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## 2-[(1,3-Benzodioxol-5-ylmethylidene)-amino]-4,5,6,7-tetrahydro-1-benzo-thiophene-3-carbonitrile

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Key indicators: single-crystal X-ray study; $T=296 \mathrm{~K}$; mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$; $R$ factor $=0.042 ; w R$ factor $=0.119 ;$ data-to-parameter ratio $=13.4$.

The title compound, $\mathrm{C}_{17} \mathrm{H}_{14} \mathrm{~N}_{2} \mathrm{O}_{2} \mathrm{~S}$, crystallizes with two roughly planar molecules in the asymmetric unit, in which the dihedral angles between the 1,3-benzodioxole-5-carbaldehyde moiety and the heterocyclic five-membered ring are 3.76 (5) and 5.33 (12) ${ }^{\circ}$. In each molecule, a short $\mathrm{C}-\mathrm{H} \cdots \mathrm{S}$ contact generates an $S(5)$ ring. In the crystal, pairs of molecules are linked by a weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{N}$ interaction, forming dimers.

## Related literature

For a related structure, see: Elerman \& Elmali, (1998). For graph-set notation, see: Bernstein et al. (1995).
$\gamma=92.854(1)^{\circ} \quad \mu=0.22 \mathrm{~mm}^{-1}$
$V=1506.77(7) \AA^{3} \quad T=296 \mathrm{~K}$
$Z=4$
$0.32 \times 0.23 \times 0.20 \mathrm{~mm}$
Mo $K \alpha$ radiation
Data collection
Bruker Kappa APEXII CCD
diffractometer
Absorption correction: multi-scan
(SADABS; Bruker, 2005)
$T_{\text {min }}=0.947, T_{\max }=0.962$

## Refinement

$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.042 \quad 397$ parameters
$w R\left(F^{2}\right)=0.119 \quad \mathrm{H}$-atom parameters constrained
$S=1.01$
5331 reflections
$\Delta \rho_{\text {max }}=0.47 \mathrm{e} \AA^{-3}$
$\Delta \rho_{\text {min }}=-0.19 \mathrm{e}^{-3}$

Table 1
Hydrogen-bond geometry ( $\AA \AA^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{C} 8-\mathrm{H} 8 \cdots \mathrm{~S} 1$ | 0.93 | 2.65 | $3.081(2)$ | 109 |
| $\mathrm{C} 25-\mathrm{H} 25 \cdots \mathrm{~S} 2$ | 0.93 | 2.61 | $3.060(2)$ | 110 |
| $\mathrm{C} 7-\mathrm{H} 7 A \cdots \mathrm{~N} 4$ | 0.97 | 2.62 | $3.190(3)$ | 118 |

Symmetry code: (i) $x, y-1, z$.

Data collection: APEX2 (Bruker, 2009); cell refinement: SAINT (Bruker, 2009); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997) and PLATON (Spek, 2009); software used to prepare material for publication: WinGX (Farrugia, 1999) and PLATON

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6325).

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