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1-Allyl-3,3-di-p-tolylindolin-2-one

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Key indicators: single-crystal X-ray study; T = 293 K; mean σ (C–C) = 0.002 Å; R factor = 0.052; wR factor = 0.156; data-to-parameter ratio = 25.7.

In the title compound, $C_{25}H_{23}NO$, the indoline system is essentially planar. The molecular structure is stabilized by weak intramolecular $C-H \cdot \cdot \cdot N$ interactions and the crystal packing is determined by intermolecular $C-H \cdot \cdot \cdot \pi$ interactions.

Related literature

For related literature, see: Harris & Uhle (1960); Ho et al. (1986); Rajeswaran et al. (1999); Stevenson et al. (2000); Sethusankar et al. (2002).



Experimental

Crystal data	
$C_{25}H_{23}NO$ $M_r = 353.44$ Triclinic, $P\overline{1}$	a = 9.3311 (2) Å b = 9.5793 (2) Å c = 11.5736 (2) Å

$\alpha = 92.163 \ (1)^{\circ}$	
$\beta = 103.192 \ (1)^{\circ}$	
$\gamma = 101.520 \ (1)^{\circ}$	
V = 983.15 (3) Å ³	
$\mathbf{Z} = 2$	

Data collection

Bruker Kappa APEXII	25969 measured reflections
diffractometer	6260 independent reflections
Absorption correction: multi-scan	4310 reflections with $I > 2\sigma(I)$
(Blessing, 1995)	$R_{\rm int} = 0.023$
$T_{\min} = 0.982, \ T_{\max} = 0.986$	

Refinement

 $\begin{array}{ll} R[F^2 > 2\sigma(F^2)] = 0.051 & 244 \text{ parameters} \\ wR(F^2) = 0.156 & H\text{-atom parameters constrained} \\ S = 0.99 & \Delta\rho_{\max} = 0.24 \text{ e } \text{\AA}^{-3} \\ 6260 \text{ reflections} & \Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3} \end{array}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C5-H5\cdots Cg^{i}$ $C12-H12A\cdots N1$	0.93	2.94	3.740 (2)	145
	0.93	2.54	2.858 (2)	100

Symmetry code: (i) -x + 1, -y, -z + 1. Cg denotes the centroid of the C20–C25 ring.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* and *SAINT* (Bruker, 2004); data reduction: *SAINT* and *XPREP* (Bruker, 2004); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997); software used to prepare material for publication: *PLATON* (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GW2036).

References

- Blessing, R. H. (1995). Acta Cryst. A51, 33-38.
- Bruker (2004). APEX2, SAINT and XPREP. Bruker AXS Inc., Madison, Wisconsin, USA.
- Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
- Harris, L. S. & Uhle, F. C. (1960). J. Pharmacol. Exp. Ther. 128, 353-363.
- Ho, C. Y., Haegman, W. E. & Perisco, F. (1986). J. Med. Chem. 29, 118-121.
- Rajeswaran, W. G., Labroo, R. B., Cohen, L. A. & King, M. M. (1999). J. Org. Chem. 64, 1369–1371.
- Sethu Sankar, K., Kannadasan, S., Velmurugan, D., Srinivasan, P. C., Shanmuga Sundara Raj, S & Fun, H.-K. (2002). Acta Cryst. C58, o277–o279. Sheldrick, G. M. (2008). Acta Cryst. A64, 112–122.
- Spek, A. L. (2003). J. Appl. Cryst. 36, 7-13.
- Stevenson, G. I., Smith, A. L., Lewis, S., Michie, S. G., Neduvelil, J. G., Patel, S., Marwood, R., Patel, S. & Castro, J. L. (2000). *Bioorg. Med. Chem. Lett.* 10, 2697–2704.

