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## Key indicators

Single-crystal X-ray study
$T=293 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.005 \AA$
$R$ factor $=0.065$
$w R$ factor $=0.206$
Data-to-parameter ratio $=13.8$
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## 5-Ethoxycarbonyl-2,6-dimethyl-4-(3-nitrophenyl)-1,4-dihydropyridine-3-carboxylic anhydride ethyl acetate solvate

In the title compound, $\mathrm{C}_{34} \mathrm{H}_{34} \mathrm{~N}_{4} \mathrm{O}_{11} \cdot \mathrm{C}_{4} \mathrm{H}_{8} \mathrm{O}_{2}$, the dihydropyridine rings display envelope configuration. The solvent molecule links to the anhydride via weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding.

## Comment

4-Aryl-1,4-dihydropyridine-3,5-dicarboxylic diesters of the nefidipine type have become almost indispensable for the treatment of cardiovascular diseases since they first appeared on the market in 1975 (Yiu \& Edward, 1999; Goldmann \& Stoltefuss, 1991). The title compound, (I), is related to the preparation of the above compounds.

(I)

The molecular structure of (I) is shown in Fig. 1. Both dihydropyridine rings display an envelope conformation, with atoms C10 and C28 displaced from the mean planes formed by the other atoms in the same ring by 0.287 (1) and 0.217 (1) $\AA$, respectively. The dihedral angle between the C11-containing benzene ring and the $\mathrm{N} 1 / \mathrm{C} 4 / \mathrm{C} 5 / \mathrm{C} 7 / \mathrm{C} 9$ plane is $87.42(5)^{\circ}$, while the dihedral angle between the C29-containing benzene ring and the $\mathrm{N} 3 / \mathrm{C} 19 / \mathrm{C} 20 / \mathrm{C} 22 / \mathrm{C} 24$ plane is $85.04(5)^{\circ}$. This is comparable to the situation found in nefidipine (Hofmann \& Cimiraglia, 1990; Ramusino \& Varì, 1999). The C17-O6 and C18-O6 bond distances (Table 1) are much longer than the C17-O5 and $\mathrm{C} 18-\mathrm{O} 7$ bond distances, respectively, confirming their single bond character.

Both classic $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ and weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonds occur in the crystal structure (Table 2). The solvent molecule links to the anhydride via weak $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ hydrogen bonding.

## Experimental

5-Ethoxycarbonyl-2,6-dimethyl-4-(3-nitrophenyl)-1,4-dihydropyridine-3-carboxylic acid ( $0.45 \mathrm{~g}, 1 \mathrm{mmol}$ ) was dissolved in $\mathrm{CH}_{2} \mathrm{Cl}_{2}(30 \mathrm{ml})$,

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The molecular structure of (I), with $30 \%$ probability displacement ellipsoids. H atoms are shown as small spheres of arbitrary radii.
and dicyclohexylcarbodiimide $(0.21 \mathrm{~g}, 1 \mathrm{mmol})$ in $\mathrm{CH}_{2} \mathrm{Cl}_{2}(10 \mathrm{ml})$ was dropped into the above solution at 278 K . The reaction mixture was stirred at $276-279 \mathrm{~K}$ for 4 h . The solvent $\mathrm{CH}_{2} \mathrm{Cl}_{2}$ was removed by vacuum evaporation at 293 K . The product was purified by chromatography on a silica gel column (eluted by ethyl acetate and petroleum, $3: 7$ ) at room temperature. The product $(0.41 \mathrm{~g})$ was obtained in a yield of $94 \%$. Recrystallization from a dichloromethane/ethyl acetate solution gave single crystals of (I).

