

- Filippakis, S. E., Leiserowitz, L., Rabinovich, D. & Schmidt, G. M. J. (1972). *J. Chem. Soc. Perkin Trans. 2*, pp. 1750–1756.
- Hall, S. R., Flack, H. D. & Stewart, J. M. (1992). Editors. *Xtal3.2 Reference Manual*. Universities of Western Australia, Australia, Geneva, Switzerland, and Maryland, USA.
- He, Y., Shi, J. & Su, G. (1993). *Acta Cryst. C* **49**, 1432–1434.
- Johnson, C. K. (1965). *ORTEP*. Report ORNL-3794. Oak Ridge National Laboratory, Tennessee, USA.
- Keller, E. (1988). *SCHAKAL88. Fortran Program for the Graphic Representation of Molecular and Crystallographic Models*. University of Freiburg, Germany.
- Kim, M. J., Mang, J. Y. & Suh, I. H. (1988). *J. Korean Phys. Soc.* **21**, 405.
- Liangfeng, Z., Yonghua, L., Baoling, L., Biyao, L. & Nianhe, X. (1993). In *Aromatic Plants and Essential Constituents*. South China Institute of Botany, Chinese Academy of Sciences. Hai Feng Publishing Co., Hong Kong. (Chinese National Node for APINMAP.)
- Wierda, D. A., Feng, T. L. & Barron, A. R. (1989). *Acta Cryst. C* **45**, 338–339.
- Zachariasen, W. H. (1967). *Acta Cryst.* **23**, 558–564.

*Acta Cryst.* (1996). **C52**, 1257–1258

## 4-Hydroxyantipyrine

KALIYAMOORTHY PANNEERSELVAM, NARAYANAN JAYANTHI, ENRIQUE RUDIÑO-PIÑERA AND MANUEL SORIANO-GARCÍA\*

*Instituto de Química, † Circuito Exterior, Ciudad Universitaria, Delegación Coyoacán, México DF 04510, México.*  
E-mail: soriano@servidor.unam.mx

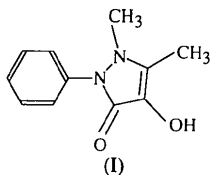
(Received 26 June 1995; accepted 13 November 1995)

### Abstract

The molecular structure of the title compound, 1,2-dihydro-4-hydroxy-1,5-dimethyl-2-phenyl-3*H*-pyrazol-3-one,  $C_{11}H_{12}N_2O_2$ , has been determined. Both rings are planar and make a dihedral angle of  $42.5(1)^\circ$  with each other. In the crystal structure, the molecule is stabilized as a centrosymmetric hydrogen-bonded dimer. There is no conjugation between the phenyl and pyrazole rings.

### Comment

The crystal and molecular structure of the title compound, (I), has been investigated in order to determine the conformation and crystal packing, and also to confirm its stereochemistry.



† Contribution No. 1356 of the Instituto de Química, UNAM.

A view of (I) with the numbering scheme is shown in Fig. 1. Bond lengths within the molecule correspond with the average C—C distance for a phenyl ring [1.378(3) Å] and the angles are normal. The molecule consists of a phenyl (C1–C6) and a pyrazole ring (C7–C9, N1, N2), which are planar within 0.007 and 0.026 Å, respectively, and form an interplanar angle of  $42.5(1)^\circ$ .

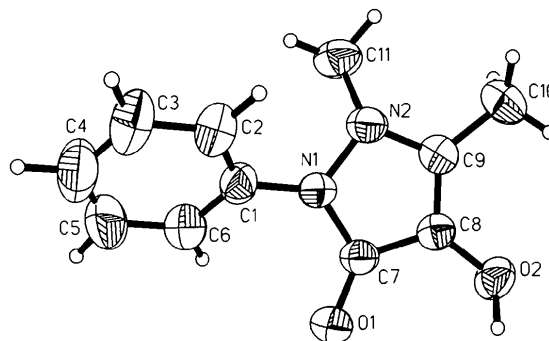


Fig. 1. The molecular structure of the title compound with the atom-labelling scheme and 50% probability displacement ellipsoids.