Mo $K\alpha$ radiation $\mu = 0.07 \text{ mm}^{-1}$

 $0.26 \times 0.20 \times 0.20$ mm

T = 293 (2) K

supporting information

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1-Allyl-3,3-di-p-tolylindolin-2-one

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S1. Comment

Indole compounds can be used as bioactive drugs (Stevenson *et al.*, 2000). Indole derivatives exhibit anti-allergic, central nervous system depressant and muscle relaxant properties (Harris & Uhle, 1960; Ho *et al.*, 1986). Indoles have also been proved to display high aldose reductase inhibitory activity (Rajeswaran *et al.*, 1999). In view of this biological importance, an X-ray study of the title compound, (I), was carried out.

An *ORTEP* (Farrugia,1997) plot of the molecular is shown in Fig.1. The indole moiety is planar [maximum deviation of 0.038 (1) from the least square plane defined by all non hydrogen atoms in the molecule] and is nearly orthogonal to methylphenyl rings A and B, and makes a dihedral angle of 72.3 (4)° with the ring A, 76.7 (3)° with the ring B. Both the *p*-tolyl rings A and B are oriented at an angle of 72.6 (4)° with respect to each other. The sum of angles around N1 [360.0]° is in accordance with *sp*² hybridization. The endocyclic angles around C4 is narrowed while those at C9 is widened from 120°. This may be caused by fusion of the smaller pyrrole ring to the six membered benzene ring of oxindole. A similar effect has also been observed by Sethu Sankar *et al.* (2002). The bond lengths in the oxindole ring systems indicate electron delocalization. The torsion angles C19—C16—C15—C14 [–179.5 (2)°], C19—C16—C17—C18 [179.7 (2)°] and C26—C23—C24—C25 [–179.6 (2)°], C26—C23—C24—C21 [179.8 (2)°] indicates that the methyl groups are coplanar with the plane of the attached benzene rings A and B. The allyl group deviates significantly from the plane of the indole moiety [C12—C11—C10—N1 = – 4.5 (3)°].

Weak intramolecular C—H···N and intermolecular C—H··· Cg interactions, with C5···Cg = 3.740 (2) Å, [Cg denotes centroid of C20—C25 ring] are observed in the molecular structure. In addition the packing is stabilized by van der Waals forces.

S2. Experimental

To a solution of *p*-methyl phenyl magnesium bromide in dry THF, at 0° C under N₂ atm.,1-*N*-allyl isatin (0.0125 mol, 2.34 g), in dry THF, was added dropwise. After the complete addition, the mixture was stirred at 0°C for 1 hr and then it was stirred at room temperature for 5 hrs. On completion of the reaction, a saturated solution of NH₄Cl was added slowly at 0°C. The aqueous layer was extracted with ether, and the combined organic layer was extracted with ether. The crude mass was obtained., which was purified over a column of silica gel using hexane/ethyl acetate as eluent. Compound was recrystallized from methanol.

S3. Refinement

H atoms were positioned geometrically and were treated as riding on their parent C atoms, with aromatic C—H distances of 0.93 Å, methyl C—H distances of 0.96 Å and methylene C—H distances of 0.97 Å, and with $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H and $1.2U_{eq}(C)$ for other H atoms.



Figure 1

The molecular structure of (I) with 30% probability displacement ellipsoids



Figure 2

The packing of the molecules viewed down b axis.

1-Allyl-3,3-di-p-tolylindolin-2-one

Crystal data C₂₅H₂₃NO Z = 2 $M_r = 353.44$ Triclinic, $P\overline{1}$ Hall symbol: -P 1 a = 9.3311 (2) Åb = 9.5793 (2) Å *c* = 11.5736 (2) Å $\alpha = 92.163 (1)^{\circ}$ $\beta = 103.192 (1)^{\circ}$ $\gamma = 101.520 (1)^{\circ}$ $V = 983.15 (3) \text{ Å}^3$ Data collection Bruker Kappa APEXII diffractometer Radiation source: fine-focus sealed tube Graphite monochromator ω and φ scan Absorption correction: multi-scan (Blessing, 1995) $T_{\rm min} = 0.982, \ T_{\rm max} = 0.986$

