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rac-2-(2-Chloro-6-methylquinolin-3-yl)-2,3-dihydroquinolin-4(1*H*)-one

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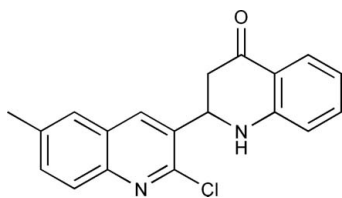
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.062; wR factor = 0.169; data-to-parameter ratio = 16.7.

In the title compound, $\text{C}_{19}\text{H}_{15}\text{ClN}_2\text{O}$, the quinoline ring forms a dihedral angle of $43.24(1)^\circ$ with the benzene ring of the dihydroquinolinyl system. In the crystal, molecules are linked through a single weak $\text{C}-\text{H}\cdots\text{O}$ hydrogen bond, forming ribbons which extend along (100), giving alternating zigzag molecular layers which stack down the b -axis direction.

Related literature

For applications of similar structures see: Chandrasekhar *et al.* (2007); Varma & Saini (1997); Donnelly & Farrell (1990); Hemanth Kumar *et al.* (2004). For the synthesis of the 2-aminochalcone, see: Gao *et al.* (1996). For related structures, see: Bouraiou *et al.* (2008, 2011); Belfaitah *et al.* (2006); Benzerka *et al.* (2011).



Experimental

Crystal data

$\text{C}_{19}\text{H}_{15}\text{ClN}_2\text{O}$
 $M_r = 322.78$
 Orthorhombic, $Pbca$
 $a = 13.8912(8)$ Å
 $b = 12.4572(4)$ Å
 $c = 17.8617(11)$ Å

$V = 3090.9(3)$ Å³
 $Z = 8$
 Mo $K\alpha$ radiation
 $\mu = 0.25$ mm⁻¹
 $T = 295$ K
 $0.15 \times 0.06 \times 0.05$ mm

Data collection

Nonius KappaCCD diffractometer
 6664 measured reflections
 3537 independent reflections
 1696 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.072$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.169$
 $S = 1.00$
 3537 reflections
 212 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.21$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.27$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C17}-\text{H17}\cdots\text{O1}^i$	0.93	2.49	3.243 (5)	138

 Symmetry code: (i) $x - \frac{1}{2}, y, -z + \frac{1}{2}$.

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg & Berndt, 2001); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: ZS2200).

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supporting information

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rac*-2-(2-Chloro-6-methylquinolin-3-yl)-2,3-dihydroquinolin-4(1*H*)-one*Abdelmalek Bouraiou, Sofiane Bouacida, Carboni Bertrand, Thierry Roisnel and Ali Belfaitah****S1. Comment**

2-Substituted dihydroquinolinones have important medicinal properties as new chemical entities and also serve as building blocks for creating further diversity in SAR studies in various therapeutic areas (Chandrasekhar *et al.*, 2007). Convenient synthesis of 2-aminochalcone and its amide derivatives and the ready cyclization of these compounds to 2-aryl-2,3-dihydroquinolin-4(1*H*)-ones have been widely explored (Varma & Saini, 1997; Donnelly & Farrell, 1990). Silica gel supported InCl₃ (20 mol %) is a new solid-support catalyst that can be used under solvent-free conditions for the facile and efficient isomerization of 2-aminochalcones to the corresponding 2-aryl-2,3-dihydroquinolin-4(1*H*)-ones (Hemanth Kumar *et al.*, 2004). As a part of a program directed toward the synthesis of new suitably functionalized heterocyclic compounds of potential biological activity (Bouraiou *et al.*, 2008, 2011; Benzerka *et al.*, 2011), we report herein the synthesis and structure determination of the title compound, C₁₉H₁₅ClN₂O.

