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**Key indicators**

Single-crystal X-ray study  
 $T = 298$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å  
 $R$  factor = 0.041  
 $wR$  factor = 0.140  
Data-to-parameter ratio = 13.7

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

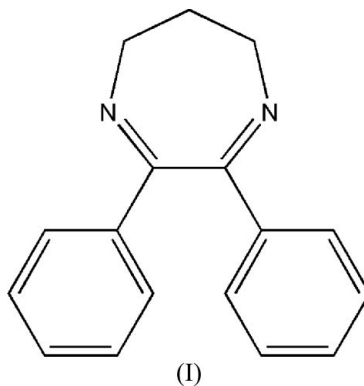
## 2,3-Diphenyl-6,7-dihydro-5H-1,4-diazepine

In the title compound,  $\text{C}_{17}\text{H}_{16}\text{N}_2$ , the 1,4-diazepine ring adopts a distorted boat conformation, in which the sum of the internal angles is  $809^\circ$ . The molecules are linked by a pair of  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds into a  $\text{C}_2^2(14)$  chain along the  $[10\bar{1}]$  direction.

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**Comment**

1,4-Diazepine derivatives are known to have a broad range of biological activities, such as selective inhibition of Rho-kinase (Tamura *et al.*, 2005), and as HDM2 antagonists used in the treatment of certain cancers (Raboisson *et al.*, 2005). The crystal structures of some 1,4-diazepine derivatives have been reported (Nesterov, 2002; Clark *et al.*, 1999). We here report the crystal structure of a 1,4-diazepine derivative, 6,7-hydro-2,3-diphenyl-1,4-diazepine, (I).

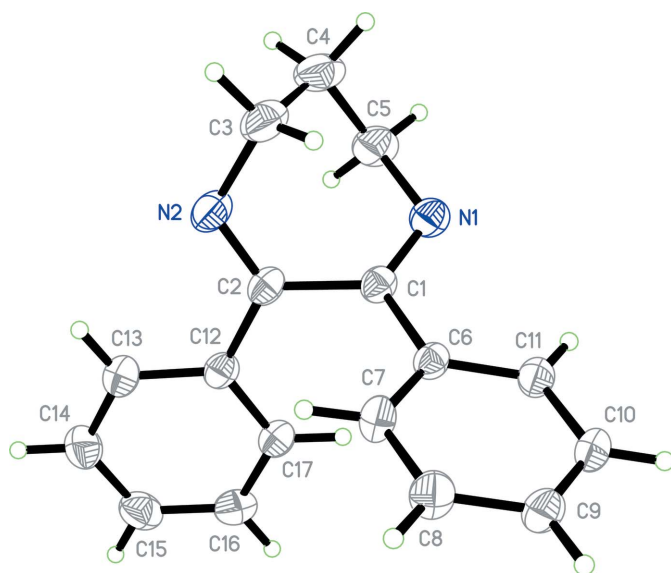


In the molecule of compound (I) (Fig. 1), the 1,4-diazepine ring adopts a distorted boat conformation, in which the sum of the internal angles is  $809^\circ$  (Table 1). The two phenyl rings are located on opposite sides of the seven-membered ring and they enclose a dihedral angle of  $82.58(6)^\circ$ .

In the crystal structure of (I), the molecules are linked by a pair of  $\text{C}-\text{H}\cdots\text{N}$  hydrogen bonds into a  $\text{C}_2^2(14)$  chain (Bernstein *et al.*, 1995) along the  $[10\bar{1}]$  direction (Fig. 2 and Table 2). Atom C16 in the molecule at  $(x, y, z)$  act as hydrogen-bond donor to atom N1 in the molecule at  $(\frac{1}{2} + x, \frac{1}{2} - y, -\frac{1}{2} + z)$ . There are no significant intermolecular interaction between the chains.

**Experimental**

The title compound, (I), was synthesized by the condensation reaction between propane-1,3-diamine and benzoin in ethanol. Colourless crystals of (I) suitable for X-ray structure analysis were obtained by recrystallizing the crude product from ethanol (m.p. 392–393 K).



