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# Racemic isopropyl 1,4-dihydro-2,6-dimethyl-4-(6-methyl-2-pyridyl)-3-nitropyridine-5-carboxylate hemibenzene solvate 

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This analysis establishes the rotameric orientation of the pyridyl-ring N atom of the title compound, $\mathrm{C}_{17} \mathrm{H}_{21} \mathrm{~N}_{3} \mathrm{O}_{4} \cdot 0.5 \mathrm{C}_{6} \mathrm{H}_{6}$, as antiperiplanar (ap) to the 1,4dihydropyridine $\mathrm{H}-4$, the absence of an intramolecular hydrogen bond between the 1,4-dihydropyridine NH and the pyridyl-N atom, and the unusual planarity of the 1,4dihydropyridine ring.

## Comment

The design of cardioselective Hantzsch 1,4-dihydropyridine (DHP) L-type voltage-sensitive calcium channel agonist positive inotropes has presented a significant challenge in drug design (Langs et al., 1990, 1991). Although there have been extensive efforts to understand the molecular basis of action (Peterson et al., 1996) and structure-activity relationships (Triggle, 1996), the development of a cardioselective calcium channel stimulant has not been reported (Rampe \& Kano, 1994). Recently, we discovered a novel third generation class of isomeric alkyl 1,4-dihydro-2,6-dimethyl-3-nitro-4-(pyridyl)-5-pyridinecarboxylates where the 4 -(2-pyridyl) isomer acted as a dual cardioselective calcium channel agonist/smooth muscle selective calcium channel antagonist (Iqbal et al., 1998). ${ }^{1} \mathrm{H}$ NMR nuclear Overhauser enhancement (NOE) studies, for a group of isopropyl or phenethyl 1,4-dihydro-3-nitro-4-(3- or 6-substituted-2-pyridyl)-5-pyridinecarboxylates, clearly indicated that a significant rotamer fraction is present in which the pyridyl-N atom is antiperiplanar (ap) to the $1,4-$ DHP H-4 in all cases irrespective of whether the substituent ( $\mathrm{H}, \mathrm{Me}, \mathrm{Ph}$ ) is located at the $\mathrm{C}-3$ or C-6 position of the 2pyridyl moiety. In addition, variable-temperature ${ }^{1} \mathrm{H}$ NMR

[^0]studies indicated the 1,4-DHP NH is hydrogen-bonded. The orientation of the 6-substituted-2-pyridyl group differs from that of an ortho-substituted-phenyl ring in Hantzsch 1,4-DHP calcium channel antagonists since the rotamer where the ortho-phenyl ring substituent is synperiplanar ( $s p$ ) to the 1,4 DHP H-4 is generally thermodynamically more preferential (Rovnyak et al., 1991).

We now describe the X-ray analysis of racemic isopropyl 1,4-dihydro-2,6-dimethyl-3-nitro-4-(6-methyl-2-pyridyl)-pyri-dine-5-carboxylate, (I) (Iqbal et al., 1998), to establish the rotameric orientation of the pyridyl ring unambiguously and to determine whether a potential intramolecular hydrogenbonding interaction involving the amine NH moiety exists. These novel third generation calcium channel modulators offer a new drug-design approach directed to the treatment of congestive heart failure, and may be useful as probes to study the structure-function relationships of calcium channels.

(I)

Each unit cell contains four molecules of the title compound and two benzene molecules of crystallization. The benzene is packed between the isopropyl groups of two neighbouring molecules of the title compound, and lies on a crystallographic centre of symmetry.

The torsion angles indicate that the $1,4-\mathrm{DHP}$ ring is relatively flat; however, angles of the greatest magnitude are found around the N1 and C4 atoms. The deviation from strict


Figure 1
ZORTEP (Zsolnai \& Huttner, 1994) diagram of the title compound showing the atom-labelling scheme. Two molecules are shown, with a symmetry-related molecule shown in grey. Hydrogen bonds between N7 and $\mathrm{N} 1-\mathrm{H}$ are shown as dotted lines. The benzene which co-crystallized is not included. Displacement ellipsoids are at the $50 \%$ probability level.
planarity takes the form of a slight boat conformation with the N 1 and C 4 atoms being displaced towards the pyridyl side of the ring. The sum of the absolute values of all the torsion values around the $1,4-\mathrm{DHP}$ ring is $28(5)^{\circ}$, a value much smaller than the reported range of 39 to $78^{\circ}$ for other 1,4-DHP ring structures (Langs \& Triggle, 1985; Fossheim, 1987). Increased planarity of this ring has been suggested to be associated with greater pharmacological activity (Triggle et al., 1980; Fossheim et al., 1982).