In the title compound (Fig. 1), the quinoline ring forms a dihedral angle of 43.24 (1)° with the phenyl ring of the 2,3-dihydroquinolin-4(1*H*)-one moiety. The geometric parameters are in agreement with those of other structures possessing a quinolyl substituent, previously reported in the literature (Belfaitah *et al.*, 2006; Benzerka *et al.*, 2011). The crystal structure can be described as alternating zigzag ribbons which stack down the *b* axis of the unit cell (Fig. 2), these ribbons comprising molecules linked through a single weak intermolecular C17—H···O1 hydrogen bond (Table 1), and extending down *a* (Fig. 3). The hetero N2—H2 group has no acceptor in the crystal structure.

S2. Experimental

2-Aminoacetophenone (1mmol) was first condensed with 2-chloro-3-formyl-6-methylquinoline (2 mmol) to give the corresponding 2-aminochalcone in 74% yield, according to the procedure described by Gao *et al.* (1996). In the next step, a mixture of 2-aminochalcone (100 mg) and 1 g of silica gel impregnated with indium(III) chloride (20 mol%, 13.6 mg) was irradiated in a domestic microwave oven at 360 W for 5 minutes. Under these conditions, the title compound was successfully synthesized in good yield (69%). Single crystals suitable for the X-ray diffraction analysis were obtained by dissolving the compound in a diisopropyl ether/CHCl₃ solvent mixture and allowing the solution to slowly evaporate at room temperature.

S3. Refinement

The N-bound H-atom (H2) was located in a difference-Fourier map and its positional parameters were refined isotropically. All other H atoms were introduced in calculated positions and treated as riding on their parent C atom, with C—H = 0.93, 0.96, 0.97 or 0.98 Å, with $U_{\text{iso}}(\text{H}) = 1.2$ or $1.5 U_{\text{eq}}(\text{C})$. No H-bond acceptor could be located for the N2—H2 group.

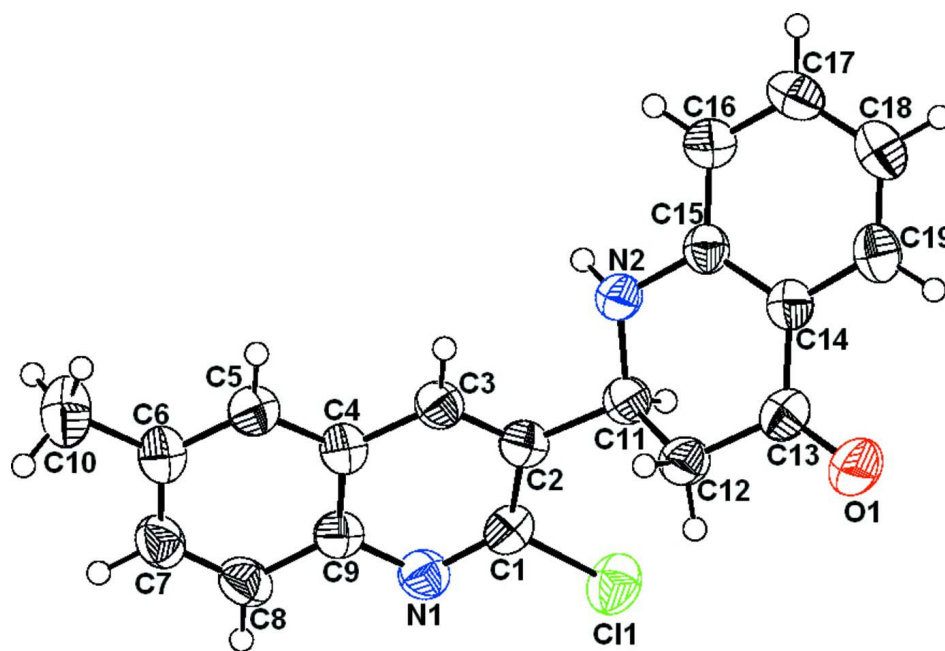


Figure 1

The structure of the title compound with the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.

