

3-Acetyl-2-methyl-4-(pyridin-3-yl)-1,4-dihydroindeno[1,2-*b*]pyridin-5-one

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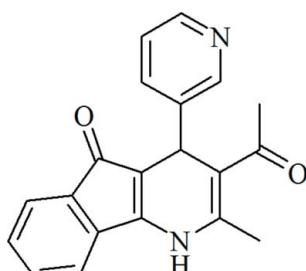
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Key indicators: single-crystal X-ray study; $T = 190\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$;
 R factor = 0.047; wR factor = 0.124; data-to-parameter ratio = 16.5.

In the title compound, $\text{C}_{20}\text{H}_{16}\text{N}_2\text{O}_2$, the condensed tricyclic fragment is near to planar, with an r.m.s. deviation of 0.0531 \AA . The 1,4-dihydropyridine (1,4-DHP) ring adopts a slightly puckered boat-like conformation. The N and opposite C atoms deviate from the least-squares plane of the four other ring atoms by $0.039(3)$ and $0.141(3)\text{ \AA}$, respectively. The $\text{C}=\text{O}$ group located at the tricyclic fragment is fixed in an *s-trans* orientation, while the second $\text{C}=\text{O}$ group adopts an *s-cis* orientation with respect to the double bonds of the 1,4-DHP ring. The pyridine ring has a pseudo-axial orientation with respect to the 1,4-DHP ring. The dihedral angle between the tricyclic system and the pyridine ring is $77.3(3)^\circ$. In the crystal, the pyridine N atom accepts a hydrogen bond from the N—H group of the 1,4-DHP ring. The hydrogen bonds link the molecules into infinite *C*(8) chains along the *b*-axis direction.

Related literature

For general information on the relationship between 1,4-dihydropyridine ring substituents and pharmaceutical effects, see: Christopher *et al.* (2010); Bisenieks *et al.* (1987, 1995); Ivanov *et al.* (1989). For the synthesis of 1,4-DHP-containing tricyclic derivatives, see: Bisenieks *et al.* (1982).



Experimental

Crystal data

$\text{C}_{20}\text{H}_{16}\text{N}_2\text{O}_2$

$M_r = 316.35$

Triclinic, $P\bar{1}$	$V = 797.00(5)\text{ \AA}^3$
$a = 8.4361(3)\text{ \AA}$	$Z = 2$
$b = 8.8812(3)\text{ \AA}$	Mo $K\alpha$ radiation
$c = 11.2582(4)\text{ \AA}$	$\mu = 0.09\text{ mm}^{-1}$
$\alpha = 87.692(2)^\circ$	$T = 190\text{ K}$
$\beta = 71.022(2)^\circ$	$0.30 \times 0.28 \times 0.13\text{ mm}$
$\gamma = 89.210(2)^\circ$	

Data collection

Nonius KappaCCD diffractometer	2753 reflections with $I > 2\sigma(I)$
5442 measured reflections	$R_{\text{int}} = 0.020$
3615 independent reflections	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$	219 parameters
$wR(F^2) = 0.124$	H-atom parameters constrained
$S = 1.04$	$\Delta\rho_{\text{max}} = 0.23\text{ e \AA}^{-3}$
3615 reflections	$\Delta\rho_{\text{min}} = -0.23\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H} \cdots A$	$D-\text{H}$	$\text{H} \cdots A$	$D \cdots A$	$D-\text{H} \cdots A$
N1—H1A \cdots N22 ⁱ	0.86	2.06	2.916 (2)	175

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

Data collection: *COLLECT* (Hooft, 1998); cell refinement: *SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO* (Otwinowski & Minor, 1997) and *SCALEPACK*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FY2091).

