

1,10-Dihydro-3,9-dimethylpyrazolo-[3,4-a]carbazole

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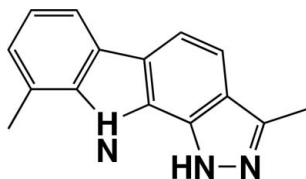
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Key indicators: single-crystal X-ray study; $T = 203$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; disorder in main residue; R factor = 0.043; wR factor = 0.096; data-to-parameter ratio = 22.9.

The heterofused carbazole unit of the title compound, $\text{C}_{15}\text{H}_{13}\text{N}_3$, is planar. The planar pyrazole ring forms dihedral angles of 1.08 (7), 0.64 (6) and 2.60 (7)° with the pyrrole, fused benzene and methyl-substituted benzene rings, respectively. The methyl group on the five-membered ring is disordered equally over two positions. In the crystal structure, the molecules are stabilized by intermolecular $\text{N}-\text{H}\cdots\text{N}$ hydrogen bonds.

Related literature

For related literature, see: Di Fabio *et al.* (2006); Gunaseelan *et al.* (2007a,b,c); Gunaseelan *et al.* (2007); Haider *et al.* (1998); Hedin *et al.* (2000); Hirata *et al.* (1999); Knolker & Reddy (2002); Thiruvalluvar *et al.* (2007).



Experimental

Crystal data

$\text{C}_{15}\text{H}_{13}\text{N}_3$
 $M_r = 235.28$
 Monoclinic, $P2_1/c$
 $a = 11.6553$ (8) Å
 $b = 5.7418$ (3) Å
 $c = 18.0804$ (12) Å
 $\beta = 101.796$ (6)°

$V = 1184.43$ (13) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 203$ (2) K
 $0.51 \times 0.43 \times 0.37$ mm

Data collection

Oxford Diffraction Gemini diffractometer
 Absorption correction: multi-scan (*CrysAlis RED*; Oxford Diffraction, 2007)
 $T_{\min} = 0.868$, $T_{\max} = 1.000$
 11247 measured reflections
 3939 independent reflections
 1587 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.047$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.043$
 $wR(F^2) = 0.096$
 $S = 0.81$
 3939 reflections
 172 parameters
 H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\max} = 0.18$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N1}-\text{H1}\cdots\text{N2}^i$	0.906 (13)	2.188 (12)	2.9878 (14)	147.0 (10)
$\text{N10}-\text{H10}\cdots\text{N2}^i$	0.906 (13)	2.263 (15)	3.0558 (15)	145.9 (12)

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

Data collection: *CrysAlis CCD* (Oxford Diffraction, 2007); cell refinement: *CrysAlis RED* (Oxford Diffraction, 2007); data reduction: *CrysAlis RED*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); 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: BQ2027).

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supplementary materials

Acta Cryst. (2007). E63, o3471 [doi:10.1107/S160053680703351X]

1,10-Dihydro-3,9-dimethylpyrazolo[3,4-*a*]carbazole

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Comment

Our recent work on the *x*-ray structural studies of simple substituted carbazoles (Gunaseelan *et al.*, 2007 a, b & c; Thiruvalluvar *et al.*, 2007) directed us towards the structural studies of heterofused carbazoles (Gunaseelan *et al.*, 2007). The heteroring fused carbazoles have found an important place in the medicinal chemistry because of their high pharmacological properties (Hedin *et al.*, 2000, Hirata *et al.*, 1999, Haider *et al.*, 1998, Knolker & Reddy, 2002). The study of three dimensional arrangement of such compounds will provide some valuable information on synthetic as well as structure activity relationship (SAR) studies (Knolker & Reddy, 2002, Di Fabio *et al.*, 2006). In view of these discussions, here, we present the crystal structure of such a heterofused carbazole, the title compound, (I).