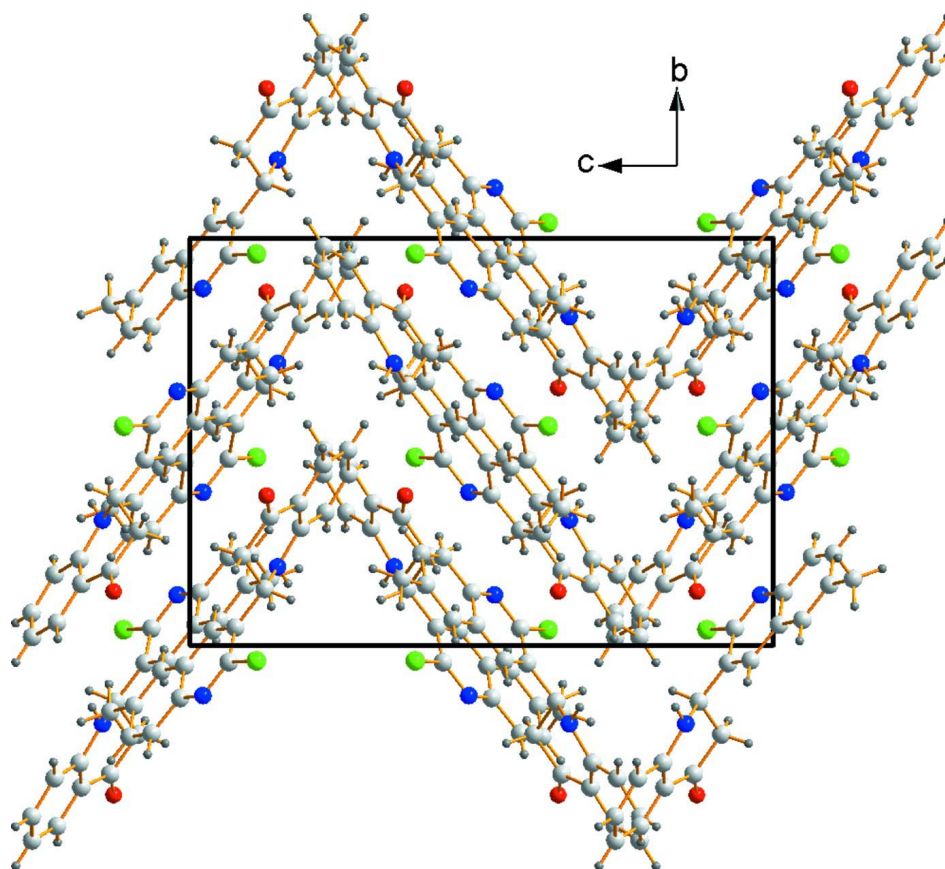


Figure 2

A diagram of the layered zigzag packing in the crystal viewed down the a axis.