F(000) = 376 $D_x = 1.194 \text{ Mg m}^{-3}$ Mo K α radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 9112 reflections $\theta = 2.3-30.1^{\circ}$ $\mu = 0.07 \text{ mm}^{-1}$ T = 293 KPrism, colourless $0.26 \times 0.20 \times 0.20 \text{ mm}$

25969 measured reflections 6260 independent reflections 4310 reflections with $I > 2\sigma(I)$ $R_{int} = 0.023$ $\theta_{max} = 31.0^{\circ}, \ \theta_{min} = 2.2^{\circ}$ $h = -13 \rightarrow 13$ $k = -13 \rightarrow 13$ $l = -16 \rightarrow 16$ Refinement

Refinement on F^2	Secondary atom site location: difference Fourier
Least-squares matrix: full	map
$R[F^2 > 2\sigma(F^2)] = 0.051$	Hydrogen site location: inferred from
$wR(F^2) = 0.156$	neighbouring sites
S = 1.00	H-atom parameters constrained
6260 reflections	$w = 1/[\sigma^2(F_o^2) + (0.0731P)^2 + 0.1889P]$
244 parameters	where $P = (F_o^2 + 2F_c^2)/3$
0 restraints	$(\Delta/\sigma)_{\rm max} < 0.001$
Primary atom site location: structure-invariant	$\Delta \rho_{\rm max} = 0.25 \ {\rm e} \ {\rm \AA}^{-3}$
direct methods	$\Delta \rho_{\rm min} = -0.21 \text{ e} \text{ Å}^{-3}$

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 (A^2)

	x	У	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
C2	0.21981 (13)	0.10769 (13)	0.14416 (10)	0.0403 (2)	
C3	0.29661 (11)	0.03653 (12)	0.25302 (9)	0.0361 (2)	
C4	0.45434 (12)	0.05217 (12)	0.23447 (10)	0.0391 (2)	
C5	0.57669 (13)	0.00390 (15)	0.29710 (12)	0.0497 (3)	
H5	0.5699	-0.0478	0.3628	0.060*	
C6	0.71037 (15)	0.03411 (17)	0.26019 (15)	0.0602 (4)	
H6	0.7938	0.0024	0.3018	0.072*	
C7	0.72086 (16)	0.10999 (18)	0.16332 (16)	0.0629 (4)	
H7	0.8116	0.1295	0.1405	0.075*	
C8	0.59809 (17)	0.15830 (17)	0.09859 (14)	0.0587 (4)	
H8	0.6049	0.2096	0.0327	0.070*	
C9	0.46580 (14)	0.12724 (13)	0.13582 (11)	0.0443 (3)	
C10	0.29599 (19)	0.24323 (18)	-0.01709 (12)	0.0614 (4)	
H10A	0.1917	0.2102	-0.0604	0.074*	
H10B	0.3589	0.2267	-0.0701	0.074*	
C11	0.32409 (19)	0.39967 (19)	0.01642 (16)	0.0658 (4)	
H11	0.3133	0.4583	-0.0458	0.079*	
C12	0.3620(2)	0.4623 (2)	0.12318 (19)	0.0749 (5)	
H12A	0.3742	0.4084	0.1885	0.090*	
H12B	0.3770	0.5612	0.1347	0.090*	
C13	0.22090 (12)	-0.12027 (12)	0.25034 (10)	0.0391 (2)	
C14	0.24490 (16)	-0.19198 (15)	0.35239 (13)	0.0541 (3)	
H14	0.3029	-0.1423	0.4237	0.065*	
C15	0.18392 (18)	-0.33608 (16)	0.34970 (16)	0.0626 (4)	
H15	0.2020	-0.3818	0.4194	0.075*	

