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Key indicators

Single-crystal X-ray study T = 293 KMean σ (C–C) = 0.002 Å R factor = 0.037 wR factor = 0.112 Data-to-parameter ratio = 13.4

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

4-p-Tolyl-1,3,4,4a,5,6,7,8a-octahydro-2H-pyrano[2,3-d]pyrimidin-2-one

The title compound, $C_{14}H_{18}N_2O_2$, which was obtained from an extended Biginelli reaction, contains a pyran ring with a chair conformation, a pyrimidine ring with a half-chair conformation and a benzene ring. The crystal structure is stabilized by an intermolecular hydrogen-bond network involving both NH groups and the carbonyl group.

Comment

The Biginelli reaction is a classic multicomponent reaction (Biginelli, 1893). It is increasingly attracting the interest of chemists because it shows great advantages of speed, diversity and efficiency in the drug discovery process. Its products, *viz*. functionalized dihydropyrimidines, originally obtained by cyclocondensation of ethyl acetoacetate, benzaldehyde and urea, represent a heterocylic system of remarkable pharmacological efficiency (Kappe, 2000). The aim of the present work was to study classical and extended Biginelli reactions. Recently, we synthesized a tetrahydropyrimidine scaffold instead of the dihydropyrimidine through an extended Biginelli reaction. We report here the cystal and molecular structure of the title compound, (I).



Fig. 1 shows the molecular structure of (I). It contains two fused six-membered rings and a benzene ring. The pyran ring, one of the two fused rings, adopts a chair conformation, while the pyrimidine ring adopts a half-chair conformation due to the two Nsp³ atoms and the C=O at C12. The O8-C9-C10-C5 torsion angle of -54.76 (13)° describes a *cis* configuration of the two fused rings. Also, the C11-C4-C10-C5 torsion angle of -73.58 (14)° suggests a *trans*-C4,C10 configuration. Selected bond distances and angles are listed in Table 1.

The H atoms at both N1 and N3 participate in intermolecular hydrogen bonding, which stabilizes the molecular Received 28 February 2005 Accepted 14 March 2005 Online 25 March 2005

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Figure 1

A view of the title molecule, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented as small spheres of arbitrary radii.



Figure 2

Four molecules of the title compound. For clarity, H atoms bonded to C atoms have been omitted. Dashed lines indicate hydrogen bonds.

packing in the crystal structure of (I) (Fig. 2). In the hydrogenbond system, the N atoms of the amine group act as hydrogenbond donors and the O atom of the carbonyl group acts as a hydrogen-bond acceptor (Table 2). A view of the molecular packing down the b axis is shown in Fig. 3.



Figure 3

The molecular packing of (I), viewed along the *b* axis. H atoms bonded to C atoms have been omitted for clarity. Dashed lines indicate the hydrogen-bonding interactions.

Experimental

Details of the synthesis and purification of compound (I) will be described elsewhere (Zhu *et al.*, 2005). Crystals appropriate for data collection were obtained by slow evaporation of an N,N-dimethyl-formide solution at room temperature over a period of a week.

Crystal data

С Л Т

$C_{14}H_{18}N_2O_2$	Z = 2
$A_r = 246.30$	$D_x = 1.322 \text{ Mg m}^{-3}$
Friclinic, $P\overline{1}$	Mo $K\alpha$ radiation
= 7.7151 (10) Å	Cell parameters from 1671
= 8.5541 (11) Å	reflections
= 9.6887 (13) Å	$\theta = 2.7 - 28.4^{\circ}$
$x = 88.516 \ (2)^{\circ}$	$\mu = 0.09 \text{ mm}^{-1}$
$B = 76.964 \ (2)^{\circ}$	T = 293 (2) K
$r = 83.401 \ (2)^{\circ}$	Prism, colourless
$V = 618.81 (14) \text{ Å}^3$	$0.34 \times 0.34 \times 0.30 \text{ mm}$

Data collection

Bruker SMART CCD area-detector1860diffractometer R_{int} φ and ω scans θ_{max} Absorption correction: noneh =3223 measured reflectionsk =204 independent reflectionsl =

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.037$ $wR(F^2) = 0.112$ S = 1.112204 reflections 164 parameters H-atom parameters constrained $\mu = 0.09 \text{ mm}$ T = 293 (2) KPrism, colourless $0.34 \times 0.34 \times 0.30 \text{ mm}$ $1865 \text{ reflections with } I > 2\sigma(I)$ $R_{\text{int}} = 0.013$

 $\theta_{\text{max}} = 25.3^{\circ}$ $h = -8 \rightarrow 9$ $k = -6 \rightarrow 10$ $l = -11 \rightarrow 11$

$$\begin{split} &w = 1/[\sigma^2(F_o^2) + (0.0609P)^2 \\ &+ 0.0889P] \\ &\text{where } P = (F_o^2 + 2F_c^2)/3 \\ (\Delta/\sigma)_{\text{max}} < 0.001 \\ \Delta\rho_{\text{max}} = 0.20 \text{ e } \text{ Å}^{-3} \\ \Delta\rho_{\text{min}} = -0.14 \text{ e } \text{ Å}^{-3} \\ &\text{Extinction correction: SHELXL97} \\ &\text{Extinction coefficient: } 0.032 (7) \end{split}$$

 Table 1

 Selected geometric parameters (Å, °).

N1-C2	1.3573 (18)	N3-C4	1.4617 (16)
N1-C9	1.4251 (17)	C4-C10	1.5339 (18)
N3-C2	1.3453 (17)	C5-C10	1.5337 (18)
C2 N1 C9	122 01 (11)	C5 C10 C4	111 88 (11)
$C_2 = N_1 = C_3$ $C_2 = N_3 = C_4$	122.91(11) 127.05(11)	$C_{12} = C_{11} = C_{16}$	117 32 (13)
N3-C2-N1	117.01 (11)	C12-C11-C10 C13-C14-C15	116.75 (14)
C6-C5-C10	112.17 (11)		
08-C9-C10-C5	-54.76 (13)	C11-C4-C10-C9	163.51 (11)
O8-C9-C10-C4	69.22 (13)	C11-C4-C10-C5	-73.58 (14)

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

$D - H \cdot \cdot \cdot A$	$D-\mathrm{H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1-H1\cdots O^i$	0.86	2.26	3.1177 (16)	176
$N3{-}H3{\cdots}O^{ii}$	0.86	2.33	3.1096 (16)	151

Symmetry codes: (i) 2 - x, -y, -z; (ii) 1 - x, -y, -z.

N-bound H atoms were located in a difference Fourier map and their parameters refined, with N–H distances restrained to 0.86(1) Å. The positions of the C-bound H atoms were calculated

geometrically and refined using a riding model [C-H = 0.93–0.98 Å], with U_{iso} (H) = 1.2 U_{eq} (C).

Data collection: *SMART* (Bruker, 1999); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997; program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1998); software used to prepare material for publication: *SHELXTL*.

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