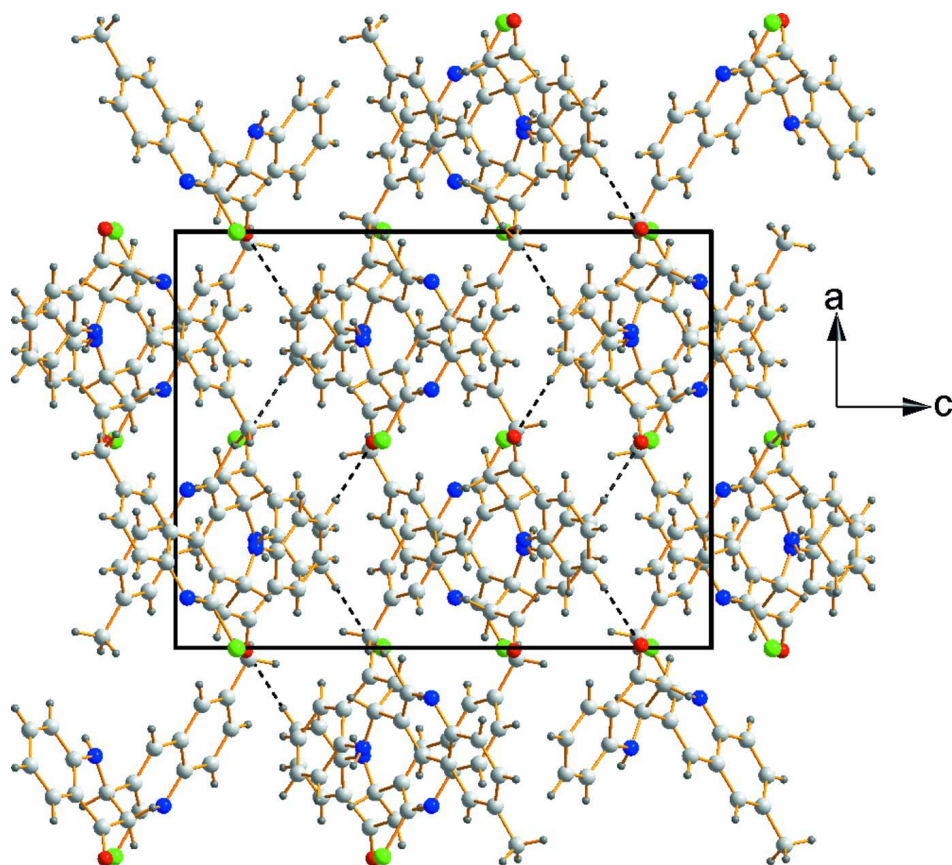


Figure 3

A part of crystal packing viewed down the b axis showing hydrogen-bond interactions as dashed lines.

rac-2-(2-Chloro-6-methylquinolin-3-yl)-2,3-dihydroquinolin-4(1H)-one

Crystal data

$C_{19}H_{15}ClN_2O$

$M_r = 322.78$

Orthorhombic, $Pbca$

Hall symbol: $-P\ 2ac\ 2ab$

$a = 13.8912\ (8)\ \text{\AA}$

$b = 12.4572\ (4)\ \text{\AA}$

$c = 17.8617\ (11)\ \text{\AA}$

$V = 3090.9\ (3)\ \text{\AA}^3$

$Z = 8$

$F(000) = 1344$

$D_x = 1.387\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 3949 reflections

$\theta = 2.9\text{--}27.5^\circ$

$\mu = 0.25\ \text{mm}^{-1}$

$T = 295\ \text{K}$

Needle, colourless

$0.15 \times 0.06 \times 0.05\ \text{mm}$

Data collection

Nonius KappaCCD

diffractometer

Radiation source: Enraf Nonius FR590

diffractometer

Graphite monochromator

Detector resolution: 9 pixels mm^{-1}

CCD rotation images, thick slices scans

6664 measured reflections

3537 independent reflections

1696 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.072$

$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 2.9^\circ$

$h = -18 \rightarrow 17$

$k = -16 \rightarrow 16$

$l = -23 \rightarrow 23$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.062$
 $wR(F^2) = 0.169$
 $S = 1.00$
 3537 reflections
 212 parameters
 0 restraints
 Primary atom site location: structure-invariant
 direct methods

Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H atoms treated by a mixture of independent
 and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0676P)^2 + 0.5198P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.21 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	1.00123 (7)	0.53833 (7)	0.38618 (6)	0.0598 (3)
O1	1.00620 (19)	0.1322 (2)	0.36882 (17)	0.0724 (9)
N1	0.8795 (2)	0.6237 (2)	0.47797 (17)	0.0467 (7)
N2	0.7597 (2)	0.3073 (2)	0.34987 (16)	0.0461 (7)
C1	0.8911 (2)	0.5416 (2)	0.43385 (19)	0.0430 (8)
C2	0.8250 (2)	0.4560 (2)	0.42234 (18)	0.0408 (7)
C3	0.7405 (2)	0.4630 (2)	0.46099 (18)	0.0430 (8)
H3	0.6944	0.4096	0.4554	0.052*
C4	0.7221 (2)	0.5504 (2)	0.50953 (18)	0.0411 (8)
C5	0.6356 (2)	0.5626 (2)	0.54971 (19)	0.0453 (8)
H5	0.5879	0.5108	0.5447	0.054*
C6	0.6196 (3)	0.6486 (2)	0.59604 (19)	0.0474 (9)
C7	0.6938 (3)	0.7251 (3)	0.60359 (19)	0.0515 (9)
H7	0.6842	0.7834	0.6353	0.062*
C8	0.7791 (3)	0.7169 (2)	0.56608 (19)	0.0508 (9)
H8	0.8269	0.7682	0.5728	0.061*
C9	0.7940 (2)	0.6298 (2)	0.51708 (19)	0.0424 (8)
C10	0.5261 (3)	0.6631 (3)	0.6368 (2)	0.0687 (12)
H10A	0.4871	0.7146	0.6107	0.103*
H10B	0.5385	0.6883	0.6867	0.103*
H10C	0.4927	0.5957	0.6391	0.103*
C11	0.8477 (2)	0.3617 (2)	0.37226 (19)	0.0442 (8)
H11	0.8803	0.3883	0.3273	0.053*
C12	0.9131 (3)	0.2810 (2)	0.41094 (19)	0.0479 (9)