The molecular structure of (I), with atomic numbering scheme, is shown in Fig. 1. The heterofused carbazole unit is planar. The planar pyrazole ring forms dihedral angles of 1.08 (7)°, 0.64 (6)° and 2.60 (7)° with pyrrole, fused benzene and methyl substituted benzene ring, respectively. The methyl group at position 3 is disordered over two positions. In the crystal structure, the molecules are stabilized by intermolecular N—H⋯N ($-x + 1, y - 1/2, -z + 1/2$) hydrogen bonds (see Fig. 2).

Experimental

Hydrazine hydrate (0.1 ml, 0.002 mol) was added to the solution of 2-acetyl-2,3,4,9-tetrahydro-1*H*-carbazol-1-one (240 mg, 0.001 mol) in glacial acetic acid (15 ml) and refluxed on oil bath for 1 h. The reaction was monitored by thin-layer chromatography. After the completion of the reaction it was poured into crushed ice. The precipitate was filtered, washed with water and dried to get the brown residue of crude compound (I), which was purified as a white powder by column chromatography over silica gel using petroleum ether-ethyl acetate (98:2 *v/v*) as eluant (50 mg, 20%) and recrystallized from glacial acetic.

Refinement

H atoms bonded to N1 and N10 were located in a difference Fourier map and refined isotropically. Other H atoms were positioned geometrically and allowed to ride on their parent atoms, with C—H = 0.94–0.97 Å and $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5$ times $U_{\text{eq}}(\text{C})$. The methyl group at position 3 was found to be disordered over two positions.

Figures

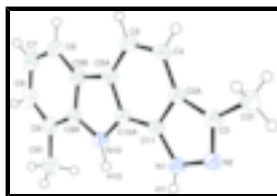


Fig. 1. The molecular structure of (I), showing the atom-numbering scheme and displacement ellipsoids drawn at the 50% probability level.

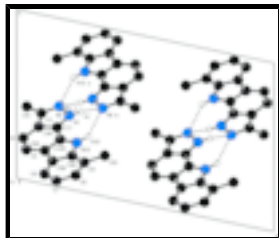


Fig. 2. The molecular packing of (I), viewed down the *b* axis. Dashed lines indicate hydrogen bonds. Symmetry code for both *a* and *b*: $-x + 1, y - 1/2, -z + 1/2$

1,10-Dihydro-3,9-dimethylpyrazolo[3,4-a]carbazole

Crystal data

$C_{15}H_{13}N_3$

$M_r = 235.28$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2ybc$

$a = 11.6553$ (8) Å

$b = 5.7418$ (3) Å

$c = 18.0804$ (12) Å

$\beta = 101.796$ (6)°

$V = 1184.43$ (13) Å³

$Z = 4$

$F_{000} = 496$

$D_x = 1.319$ Mg m⁻³

Melting point: 416(1) K

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 2712 reflections

$\theta = 4.8\text{--}32.4^\circ$

$\mu = 0.08$ mm⁻¹

$T = 203$ (2) K

Plate, light_brown

$0.51 \times 0.43 \times 0.37$ mm

Data collection

Oxford Diffraction Gemini
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 203$ (2) K

φ and ω scans

Absorption correction: multi-scan
(CrysAlis RED; Oxford Diffraction, 2007)

$T_{\min} = 0.868$, $T_{\max} = 1.000$

11247 measured reflections

3939 independent reflections

1587 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.047$

$\theta_{\max} = 32.4^\circ$

$\theta_{\min} = 4.8^\circ$

$h = -16 \rightarrow 12$

$k = -8 \rightarrow 8$

$l = -26 \rightarrow 27$

Standard reflections: .;

every . reflections

intensity decay: .

