# organic papers

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#### **Key indicators**

Single-crystal X-ray study T = 293 K Mean  $\sigma$ (C–C) = 0.004 Å Disorder in main residue R factor = 0.043 wR factor = 0.079 Data-to-parameter ratio = 12.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

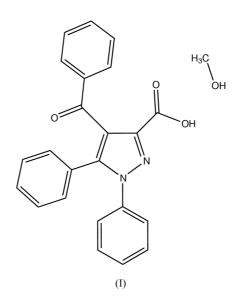
# 4-Benzoyl-1,5-diphenyl-1*H*-pyrazole-3carboxylic acid methanol solvate

In the title compound,  $C_{23}H_{16}N_2O_3 \cdot CH_4O$ , the N–N bond distance in the pyrazole ring, which is planar within 0.008 Å, is 1.3634 (18) Å. The crystal packing is stabilized by O–H···O, O–H···N and C–H···O intermolecular hydrogen bonds.

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## Comment

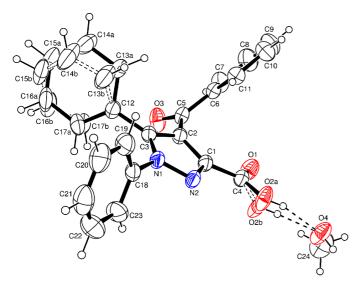
Derivatives of pyrazole continue to attract interest because of their wide spectrum of biological, medicinal (Badawey & El-Ashmawey, 1998) and agricultural (Thomson, 1997) activities. Some pyrazoles have been reported to possess significant antimicrobial (Mahajan et al., 1991), antiviral (Baraldi et al., 1998), antitumor (Hatheway et al., 1978; Katayama & Oshiyama, 1997), antifungal (Chen & Li, 1998), pesticidal (Londershausen, 1996), anticonvulsant (Lepage & Hublot, 1992), antihistaminic (Mishra et al., 1998) and antidepressant activities (Bailley et al., 1985), as well as interesting properties in commercially important dyestuffs (Baroni & Kovyrzina, 1961; Neunhoeffer et al., 1959). Recyclization reactions of 4benzoyl-5-phenyl-2,3-dihydro-2,3-furandione with various phenylhydrazones and phenylhydrazine lead to pyrazolecarboxylic acid and pyridazinones (Akçamur et al., 1986, 1997; Sener et al., 2002, 2004; Verirsen & Erturan, 1998). As part of our ongoing study of the relationship between the molecular and crystal structures of pyrazole derivatives, a crystal structure determination of the title compound, (I), has been undertaken and the results are presented here.



Previously, we have reported closely related compounds, namely methyl 4-benzoyl-1,5-diphenyl-1*H*-pyrazole-3-carboxylate, (II) (Dinçer, Özdemir, Yıldırım, Demir, Akçamur & Işık, 2004), and 4-benzoyl-*N*-methyl-1,5-diphenyl-1*H*-pyra-

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**0256** Yıldırım et al. • C<sub>23</sub>H<sub>16</sub>N<sub>2</sub>O<sub>3</sub>·CH<sub>4</sub>O





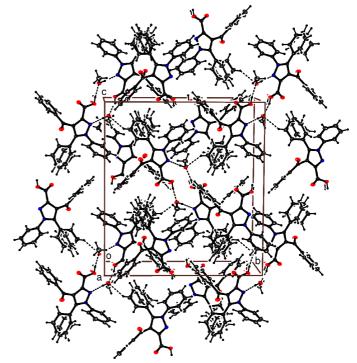
An ORTEP-3 (Farrugia, 1997) drawing of the title compound, (I), with the atomic numbering scheme; both disordered components are shown. Displacement ellipsoids of non-H atoms are drawn at the 30% probability level. The intermolecular O-H···O hydrogen bonds are represented by dashed lines.

zole-3-carboxamide, (III) (Dinçer, Özdemir, Yıldırım, Demir & Işık, 2004). The main aim of the present investigation is to study the differences between the structures of (I), (II) and (III), and to determine the type of hydrogen bonding.

