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#### **Key indicators**

Single-crystal X-ray study T = 298 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.036 wR factor = 0.091 Data-to-parameter ratio = 8.1

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

# 5-(2-Furyl)-3-methyl-1-(4-nitrophenyl)-2-pyrazoline

In the title compound,  $C_{14}H_{13}N_3O_3$ , the pyrazoline ring and the 4-nitrophenyl group are nearly coplanar, whereas the furyl and pyrazoline rings are roughly perpendicular. The occurrence of weak  $C-H\cdots O$  hydrogen-bonding interactions results in the formation of an  $R_3^3(15)$  ring. Received 30 October 2006 Accepted 6 November 2006

## Comment

The derivatives of pyrazoline are mostly used in medicine, for example as antitumor (Hatheway *et al.*, 1978), analgesic (Sobczak & Pawlaczyk, 1998), and antimicrobial (Mahajan *et al.*, 1991) agents. As part of our research, we have synthesized the new pyrazoline compound, (I).



The 4-nitrophenyl group and the pyrazoline ring are almost coplanar, making a dihedral angle of  $3.52 (14)^{\circ}$ , while the furyl ring is nearly perpendicular to the pyrazoline ring, with a dihedral angle of 89.61 (15)°. The dihedral angle between the furyl ring and the 4-nitrophenyl group is 87.93 (14)°. The N2–N3 bond length [1.400 (3) Å] agrees with that of a single bond (Burke-Laing & Laing, 1976).

An interesting feature is the occurrence of weak C–H···O hydrogen bonds involving two O atoms of the nitro group, leading to the formation of an  $R_3^3(15)$  graph-set motif (Etter *et al.*, 1990) (Table 1 and Fig. 2).

### **Experimental**

4-Nitrophenylhydrazine (1 mmol, 0.153 g) was dissolved in anhydrous methanol (15 ml), and  $H_2SO_4$  (98%, 0.5 ml) was added. The mixture was stirred for several minitutes at 351 K, furylideneacetone (1 mmol, 0.136 g) in methanol (8 ml) was added dropwise and the mixture was stirred at refluxing temperature for 2 h. The product was isolated and recrystallized from dichloromethane, brown single crystals of (I) being obtained after 1 d.

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#### Crystal data

 $C_{14}H_{13}N_{3}O_{3}$   $M_{r} = 271.27$ Monoclinic, P2<sub>1</sub> a = 8.1504 (13) Å b = 8.4608 (14) Å c = 9.6959 (16) Å  $\beta = 98.036$  (2)° V = 662.05 (19) Å<sup>3</sup>

#### Data collection

Bruker SMART CCD area-detector
diffractometer
$\omega$ scans
Absorption correction: multi-scan
SADABS(Bruker, 2000)
$T_{\min} = 0.974, \ T_{\max} = 0.981$

#### Refinement

Refinement on $F^2$	$w = 1/[\sigma^2(F_0^2) + (0.0428P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.036$	+ 0.0117P]
$wR(F^2) = 0.091$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.00	$(\Delta/\sigma)_{\rm max} < 0.001$
1470 reflections	$\Delta \rho_{\rm max} = 0.10 \ {\rm e} \ {\rm \AA}^{-3}$
182 parameters	$\Delta \rho_{\rm min} = -0.15 \text{ e } \text{\AA}^{-3}$
H-atom parameters constrained	

Та	ble	1	
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Hydrogen-bond	geometry	(A,	°).
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$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$\begin{array}{c} C2 - H2 \cdots O3^{i} \\ C10 - H10 \cdots O2^{ii} \end{array}$	0.93 0.98	2.51 2.47	3.233 (4) 3.369 (4)	135 153
	1 .			

Z = 2

 $D_x = 1.361 \text{ Mg m}^{-3}$ 

 $0.27 \times 0.23 \times 0.20$  mm

5482 measured reflections 1470 independent reflections 1047 reflections with  $I > 2\sigma(I)$ 

Mo  $K\alpha$  radiation

 $\mu = 0.10 \text{ mm}^{-1}$ 

T = 298 (2) K

Block, brown

 $R_{\rm int} = 0.027$  $\theta_{\rm max} = 26.5^{\circ}$ 

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

All H atoms were positioned geometrically and refined as riding with C–H = 0.93 (aromatic), 0.97 (methylene), 0.98 (methine) and 0.96 Å (methyl), with  $U_{\rm iso}(\rm H) = 1.2U_{eq}(\rm CH \ or \ CH_2)$  and  $U_{\rm iso}(\rm H) = 1.5U_{eq}(\rm CH_3)$ . In the absence of significant anomalous scattering, Friedel pairs were merged.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1998); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPIII* (Burnett & Johnson, 1996); *ORTEP-3 for Windows* (Farrugia, 1997); *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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#### Figure 1

The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level. H atoms are represented as small spheres of arbitrary radii.



#### Figure 2

Partial packing view, showing the formation of a pseudo-trimer through  $C-H\cdots O$  hydrogen bonding. Hydrogen bonds are shown as dashed lines. H atoms not involved in hydrogen bonding have been omitted for clarity. [Symmetry codes: (i) -x,  $y - \frac{1}{2}$ , 2 - z; (ii) x, y - 1, z].

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