## Crystal data

$\mathrm{C}_{34} \mathrm{H}_{34} \mathrm{~N}_{4} \mathrm{O}_{11} \cdot \mathrm{C}_{4} \mathrm{H}_{8} \mathrm{O}_{2}$
$M_{r}=762.76$
Triclinic, $P \overline{1}$
$a=10.896$ (2) $\AA$
$b=12.017$ (2) $\AA$
$c=16.162$ (3) $\AA$
$\alpha=99.59$ (3) ${ }^{\circ}$
$\beta=106.29(3)^{\circ}$
$\gamma=105.69$ (3) ${ }^{\circ}$
$V=1887.1(9) \AA^{3}$

$$
\begin{aligned}
& Z=2 \\
& D_{x}=1.342 \mathrm{Mg} \mathrm{~m}^{-3}
\end{aligned}
$$

Mo $K \alpha$ radiation
Cell parameters from 15558 reflections
$\theta=3.2-25.5^{\circ}$
$\mu=0.10 \mathrm{~mm}^{-1}$
$T=293$ (2) K
Block, yellow
$0.28 \times 0.22 \times 0.10 \mathrm{~mm}$

## Data collection

Rigaku R-AXIS RAPID diffractometer $\omega$ scans
Absorption correction: none 18422 measured reflections 6953 independent reflections

4362 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.038$
$\theta_{\text {max }}=25.5^{\circ}$
$h=-13 \rightarrow 13$
$k=-14 \rightarrow 14$
$l=-19 \rightarrow 19$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.065$
$w R\left(F^{2}\right)=0.206$
$S=1.08$
6953 reflections
504 parameters

> H -atom parameters constrained
> $w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.1184 P)^{2}\right]$
> where $P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3$
> $(\Delta / \sigma)_{\max }=0.001$
> $\Delta \rho_{\max }=0.57 \mathrm{e} \AA^{-3}$
> $\Delta \rho_{\min }=-0.45 \mathrm{e}^{-3}$

Table 1
Selected bond lengths ( A ).

| O1-C2 | $1.453(4)$ | O10-C25 | $1.204(4)$ |
| :--- | :--- | :--- | :--- |
| O1-C3 | $1.345(4)$ | O11-C25 | $1.347(3)$ |
| O2-C3 | $1.212(3)$ | O11-C26 | $1.451(3)$ |
| O5-C17 | $1.199(3)$ | O12-C36 | $1.269(6)$ |
| O6-C17 | $1.391(3)$ | O13-C36 | $1.256(6)$ |
| O6-C18 | $1.399(3)$ | O13-C37 | $1.482(6)$ |
| O7-C18 | $1.203(3)$ |  |  |

Table 2
Hydrogen-bond geometry ( $\AA^{\circ},{ }^{\circ}$ ).

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 1-\mathrm{H} 1 D \cdots \mathrm{O} 4^{\mathrm{i}}$ | 0.86 | 2.39 | $3.121(4)$ | 143 |
| $\mathrm{~N} 3-\mathrm{H} 3 A \cdots \mathrm{O} 9^{\text {ii }}$ | 0.86 | 2.37 | $3.131(4)$ | 147 |
| $\mathrm{C} 6-\mathrm{H} 6 C \cdots \mathrm{O} 2^{\mathrm{i}}$ | 0.96 | 2.36 | $3.317(6)$ | 178 |
| $\mathrm{C} 23-\mathrm{H} 23 C \cdots \mathrm{O} 2^{\mathrm{iii}}$ | 0.96 | 2.44 | $3.348(5)$ | 157 |

Symmetry codes: (i) $x+1, y, z$; (ii) $x-1, y, z$; (iii) $x-1, y-1, z$.

Methyl H atoms were placed in calculated positions with $\mathrm{C}-\mathrm{H}=$ $0.96 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.5 U_{\text {eq }}(\mathrm{C})$; the torsion angles were refined to fit the electron density. Other H atoms were placed in calculated positions with $\mathrm{C}-\mathrm{H}=0.93-0.98 \AA$ and $\mathrm{N}-\mathrm{H}=0.86 \AA$, and constrained to ride on their parent atoms with $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}$ (carrier).

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC, 2002); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2002); software used to prepare material for publication: SHELXTL.

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