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## Structure Reports

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## Key indicators

Single-crystal X-ray study
$T=120 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$
$R$ factor $=0.055$
$w R$ factor $=0.146$
Data-to-parameter ratio $=9.1$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 10-Methoxy-5H-dibenz[b,f]azepine

The structure of the title compound, $\mathrm{C}_{15} \mathrm{H}_{13} \mathrm{NO}$, has six independent molecules in the asymmetric unit; in each case, the seven-membered ring adopts a boat conformation and the overall molecular shape is that of a butterfly. All molecules display $\mathrm{N}-\mathrm{H} \cdots \mathrm{C}=\mathrm{C}$ close contacts, instead of $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ interactions. The intramolecular dihedral angles between the benzene rings are within the range 43.7 (1)-46.4 (1) ${ }^{\circ}$ for the six molecules.

## Comment

The title compound, (I), is used as an intermediate for the synthesis of the registered anticonvulsant drug oxcarbazepine (Kricka \& Ledwith, 1974), the structure of which has recently been reported (Hempel et al., 2005). As part of a series of studies into the structural aspects of iminostilbene analogues, the structure of (I) was determined and is reported here.

(I)

A search of the Cambridge Structural Database (CSD, Version 5.26; Allen, 2002) reveals 29 compounds that contain a dibenz $[b, f]$ azepine moiety, with all bar four being either structures or adducts of 5 H -dibenz $[b, f]$ azepine- 5 -carboxamide (carbamazepine). The structure of (I) has six unique molecules in the asymmetric unit (Fig. 1); in each case, the sevenmembered ring adopts a boat conformation (Cremer \& Pople, 1975) and the overall molecular shape is that of a butterfly. The intramolecular dihedral angles between the benzene rings are 43.7 (1), 45.1 (1), 46.4 (1), 44.7 (1), 44.7 (1) and 45.2 (1) ${ }^{\circ}$ for molecules $A$ to $F$, respectively. All molecules display $\mathrm{N}-$ $\mathrm{H} \cdots \mathrm{C}=\mathrm{C}$ close contacts, listed in Table 1, instead of $\mathrm{N}-$ $\mathrm{H} \cdots \mathrm{O}$ interactions. These close contacts all occur between equivalent molecules in the $b$-cell direction. The unit-cell packing of (I) is shown in Fig. 2.

## Experimental

The title compound was prepared by brominating N -acetyl-5 H dibenz[ $b, f]$ azepine ( $2.35 \mathrm{~g}, 10 \mathrm{mmol}$ ) using bromine ( $3.2 \mathrm{~g}, 20 \mathrm{mmol}$ )

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Figure 1
The molecular configuration and atom-numbering scheme for the six molecules in the asymmetric unit of (I). Displacement ellipsoids are drawn at the $50 \%$ probability level and H atoms are drawn as spheres of arbitrary radius. All molecules have been separately plotted in comparable orientations.
in dichloromethane ( 5 ml ) to obtain the dibromo derivative, which was further refluxed with $\mathrm{KOH}(1.12 \mathrm{~g}, 20 \mathrm{mmol})$ in $\mathrm{CH}_{3} \mathrm{OH}(5 \mathrm{ml})$ to yield the product. Crystals were grown from a dichloromethaneethanol ( $1: 1 \mathrm{v} / \mathrm{v}$ ) solution.

## Crystal data

## $\mathrm{C}_{15} \mathrm{H}_{13} \mathrm{NO}$

$M_{r}=223.26$
Monoclinic, C2
$a=54.925$ (11) $\AA$
$b=5.8189$ (12) $\AA$
$c=21.628$ (4) A
$\beta=98.14$ (3) ${ }^{\circ}$
$V=6843(2) \AA^{3}$
$Z=24$

## Data collection

Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 2003) $T_{\text {min }}=0.970, T_{\text {max }}=0.989$ 38719 measured reflections
8443 independent reflections
$D_{x}=1.300 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 7575
reflections
$\theta=2.9-27.5^{\circ}$
$\mu=0.08 \mathrm{~mm}^{-1}$
$T=120$ (2) K
Prism, yellow
$0.38 \times 0.34 \times 0.14 \mathrm{~mm}$
$R_{\text {int }}=0.062$
$\theta_{\text {max }}=27.5^{\circ}$
$h=-70 \rightarrow 69$
$k=-7 \rightarrow 7$
$l=-26 \rightarrow 28$

8443 reflections
926 parameters
H -atom parameters constrained

Table 1
Hydrogen-bonding geometry $\left(\AA{ }^{\circ},{ }^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| N5 $A-\mathrm{H} 5 A \cdots \mathrm{C} 11 A^{\mathrm{i}}$ | 0.88 | 2.54 | $3.407(5)$ | 167 |
| N5 $B-\mathrm{H} 5 B \cdots \mathrm{C} 11 B^{\mathrm{i}}$ | 0.88 | 2.52 | $3.388(5)$ | 167 |
| N5C-H5C C $11 C^{\mathrm{i}}$ | 0.88 | 2.54 | $3.403(5)$ | 167 |
| N5D-H5D $\cdots \mathrm{C} 11 D^{\mathrm{i}}$ | 0.88 | 2.53 | $3.388(5)$ | 166 |
| N5E-H5E $11 E^{\mathrm{i}}$ | 0.88 | 2.54 | $3.405(5)$ | 167 |
| N5 $F-\mathrm{H} 5 F \cdots \mathrm{C} 11 F^{\mathrm{i}}$ | 0.88 | 2.53 | $3.391(5)$ | 167 |

Symmetry code: (i) $x, 1+y, z$.
All H atoms were included in the refinement in calculated positions, in the riding-model approximation, with $\mathrm{C}-\mathrm{H}$ distances of 0.95


Figure 2
The unit cell contents of (I), viewed down the $b$ cell axis. All atoms are drawn as circles of arbitrary radii. For clarity, all H atoms except those of the NH groups have been omitted.

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.055$
$w R\left(F^{2}\right)=0.146$
$S=1.18$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0668 P)^{2}\right] \\
& \text { where } P=\left(F_{o}^{2}+2 F_{c}^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.37 \mathrm{e}^{-3} \\
& \Delta \rho_{\min }=-0.43 \mathrm{e} \AA^{-3} \\
& \text { Extinction correction: SHELXL97 } \\
& \text { Extinction coefficient: } 0.0014(2)
\end{aligned}
$$

$(\mathrm{ArH})$ and $0.98 \AA\left(\mathrm{CH}_{3}\right)$ and an $\mathrm{N}-\mathrm{H}$ distance of $0.88 \AA$. The isotropic displacement parameters for all H atoms were set equal to $1.25 U_{\text {eq }}$ of the carrier atom. In the absence of significant anomalous scattering, 6006 measured Friedel pairs were merged.

Data collection: COLLECT (Hooft, 1998); cell refinement: DENZO (Otwinowski \& Minor, 1997) and COLLECT; data reduction: DENZO and COLLECT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

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