The nitro group attached to C3 is almost coplanar with the 1,4-DHP ring, with a torsion of $13.4(2)^{\circ}$ relative to the $\mathrm{C} 2=\mathrm{C} 3$ bond. This rotation relieves the potential steric clash between the pyridyl group and the proximal oxygen of the nitro group. Similarly the ester group attached to C5 is rotated approximately $18(2)^{\circ}$ out of the plane of the $1,4-\mathrm{DHP}$ ring. The shortness of the C3-N31 (nitro) bond [1.430 (2) A] relative to the canonical length ( $1.471 \AA$ ) found in nifedipine structures indicates a greater degree of electron delocalization, this was previously observed in another nitro-substituted 1,4-DHP ring (Langs \& Triggle, 1985). Similarly the N1-C2 bond is shorter than in nifedipine structures [1.363 (2) $\AA$ compared to $1.380 \AA$ ], but longer than that seen in methyl 2,6-dimethyl-5-nitro-4-(2-trifluoromethylphenyl)-1,4-dihydropyr-idine-3-carboxylate $(1.350 \AA)$. Despite the differences, both compounds appear to share more electron delocalization over the nitro groups which may explain the greater degree of planarity of the $1,4-\mathrm{DHP}$ ring.

The pyridine group is oriented approximately perpendicular to the plane of the $1,4-\mathrm{DHP}$ ring, with the N atom of the pyridyl group antiperiplanar to the 1,4 -DHP H4. The pyridylN atom is turned slightly toward C 5 (approximately $3.7^{\circ}$ relative to a line bisecting the $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 5$ angle). No intramolecular hydrogen bond is formed between the NH of the 1,4-DHP ring and the pyridyl-N atom since the $\mathrm{N}-\mathrm{H}$ bond vector is oriented in the same direction as the pyridyl- N atom lone pair; however, a hydrogen bond is formed between these atoms in the intermolecular packing interactions (Fig. 1). The possibility of a dimer persisting in solution similar to that found in the solid-state structure is supported by the NMR data and is worth investigating further. The ap orientation of the pyridyl-N atom may be due to the smaller relative size of the N atom to the $\mathrm{C} 3-\mathrm{H}$ atoms; alternately, there may be an electrostatic explanation because in the $s p$ orientation the pyridyl-N atom lone pair would be approximately $3.0 \AA$ from the nitro oxygen compared to $3.8 \AA$ in the $a p$ orientation.

The hydrogen positions that were defined for the methyl group attached to C 2 on the $1,4-\mathrm{DHP}$ ring contrast with the methyl group on C6 which does not show clearly defined Hatom positions. The explanation for this lies in the closer approach of a nitro-O atom to the methyl at $\mathrm{C} 2(2.70 \AA)$ relative to an ester O atom to C 6 methyl distance ( $2.85 \AA$ ); this shorter distance forces the H atoms to adopt a staggered conformation relative to the nitro-O atom.

## Experimental

The title compound was dissolved in benzene and recrystallized by slow equilibration by placing the solution inside another chamber
containing hexane. Yellow crystals of appropriate size were obtained after 1-2 d.

## Crystal data

$\mathrm{C}_{17} \mathrm{H}_{21} \mathrm{~N}_{3} \mathrm{O}_{4} \cdot 0.5 \mathrm{C}_{6} \mathrm{H}_{6}$
$M_{r}=370.42$
Monoclinic, $P 2_{1 /} / n$
$a=7.9436$ (6) A
$b=15.1440$ (11) £
$c=16.3075$ (11) $\AA$
$\beta=102.2083(13)^{\circ}$
$V=1917.4(2) \AA^{3}$
$Z=4$
$D_{x}=1.283 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
Cell parameters from 4348 reflections
$\theta=1.85-27.49^{\circ}$
$\mu=0.090 \mathrm{~mm}^{-1}$
$T=193$ (2) K
Prism, clear pale yellow
$0.38 \times 0.28 \times 0.18 \mathrm{~mm}$

## Data collection

CCD area detector diffractometer
$R_{\text {int }}=0.035$
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 1997a)
$T_{\text {min }}=0.966, T_{\text {max }}=0.984$
9844 measured reflections
4293 independent reflections
2898 reflections with $I>2 \sigma(I)$
$\theta_{\text {max }}=27.49^{\circ}$
$h=-10 \rightarrow 9$
$k=-19 \rightarrow 13$
$l=-13 \rightarrow 21$
1 standard reflection every 1200 reflections intensity decay: $<1 \%$

