

3,5-Dimethyl-1,7-diphenyl-4-(2-pyridyl)-
1*H*,7*H*-pyrazolo[3,4-*b*:4',3'-*e*]pyridineJohn Nicolson Low,^{a*†}
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Jairo Quiroga^c^aDepartment of Chemistry, University of Aberdeen, Meston Walk, Old Aberdeen AB24 3UE, Scotland, ^bDepartamento de Química Inorgánica y Orgánica, Universidad de Jaén, 23071 Jaén, Spain, and ^cGrupo de Investigación de Compuestos Heterocíclicos, Departamento de Química, Universidad de Valle, AA 25360, Cali, Colombia

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Key indicators

Single-crystal X-ray study

T = 120 K

Mean $\sigma(\text{C}-\text{C}) = 0.002 \text{ \AA}$

R factor = 0.040

wR factor = 0.110

Data-to-parameter ratio = 15.9

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.In the title compound, $\text{C}_{26}\text{H}_{20}\text{N}_6$, the supramolecular structure is determined by a weak $\text{C}-\text{H}\cdots\text{N}$ hydrogen bond which links the molecules into chains running parallel to [110].Received 18 July 2003
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Comment

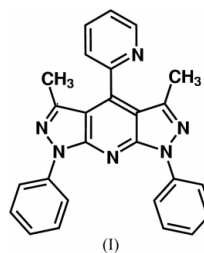
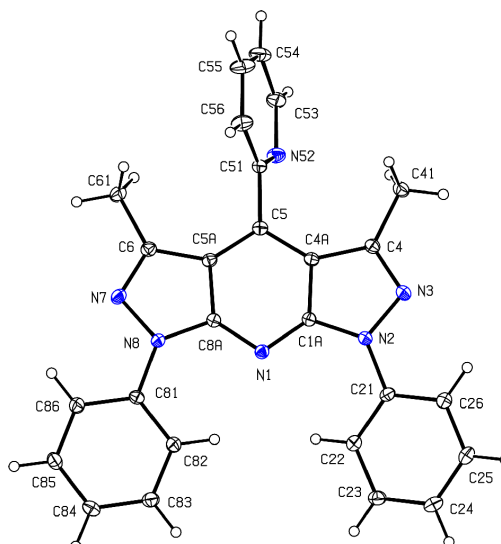
In our ongoing application of microwave irradiation to heterocyclic synthesis by multicomponent condensation, we have prepared compound (I) from 5-amino-3-methyl-1-phenyl-1*H*-pyrazole and 2-pyridinecarbaldehyde.There are no unusual bonds and angles in the structure (Table 1 and Fig. 1). The 12-membered fused-ring system defined by atoms N1 to C8A is essentially planar and symmetrical about the $\text{N1}\cdots\text{C5}$ line, as shown by the bonds and angles in Table 1. However, the angle of tilt of the phenyl rings out of the plane of the 12-membered ring system varies as is shown by the torsion angles listed in Table 1 so that this

Figure 1

A view of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

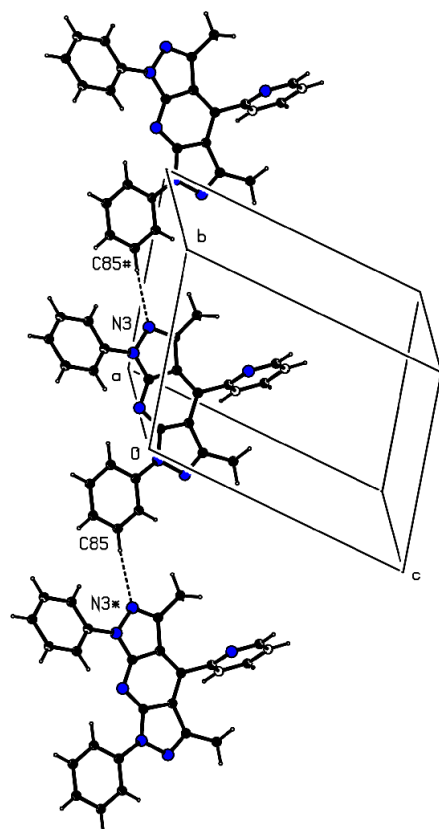


Figure 2

A view of the $C(10)$ chains running parallel to $[110]$. The atom labelled with an asterisk (*) is in the molecule at $(x - 1, y - 1, z)$ while that labelled with a hash (#) is in the molecule at $(1 + x, 1 + y, z)$.

symmetry does not extend to the whole molecule. There are two short intramolecular contacts $C22-H22 \cdots N1$ and $C82-H82 \cdots N1$, *i.e.* between the two phenyl rings defined by atoms $C21-C26$ and $C81-C86$ to atom $N1$ (Table 2). The $C8-H85 \cdots N3$ weak hydrogen bond links the molecules into $C(10)$ chains (Bernstein *et al.*, 1995) which run parallel to $[110]$ (Fig. 2).

Experimental

A mixture of 5-amino-3-methyl-1-phenylpyrazole (10 mmol) and 2-pyridinecarbaldehyde (20 mmol) was placed into Pyrex-glass open vessels and irradiated in a domestic microwave oven for 1.5 min (at 600 W). The solid was filtered, washed with ethanol, dried and recrystallized from ethanol. Yield 68%, m.p. 517 K. Analysis calculated for $C_{26}H_{20}N_6$: C 74.98, H 4.84, N 20.18%; found: C 74.93, H 4.78, N 20.15%.