References

- Bisenieks, E. A., Uldrikis, J. R., Duburs, G. J., Kozuhov, A. N., Neimane, D. U., Ivanov, E. B. & Merkusev, G. N. (1987). USSR Inventors Certif. No. 1324260.
- Bisenieks, E. A., Uldrikis, J. R., Duburs, G. J., Tirzitis, G. D., Dauvarate, A. D., Zidermane, A. A. & Ponomareva, A. B. (1995). USSR Inventors Certif. No. 1050261.
- Bisenieks, E. A., Uldrikis, J. R., Kirule, I. E., Tirzitis, G. D. & Duburs, G. J. (1982). *Chem. Heterocycl. Compd.*, **18**, 1187–1189.
- Christopher, G. E., Umesh, K. J., Leah, N. M., Dickey, A. C. & Gestwicki, J. E. (2010). *Chem. Commun.*, **47**, 529–531.
- Farrugia, L. J. (2012). *J. Appl. Cryst.*, **45**, 849–854.
- Hooft, R. (1998). *COLLECT*. Nonius BV, Delft, The Netherlands.
- Ivanov, E. I., Ponamareva, T. V., Merkushov, G. N., Duburs, G. J., Bisenieks, E. A., Uldrikis, J. R. & Makarova, N. V. (1989). Abstract Book of the IXth Soviet–Indian Symposium on the Chemistry of Natural Products, Riga, May 15–18, 1989, pp. 87–88.
- Otwinowski, Z. & Minor, W. (1997). *Methods in Enzymology*, Vol. 276, *Macromolecular Crystallography*, Part A, edited by C. W. Carter Jr & R. M. Sweet, pp. 307–326. New York: Academic Press.
- Sheldrick, G. M. (2008). *Acta Cryst. A*, **64**, 112–122.

supporting information

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3-Acetyl-2-methyl-4-(pyridin-3-yl)-1,4-dihydroindeno[1,2-*b*]pyridin-5-one

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S1. Comment

3-Acetyl-2-methyl-4-pyridin-3-yl-1,4-dihydro-indeno[1,2-*b*]pyridin-5-one is a condensed tricyclic 1,4-dihydropyridine derivative containing a conjugated system, with an intense red colour which is common for this type of compounds. In this particular compound, the 1,4-dihydropyridine ring acts as a pharmacophore, and gives potentially useful pharmacological activities. Depending on the particular set of substituents, similar types of 1,4-dihydropyridine derivatives have been identified as compounds possessing an effect on the accumulation of tau, a microtubule-binding protein, which plays a major role in many neurodegenerative disorders, including Alzheimer's disease (Christopher *et al.*, 2010). It has been reported that similar types of 1,4-dihydropyridine derivatives have anti-cancer and anti-radiation activity (Bisenieks *et al.*, 1995; Ivanov *et al.*, 1989), and also skin regenerating and radioprotection properties (Bisenieks *et al.*, 1987). Some of these types of compounds act as antioxidants (Bisenieks *et al.*, 1982). Fig. 1 shows a view of the crystal structure of the title compound. In the crystal structure the 1,4-dihydropyridine (1,4-DHP) ring adopts a slightly puckered boat conformation. The N1 and C4 atoms deviate from the least-squares plane calculated through the four other ring atoms by 0.039 (3) Å and 0.141 (3) Å, respectively. The carbonyl group C13=O14 is fixed in *trans* orientation with respect to the double bonds of the 1,4-DHP ring while the second carbonyl group C15=O16 assumes *cis* orientation. The condensed tricyclic fragment is planar with an r.m.s. deviation of 0.0531 Å. The pyridine ring has an axial orientation with respect to the 1,4-DHP ring. A dihedral angle between the tricyclic system and the pyridine fragment is 77.3 (3)°. In the crystal, the pyridine nitrogen accepts a hydrogen bond from the N—H group of the 1,4-DHP cycle (Table 1). The hydrogen bonds essembles the molecules in infinite chains, C¹(8), along the *b* axis.

