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3-(1-Hydroxyethylidene)-1-phenylpyrrolidine-2,4-dione

Dianne D. Ellis* and Anthony L. Spek

Bijvoet Center for Biomolecular Research, Department of Crystal and Structural Chemistry, Utrecht University, Padualaan 8, 3584 CH Utrecht, The Netherlands Correspondence e-mail: d.d.ellis@chem.uu.nl

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Analysis of $C_{12}H_{11}NO_3$ revealed a coplanar *N*-substituted phenyl group on a pyrrolidine ring with two keto moieties and a hydroxyethylidene functionality. The hydroxy group forms part of a hydrogen-bonding network characterized by a short intramolecular $H \cdots O$ distance of 1.81 (3) Å, and a longer intermolecular interaction with an $H \cdots O$ distance of 2.38 (3) Å. Both keto groups form additional intra- and intermolecular $C-H \cdots O$ contacts with $H \cdots O$ distances ranging from 2.26 to 2.41 Å.

Comment

The pyrrolidine skeleton occurs in many families of biologically important compounds. The resulting functionality, due to ease of substitution and therefore modification at several positions (Baldwin *et al.*, 1994a,b), has been utilized to synthesize compounds with varying properties. For example, several unusual amino acids which contain the pyrrolidine motif were investigated (Galeazzi *et al.*, 1999) and optically active pyrrolidines have been used as intermediates in controlled asymmetric synthesis (Suzuki *et al.*, 1994).



The analysis of crystals grown from a solution of *N*-phenyl-3-acetylpyrrolidine-2,4-dione showed that we had obtained crystals of the related tautomeric form 3-(1-hydroxyethylidene)-1-phenylpyrrolidine-2,4-dione, (I). The title compound (Fig. 1) shows a coplanar pyrrolidine ring with an *N*-substituted phenyl group [dihedral angle between the rings of 0.6 (1)°], substitution at two positions with keto groups and at a third with a hydroxyethylidene moiety. The pyrrolidine atoms do not deviate significantly from their least-squares plane, nor do any of the other non-H atoms [mean deviation, using unit weights, of all non-H atom is 0.02 (3) Å (*PLATON*; Spek, 2000)]. The bond distances around the N1 atom have a wide range (Table 1) and depend on the nature of the carbon. The shortest distance involves an $N-Csp^2$ bond, followed by an $N-C_{ar}$ of intermediate length with the longest belonging to an $N-Csp^3$ interaction. This results in an asymmetric planar geometry around the nitrogen. The bond geometries correlate well with a variety of *N*-phenyl-substituted pyrrolidin-2-one systems (for example, see Billing *et al.*, 1991).





PLATON/ORTEP representation (Spek, 2000) of (I) with displacement ellipsoids at the 50% probability level.

The nature of the hydroxyethylidene side group is confirmed by the C2=C3 distance of 1.368 (2) Å, the C2-O1 distance of 1.3258 (19) Å, and refinement of H1. The C-C bonds connecting the carbonyl groups show partial doublebond characteristics but are within the range of related bond types observed which have adjoining amino and carbon environments (Allen *et al.*, 1987). The two keto groups also have differing surroundings; thus, O2=C6 is involved in a cyclic amino linkage and is slightly longer than O3=C4 which has only adjacent C atoms. The geometric parameters concerning both keto functionalities agree with the tabulated compilation.



Figure 2

PLATON/PLUTON plot (Spek, 2000) showing part of the hydrogenbond network and the stacking of molecules.

The hydroxyl group forms an intramolecular contact with O2 and an intermolecular contact to the second keto group O3; H1···O distances of 1.81 (3) Å to O2 and 2.38 (3) Å to $O3^{i}$ [symmetry code: (i) 1 + x, y, z]. The O1-H1 distance is slightly elongated at 0.99 (2) Å, presumably due to the threecentre (bifurcated) hydrogen-bonding interaction. The geometries are similar to those observed in 2-(1-hydroxyethylidene)indan-1-one (Garcia et al., 1993) and α -cyclopiazonic acid (van Rooyen, 1992). The molecules form chains along the [100] direction in the crystal and pack in a head-totail fashion, due to partial π - π stacking of the hydroxyethylidene group over the phenyl ring. The hydrogen-bonding network is supplemented by weak $C-H \cdot \cdot \cdot O$ intermolecular contacts between neighbouring molecules (Fig. 2) and extended by $C-H \cdots O$ intramolecular contacts in the plane of the molecule (Table 2).