Crystal data

$C_{26}H_{20}N_6$
 $M_r = 416.48$
 Triclinic, $P\bar{1}$
 $a = 8.4576$ (2) Å
 $b = 10.6599$ (3) Å
 $c = 12.3642$ (4) Å
 $\alpha = 106.9170$ (14)°
 $\beta = 98.6240$ (16)°
 $\gamma = 100.803$ (2)°
 $V = 1022.48$ (5) Å³

$Z = 2$
 $D_x = 1.353$ Mg m⁻³
 Mo $K\alpha$ radiation
 Cell parameters from 4637 reflections
 $\theta = 3.1-27.5^\circ$
 $\mu = 0.08$ mm⁻¹
 $T = 120.0$ (2) K
 Block, brown
 $0.28 \times 0.22 \times 0.18$ mm

Data collection

Nonius KappaCCD diffractometer
 φ scans and ω scans with κ offsets
 Absorption correction: multi-scan
 (DENZO-SMN; Otwinowski & Minor, 1997)
 $T_{\min} = 0.977$, $T_{\max} = 0.985$
 19 117 measured reflections

4637 independent reflections
 3732 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.050$
 $\theta_{\max} = 27.5^\circ$
 $h = -10 \rightarrow 10$
 $k = -13 \rightarrow 13$
 $l = -16 \rightarrow 15$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.110$
 $S = 1.03$
 4637 reflections
 291 parameters
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0603P)^2 + 0.1734P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.25$ e Å⁻³
 $\Delta\rho_{\min} = -0.25$ e Å⁻³

Table 1

Selected geometric parameters (Å, °).

N1—C1A	1.3347 (15)	N52—C53	1.3400 (16)
N1—C8A	1.3364 (15)	N52—C51	1.3420 (15)
N2—N3	1.3879 (13)	C1A—C4A	1.4206 (15)
N3—C4	1.3139 (15)	C4—C4A	1.4375 (16)
N7—C6	1.3147 (15)	C4A—C5	1.3953 (17)
N7—N8	1.3896 (13)	C5—C5A	1.3954 (16)
N8—C8A	1.3717 (15)	C5A—C8A	1.4240 (15)
C1A—N1—C8A	110.97 (9)	C5A—C5—C4A	115.03 (10)
C1A—N2—N3	110.74 (9)	C5A—C5—C51	122.83 (10)
C1A—N2—C21	130.19 (10)	C4A—C5—C51	122.07 (10)
N3—N2—C21	119.06 (9)	C5—C5A—C8A	119.47 (10)
C4—N3—N2	107.25 (9)	C5—C5A—C6	135.71 (11)
C6—N7—N8	107.25 (10)	C8A—C5A—C6	104.67 (10)
C8A—N8—N7	110.61 (9)	N7—C6—C5A	110.96 (10)
C8A—N8—C81	130.72 (10)	N7—C6—C61	119.88 (11)
N7—N8—C81	118.49 (9)	C5A—C6—C61	129.15 (11)
C53—N52—C51	117.14 (10)	N1—C8A—N8	126.07 (10)
N1—C1A—N2	125.73 (10)	N1—C8A—C5A	127.41 (10)
N1—C1A—C4A	127.79 (10)	N8—C8A—C5A	106.49 (10)
N2—C1A—C4A	106.48 (10)	N52—C51—C56	122.88 (11)
N3—C4—C4A	110.79 (10)	N52—C51—C5	115.80 (10)
N3—C4—C41	120.45 (11)	N52—C53—C54	123.90 (12)
C4A—C4—C41	128.74 (11)	C53—C54—C55	118.28 (12)
C5—C4A—C1A	119.31 (10)	C54—C55—C56	119.28 (12)
C5—C4A—C4	135.98 (11)	C55—C56—C51	118.50 (12)
C1A—C4A—C4	104.71 (10)		
C1A—N2—C21—C22	22.81 (18)	C5A—C5—C51—C56	70.95 (16)
N3—N2—C21—C22	-156.16 (10)	C4A—C5—C51—C56	-112.34 (14)
C1A—N2—C21—C26	-158.64 (12)	C8A—N8—C81—C82	-3.78 (18)
N3—N2—C21—C26	22.40 (15)	N7—N8—C81—C82	170.86 (10)
C5A—C5—C51—N52	-108.40 (13)	C8A—N8—C81—C86	177.07 (11)
C4A—C5—C51—N52	68.30 (14)	N7—N8—C81—C86	-8.29 (15)

Table 2

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$C22-H22 \cdots N1$	0.95	2.43	3.0029 (16)	119
$C82-H82 \cdots N1$	0.95	2.31	2.9721 (16)	126
$C85-H85 \cdots N3^i$	0.95	2.55	3.4922 (18)	173

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

H atoms were treated as riding atoms, with $C-H = 0.95$ Å (aromatic) and 0.98 Å (methyl), and U_{iso} values of $1.2U_{\text{eq}}$ (aromatic C) and $1.5U_{\text{eq}}$ (methyl C).

Data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data

reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *WordPerfect* macro *PRPKAPPA* (Ferguson, 1999).

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