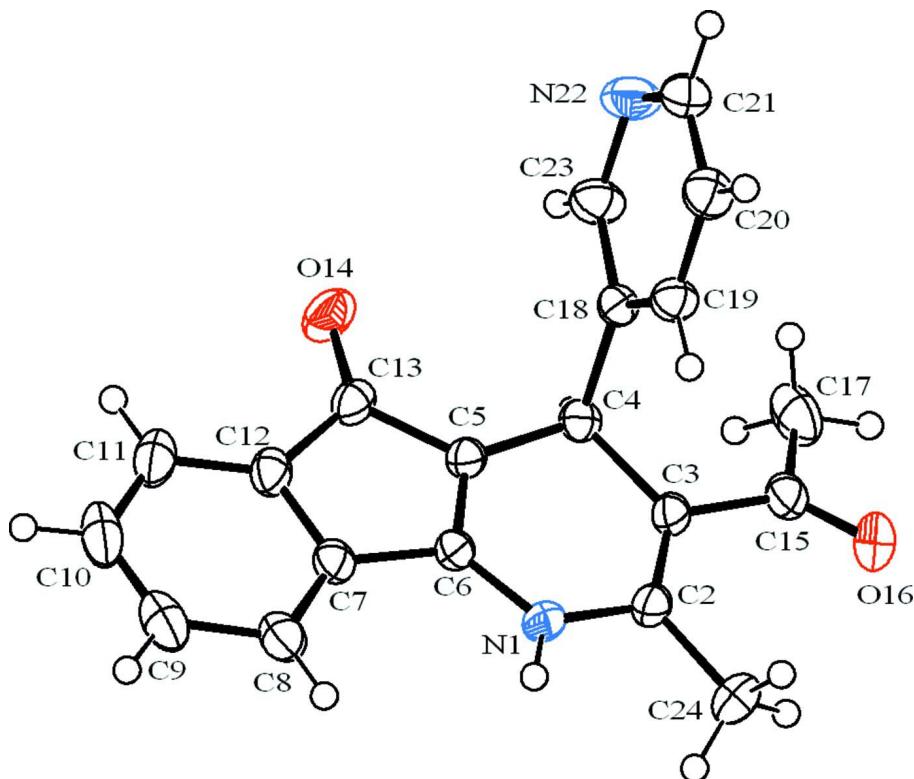
S2. Experimental

3-Acetyl-2-methyl-4-pyridin-3-yl-1,4-dihydro-indeno[1,2-*b*]pyridin-5-one was synthesized by the method described by Bisenieks *et al.* (1982). The resulting compound was dissolved in an acetic acid, DMAA and water (2:1:1) mixture while heating at 100 °C. Then the solution was kept at -5 °C until block crystals were obtained.

¹H-NMR (400 MHz, DMSO-d₆), δ/p.p.m.: 10.19 (s, 1H, N—H), 8.48–8.49 (m, 1H, py-2-H), 8.34–8.35 (m, 1H, py-6-H), 7.25–7.61 (m, 6H, py-4,5-H and –C₆H₄CO–), 4.96 (s, 1H, 4-H), 2.4 (s, 3H, COCH₃), 2.12 (s, 3H, 2-CH₃)

S3. Refinement

The H-atoms were included in the refinement at calculated positions (N—H = 0.86 Å, C—H = 0.93 to 0.98 Å) and refined using a riding-model approximation with $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{NH,CH})$ and $1.5U_{\text{eq}}(\text{CH}_3)$.

**Figure 1**

Molecular structure of the title compound showing 50% probability ellipsoids.

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Crystal data

$C_{20}H_{16}N_2O_2$
 $M_r = 316.35$
Triclinic, $P\bar{1}$
Hall symbol: -P 1
 $a = 8.4361 (3) \text{ \AA}$
 $b = 8.8812 (3) \text{ \AA}$
 $c = 11.2582 (4) \text{ \AA}$
 $\alpha = 87.692 (2)^\circ$
 $\beta = 71.022 (2)^\circ$
 $\gamma = 89.210 (2)^\circ$
 $V = 797.00 (5) \text{ \AA}^3$

$Z = 2$
 $F(000) = 332$
 $D_x = 1.318 \text{ Mg m}^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Cell parameters from 8654 reflections
 $\theta = 1.0\text{--}27.5^\circ$
 $\mu = 0.09 \text{ mm}^{-1}$
 $T = 190 \text{ K}$
Plate, red
 $0.30 \times 0.28 \times 0.13 \text{ mm}$