The conformation of (I), together with the atom-numbering scheme, is shown in Fig. 1. In the structure, the phenyl ring at the 5-position and the O atom (O2) belonging to the carboxylic acid group show positional disorder and the refined site-occupancy factors of the disordered components (C13a-C17a and C13b-C17b, and O2a and O2b) are 0.51 (1) and 0.49 (1), respectively. The planes of the C13a-C17a and C13b-C17b parts seem to be rotated with respect to each other about the C12 $\cdots$ C15 vector. The dihedral angle between these two planes, including atom C12, is  $36.7 (8)^{\circ}$ .

As expected, all five rings [A (N1/N2/C1-C3), B (C6-C11), C (C12–C17a), C' (C12–C17b) and D (C18–C23)] are essentially planar. The maximum deviations from the leastsquares plane of the pyrazole ring is 0.008 (1) Å for atom N2. However, all fragments in the structure do not share a common plane. The dihedral angles between the mean planes of the rings are 71.45 (A/B), 65.67 (A/C), 77.90 (A/C'), 66.70 (A/D), 85.48 (B/C), 78.53 (B/C'), 43.37 (B/D), 67.59 (C/D) and 77.17° (C'/D). The  $\Phi_{CN}$  torsion angle (C12–C3–N1–C18) is  $4.0 (4)^{\circ}$ , which shows that the conformation about the C3–N1 bond is (+)synperiplanar. The  $\Phi_{CC}$  (C5–C2–C3–C12 and C4– C1–C2–C5) torsion angles are -4.3(4) and  $-1.8(4)^{\circ}$ , respectively, showing that the conformations about the C2–C3 and C1–C2 bonds are (-)synperiplanar.

The N-N bond distance is 1.363 (2) Å. Comparison of the bond lengths and angles of the pyrazole rings in (I), (II) and (III) shows that there are no important differences in (I) with respect to the others. The crystal structure contains intermolecular hydrogen bonds (Table 2). As can be seen in Fig. 2, the methanol solvent molecule plays an active role in linking



### Figure 2

A projection of the crystal structure of (I) approximately along the a axis. Dashed lines show the intermolecular interactions.

the molecules. The hydrogen bonds between atoms O2 and O4, acting both as donors and acceptors, together with O-H···N hydrogen bonds form an R(14) ring. These interactions along with the C-H···O interactions (Table 2) generate a two-dimensional network.

## **Experimental**

4-Benzoyl-5-phenyl-2,3-dihydro-2,3-furandione (2 g, 7.15 mmol) (Ziegler et al., 1967) and phenylhydrazine (0.81 g, 7.16 mmol), as described by Akçamur et al. (1986), were refluxed in boiling benzene (30 ml) for 1-1.5 h. The solvent was removed by evaporation and the oily residue was triturated with dry diethyl ether to give a crude solid, which was then recrystallized from methanol, yielding pure 4benzoyl-1,5-diphenyl-1H-pyrazole-3-carboxylic acid, (I) (1.31 g, 49%; m.p. 468-470 K). Solvents were dried by refluxing with the appropriate drying agents and distilled before use. All other reagents were purchased from Merck, Fluka, Aldrich or the Acros Chemical Co., and used without further purification. Analysis calculated for C23H16N2O3: C 74.99, H 4.38, N 7.60%; found: C 75.20, H 4.50, N 7.47%.

$C_{23}H_{16}N_2O_3 \cdot CH_4O$	Mo $K\alpha$ radiation		
$M_r = 400.42$	Cell parameters from 8043		
Orthorhombic, Pcab	reflections		
a = 10.8731 (9) Å	$\theta = 2.0-24.5^{\circ}$		
b = 18.7861 (14) Å	$\mu = 0.09 \text{ mm}^{-1}$		
$c = 20.5836 (19) \text{\AA}$	T = 293 (2) K		
V = 4204.5 (6) Å <sup>3</sup>	Prism, colorless		
Z = 8	$0.50 \times 0.33 \times 0.21 \text{ mm}$		
$D_x = 1.265 \text{ Mg m}^{-3}$			

