rm Extinction \ correction: \ SHELXL,} \\ & {\rm Fc}^* = {\rm kFc} [1 + 0.001 {\rm xFc}^2 \lambda^3 / {\rm sin}(2\theta)]^{-1/4} \\ & {\rm Extinction \ coefficient: \ } 0.0024 \ (2) \end{split}$$

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.

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
Cl1	0.86090 (9)	0.51416 (3)	0.60657 (12)	0.0256 (2)	
C12	0.73128 (9)	0.32459 (3)	0.75788 (12)	0.0262 (2)	
01	0.3936 (2)	0.32320 (9)	0.2910 (3)	0.0280 (5)	
O2	0.5668 (3)	0.34472 (9)	0.1210 (3)	0.0286 (6)	
H2	0.502 (4)	0.3295 (15)	0.155 (6)	0.043*	
N1	0.3832 (3)	0.36887 (10)	0.5389 (4)	0.0190 (6)	
N2	0.6028 (3)	0.51570 (10)	0.8819 (4)	0.0213 (6)	
N3	0.3468 (3)	0.41829 (11)	0.7719 (4)	0.0216 (6)	
H3A	0.376 (4)	0.4411 (13)	0.857 (5)	0.026*	
H3B	0.275 (4)	0.3976 (13)	0.763 (5)	0.026*	
C1	0.6193 (3)	0.38087 (12)	0.2516 (4)	0.0222 (7)	
C2	0.4484 (3)	0.35814 (12)	0.4048 (4)	0.0215 (7)	
C3	0.5700 (3)	0.38792 (12)	0.3994 (4)	0.0190 (7)	
C4	0.6393 (3)	0.42841 (12)	0.5454 (4)	0.0185 (7)	
H4	0.6402	0.4592	0.4717	0.022*	
C5	0.5438 (3)	0.43903 (11)	0.6633 (4)	0.0178 (6)	
C6	0.4288 (3)	0.40975 (12)	0.6622 (4)	0.0190 (6)	
C7	0.2636 (3)	0.33754 (11)	0.5457 (4)	0.0182 (6)	
C8	0.2954 (3)	0.29874 (12)	0.6757 (4)	0.0225 (7)	
H8	0.3954	0.2923	0.7572	0.027*	
C9	0.1808 (4)	0.26931 (12)	0.6866 (5)	0.0247 (7)	
H9	0.2018	0.2428	0.7762	0.030*	
C10	0.0355 (4)	0.27893 (12)	0.5655 (5)	0.0254 (7)	
H10	-0.0431	0.2588	0.5722	0.030*	
C11	0.0039 (3)	0.31762 (12)	0.4351 (5)	0.0233 (7)	
H11	-0.0959	0.3239	0.3526	0.028*	
C12	0.1182 (3)	0.34720 (12)	0.4251 (4)	0.0219 (7)	
H12	0.0970	0.3739	0.3362	0.026*	
C13	0.7339 (4)	0.41270 (13)	0.2175 (5)	0.0262 (7)	
H13A	0.7588	0.3988	0.1121	0.039*	
H13B	0.8235	0.4139	0.3346	0.039*	
H13C	0.6943	0.4462	0.1834	0.039*	
C14	0.8003 (3)	0.41904 (12)	0.6829 (4)	0.0168 (6)	
C15	0.9070 (3)	0.45717 (12)	0.7254 (4)	0.0206 (7)	

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

C16	1.0502 (3)	0.45270 (13)	0.8591 (5)	0.0237 (7)	
H16	1.1177	0.4797	0.8851	0.028*	
C17	1.0934 (3)	0.40785 (13)	0.9546 (4)	0.0252 (7)	
H17	1.1915	0.4041	1.0469	0.030*	
C18	0.9949 (3)	0.36865 (13)	0.9164 (5)	0.0238 (7)	
H18	1.0255	0.3378	0.9793	0.029*	
C19	0.8510(3)	0.37491 (12)	0.7852 (4)	0.0194 (7)	
C20	0.5773 (3)	0.48144 (12)	0.7841 (4)	0.0190 (7)	