H12A	0.9741	0.3151	0.4226	0.058*
H12B	0.8838	0.2588	0.4577	0.058*
C13	0.9313 (3)	0.1834 (3)	0.3631 (2)	0.0475 (8)
C14	0.8524 (2)	0.1510 (2)	0.31357 (18)	0.0420 (8)
C15	0.7678 (2)	0.2110 (2)	0.31041 (18)	0.0412 (8)
C16	0.6899 (3)	0.1735 (3)	0.2677 (2)	0.0523 (9)
H16	0.633	0.2127	0.266	0.063*
C17	0.6976 (3)	0.0793 (3)	0.2284 (2)	0.0534 (9)
H17	0.6453	0.0546	0.2008	0.064*
C18	0.7825 (3)	0.0204 (3)	0.2292 (2)	0.0559 (9)
H18	0.7878	-0.0423	0.2012	0.067*
C19	0.8584 (3)	0.0556 (2)	0.2717 (2)	0.0505 (9)
H19	0.915	0.0157	0.2728	0.061*
H2	0.717 (3)	0.348 (2)	0.3321 (19)	0.05*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0485 (5)	0.0545 (5)	0.0763 (7)	-0.0051 (4)	0.0144 (5)	0.0027 (5)
O1	0.0539 (18)	0.0643 (16)	0.099 (2)	0.0201 (13)	-0.0134 (16)	-0.0173 (15)
N1	0.0479 (18)	0.0388 (14)	0.0534 (17)	-0.0029 (13)	-0.0024 (14)	-0.0010 (13)
N2	0.0408 (17)	0.0419 (15)	0.0554 (18)	0.0075 (12)	-0.0080 (14)	-0.0077 (13)
C1	0.0382 (19)	0.0412 (17)	0.050 (2)	0.0007 (14)	0.0014 (15)	0.0067 (15)
C2	0.0450 (19)	0.0353 (16)	0.0421 (18)	0.0008 (14)	-0.0046 (16)	0.0041 (14)
C3	0.045 (2)	0.0360 (16)	0.0482 (19)	-0.0027 (14)	-0.0011 (16)	-0.0015 (15)
C4	0.047 (2)	0.0352 (15)	0.0412 (18)	0.0007 (15)	0.0025 (15)	0.0043 (14)
C5	0.047 (2)	0.0378 (16)	0.051 (2)	-0.0021 (14)	-0.0016 (17)	0.0033 (15)
C6	0.057 (2)	0.0407 (17)	0.045 (2)	0.0054 (16)	0.0057 (17)	0.0048 (15)
C7	0.062 (2)	0.0438 (18)	0.049 (2)	0.0108 (17)	-0.0084 (19)	-0.0058 (16)
C8	0.059 (2)	0.0394 (17)	0.053 (2)	-0.0029 (16)	-0.0091 (19)	-0.0063 (16)
C9	0.043 (2)	0.0382 (16)	0.0454 (19)	-0.0004 (14)	-0.0040 (16)	0.0028 (14)
C10	0.075 (3)	0.054 (2)	0.078 (3)	0.007 (2)	0.022 (2)	-0.004 (2)
C11	0.042 (2)	0.0401 (17)	0.050 (2)	0.0005 (14)	-0.0020 (16)	-0.0038 (15)
C12	0.047 (2)	0.0439 (18)	0.053 (2)	0.0052 (15)	-0.0083 (17)	-0.0013 (15)
C13	0.042 (2)	0.0461 (18)	0.054 (2)	0.0036 (16)	0.0013 (17)	0.0044 (16)
C14	0.044 (2)	0.0397 (16)	0.0423 (19)	-0.0002 (14)	0.0043 (15)	0.0014 (14)
C15	0.042 (2)	0.0396 (16)	0.0419 (18)	0.0000 (14)	0.0038 (15)	0.0004 (14)
C16	0.047 (2)	0.055 (2)	0.055 (2)	-0.0028 (16)	-0.0017 (17)	-0.0058 (17)
C17	0.060 (3)	0.053 (2)	0.047 (2)	-0.0127 (17)	-0.0048 (19)	-0.0062 (16)
C18	0.071 (3)	0.0454 (18)	0.051 (2)	-0.0023 (18)	0.005 (2)	-0.0066 (16)
C19	0.061 (2)	0.0423 (17)	0.048 (2)	0.0054 (16)	0.0067 (19)	0.0001 (16)