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Data collection

Stoe IPDS-II diffractometer	1743 reflections with $I > 2\sigma(I)$
$\omega$ scans	$R_{\rm int} = 0.099$
Absorption correction: integration	$\theta_{\rm max} = 25.5^{\circ}$
(X-RED32; Stoe & Cie, 2002)	$h = -13 \rightarrow 13$
$T_{\min} = 0.965, T_{\max} = 0.984$	$k = -22 \rightarrow 22$
27 164 measured reflections	$l = -24 \rightarrow 24$
3914 independent reflections	

### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.043$	$w = 1/[\sigma^2(F_o^2) + (0.0154P)^2]$
$wR(F^2) = 0.079$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
3914 reflections	$\Delta \rho_{\rm max} = 0.11 \text{ e } \text{\AA}^{-3}$
327 parameters	$\Delta \rho_{\rm min} = -0.20 \text{ e } \text{\AA}^{-3}$

### Table 1

Selected geometric parameters (Å, °).

0 1			
01-C4	1.186 (2)	N1-C18	1.449 (2)
O2a - C4	1.300 (6)	N2-C1	1.331 (2)
O2b-C4	1.308 (8)	C1-C2	1.398 (3)
O3-C5	1.218 (2)	C1-C4	1.502 (3)
O4-C24	1.385 (3)	C2-C3	1.388 (2)
N1-C3	1.346 (2)	C2-C5	1.475 (3)
N1-N2	1.3634 (18)	C3-C12	1.483 (3)
C3-N1-N2	112.36 (15)	N1-C3-C2	107.07 (17)
C3-N1-C18	130.22 (16)	N1-C3-C12	123.02 (17)
N2-N1-C18	117.33 (18)	C2-C3-C12	129.9 (2)
C1-N2-N1	103.84 (16)	O1-C4-O2a	124.5 (4)
N2-C1-C2	112.77 (15)	O1 - C4 - O2b	125.5 (5)
N2-C1-C4	119.16 (19)	O1-C4-C1	122.7 (2)
C2-C1-C4	127.9 (2)	O2a - C4 - C1	112.3 (4)
C3-C2-C1	103.93 (18)	O2b-C4-C1	111.0 (5)
C3-C2-C5	126.19 (18)	O3-C5-C2	119.6 (2)
C1-C2-C5	129.81 (17)	O3-C5-C6	121.0 (2)
C4-C1-C2-C5	-1.8(4)	C5-C2-C3-C12	-4.3 (4)
C18-N1-C3-C12	4.0 (4)	00 02 05 012	1.5 (1)

Table 2

Hydrogen-bond geometry (Å, °).

$\overline{D - \mathbf{H} \cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O4-H4\cdots N2^i$	0.82	1.99	2.804 (2)	176
$O2b - H2b \cdots O4$	0.82	1.77	2.583 (10)	171
$O2a - H2a \cdot \cdot \cdot O4$	0.82	1.77	2.576 (8)	165
$C10-H10\cdots O1^{ii}$	0.93	2.46	3.339 (3)	158
$C14a - H14a \cdot \cdot \cdot O4^{iii}$	0.93	2.53	3.334 (6)	145

Symmetry codes: (i) -x + 1, -y, -z; (ii)  $x + \frac{1}{2}$ ,  $-y + \frac{1}{2}$ , z; (iii) -x + 1,  $-y + \frac{1}{2}$ ,  $z + \frac{1}{2}$ .

H atoms were positioned geometrically and treated using a riding model, fixing the bond lengths at 0.82 Å for O-bound H atoms, 0.93 Å

for the benzene ring H atoms and 0.96 Å for the methyl group H atoms. The displacement parameters of the H atoms were constrained as  $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}$  (1.5 $U_{\rm eq}$  for methyl atoms) of the carrier atom.

Data collection: X-AREA (Stoe & Cie, 2002); cell refinement: X-AREA; data reduction: X-RED32 (Stoe & Cie, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999) and PLATON (Spek, 2003).

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