C16	0.09700 (17)	-0.41371 (15)	0.24620 (16)	0.0592 (4)
C17	0.07309 (17)	-0.34194 (16)	0.14492 (15)	0.0603 (4)
H17	0.0145	-0.3920	0.0739	0.072*
C18	0.13362 (15)	-0.19778 (14)	0.14583 (12)	0.0490 (3)
H18	0.1156	-0.1526	0.0759	0.059*
C19	0.0294 (3)	-0.57073 (17)	0.2435 (2)	0.0907 (6)
H19A	-0.0262	-0.6064	0.1639	0.136*
H19B	-0.0372	-0.5842	0.2963	0.136*
H19C	0.1083	-0.6216	0.2686	0.136*
C20	0.28928 (12)	0.12652 (12)	0.36267 (10)	0.0372 (2)
C21	0.40937 (13)	0.23204 (14)	0.42458 (11)	0.0445 (3)
H21	0.5003	0.2467	0.4020	0.053*
C22	0.39541 (16)	0.31600 (15)	0.51984 (12)	0.0530 (3)
H22	0.4778	0.3858	0.5605	0.064*
C23	0.26275 (17)	0.29871 (17)	0.55575 (12)	0.0553 (3)
C24	0.14184 (16)	0.19372 (17)	0.49283 (13)	0.0561 (3)
H24	0.0506	0.1801	0.5150	0.067*
C25	0.15471 (14)	0.10923 (15)	0.39803 (12)	0.0483 (3)
H25	0.0721	0.0398	0.3573	0.058*
C26	0.2485 (3)	0.3901 (3)	0.65992 (18)	0.0932 (7)
H26A	0.1485	0.3626	0.6718	0.140*
H26B	0.2675	0.4889	0.6439	0.140*
H26C	0.3204	0.3771	0.7304	0.140*
N1	0.32609 (13)	0.15965 (12)	0.08421 (9)	0.0484 (3)
01	0.08971 (10)	0.11949 (10)	0.11825 (8)	0.0515 (2)

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

		T 72	T 70	T 710	T 710	
	U^{II}	U^{22}	U^{ss}	U^{12}	U^{13}	U^{23}
C2	0.0417 (5)	0.0399 (6)	0.0385 (5)	0.0124 (4)	0.0053 (4)	0.0024 (4)
C3	0.0306 (5)	0.0386 (6)	0.0381 (5)	0.0075 (4)	0.0059 (4)	0.0061 (4)
C4	0.0342 (5)	0.0399 (6)	0.0429 (6)	0.0077 (4)	0.0094 (4)	0.0027 (4)
C5	0.0372 (6)	0.0554 (8)	0.0569 (7)	0.0132 (5)	0.0086 (5)	0.0103 (6)
C6	0.0364 (6)	0.0650 (9)	0.0805 (10)	0.0145 (6)	0.0135 (6)	0.0057 (8)
C7	0.0444 (7)	0.0637 (9)	0.0859 (11)	0.0079 (6)	0.0305 (7)	0.0025 (8)
C8	0.0586 (8)	0.0602 (9)	0.0656 (9)	0.0114 (7)	0.0317 (7)	0.0128 (7)
C9	0.0444 (6)	0.0442 (6)	0.0468 (6)	0.0109 (5)	0.0148 (5)	0.0051 (5)
C10	0.0757 (10)	0.0732 (10)	0.0455 (7)	0.0298 (8)	0.0202 (7)	0.0229 (7)
C11	0.0697 (9)	0.0666 (10)	0.0719 (10)	0.0222 (8)	0.0280 (8)	0.0323 (8)
C12	0.0729 (11)	0.0612 (10)	0.0954 (13)	0.0132 (8)	0.0299 (10)	0.0153 (9)
C13	0.0318 (5)	0.0389 (6)	0.0461 (6)	0.0085 (4)	0.0074 (4)	0.0048 (5)
C14	0.0522 (7)	0.0470 (7)	0.0556 (7)	0.0060 (6)	0.0005 (6)	0.0122 (6)
C15	0.0626 (9)	0.0476 (8)	0.0785 (10)	0.0132 (6)	0.0152 (7)	0.0224 (7)
C16	0.0542 (7)	0.0368 (7)	0.0920 (11)	0.0108 (6)	0.0280 (7)	0.0023 (7)
C17	0.0582 (8)	0.0462 (8)	0.0716 (9)	0.0058 (6)	0.0133 (7)	-0.0137 (7)
C18	0.0499 (7)	0.0470 (7)	0.0480 (6)	0.0096 (5)	0.0092 (5)	-0.0022 (5)
C19	0.0989 (14)	0.0380 (8)	0.1420 (19)	0.0073 (8)	0.0500 (14)	0.0009 (10)
C20	0.0339 (5)	0.0404 (6)	0.0373 (5)	0.0077 (4)	0.0078 (4)	0.0079 (4)