Data collection

Nonius KappaCCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
CCD scans
5442 measured reflections
3615 independent reflections

2753 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$
 $\theta_{\text{max}} = 27.5^\circ, \theta_{\text{min}} = 2.7^\circ$
 $h = -10 \rightarrow 10$
 $k = -11 \rightarrow 10$
 $l = -14 \rightarrow 14$

*Refinement*Refinement on F^2

Least-squares matrix: full

$$R[F^2 > 2\sigma(F^2)] = 0.047$$

$$wR(F^2) = 0.124$$

$$S = 1.04$$

3615 reflections

219 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0583P)^2 + 0.1857P]$$
$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

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

$$\Delta\rho_{\max} = 0.23 \text{ e \AA}^{-3}$$

$$\Delta\rho_{\min} = -0.23 \text{ e \AA}^{-3}$$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.95378 (15)	0.05051 (14)	0.30285 (11)	0.0282 (3)
H1A	0.9247	0.1425	0.2948	0.034*
O14	0.77556 (15)	-0.36855 (13)	0.58601 (11)	0.0409 (3)
C5	0.90043 (17)	-0.18330 (16)	0.41963 (13)	0.0245 (3)
C6	0.86632 (17)	-0.03576 (16)	0.40434 (13)	0.0243 (3)
C23	0.9277 (2)	-0.53299 (17)	0.30646 (16)	0.0355 (4)
H3	0.9676	-0.5603	0.3722	0.043*
N22	0.85901 (19)	-0.64053 (15)	0.25866 (14)	0.0424 (4)
C18	0.94286 (16)	-0.38372 (16)	0.26401 (13)	0.0237 (3)
C3	1.13273 (17)	-0.15786 (17)	0.21807 (13)	0.0258 (3)
C19	0.87866 (18)	-0.34450 (17)	0.16849 (14)	0.0287 (3)
H3A	0.8850	-0.2454	0.1373	0.034*
C15	1.27713 (18)	-0.22251 (19)	0.11903 (15)	0.0336 (4)
C4	1.02827 (17)	-0.26829 (16)	0.32047 (13)	0.0245 (3)
H7	1.1027	-0.3218	0.3589	0.029*
C2	1.09093 (18)	-0.01006 (17)	0.21154 (13)	0.0279 (3)
C13	0.78546 (18)	-0.24006 (18)	0.53979 (13)	0.0287 (3)
O16	1.33862 (17)	-0.16140 (15)	0.01533 (11)	0.0523 (4)
C24	1.1816 (2)	0.1054 (2)	0.11276 (16)	0.0412 (4)
H11A	1.2993	0.0830	0.0844	0.062*
H11B	1.1638	0.2036	0.1475	0.062*
H11C	1.1397	0.1034	0.0431	0.062*
C20	0.80500 (19)	-0.45370 (19)	0.11972 (15)	0.0346 (4)
H12	0.7604	-0.4287	0.0560	0.042*
C7	0.72519 (17)	0.01618 (17)	0.51290 (13)	0.0274 (3)