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.096$

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

$$w = 1/[\sigma^2(F_o^2) + (0.045P)^2]$$

where $P = (F_o^2 + 2F_c^2)/3$

$S = 0.81$ $(\Delta/\sigma)_{\max} < 0.001$
 3939 reflections $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$
 172 parameters $\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$
 Primary atom site location: structure-invariant direct methods Extinction correction: none

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

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 $F^2 > 2\sigma(F^2)$ 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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
N1	0.43672 (9)	0.39900 (18)	0.20431 (6)	0.0308 (4)	
N2	0.47899 (9)	0.57711 (17)	0.16614 (5)	0.0313 (3)	
N10	0.28403 (9)	-0.04257 (18)	0.23365 (6)	0.0301 (3)	
C3	0.41609 (11)	0.5733 (2)	0.09612 (6)	0.0289 (4)	
C3A	0.33286 (10)	0.3874 (2)	0.08653 (6)	0.0273 (4)	
C4	0.24616 (11)	0.2997 (2)	0.02588 (6)	0.0321 (4)	
C5	0.18152 (11)	0.1101 (2)	0.03769 (6)	0.0324 (4)	
C5A	0.19858 (10)	0.0022 (2)	0.10985 (6)	0.0283 (4)	
C5B	0.14567 (10)	-0.1954 (2)	0.13948 (6)	0.0300 (4)	
C6	0.05963 (11)	-0.3574 (2)	0.10872 (7)	0.0377 (4)	
C7	0.03272 (12)	-0.5330 (2)	0.15464 (8)	0.0426 (5)	
C8	0.09011 (12)	-0.5508 (2)	0.22997 (8)	0.0417 (5)	
C9	0.17617 (11)	-0.3965 (2)	0.26261 (7)	0.0344 (4)	
C9A	0.20113 (10)	-0.2182 (2)	0.21583 (7)	0.0291 (4)	
C10A	0.28287 (10)	0.0885 (2)	0.16958 (6)	0.0261 (4)	
C11	0.35017 (10)	0.2806 (2)	0.15764 (6)	0.0263 (4)	
C31	0.43731 (12)	0.7499 (2)	0.03983 (7)	0.0374 (4)	
C91	0.24276 (13)	-0.4178 (3)	0.34279 (7)	0.0503 (6)	
H1	0.4739 (12)	0.358 (2)	0.2515 (7)	0.048 (4)*	
H4	0.23351	0.37154	-0.02181	0.0385*	
H5	0.12498	0.05020	-0.00244	0.0388*	
H6	0.02084	-0.34717	0.05786	0.0452*	
H7	-0.02538	-0.64237	0.13466	0.0510*	
H8	0.06942	-0.67235	0.25954	0.0500*	
H10	0.3347 (13)	-0.015 (2)	0.2779 (7)	0.053 (4)*	
H31A	0.49975	0.85390	0.06335	0.0560*	0.500
H31B	0.36626	0.83888	0.02215	0.0560*	0.500
H31C	0.45969	0.67151	-0.00261	0.0560*	0.500

