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Key indicators

Single-crystal X-ray study
 $T = 173$ K
Mean $\sigma(\text{C}-\text{C}) = 0.002$ Å
 R factor = 0.034
 wR factor = 0.098
Data-to-parameter ratio = 12.2

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

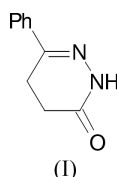
6-Phenyl-4,5-dihydropyridazin-3(2H)-one

The title compound, $\text{C}_{10}\text{H}_{10}\text{N}_2\text{O}$, is of pharmacological interest. It contains a keto-tetrahydropyridazine ring carrying a phenyl ring *para* to the carbonyl group. In the crystal structure, hydrogen-bonded centrosymmetric dimers are observed.

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Comment

Pyridazine compounds homologous to the title compound, (I), are known and are of particular interest as antidepressive agents (Biziere *et al.*, 1988). This kind of compound has also been used for the treatment of Alzheimer's disease (Passeri *et al.*, 1985).



The geometric parameters of (I) are normal. The dihydropyridazine ring adopts a half-chair conformation, with atoms C1, N2, N3 and C4 in a common plane and C5 0.222 (2) Å and C6 0.262 (2) Å on opposite sides of this plane. The plane is almost coplanar with the phenyl ring; the dihedral angle between the two planes is 1.73 (9) Å. In the crystal structure, hydrogen-bonded centrosymmetric dimers are observed.

Experimental

142.4 g (8 mmol) of benzoylpropionic acid was dissolved in 170 ml ethanol. 43 g (8.6 mmol) of hydrazine hydrate was then added to this mixture and it was refluxed for 1 h. After cooling to room temperature, yellow crystals grew from the solution and were filtered off and dried.

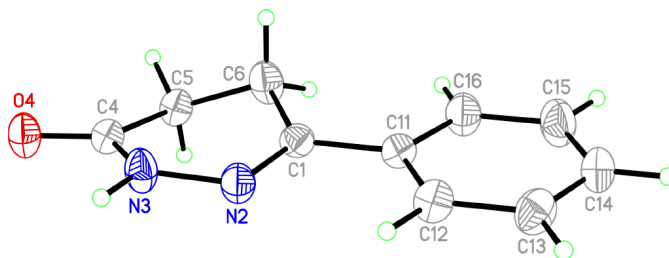


Figure 1
Perspective view of the asymmetric unit of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

Crystal data

C₁₀H₁₀N₂O
M_r = 174.20
 Triclinic, *P* $\bar{1}$
a = 5.6712 (10) Å
b = 8.1749 (14) Å
c = 10.1055 (17) Å
 α = 104.740 (14)°
 β = 104.392 (13)°
 γ = 98.097 (14)°
V = 428.33 (13) Å³

Z = 2
D_x = 1.351 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 10040 reflections
 θ = 3.7–25.1°
 μ = 0.09 mm⁻¹
T = 173 (2) K
 Block, colourless
 0.44 × 0.38 × 0.36 mm

Data collection

Stoe IPDS II two-circle diffractometer
 ω scans
 Absorption correction: none
 5318 measured reflections
 1483 independent reflections

1318 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.025
 θ_{\max} = 25.0°
h = -6 → 6
k = -9 → 9
l = -11 → 11

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.034
wR(*F*²) = 0.098
S = 1.08
 1483 reflections
 122 parameters
 H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0639P)^2 + 0.0435P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.18 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.16 \text{ e \AA}^{-3}$

Table 1

Selected geometric parameters (Å, °).

C1–N2	1.2830 (15)	N3–C4	1.3464 (15)
N2–N3	1.3877 (14)	C4–O4	1.2316 (14)
C1–N2–N3	117.98 (9)	C4–N3–N2	127.23 (9)
C1–N2–N3–C4	15.45 (18)		

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> –H··· <i>A</i>	<i>D</i> –H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> –H··· <i>A</i>
N3–H3···O4 ⁱ	0.870 (16)	1.990 (16)	2.8587 (13)	177.1 (14)

Symmetry code: (i) 1 – *x*, 1 – *y*, 2 – *z*.

H atoms bonded to C atoms were refined with fixed individual displacement parameters [*U*_{iso}(H) = 1.2*U*_{eq}(C)], using a riding model with aromatic C–H = 0.99 Å and methylene C–H = 0.98 Å. The H atom bonded to N was refined freely.

Data collection: *X*-AREA (Stoe & Cie, 2001); cell refinement: *X*-AREA; data reduction: *X*-AREA; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *XP* in *SHELXTL-Plus* (Sheldrick, 1991); software used to prepare material for publication: *SHELXL97*.

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