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

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	<i>U</i> ²³
Cl1	0.0223 (4)	0.0243 (4)	0.0291 (4)	-0.0013 (3)	0.0083 (3)	0.0025 (4)
Cl2	0.0236 (4)	0.0258 (4)	0.0260 (4)	-0.0006 (3)	0.0056 (3)	0.0049 (4)
01	0.0271 (12)	0.0342 (14)	0.0240 (12)	-0.0085 (11)	0.0108 (10)	-0.0124 (11)
O2	0.0296 (13)	0.0394 (15)	0.0171 (12)	-0.0043 (11)	0.0091 (10)	-0.0072 (11)
N1	0.0169 (13)	0.0243 (15)	0.0161 (13)	-0.0028 (11)	0.0063 (10)	-0.0035 (11)
N2	0.0213 (14)	0.0210 (14)	0.0200 (14)	-0.0026 (11)	0.0058 (11)	-0.0030 (12)
N3	0.0199 (14)	0.0270 (15)	0.0196 (14)	-0.0071 (12)	0.0095 (11)	-0.0081 (12)
C1	0.0210 (16)	0.0286 (18)	0.0135 (15)	0.0038 (13)	0.0022 (12)	-0.0012 (14)
C2	0.0200 (15)	0.0275 (18)	0.0160 (15)	0.0009 (13)	0.0053 (12)	-0.0023 (14)
C3	0.0174 (15)	0.0225 (17)	0.0152 (15)	-0.0002 (12)	0.0038 (12)	-0.0001 (13)
C4	0.0157 (15)	0.0228 (17)	0.0166 (15)	0.0014 (12)	0.0055 (12)	-0.0020 (13)
C5	0.0172 (15)	0.0207 (16)	0.0139 (15)	0.0005 (12)	0.0037 (12)	-0.0014 (13)
C6	0.0162 (14)	0.0237 (17)	0.0135 (15)	0.0036 (12)	0.0012 (12)	-0.0002 (13)
C7	0.0175 (15)	0.0185 (16)	0.0184 (15)	-0.0037 (12)	0.0062 (12)	-0.0048 (13)
C8	0.0201 (16)	0.0264 (18)	0.0174 (16)	0.0015 (13)	0.0026 (13)	-0.0003 (14)
C9	0.0275 (17)	0.0219 (17)	0.0254 (18)	0.0030 (14)	0.0106 (14)	0.0049 (14)
C10	0.0228 (16)	0.0245 (18)	0.0310 (19)	-0.0028 (14)	0.0124 (14)	-0.0042 (15)
C11	0.0177 (15)	0.0259 (18)	0.0223 (17)	0.0019 (13)	0.0024 (13)	0.0013 (14)
C12	0.0230 (16)	0.0224 (17)	0.0177 (16)	0.0031 (13)	0.0044 (13)	0.0001 (13)
C13	0.0264 (17)	0.033 (2)	0.0206 (17)	0.0037 (15)	0.0107 (14)	0.0041 (15)
C14	0.0152 (14)	0.0241 (16)	0.0126 (14)	0.0025 (12)	0.0070 (11)	-0.0011 (12)
C15	0.0200 (15)	0.0254 (17)	0.0179 (16)	0.0040 (13)	0.0088 (13)	0.0024 (14)
C16	0.0185 (15)	0.0315 (19)	0.0222 (17)	-0.0020 (13)	0.0089 (13)	-0.0035 (14)
C17	0.0162 (15)	0.042 (2)	0.0173 (16)	0.0027 (14)	0.0063 (13)	-0.0010 (15)
C18	0.0219 (16)	0.0307 (19)	0.0196 (16)	0.0047 (14)	0.0087 (13)	0.0047 (14)
C19	0.0174 (15)	0.0266 (17)	0.0150 (15)	-0.0009 (13)	0.0066 (12)	-0.0022 (13)
C20	0.0110 (14)	0.0270 (18)	0.0193 (16)	0.0044 (12)	0.0058 (12)	0.0062 (14)

Geometric parameters (Å, °)