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H31D	0.38405	0.72230	-0.00809	0.0560*	0.500
H31E	0.51754	0.73732	0.03311	0.0560*	0.500
H31F	0.42411	0.90468	0.05787	0.0560*	0.500
H91A	0.21018	-0.54402	0.36762	0.0754*	
H91B	0.23653	-0.27322	0.36946	0.0754*	
H91C	0.32456	-0.44983	0.34305	0.0754*	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0319 (6)	0.0337 (7)	0.0245 (6)	-0.0023 (5)	0.0006 (5)	0.0018 (5)
N2	0.0321 (6)	0.0309 (6)	0.0301 (6)	-0.0016 (5)	0.0043 (5)	0.0025 (5)
N10	0.0270 (6)	0.0348 (6)	0.0274 (6)	-0.0005 (5)	0.0027 (5)	0.0021 (5)
C3	0.0299 (7)	0.0297 (7)	0.0273 (7)	0.0069 (6)	0.0060 (5)	0.0003 (6)
C3A	0.0276 (7)	0.0287 (7)	0.0259 (6)	0.0058 (6)	0.0059 (5)	-0.0022 (5)
C4	0.0340 (7)	0.0383 (8)	0.0221 (6)	0.0067 (7)	0.0016 (5)	-0.0004 (6)
C5	0.0293 (7)	0.0367 (8)	0.0283 (7)	0.0027 (6)	-0.0008 (5)	-0.0057 (6)
C5A	0.0258 (7)	0.0289 (7)	0.0291 (7)	0.0049 (6)	0.0029 (5)	-0.0031 (6)
C5B	0.0244 (7)	0.0298 (7)	0.0360 (7)	0.0024 (6)	0.0066 (5)	-0.0049 (6)
C6	0.0289 (7)	0.0399 (8)	0.0426 (8)	0.0008 (6)	0.0037 (6)	-0.0087 (7)
C7	0.0336 (8)	0.0353 (8)	0.0606 (10)	-0.0067 (7)	0.0139 (7)	-0.0103 (7)
C8	0.0370 (8)	0.0338 (8)	0.0595 (10)	0.0011 (7)	0.0219 (7)	0.0033 (7)
C9	0.0313 (7)	0.0329 (8)	0.0419 (8)	0.0032 (6)	0.0143 (6)	0.0029 (6)
C9A	0.0253 (7)	0.0289 (7)	0.0340 (7)	0.0035 (6)	0.0083 (5)	-0.0020 (6)
C10A	0.0269 (7)	0.0273 (7)	0.0241 (6)	0.0048 (6)	0.0049 (5)	-0.0001 (5)
C11	0.0246 (6)	0.0288 (7)	0.0245 (6)	0.0035 (6)	0.0024 (5)	-0.0033 (6)
C31	0.0419 (8)	0.0352 (8)	0.0359 (7)	0.0020 (7)	0.0100 (6)	0.0042 (6)
C91	0.0505 (10)	0.0557 (10)	0.0467 (9)	0.0045 (8)	0.0147 (7)	0.0151 (7)

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

N1—N2	1.3798 (14)	C8—C9	1.3774 (18)
N1—C11	1.3579 (16)	C9—C9A	1.3960 (17)
N2—C3	1.3271 (14)	C9—C91	1.5030 (18)
N10—C9A	1.3882 (16)	C10A—C11	1.3956 (16)
N10—C10A	1.3792 (15)	C4—H4	0.9400
N1—H1	0.906 (13)	C5—H5	0.9400
N10—H10	0.906 (13)	C6—H6	0.9400
C3—C3A	1.4289 (17)	C7—H7	0.9400
C3—C31	1.4927 (17)	C8—H8	0.9400
C3A—C4	1.4227 (16)	C31—H31A	0.9700
C3A—C11	1.4013 (15)	C31—H31B	0.9700
C4—C5	1.3653 (17)	C31—H31C	0.9700
C5—C5A	1.4211 (15)	C31—H31D	0.9700
C5A—C5B	1.4456 (16)	C31—H31E	0.9700
C5A—C10A	1.3940 (16)	C31—H31F	0.9700
C5B—C6	1.3972 (17)	C91—H91A	0.9700
C5B—C9A	1.4056 (16)	C91—H91B	0.9700
C6—C7	1.3823 (18)	C91—H91C	0.9700