Geometric parameters (Å, °)

Cl1—C1	1.751 (3)	C8—H8	0.93
O1—C13	1.224 (4)	C10—H10A	0.96
N1—C1	1.301 (4)	C10—H10B	0.96
N1—C9	1.380 (4)	C10—H10C	0.96

N2—C15	1.395 (4)	C11—C12	1.522 (4)
N2—C11	1.454 (4)	C11—H11	0.98
N2—H2	0.85 (3)	C12—C13	1.507 (4)
C1—C2	1.423 (4)	C12—H12A	0.97
C2—C3	1.364 (5)	C12—H12B	0.97
C2—C11	1.509 (4)	C13—C14	1.466 (5)
C3—C4	1.415 (4)	C14—C15	1.395 (4)
C3—H3	0.93	C14—C19	1.406 (4)
C4—C5	1.407 (5)	C15—C16	1.404 (4)
C4—C9	1.412 (4)	C16—C17	1.372 (4)
C5—C6	1.372 (4)	C16—H16	0.93
C5—H5	0.93	C17—C18	1.389 (5)
C6—C7	1.410 (5)	C17—H17	0.93
C6—C10	1.500 (5)	C18—C19	1.372 (5)
C7—C8	1.365 (5)	C18—H18	0.93
C7—H7	0.93	C19—H19	0.93
C8—C9	1.409 (4)		
C1—N1—C9	117.2 (3)	H10A—C10—H10C	109.5
C15—N2—C11	118.2 (3)	H10B—C10—H10C	109.5
C15—N2—H2	112 (2)	N2—C11—C2	110.5 (3)
C11—N2—H2	115 (2)	N2—C11—C12	108.6 (3)
N1—C1—C2	126.6 (3)	C2—C11—C12	111.7 (3)
N1—C1—C11	114.9 (2)	N2—C11—H11	108.7
C2—C1—C11	118.4 (2)	C2—C11—H11	108.7
C3—C2—C1	115.7 (3)	C12—C11—H11	108.7
C3—C2—C11	122.0 (3)	C13—C12—C11	112.1 (3)
C1—C2—C11	122.3 (3)	C13—C12—H12A	109.2
C2—C3—C4	121.1 (3)	C11—C12—H12A	109.2
C2—C3—H3	119.5	C13—C12—H12B	109.2
C4—C3—H3	119.5	C11—C12—H12B	109.2
C5—C4—C9	118.7 (3)	H12A—C12—H12B	107.9
C5—C4—C3	123.4 (3)	O1—C13—C14	122.8 (3)
C9—C4—C3	118.0 (3)	O1—C13—C12	121.0 (3)
C6—C5—C4	122.0 (3)	C14—C13—C12	116.1 (3)
C6—C5—H5	119	C15—C14—C19	118.8 (3)
C4—C5—H5	119	C15—C14—C13	120.5 (3)
C5—C6—C7	117.9 (3)	C19—C14—C13	120.6 (3)
C5—C6—C10	121.8 (3)	C14—C15—N2	120.5 (3)
C7—C6—C10	120.3 (3)	C14—C15—C16	119.5 (3)
C8—C7—C6	122.4 (3)	N2—C15—C16	119.9 (3)
C8—C7—H7	118.8	C17—C16—C15	120.2 (3)
C6—C7—H7	118.8	C17—C16—H16	119.9
C7—C8—C9	119.4 (3)	C15—C16—H16	119.9
C7—C8—H8	120.3	C16—C17—C18	120.9 (3)
C9—C8—H8	120.3	C16—C17—H17	119.6
N1—C9—C8	118.9 (3)	C18—C17—H17	119.6
N1—C9—C4	121.5 (3)	C19—C18—C17	119.3 (3)