Cl1—C15	1.751 (3)	C7—C12	1.387 (4)	
Cl2—C19	1.747 (3)	C8—C9	1.390 (4)	
O1—C2	1.250 (4)	C8—H8	0.9500	
O2—C1	1.342 (4)	C9—C10	1.387 (4)	
O2—H2	0.86 (4)	С9—Н9	0.9500	
N1—C2	1.397 (4)	C10—C11	1.385 (5)	

N1—C6	1.401 (4)	C10—H10	0.9500
N1—C7	1.448 (4)	C11—C12	1.387 (4)
N2—C20	1.149 (4)	C11—H11	0.9500
N3—C6	1.354 (4)	C12—H12	0.9500
N3—H3A	0.86 (4)	С13—Н13А	0.9800
N3—H3B	0.88 (4)	С13—Н13В	0.9800
C1—C3	1.368 (4)	С13—Н13С	0.9800
C1—C13	1.496 (4)	C14—C19	1.404 (4)
C2—C3	1.437 (4)	C14—C15	1.408 (4)
C3—C4	1.516 (4)	C15—C16	1.382 (4)
C4—C5	1.519 (4)	C16—C17	1.390 (5)
C4—C14	1.538 (4)	С16—Н16	0.9500
C4—H4	1.0000	C17—C18	1.382 (5)
C5—C6	1.361 (4)	С17—Н17	0.9500
C5-C20	1 421 (4)	C18 - C19	1 386 (4)
C7—C8	1 385 (4)	C18—H18	0.9500
0, 00	1.505 (1)		0.9500
C1—O2—H2	105 (3)	С8—С9—Н9	120.3
C_2 —N1—C6	121.3 (3)	C11—C10—C9	120.6(3)
C2—N1—C7	118.7 (2)	C11—C10—H10	119.7
C6—N1—C7	120.0 (2)	C9—C10—H10	119.7
C6—N3—H3A	118 (2)	C10—C11—C12	120.0 (3)
C6—N3—H3B	118 (2)	C10—C11—H11	120.0
H3A—N3—H3B	123 (3)	C12—C11—H11	120.0
O2—C1—C3	122.6 (3)	C11—C12—C7	119.5 (3)
O2—C1—C13	113.5 (3)	C11—C12—H12	120.2
C3—C1—C13	123.8 (3)	C7—C12—H12	120.2
01—C2—N1	117.1 (3)	C1—C13—H13A	109.5
O1—C2—C3	123.3 (3)	C1—C13—H13B	109.5
N1—C2—C3	119.6 (3)	H13A—C13—H13B	109.5
C1—C3—C2	118.4 (3)	C1—C13—H13C	109.5
C1—C3—C4	119.4 (3)	H13A—C13—H13C	109.5
C2—C3—C4	122.1 (3)	H13B—C13—H13C	109.5
C3—C4—C5	110.7 (2)	C19—C14—C15	114.7 (3)
C3—C4—C14	115.6 (2)	C19—C14—C4	124.6 (3)
C5—C4—C14	108.9 (2)	C15—C14—C4	120.5 (3)
C3—C4—H4	107.1	C16—C15—C14	123.7 (3)
C5—C4—H4	107.1	C16—C15—Cl1	116.4 (3)
C14—C4—H4	107.1	C14—C15—Cl1	120.0 (2)
C6—C5—C20	117.8 (3)	C15—C16—C17	118.7 (3)
C6—C5—C4	123.7 (3)	C15—C16—H16	120.7
C20—C5—C4	118.4 (3)	C17—C16—H16	120.7
N3—C6—C5	123.7 (3)	C18—C17—C16	120.5 (3)
N3—C6—N1	114.8 (3)	C18—C17—H17	119.8
C5—C6—N1	121.4 (3)	C16—C17—H17	119.8
C8—C7—C12	120.6 (3)	C17—C18—C19	119.2 (3)
C8—C7—N1	119.5 (3)	C17—C18—H18	120.4
C12—C7—N1	119.9 (3)	C19—C18—H18	120.4