C7—C8	1.393 (2)		
N1…N10	3.2036 (15)	C10A…H31F ^{iv}	3.0400
N1…N2 ⁱ	2.9878 (14)	C11…H31F ^{iv}	3.0500
N1…N10 ⁱⁱ	3.2360 (15)	C31…H31C ^v	2.8400
N2…N10 ⁱⁱ	3.0558 (15)	C91…H10	2.896 (13)
N2…N1 ⁱⁱ	2.9878 (14)	C91…H7 ^{vii}	3.0800
N10…N1 ⁱ	3.2360 (15)	C91…H31D ^{viii}	3.0700
N10…N1	3.2036 (15)	H1…N1 ⁱ	2.886 (12)
N10…N2 ⁱ	3.0558 (15)	H1…N2 ⁱ	2.188 (12)
N1…H10 ⁱⁱ	2.664 (15)	H1…N10 ⁱⁱ	2.837 (14)
N1…H1 ⁱⁱ	2.886 (12)	H1…H10 ⁱⁱ	2.51 (2)
N2…H1 ⁱⁱ	2.188 (12)	H5…H6 ^{vi}	2.4700
N2…H10 ⁱⁱ	2.263 (15)	H6…C5 ^{vi}	2.9600
N10…H91B	2.9400	H6…H5 ^{vi}	2.4700
N10…H1 ⁱ	2.837 (14)	H7…C91 ^{ix}	3.0800
C3…C5A ⁱⁱⁱ	3.5792 (17)	H7…H91B ^{ix}	2.5600
C3A…C5B ⁱⁱⁱ	3.5027 (17)	H8…H91A	2.4000
C4…C6 ⁱⁱⁱ	3.4897 (18)	H8…C7 ^{ix}	2.9800
C5A…C31 ^{iv}	3.5894 (18)	H8…C8 ^{ix}	2.8900
C5A…C7 ⁱⁱⁱ	3.4865 (17)	H8…C9 ^{ix}	3.0900
C5A…C3 ^{iv}	3.5792 (17)	H10…C91	2.896 (13)
C5B…C3A ^{iv}	3.5027 (17)	H10…N1 ⁱ	2.664 (15)
C6…C4 ^{iv}	3.4897 (18)	H10…N2 ⁱ	2.263 (15)
C7…C5A ^{iv}	3.4865 (17)	H10…H1 ⁱ	2.51 (2)
C8…C10A ^{iv}	3.3959 (18)	H31B…C4 ⁱⁱⁱ	3.0000
C9…C11 ^{iv}	3.5674 (17)	H31B…C5 ⁱⁱⁱ	2.7200
C10A…C8 ⁱⁱⁱ	3.3959 (18)	H31B…C5A ⁱⁱⁱ	2.9100
C11…C9 ⁱⁱⁱ	3.5674 (17)	H31C…C3 ^v	2.8200
C31…C5A ⁱⁱⁱ	3.5894 (18)	H31C…C31 ^v	2.8400
C3…H31C ^v	2.8200	H31D…C4	3.0400
C3A…H31F ^{iv}	3.0500	H31D…C91 ^x	3.0700
C4…H31F ^{iv}	3.0500	H31D…H91B ^x	2.5300
C4…H31B ^{iv}	3.0000	H31F…C3A ⁱⁱⁱ	3.0500
C4…H31D	3.0400	H31F…C4 ⁱⁱⁱ	3.0500
C5…H31B ^{iv}	2.7200	H31F…C5 ⁱⁱⁱ	3.0200
C5…H6 ^{vi}	2.9600	H31F…C5A ⁱⁱⁱ	3.0200
C5…H31F ^{iv}	3.0200	H31F…C10A ⁱⁱⁱ	3.0400
C5A…H31F ^{iv}	3.0200	H31F…C11 ⁱⁱⁱ	3.0500
C5A…H31B ^{iv}	2.9100	H91A…H8	2.4000
C7…H8 ^{vii}	2.9800	H91B…N10	2.9400
C8…H8 ^{vii}	2.8900	H91B…H7 ^{vii}	2.5600