supporting information

C21	0.0372 (5)	0.0475 (7)	0.0461 (6)	0.0038 (5)	0.0097 (5)	0.0032 (5)
C22	0.0521 (7)	0.0519 (8)	0.0491 (7)	0.0055 (6)	0.0063 (6)	-0.0046 (6)
C23	0.0607 (8)	0.0625 (9)	0.0456 (7)	0.0186 (7)	0.0145 (6)	0.0007 (6)
C24	0.0486 (7)	0.0701 (9)	0.0558 (7)	0.0148 (6)	0.0229 (6)	0.0056 (7)
C25	0.0361 (5)	0.0550(7)	0.0522 (7)	0.0046 (5)	0.0121 (5)	0.0027 (6)
C26	0.0965 (14)	0.1150 (17)	0.0707 (11)	0.0262 (12)	0.0279 (10)	-0.0268 (11)
N1	0.0520 (6)	0.0551 (6)	0.0434 (5)	0.0188 (5)	0.0140 (5)	0.0154 (5)
O1	0.0433 (4)	0.0595 (6)	0.0515 (5)	0.0207 (4)	0.0023 (4)	0.0074 (4)

Geometric parameters (Å, °)

C2—01	1.2114 (14)	C14—C15	1.380 (2)	
C2—N1	1.3621 (16)	C14—H14	0.9300	
С2—С3	1.5535 (15)	C15—C16	1.376 (2)	
С3—С4	1.5143 (15)	C15—H15	0.9300	
C3—C13	1.5228 (16)	C16—C17	1.378 (2)	
C3—C20	1.5294 (16)	C16—C19	1.507 (2)	
C4—C5	1.3788 (16)	C17—C18	1.382 (2)	
С4—С9	1.3853 (17)	C17—H17	0.9300	
С5—С6	1.3895 (19)	C18—H18	0.9300	
С5—Н5	0.9300	C19—H19A	0.9600	
С6—С7	1.371 (2)	C19—H19B	0.9600	
С6—Н6	0.9300	C19—H19C	0.9600	
С7—С8	1.391 (2)	C20—C21	1.3847 (16)	
С7—Н7	0.9300	C20—C25	1.3885 (16)	
С8—С9	1.3798 (19)	C21—C22	1.3857 (19)	
С8—Н8	0.9300	C21—H21	0.9300	
C9—N1	1.4046 (16)	C22—C23	1.375 (2)	
C10—N1	1.4508 (16)	C22—H22	0.9300	
C10-C11	1.489 (2)	C23—C24	1.390 (2)	
C10—H10A	0.9700	C23—C26	1.509 (2)	
C10—H10B	0.9700	C24—C25	1.380 (2)	
C11—C12	1.293 (3)	C24—H24	0.9300	
C11—H11	0.9300	C25—H25	0.9300	
C12—H12A	0.9300	C26—H26A	0.9600	
C12—H12B	0.9300	C26—H26B	0.9600	
C13—C18	1.3852 (17)	C26—H26C	0.9600	
C13—C14	1.3853 (17)			
01—C2—N1	125.45 (11)	C16—C15—C14	121.51 (14)	
O1—C2—C3	126.57 (11)	C16—C15—H15	119.2	
N1-C2-C3	107.95 (9)	C14—C15—H15	119.2	
C4—C3—C13	111.14 (9)	C15-C16-C17	117.38 (13)	
C4—C3—C20	113.55 (9)	C15-C16-C19	121.49 (17)	
C13—C3—C20	112.94 (9)	C17—C16—C19	121.13 (17)	
C4—C3—C2	101.28 (9)	C16—C17—C18	121.93 (14)	
С13—С3—С2	111.74 (9)	C16—C17—H17	119.0	
С20—С3—С2	105.46 (9)	C18—C17—H17	119.0	