**Figure 1**  
The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 30% probability level.

#### Crystal data

$C_{17}H_{16}N_2$   
 $M_r = 248.32$   
 Monoclinic,  $P2_1/n$   
 $a = 8.708$  (2) Å  
 $b = 17.649$  (3) Å  
 $c = 9.171$  (2) Å  
 $\beta = 105.115$  (2)°  
 $V = 1360.7$  (5) Å<sup>3</sup>

$Z = 4$   
 $D_x = 1.212$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 $\mu = 0.07$  mm<sup>-1</sup>  
 $T = 298$  (2) K  
 Block, colourless  
 $0.63 \times 0.58 \times 0.52$  mm

#### Data collection

Siemens SMART 1000 CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
 $T_{\min} = 0.956$ ,  $T_{\max} = 0.964$

6849 measured reflections  
 2375 independent reflections  
 1671 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.054$   
 $\theta_{\text{max}} = 25.0^\circ$

#### Refinement

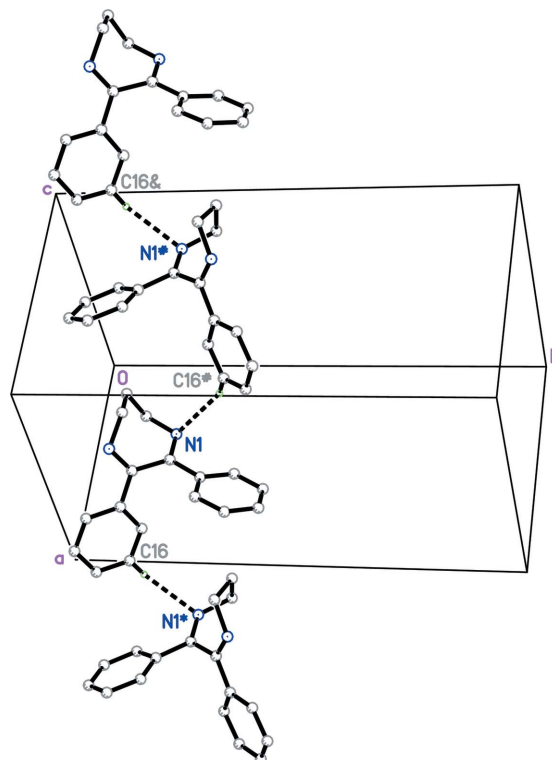
Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.140$   
 $S = 1.01$   
 2375 reflections  
 173 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0742P)^2 + 0.1494P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\text{max}} < 0.001$   
 $\Delta\rho_{\text{max}} = 0.11$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.15$  e Å<sup>-3</sup>  
 Extinction correction: SHELXL97  
 Extinction coefficient: 0.44 (3)

**Table 1**

Selected geometric parameters (Å, °).

N1—C1	1.276 (2)	N2—C2	1.274 (2)
C1—N1—C5	116.47 (16)	C3—C4—C5	112.24 (16)
C2—N2—C3	116.18 (16)	N1—C5—C4	111.32 (16)
N2—C3—C4	110.48 (17)		
N2—C3—C4—C5	-47.0 (2)	C3—C4—C5—N1	-43.1 (2)



**Figure 2**

Part of the crystal structure of (I), showing the formation of a  $C_2^2(14)$  chain along the  $[10\bar{1}]$  direction. For clarity, H atoms not involved in the motifs shown have been omitted. [Symmetry codes: (\*)  $\frac{1}{2} + x, \frac{1}{2} - y, -\frac{1}{2} + z$ ; (#)  $-\frac{1}{2} + x, \frac{1}{2} - y, \frac{1}{2} + z$ ; (&#)  $-1 + x, y, 1 + z$ ]. Dashed lines indicate hydrogen bonds.

**Table 2**

Hydrogen-bond geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
C16—H16 <sup>#</sup> ···N1 <sup>i</sup>	0.93	2.62	3.454 (2)	149

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

All H atoms were positioned geometrically and refined as riding on their parent atoms, with C—H = 0.93 Å for aromatic H atoms, C—H = 0.97 Å for methylene H atoms and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  for all H atoms.

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

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