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C9...H8 ^{vii}	3.0900	H91B...H31D ^{viii}	2.5300
N2—N1—C11	110.86 (10)	C5—C4—H4	120.00
N1—N2—C3	106.31 (10)	C4—C5—H5	120.00
C9A—N10—C10A	108.34 (10)	C5A—C5—H5	120.00
C11—N1—H1	127.4 (8)	C5B—C6—H6	121.00
N2—N1—H1	120.9 (8)	C7—C6—H6	121.00
C9A—N10—H10	129.0 (8)	C6—C7—H7	119.00
C10A—N10—H10	122.6 (8)	C8—C7—H7	119.00
N2—C3—C31	120.20 (11)	C7—C8—H8	119.00
C3A—C3—C31	128.88 (10)	C9—C8—H8	119.00
N2—C3—C3A	110.92 (10)	C3—C31—H31A	109.00
C3—C3A—C11	104.40 (10)	C3—C31—H31B	109.00
C4—C3A—C11	119.91 (11)	C3—C31—H31C	109.00
C3—C3A—C4	135.69 (10)	C3—C31—H31D	109.00
C3A—C4—C5	119.30 (10)	C3—C31—H31E	109.00
C4—C5—C5A	120.87 (10)	C3—C31—H31F	109.00
C5—C5A—C5B	133.80 (11)	H31A—C31—H31B	109.00
C5—C5A—C10A	120.03 (11)	H31A—C31—H31C	109.00
C5B—C5A—C10A	106.16 (9)	H31A—C31—H31D	141.00
C5A—C5B—C9A	106.96 (10)	H31A—C31—H31E	56.00
C6—C5B—C9A	118.52 (11)	H31A—C31—H31F	56.00
C5A—C5B—C6	134.50 (10)	H31B—C31—H31C	109.00
C5B—C6—C7	118.62 (11)	H31B—C31—H31D	56.00
C6—C7—C8	121.21 (12)	H31B—C31—H31E	141.00
C7—C8—C9	122.26 (12)	H31B—C31—H31F	56.00
C8—C9—C91	123.24 (12)	H31C—C31—H31D	56.00
C9A—C9—C91	120.90 (12)	H31C—C31—H31E	56.00
C8—C9—C9A	115.84 (12)	H31C—C31—H31F	141.00
N10—C9A—C5B	108.59 (10)	H31D—C31—H31E	109.00
N10—C9A—C9	127.84 (11)	H31D—C31—H31F	109.00
C5B—C9A—C9	123.53 (11)	H31E—C31—H31F	109.00
N10—C10A—C5A	109.94 (10)	C9—C91—H91A	109.00
N10—C10A—C11	130.68 (10)	C9—C91—H91B	109.00
C5A—C10A—C11	119.37 (10)	C9—C91—H91C	109.00
N1—C11—C10A	132.01 (10)	H91A—C91—H91B	109.00
C3A—C11—C10A	120.51 (10)	H91A—C91—H91C	109.00
N1—C11—C3A	107.48 (10)	H91B—C91—H91C	109.00
C3A—C4—H4	120.00		
C11—N1—N2—C3	1.77 (13)	C10A—C5A—C5B—C6	-178.14 (13)
N2—N1—C11—C3A	-1.32 (13)	C10A—C5A—C5B—C9A	-0.06 (14)
N2—N1—C11—C10A	179.04 (12)	C5—C5A—C10A—N10	-179.96 (11)
N1—N2—C3—C3A	-1.52 (13)	C5—C5A—C10A—C11	-0.21 (17)
N1—N2—C3—C31	178.28 (11)	C5B—C5A—C10A—N10	-0.44 (13)
C10A—N10—C9A—C5B	-0.82 (13)	C5B—C5A—C10A—C11	179.31 (11)
C10A—N10—C9A—C9	177.07 (12)	C5A—C5B—C6—C7	177.92 (13)
C9A—N10—C10A—C5A	0.79 (14)	C9A—C5B—C6—C7	0.00 (19)
C9A—N10—C10A—C11	-178.93 (12)	C5A—C5B—C9A—N10	0.54 (13)
N2—C3—C3A—C4	-179.12 (13)	C5A—C5B—C9A—C9	-177.46 (11)

N2—C3—C3A—C11	0.74 (14)	C6—C5B—C9A—N10	178.97 (11)
C31—C3—C3A—C4	1.1 (2)	C6—C5B—C9A—C9	0.97 (18)
C31—C3—C3A—C11	-179.03 (12)	C5B—C6—C7—C8	-0.41 (19)
C3—C3A—C4—C5	179.16 (13)	C6—C7—C8—C9	-0.1 (2)
C11—C3A—C4—C5	-0.69 (18)	C7—C8—C9—C9A	1.05 (19)
C3—C3A—C11—N1	0.36 (13)	C7—C8—C9—C91	-177.40 (13)
C3—C3A—C11—C10A	-179.94 (12)	C8—C9—C9A—N10	-179.07 (12)
C4—C3A—C11—N1	-179.75 (11)	C8—C9—C9A—C5B	-1.48 (18)
C4—C3A—C11—C10A	-0.06 (18)	C91—C9—C9A—N10	-0.6 (2)
C3A—C4—C5—C5A	0.98 (18)	C91—C9—C9A—C5B	177.01 (12)
C4—C5—C5A—C5B	-179.90 (13)	N10—C10A—C11—N1	-0.2 (2)
C4—C5—C5A—C10A	-0.54 (18)	N10—C10A—C11—C3A	-179.81 (12)
C5—C5A—C5B—C6	1.3 (2)	C5A—C10A—C11—N1	-179.90 (12)
C5—C5A—C5B—C9A	179.36 (13)	C5A—C10A—C11—C3A	0.49 (17)