С7—С8—С9	119.9 (3)	C18—C19—C14	123.2 (3)
С7—С8—Н8	120.1	C18—C19—Cl2	115.9 (3)
С9—С8—Н8	120.1	C14—C19—Cl2	120.9 (2)
C10—C9—C8	119.5 (3)	N2-C20-C5	179.2 (3)
С10—С9—Н9	120.3		
C6—N1—C2—O1	-174.8 (3)	C6—N1—C7—C8	-86.5 (4)
C7—N1—C2—O1	3.5 (4)	C2—N1—C7—C12	-86.2 (4)
C6—N1—C2—C3	3.4 (4)	C6—N1—C7—C12	92.2 (3)
C7—N1—C2—C3	-178.2 (3)	C12—C7—C8—C9	-0.4 (5)
O2—C1—C3—C2	-5.5 (5)	N1-C7-C8-C9	178.3 (3)
C13—C1—C3—C2	173.5 (3)	C7—C8—C9—C10	0.5 (5)
O2—C1—C3—C4	177.7 (3)	C8—C9—C10—C11	-0.3 (5)
C13—C1—C3—C4	-3.3 (5)	C9—C10—C11—C12	-0.1 (5)
O1—C2—C3—C1	6.4 (5)	C10-C11-C12-C7	0.3 (5)
N1—C2—C3—C1	-171.7 (3)	C8—C7—C12—C11	0.0 (5)
O1—C2—C3—C4	-176.9 (3)	N1-C7-C12-C11	-178.7 (3)
N1-C2-C3-C4	5.0 (4)	C3—C4—C14—C19	-48.4 (4)
C1—C3—C4—C5	165.1 (3)	C5—C4—C14—C19	76.9 (4)
C2—C3—C4—C5	-11.6 (4)	C3—C4—C14—C15	136.5 (3)
C1—C3—C4—C14	-70.5 (4)	C5-C4-C14-C15	-98.2 (3)
C2-C3-C4-C14	112.8 (3)	C19—C14—C15—C16	-1.0 (4)
C3—C4—C5—C6	11.6 (4)	C4-C14-C15-C16	174.6 (3)
C14—C4—C5—C6	-116.5 (3)	C19—C14—C15—Cl1	179.1 (2)
C3—C4—C5—C20	-169.4 (3)	C4—C14—C15—Cl1	-5.3 (4)
C14—C4—C5—C20	62.5 (4)	C14—C15—C16—C17	1.4 (5)
C20—C5—C6—N3	-1.3 (5)	Cl1—C15—C16—C17	-178.7 (2)
C4—C5—C6—N3	177.7 (3)	C15—C16—C17—C18	0.0 (5)
C20—C5—C6—N1	176.3 (3)	C16—C17—C18—C19	-1.7 (5)
C4—C5—C6—N1	-4.7 (5)	C17—C18—C19—C14	2.2 (5)
C2—N1—C6—N3	174.1 (3)	C17—C18—C19—Cl2	-176.1 (2)
C7—N1—C6—N3	-4.2 (4)	C15—C14—C19—C18	-0.8 (4)
C2—N1—C6—C5	-3.6 (4)	C4-C14-C19-C18	-176.2 (3)
C7—N1—C6—C5	178.0 (3)	C15—C14—C19—Cl2	177.3 (2)
C2—N1—C7—C8	95.1 (3)	C4—C14—C19—Cl2	1.9 (4)

Hydrogen-bond geometry (Å, °)

Cg2 is the centroid of the C7–C12 phenyl ring.

D—H···A	D—H	H···A	D····A	<i>D</i> —H··· <i>A</i>
O2—H2…O1	0.86 (4)	1.72 (4)	2.514 (3)	153 (4)
N3—H3A···N2 ⁱ	0.86 (4)	2.22 (4)	3.032 (4)	159 (3)
C16—H16…N2 ⁱⁱ	0.95	2.62	3.308 (4)	129
N3—H3 <i>B</i> ··· <i>Cg</i> 2	0.88 (4)	2.88 (4)	3.581 (3)	138 (3)
С9—Н9…Сg2 ^{ііі}	0.95	2.70	3.564 (4)	151

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