C21	0.7984 (2)	-0.59916 (19)	0.16626 (16)	0.0372 (4)
H14	0.7498	-0.6722	0.1322	0.045*
C12	0.67731 (18)	-0.10812 (18)	0.59719 (14)	0.0294 (3)
C8	0.64311 (19)	0.15161 (19)	0.53788 (15)	0.0340 (4)
H16	0.6747	0.2341	0.4818	0.041*
C10	0.4653 (2)	0.0398 (2)	0.73495 (16)	0.0434 (4)
H17	0.3782	0.0497	0.8102	0.052*
C9	0.5097 (2)	0.1612 (2)	0.65120 (16)	0.0411 (4)
H18	0.4505	0.2511	0.6699	0.049*
C17	1.3462 (2)	-0.3722 (2)	0.14648 (18)	0.0454 (5)
H19A	1.2708	-0.4510	0.1440	0.068*
H19B	1.3582	-0.3725	0.2284	0.068*
H19C	1.4536	-0.3885	0.0846	0.068*
C11	0.54884 (19)	-0.0980 (2)	0.70877 (15)	0.0376 (4)
H20	0.5183	-0.1803	0.7652	0.045*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0341 (6)	0.0202 (6)	0.0261 (6)	-0.0024 (5)	-0.0044 (5)	0.0028 (5)
O14	0.0472 (7)	0.0363 (7)	0.0346 (6)	-0.0057 (5)	-0.0084 (5)	0.0139 (5)
C5	0.0275 (7)	0.0254 (8)	0.0205 (7)	-0.0041 (6)	-0.0078 (6)	0.0001 (6)
C6	0.0267 (7)	0.0260 (8)	0.0204 (7)	-0.0037 (5)	-0.0077 (6)	-0.0022 (6)
C23	0.0484 (9)	0.0247 (8)	0.0374 (9)	-0.0005 (7)	-0.0197 (7)	0.0021 (7)
N22	0.0566 (9)	0.0241 (7)	0.0477 (9)	-0.0066 (6)	-0.0184 (7)	0.0002 (6)
C18	0.0234 (6)	0.0221 (7)	0.0243 (7)	0.0018 (5)	-0.0058 (5)	-0.0023 (6)
C3	0.0262 (7)	0.0304 (8)	0.0199 (7)	-0.0047 (6)	-0.0057 (6)	-0.0028 (6)
C19	0.0318 (7)	0.0245 (8)	0.0305 (8)	0.0000 (6)	-0.0112 (6)	0.0005 (6)
C15	0.0292 (7)	0.0397 (9)	0.0290 (8)	-0.0078 (6)	-0.0040 (6)	-0.0111 (7)
C4	0.0268 (7)	0.0247 (7)	0.0233 (7)	-0.0004 (5)	-0.0097 (6)	-0.0004 (6)
C2	0.0315 (7)	0.0306 (8)	0.0202 (7)	-0.0073 (6)	-0.0062 (6)	0.0006 (6)
C13	0.0300 (7)	0.0339 (9)	0.0231 (7)	-0.0066 (6)	-0.0101 (6)	0.0031 (6)
O16	0.0564 (8)	0.0545 (8)	0.0305 (7)	-0.0109 (6)	0.0084 (6)	-0.0084 (6)
C24	0.0468 (9)	0.0371 (10)	0.0320 (9)	-0.0098 (8)	-0.0027 (7)	0.0075 (7)
C20	0.0344 (8)	0.0404 (9)	0.0322 (8)	-0.0029 (7)	-0.0149 (7)	-0.0036 (7)
C7	0.0270 (7)	0.0321 (8)	0.0238 (7)	-0.0031 (6)	-0.0088 (6)	-0.0049 (6)
C21	0.0370 (8)	0.0359 (9)	0.0372 (9)	-0.0104 (7)	-0.0088 (7)	-0.0084 (7)
C12	0.0277 (7)	0.0388 (9)	0.0220 (7)	-0.0045 (6)	-0.0080 (6)	-0.0031 (6)
C8	0.0350 (8)	0.0355 (9)	0.0322 (8)	0.0010 (6)	-0.0111 (7)	-0.0091 (7)
C10	0.0294 (8)	0.0696 (13)	0.0284 (9)	0.0008 (8)	-0.0041 (7)	-0.0165 (9)
C9	0.0338 (8)	0.0522 (11)	0.0389 (9)	0.0079 (7)	-0.0119 (7)	-0.0198 (8)
C17	0.0326 (8)	0.0541 (12)	0.0476 (10)	0.0104 (7)	-0.0086 (8)	-0.0216 (9)
C11	0.0313 (8)	0.0558 (11)	0.0234 (8)	-0.0073 (7)	-0.0057 (6)	-0.0018 (7)

Geometric parameters (\AA , $^\circ$)

N1—C6	1.3541 (18)	C2—C24	1.500 (2)
N1—C2	1.3923 (19)	C13—C12	1.507 (2)