Symmetry codes: (i) $-x+1, y-1/2, -z+1/2$; (ii) $-x+1, y+1/2, -z+1/2$; (iii) $x, y+1, z$; (iv) $x, y-1, z$; (v) $-x+1, -y+1, -z$; (vi) $-x, -y, -z$; (vii) $-x, y+1/2, -z+1/2$; (viii) $x, -y+1/2, z+1/2$; (ix) $-x, y-1/2, -z+1/2$; (x) $x, -y+1/2, z-1/2$.

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1 \cdots N2 ⁱ	0.906 (13)	2.188 (12)	2.9878 (14)	147.0 (10)
N10—H10 \cdots N2 ⁱ	0.906 (13)	2.263 (15)	3.0558 (15)	145.9 (12)

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

Fig. 1

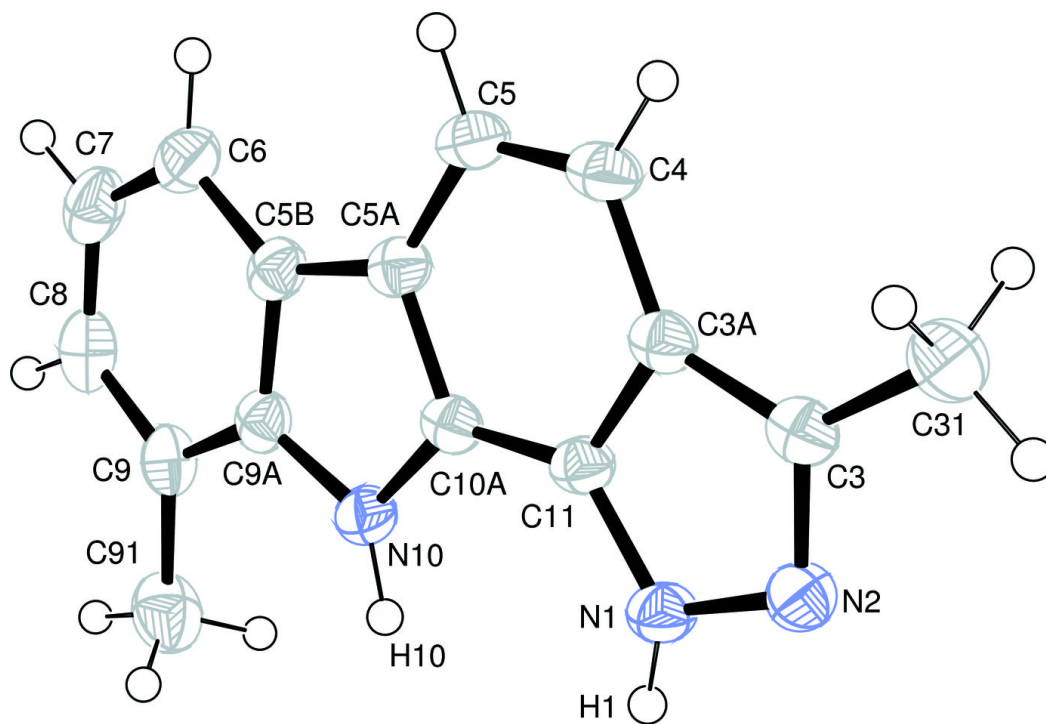


Fig. 2