C8—C9—C4	119.6 (3)	C19—C18—H18	120.4
C6—C10—H10A	109.5	C17—C18—H18	120.4
C6—C10—H10B	109.5	C18—C19—C14	121.3 (3)
H10A—C10—H10B	109.5	C18—C19—H19	119.4
C6—C10—H10C	109.5	C14—C19—H19	119.4
C9—N1—C1—C2	-1.0 (5)	C15—N2—C11—C12	-50.0 (4)
C9—N1—C1—C11	-179.8 (2)	C3—C2—C11—N2	21.5 (4)
N1—C1—C2—C3	1.2 (5)	C1—C2—C11—N2	-160.2 (3)
C11—C1—C2—C3	180.0 (2)	C3—C2—C11—C12	-99.5 (4)
N1—C1—C2—C11	-177.2 (3)	C1—C2—C11—C12	78.8 (4)
C11—C1—C2—C11	1.6 (4)	N2—C11—C12—C13	54.4 (4)
C1—C2—C3—C4	-0.5 (5)	C2—C11—C12—C13	176.5 (3)
C11—C2—C3—C4	177.9 (3)	C11—C12—C13—O1	151.2 (3)
C2—C3—C4—C5	178.9 (3)	C11—C12—C13—C14	-32.2 (4)
C2—C3—C4—C9	-0.4 (5)	O1—C13—C14—C15	178.7 (3)
C9—C4—C5—C6	-0.4 (5)	C12—C13—C14—C15	2.1 (5)
C3—C4—C5—C6	-179.6 (3)	O1—C13—C14—C19	2.3 (5)
C4—C5—C6—C7	-1.0 (5)	C12—C13—C14—C19	-174.2 (3)
C4—C5—C6—C10	177.9 (3)	C19—C14—C15—N2	-178.6 (3)
C5—C6—C7—C8	0.8 (5)	C13—C14—C15—N2	5.0 (5)
C10—C6—C7—C8	-178.2 (3)	C19—C14—C15—C16	2.0 (5)
C6—C7—C8—C9	0.9 (5)	C13—C14—C15—C16	-174.4 (3)
C1—N1—C9—C8	179.3 (3)	C11—N2—C15—C14	20.8 (4)
C1—N1—C9—C4	0.0 (5)	C11—N2—C15—C16	-159.8 (3)
C7—C8—C9—N1	178.4 (3)	C14—C15—C16—C17	-1.0 (5)
C7—C8—C9—C4	-2.3 (5)	N2—C15—C16—C17	179.6 (3)
C5—C4—C9—N1	-178.6 (3)	C15—C16—C17—C18	-1.0 (5)
C3—C4—C9—N1	0.7 (5)	C16—C17—C18—C19	1.9 (6)
C5—C4—C9—C8	2.0 (5)	C17—C18—C19—C14	-0.9 (5)
C3—C4—C9—C8	-178.7 (3)	C15—C14—C19—C18	-1.0 (5)
C15—N2—C11—C2	-172.8 (3)	C13—C14—C19—C18	175.3 (3)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C3—H3 \cdots N2	0.93	2.45	2.788 (4)	102
C11—H11 \cdots C11	0.98	2.72	3.074 (3)	102
C17—H17 \cdots O1 ⁱ	0.93	2.49	3.243 (5)	138

Symmetry code: (i) $x-1/2, y, -z+1/2$.