Experimental

N-Phenyl-3-acetylpyrrolidine-2,4-dione (available from Interbioscreen Ltd) was prepared in a solution of chloroform and ethanol. Colourless crystals of the 'enol' form 3-(1-hydroxyethylidene)-1phenylpyrrolidine-2,4-dione were obtained by recrystallization from a mixture of chloroform, dichloromethane, methanol and petroleum ether.

Crystal data

 $\begin{array}{l} C_{12}H_{11}NO_{3} \\ M_{r} = 217.22 \\ \text{Triclinic, } P\overline{1} \\ a = 6.5984 \ (5) \text{ Å} \\ b = 7.1839 \ (4) \text{ Å} \\ c = 11.2817 \ (6) \text{ Å} \\ \alpha = 103.084 \ (4)^{\circ} \\ \beta = 98.380 \ (5)^{\circ} \\ \gamma = 102.485 \ (5)^{\circ} \\ V = 497.81 \ (5) \text{ Å}^{3} \end{array}$

Data collection

Nonius CAD-4 diffractometer ω scans 4924 measured reflections 2279 independent reflections 1652 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.068$ $\theta_{\text{max}} = 27.47^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.116$ S = 1.0222279 reflections 150 parameters H atoms treated by a mixture of independent and constrained refinement Z = 2 $D_x = 1.449 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 25 reflections $\theta = 8.65 - 15.35^{\circ}$ $\mu = 0.105 \text{ mm}^{-1}$ T = 150 (2) KPlate, colourless $0.50 \times 0.50 \times 0.08 \text{ mm}$

 $h = -8 \rightarrow 8$ $k = -9 \rightarrow 9$ $l = -14 \rightarrow 14$ 3 standard reflections frequency: 60 min intensity decay: <1%

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0494P)^{2} + 0.0415P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.27 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{\text{min}} = -0.22 \text{ e} \text{ Å}^{-3}$

Aromatic H atoms were constrained and allowed to ride on their C atoms with $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$. The methyl H atoms were constrained to an ideal geometry with $U_{\rm iso}({\rm H}) = 1.5U_{\rm eq}({\rm C})$, and allowed to rotate freely about their C–C bonds. The hydroxyl H1 atom was located in the Fourier map and the displacement parameter was freely refined.

Table 1

Selected geometric parameters (Å, °).

N1-C5	1.4732 (18)	O3-C4	1.2205 (19)
N1-C6	1.375 (2)	C2-C3	1.368 (2)
N1-C7	1.412 (2)	C3-C4	1.444 (2)
O1-C2 O2-C6	1.3258 (19) 1.2398 (18)	C3-C6	1.456 (2)
C6-N1-C7 C6-N1-C5	128.11 (13) 111.22 (13)	C7-N1-C5	120.66 (12)

Table 2		
Hydrogen-bonding geometry	(Å,	°).

$D-\mathrm{H}\cdots A$	D-H	H···A	$D \cdots A$	$D-\mathrm{H}\cdots A$
$O1 - H1 \cdots O2$	0.99 (3)	1.81 (3)	2.6948 (18)	147 (2)
$O1 - H1 \cdots O3^{i}$	0.99 (3)	2.38 (3)	2.8155 (18)	106 (2)
$C5-H5B\cdots O3^{ii}$	0.99	2.41	3.328 (2)	154
$C1 - H1B \cdots O3$	0.98	2.38	3.043 (2)	124
C12−H12···O2	0.95	2.26	2.898 (2)	124

Symmetry codes: (i) 1 + x, y, z; (ii) -x, -y, -z.

Data collection: locally modified *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *SET4* (de Boer & Duisenberg, 1984); data reduction: *HELENA* (Spek, 1997); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2000); software used to prepare material for publication: *PLATON*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1499). Services for accessing these data are described at the back of the journal.

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