N1—H1A	0.8600	C24—H11A	0.9600
O14—C13	1.2274 (19)	C24—H11B	0.9600
C5—C6	1.354 (2)	C24—H11C	0.9600
C5—C13	1.459 (2)	C20—C21	1.369 (2)
C5—C4	1.497 (2)	C20—H12	0.9300
C6—C7	1.486 (2)	C7—C8	1.372 (2)
C23—N22	1.340 (2)	C7—C12	1.400 (2)
C23—C18	1.384 (2)	C21—H14	0.9300
C23—H3	0.9300	C12—C11	1.372 (2)
N22—C21	1.338 (2)	C8—C9	1.405 (2)
C18—C19	1.3837 (19)	C8—H16	0.9300
C18—C4	1.5312 (18)	C10—C9	1.375 (3)
C3—C2	1.360 (2)	C10—C11	1.398 (3)
C3—C15	1.485 (2)	C10—H17	0.9300
C3—C4	1.529 (2)	C9—H18	0.9300
C19—C20	1.382 (2)	C17—H19A	0.9600
C19—H3A	0.9300	C17—H19B	0.9600
C15—O16	1.220 (2)	C17—H19C	0.9600
C15—C17	1.505 (2)	C11—H20	0.9300
C4—H7	0.9800		
C6—N1—C2	119.88 (13)	C2—C24—H11A	109.5
C6—N1—H1A	120.1	C2—C24—H11B	109.5
C2—N1—H1A	120.1	H11A—C24—H11B	109.5
C6—C5—C13	108.51 (13)	C2—C24—H11C	109.5
C6—C5—C4	122.68 (13)	H11A—C24—H11C	109.5
C13—C5—C4	128.66 (13)	H11B—C24—H11C	109.5
C5—C6—N1	123.00 (13)	C21—C20—C19	119.23 (14)
C5—C6—C7	111.18 (13)	C21—C20—H12	120.4
N1—C6—C7	125.81 (13)	C19—C20—H12	120.4
N22—C23—C18	124.02 (15)	C8—C7—C12	121.17 (14)
N22—C23—H3	118.0	C8—C7—C6	132.56 (15)
C18—C23—H3	118.0	C12—C7—C6	106.25 (13)
C21—N22—C23	117.45 (14)	N22—C21—C20	122.67 (14)
C19—C18—C23	117.09 (13)	N22—C21—H14	118.7
C19—C18—C4	121.92 (13)	C20—C21—H14	118.7
C23—C18—C4	120.99 (13)	C11—C12—C7	121.20 (15)
C2—C3—C15	120.83 (13)	C11—C12—C13	130.70 (15)
C2—C3—C4	122.57 (13)	C7—C12—C13	108.08 (13)
C15—C3—C4	116.45 (13)	C7—C8—C9	117.61 (16)
C20—C19—C18	119.51 (14)	C7—C8—H16	121.2
C20—C19—H3A	120.2	C9—C8—H16	121.2
C18—C19—H3A	120.2	C9—C10—C11	121.08 (15)
O16—C15—C3	122.70 (16)	C9—C10—H17	119.5
O16—C15—C17	118.81 (15)	C11—C10—H17	119.5
C3—C15—C17	118.46 (14)	C10—C9—C8	121.05 (16)
C5—C4—C3	109.63 (12)	C10—C9—H18	119.5
C5—C4—C18	110.49 (11)	C8—C9—H18	119.5

C3—C4—C18	110.48 (11)	C15—C17—H19A	109.5
C5—C4—H7	108.7	C15—C17—H19B	109.5
C3—C4—H7	108.7	H19A—C17—H19B	109.5
C18—C4—H7	108.7	C15—C17—H19C	109.5
C3—C2—N1	121.09 (13)	H19A—C17—H19C	109.5
C3—C2—C24	126.54 (14)	H19B—C17—H19C	109.5
N1—C2—C24	112.36 (13)	C12—C11—C10	117.86 (16)
O14—C13—C5	127.98 (14)	C12—C11—H20	121.1
O14—C13—C12	126.06 (14)	C10—C11—H20	121.1
C5—C13—C12	105.96 (13)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···N22 ⁱ	0.86	2.06	2.916 (2)	175

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