

## 5-Hydroxy-3-phenyl-5-trifluoromethyl-4,5-dihydro-1H-pyrazole

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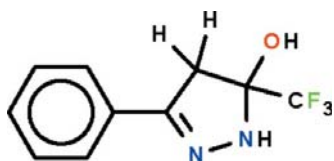
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Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å;  $R$  factor = 0.037;  $wR$  factor = 0.087; data-to-parameter ratio = 8.0.

The five-membered dihydropyrazole ring in the title compound,  $\text{C}_{10}\text{H}_9\text{F}_3\text{N}_2\text{O}$ , is approximately planar (r.m.s. deviation 0.111 Å for all non-H atoms) and its phenyl substituent is aligned at an angle of  $14.7(2)^\circ$ . Adjacent molecules are linked by  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{O}-\text{H}\cdots\text{N}$  hydrogen bonds, generating ribbons running along the  $b$  axis of the monoclinic unit cell.

### Related literature

For the synthesis, see: Yakimovich *et al.* (2002); Zelenin *et al.* (1995). For two related structures, see: Dias & Goh (2004); Yang & Raptis (2003).



### Experimental

#### Crystal data

$\text{C}_{10}\text{H}_9\text{F}_3\text{N}_2\text{O}$   
 $M_r = 230.19$   
Monoclinic,  $P2_1$

$a = 9.1000(6)$  Å  
 $b = 5.4032(3)$  Å  
 $c = 10.4515(7)$  Å

$\beta = 108.139(7)^\circ$   
 $V = 488.35(5)$  Å<sup>3</sup>  
 $Z = 2$   
Mo  $K\alpha$  radiation

$\mu = 0.14$  mm<sup>-1</sup>  
 $T = 100$  K  
 $0.20 \times 0.15 \times 0.10$  mm

#### Data collection

Agilent SuperNova Dual diffractometer with Atlas detector  
Absorption correction: multi-scan (*CrysAlis PRO*; Agilent, 2010)  
 $T_{\min} = 0.972$ ,  $T_{\max} = 0.986$

4222 measured reflections  
1230 independent reflections  
1060 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.039$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$   
 $wR(F^2) = 0.087$   
 $S = 1.05$   
1230 reflections  
153 parameters  
3 restraints

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.29$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.26$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1}\cdots\text{N1}^i$	0.84 (1)	2.03 (2)	2.833 (3)	162 (4)
$\text{N2}-\text{H2}\cdots\text{O1}^{ii}$	0.88 (1)	2.13 (2)	2.974 (3)	161 (3)

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5617).

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