C5 C4 C0	110.90 (11)	C_{17} C_{19} C_{12}	120 20 (12)
C_{3}	119.89 (11)	C17 - C18 - C13	120.39 (13)
C_{3}	130.70 (11)	C17 - C18 - H18	119.8
C9—C4—C3	109.40 (9)	C13—C18—H18	119.8
C4—C5—C6	118.68 (13)	С16—С19—Н19А	109.5
C4—C5—H5	120.7	С16—С19—Н19В	109.5
С6—С5—Н5	120.7	H19A—C19—H19B	109.5
C7—C6—C5	120.88 (13)	С16—С19—Н19С	109.5
С7—С6—Н6	119.6	H19A—C19—H19C	109.5
С5—С6—Н6	119.6	H19B—C19—H19C	109.5
C6—C7—C8	121.13 (13)	C21—C20—C25	117.95 (11)
С6—С7—Н7	119.4	C21—C20—C3	122.56 (10)
С8—С7—Н7	119.4	С25—С20—С3	119.36 (10)
C9—C8—C7	117.48 (14)	C20—C21—C22	120.67 (12)
С9—С8—Н8	121.3	C20—C21—H21	119.7
С7—С8—Н8	121.3	C22—C21—H21	119.7
C8—C9—C4	121.93 (12)	C23—C22—C21	121.62 (13)
C8—C9—N1	128.51 (12)	C23—C22—H22	119.2
C4—C9—N1	109.54 (10)	C21—C22—H22	119.2
N1-C10-C11	113.55 (13)	C22—C23—C24	117.66 (13)
N1-C10-H10A	108.9	C22—C23—C26	121.35 (15)
C11—C10—H10A	108.9	C24—C23—C26	120.99 (15)
N1-C10-H10B	108.9	C25—C24—C23	121.17 (13)
C11—C10—H10B	108.9	C25—C24—H24	119.4
H10A—C10—H10B	107.7	C23—C24—H24	119.4
C12—C11—C10	126.66 (15)	C24—C25—C20	120.91 (12)
C12—C11—H11	116.7	C_{24} C_{25} H_{25}	119.5
C10-C11-H11	116.7	C_{20} C_{25} H_{25}	119.5
C11—C12—H12A	120.0	C_{23} C_{26} H_{26A}	109.5
C_{11} C_{12} H_{12R}	120.0	$C_{23} = C_{26} = H_{26}R$	109.5
H12A - C12 - H12B	120.0	H_{26}^{-} $H_{$	109.5
C18 C13 C14	117.85 (12)	C_{23} C_{26} H_{26C}	109.5
C18 - C13 - C14	117.03(12) 121.72(11)	$H_{26}^{-} = C_{26}^{-} = H_{26}^{-} = $	109.5
$C_{10} = C_{13} = C_3$	121.72(11) 120.33(10)	H26R C26 H26C	109.5
$C_{14} = C_{13} = C_{3}$	120.35(10) 120.05(12)	1120B - C20 - 1120C	109.3 111.71 (10)
$C_{15} = C_{14} = C_{15}$	120.95 (15)	$C_2 = N_1 = C_3$	111.71(10) 122.67(11)
C13 - C14 - H14	119.5	C_2 N1 C_{10}	122.07(11) 125.57(12)
C13—C14—H14	119.5	C9—N1—C10	125.57 (12)
01 62 63 64	179 69 (12)	C14 C15 C16 C17	0.0(2)
01 - 02 - 03 - 04	1/8.08(12)	C14 - C15 - C10 - C17	0.0(2)
N1 - C2 - C3 - C4	-3.41(12)	C14 - C15 - C16 - C19	-1/9.48(16)
01 - 02 - 03 - 013	60.29 (16)		0.2(2)
N1 - C2 - C3 - C13	-121.80 (11)	C19 - C16 - C17 - C18	1/9.68 (15)
01 - C2 - C3 - C20	-62.78 (15)	C16 - C17 - C18 - C13	-0.2(2)
N1 - C2 - C3 - C20	115.14 (10)	C14—C13—C18—C17	0.1 (2)
C13—C3—C4—C5	-57.30 (16)	C3—C13—C18—C17	176.45 (12)
C20—C3—C4—C5	71.33 (16)	C4—C3—C20—C21	11.87 (15)
C2—C3—C4—C5	-176.12 (13)	C13—C3—C20—C21	139.58 (11)
C13—C3—C4—C9	121.92 (11)	C2-C3-C20-C21	-98.13 (12)
C20—C3—C4—C9	-109.45 (11)	C4—C3—C20—C25	-172.36 (10)