Fig. 2 shows a projection along the *c* axis. A hydrogen bond between the O2 hydroxy group and the carbonyl O1 atom of a neighbouring molecule constitutes the major intermolecular interaction and packing force. The two molecules are linked by a pair of O2—HO2···O1 hydrogen bonds across a crystallographic centre of inversion located at  $(-x, -y, 1-z)$ . The

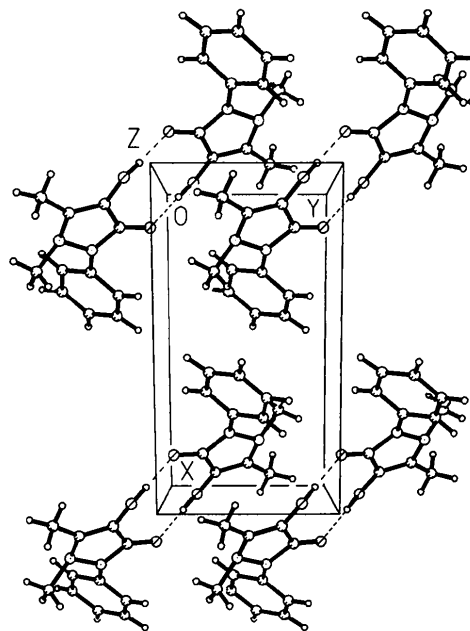


Fig. 2. A unit-cell drawing of the packing arrangement, with dashed lines indicating O—H···O intermolecular hydrogen-bonding interactions.

O2...O1 and HO2...O1 distances are 2.601 (2) and 1.66 (3) Å, respectively, and the O2—HO2...O1 angle is 161 (3)° (Allen, Kennard & Taylor, 1983). There are three intermolecular C—H...O non-bonded interactions in the range 3.40–3.58 Å (Desiraju, 1991).

## Experimental

The title compound was purchased from the Aldrich Chemical Co. Ltd and recrystallized from acetone at room temperature (m.p. 457–459 K).

### Crystal data

C<sub>11</sub>H<sub>12</sub>N<sub>2</sub>O<sub>2</sub>  
*M<sub>r</sub>* = 204.23  
 Monoclinic  
*P*2<sub>1</sub>/*n*  
*a* = 12.461 (6) Å  
*b* = 6.454 (4) Å  
*c* = 13.621 (5) Å  
 $\beta$  = 107.57 (1)°  
*V* = 1044.3 (9) Å<sup>3</sup>  
*Z* = 4  
*D<sub>x</sub>* = 1.299 Mg m<sup>-3</sup>  
*D<sub>m</sub>* = 1.300 Mg m<sup>-3</sup>  
*D<sub>m</sub>* measured by flotation in benzene/bromofom

Cu *K*α radiation  
 $\lambda$  = 1.54178 Å  
 Cell parameters from 25 reflections  
 $\theta$  = 10–25°  
 $\mu$  = 0.747 mm<sup>-1</sup>  
*T* = 293 (2) K  
 Needle  
 0.35 × 0.20 × 0.20 mm  
 Yellow

### Data collection

Siemens *P4* diffractometer  
 $\theta/2\theta$  scans  
 Absorption correction: none  
 1930 measured reflections  
 1383 independent reflections  
 1253 observed reflections  
 $[I > 2\sigma(I)]$   
*R<sub>int</sub>* = 0.0452

$\theta_{\max}$  = 56.74°  
 $h$  = -13 → 12  
 $k$  = 0 → 7  
 $l$  = 0 → 14  
 3 standard reflections monitored every 100 reflections  
 intensity decay: 3%

### Refinement

Refinement on *F*<sup>2</sup>  
*R*(*F*) = 0.0470  
 $wR(F^2)$  = 0.1226  
*S* = 1.073  
 1383 reflections  
 184 parameters  
 All H atoms refined isotropically  
 $w = 1/[\sigma^2(F_o^2) + (0.0786P)^2 + 0.2073P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max}$  = -0.002  
 $\Delta\rho_{\max}$  = 0.155 e Å<sup>-3</sup>  
 $\Delta\rho_{\min}$  = -0.311 e Å<sup>-3</sup>  
 Extinction correction: none  
 Atomic scattering factors from *International Tables for Crystallography* (1992, Vol. C, Tables 4.2.6.8 and 6.1.1.4)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å<sup>2</sup>)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U<sub>eq</sub></i>
C1	-0.2760 (2)	0.4135 (3)	0.5394 (1)	0.0509 (5)
C2	-0.2668 (2)	0.5901 (4)	0.5983 (2)	0.0639 (6)
C3	-0.3310 (2)	0.6092 (5)	0.6649 (2)	0.0824 (8)