$C^{2}-C^{3}-C^{4}-C^{9}$	310(12)	C13—C3—C20—C25	-44 66 (14)
C9—C4—C5—C6	1.0 (2)	C2—C3—C20—C25	77.64 (13)
C3—C4—C5—C6	-179.88 (12)	C25—C20—C21—C22	0.88 (18)
C4—C5—C6—C7	-0.2 (2)	C3—C20—C21—C22	176.71 (11)
C5C6C7C8	-0.4 (2)	C20—C21—C22—C23	-0.5 (2)
C6—C7—C8—C9	0.2 (2)	C21—C22—C23—C24	-0.2 (2)
C7—C8—C9—C4	0.7 (2)	C21—C22—C23—C26	179.75 (16)
C7—C8—C9—N1	-177.86 (14)	C22—C23—C24—C25	0.4 (2)
C5—C4—C9—C8	-1.2 (2)	C26—C23—C24—C25	-179.55 (16)
C3—C4—C9—C8	179.45 (12)	C23—C24—C25—C20	0.1 (2)
C5-C4-C9-N1	177.52 (12)	C21—C20—C25—C24	-0.69 (19)
C3—C4—C9—N1	-1.79 (14)	C3—C20—C25—C24	-176.65 (12)
N1-C10-C11-C12	-4.5 (3)	O1—C2—N1—C9	-179.43 (12)
C4—C3—C13—C18	-91.47 (13)	C3—C2—N1—C9	2.63 (14)
C20—C3—C13—C18	139.57 (11)	O1—C2—N1—C10	3.1 (2)
C2-C3-C13-C18	20.86 (15)	C3-C2-N1-C10	-174.88 (12)
C4—C3—C13—C14	84.82 (14)	C8—C9—N1—C2	178.07 (13)
C20—C3—C13—C14	-44.14 (14)	C4—C9—N1—C2	-0.58 (15)
C2-C3-C13-C14	-162.84 (11)	C8—C9—N1—C10	-4.5 (2)
C18—C13—C14—C15	0.1 (2)	C4-C9-N1-C10	176.83 (13)
C3—C13—C14—C15	-176.30 (13)	C11—C10—N1—C2	88.62 (17)
C13-C14-C15-C16	-0.2 (2)	C11—C10—N1—C9	-88.53 (17)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	D—H··· A
C5—H5····Cg ⁱ	0.93	2.94	3.740 (2)	145
C12—H12A…N1	0.93	2.54	2.858 (2)	100

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