C4	-0.4017 (2)	0.4526 (6)	0.6731 (2)	0.0926 (9)
C5	-0.4095 (2)	0.2764 (6)	0.6155 (3)	0.0927 (9)
C6	-0.3477 (2)	0.2553 (4)	0.5461 (2)	0.0709 (7)
C7	-0.1438 (2)	0.2283 (3)	0.4631 (1)	0.0456 (5)
C8	-0.0789 (1)	0.3010 (3)	0.3998 (1)	0.0439 (5)
C9	-0.1074 (2)	0.4987 (3)	0.3728 (1)	0.0454 (5)
C10	-0.0670 (2)	0.6359 (4)	0.3040 (2)	0.0648 (6)
C11	-0.2824 (2)	0.6875 (4)	0.3605 (2)	0.0644 (6)
N1	-0.2121 (1)	0.3914 (2)	0.4692 (1)	0.0487 (5)
N2	-0.1853 (1)	0.5665 (2)	0.4199 (1)	0.0478 (5)
O1	-0.1418 (1)	0.0589 (2)	0.5077 (1)	0.0632 (5)
O2	-0.0038 (1)	0.1854 (2)	0.3688 (1)	0.0582 (5)

Table 2. Selected geometric parameters (Å, °)

C1—C6	1.377 (3)	C7—N1	1.372 (2)
C1—C2	1.379 (3)	C7—C8	1.428 (3)
C1—N1	1.424 (3)	C8—C9	1.346 (3)
C2—C3	1.384 (3)	C9—N2	1.387 (2)
C3—C4	1.367 (4)	C9—C10	1.483 (3)
C4—C5	1.368 (5)	C11—N2	1.462 (3)
C5—C6	1.395 (4)	N1—N2	1.406 (2)
C7—O1	1.247 (2)		
C6—C1—C2	121.1 (2)	C9—C8—C7	109.0 (2)
C6—C1—N1	118.4 (2)	O2—C8—C7	125.1 (2)
C2—C1—N1	120.6 (2)	C8—C9—N2	109.7 (2)
C1—C2—C3	119.5 (2)	C8—C9—C10	128.8 (2)
C4—C3—C2	120.1 (3)	N2—C9—C10	121.5 (2)
C3—C4—C5	120.3 (2)	C7—N1—N2	110.3 (2)
C4—C5—C6	120.8 (3)	C7—N1—C1	126.9 (2)
C1—C6—C5	118.3 (3)	N2—N1—C1	120.1 (1)
O1—C7—N1	124.3 (2)	C9—N2—N1	105.6 (1)
O1—C7—C8	130.6 (2)	C9—N2—C11	119.4 (2)
N1—C7—C8	105.1 (2)	N1—N2—C11	114.7 (2)
C9—C8—O2	125.8 (2)		

Data collection: Siemens *P4* diffractometer software. Cell refinement: *XSCANS* (Siemens, 1992). Data reduction: *XSCANS*. Program(s) used to solve structure: *SHELXS86* (Sheldrick, 1990a). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *SHELXTL-Plus* (Sheldrick, 1990b). Software used to prepare material for publication: *SHELXL93*.

KP acknowledges the CONACYT (Cátedra Patrimonial Nivel II) for fellowship assistance. The authors thank the Instituto de Biotecnología, UNAM, for data collection.

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: KA1148). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

## References

- Allen, F. H., Kennard, O. & Taylor, R. (1983). *Acc. Chem. Res.* **16**, 146–153.  
 Desiraju, G. R. (1991). *Acc. Chem. Res.* **24**, 290–296.  
 Sheldrick, G. M. (1990a). *Acta Cryst.* **A46**, 467–473.  
 Sheldrick, G. M. (1990b). *SHELXTL-Plus. Structure Determination Software Programs*. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.  
 Sheldrick, G. M. (1993). *SHELXL93. Program for the Refinement of Crystal Structures*. University of Göttingen, Germany.  
 Siemens (1992). *XSCANS. X-ray Single Crystal Analysis System*. Version 2.0. Siemens Analytical X-ray Instruments Inc., Madison, Wisconsin, USA.