## Refinement

Refinement on $F^{2}$
$w=1 /\left[\sigma^{2}\left(F_{o}^{2}\right)+(0.0578 P)^{2}\right.$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.045$
$w R\left(F^{2}\right)=0.126$
$S=1.003$
4293 reflections
290 parameters
H atoms treated by a mixture of independent and constrained refinement
$+0.8765 P$ ]
where $P=\left(F_{o}{ }^{2}+2 F_{c}{ }^{2}\right) / 3$
$(\Delta / \sigma)_{\max }<0.001$
$\Delta \rho_{\max }=0.24 \mathrm{e} \AA^{-3}$
$\Delta \rho_{\max }=0.24 \mathrm{e} \mathrm{A}^{-3}$
$\Delta \rho_{\min }=-0.19 \mathrm{e}^{-3}$

Methyl-H atoms on C53, C54, C61 and C121 and one of the H atoms of the benzene solvate were allowed for as riding atoms; coordinates for methyl- H atoms on C 21 and other H atoms were refined; the $\mathrm{C}-\mathrm{H}$ distances are in the range 0.95 (2)-1.01 (3) $\AA$.

Data collection: SMART (Siemens, 1996); cell refinement: SMART; data reduction: SHELXTL (Sheldrick, 1996); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997b); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997c); molecular

Table 1
Selected geometric parameters $\left(\AA{ }^{\circ}{ }^{\circ}\right)$.

| N1-C2 | $1.3634(19)$ | C6-N1 | $1.381(2)$ |
| :--- | :--- | :--- | :---: |
| C2-C3 | $1.361(2)$ | $\mathrm{C} 3-\mathrm{N} 31$ | $1.4297(19)$ |
| C3-C4 | $1.511(2)$ | $\mathrm{N} 31-\mathrm{O} 31$ | $1.2397(17)$ |
| C4-C5 | $1.5263(19)$ | $\mathrm{N} 31-\mathrm{O} 32$ | $1.2327(17)$ |
| C5-C6 | $1.350(2)$ |  |  |
| C6-N1-C2 | $124.19(14)$ | $\mathrm{C} 2-\mathrm{C} 3-\mathrm{N} 31$ | $120.58(13)$ |
| N1-C2-C3 | $117.76(14)$ | $\mathrm{C} 4-\mathrm{C} 3-\mathrm{N} 31$ | $114.50(13)$ |
| C2-C3-C4 | $124.83(13)$ | $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 8$ | $112.59(11)$ |
| C3-C4-C5 | $110.17(12)$ | $\mathrm{C} 5-\mathrm{C} 4-\mathrm{C} 8$ | $112.35(12)$ |
| C4-C5-C6 | $121.89(13)$ | $\mathrm{C} 4-\mathrm{C} 5-\mathrm{C} 51$ | $117.34(13)$ |
| N1-C6-C5 | $120.56(14)$ | $\mathrm{C} 6-\mathrm{C} 5-\mathrm{C} 51$ | $120.77(14)$ |
| N1-C2-C21 | $113.93(14)$ | $\mathrm{C} 5-\mathrm{C} 6-\mathrm{C} 61$ | $126.63(14)$ |
| C3-C2-C21 | $128.25(14)$ | $\mathrm{N} 1-\mathrm{C} 6-\mathrm{C} 61$ | $112.63(14)$ |
|  |  |  |  |
| C6-N1-C2-C3 | $-4.6(2)$ | $\mathrm{C} 2-\mathrm{C} 3-\mathrm{N} 31-\mathrm{O} 31$ | $13.4(2)$ |
| N1-C2-C3-C4 | $-3.0(2)$ | N1-C6-C5-C51 | $179.05(13)$ |
| C2-C3-C4-C5 | $7.4(2)$ | $\mathrm{C} 6-\mathrm{C} 5-\mathrm{C} 51-\mathrm{O} 51$ | $-18.4(2)$ |
| C3-C4-C5-C6 | $-5.17(19)$ | $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 8-\mathrm{N} 7$ | $65.31(17)$ |
| C4-C5-C6-N1 | $-1.1(2)$ | $\mathrm{C} 3-\mathrm{C} 4-\mathrm{C} 8-\mathrm{C} 9$ | $-114.88(15)$ |
| C5-C6-N1-C2 | $6.7(2)$ | $\mathrm{C} 5-\mathrm{C} 4-\mathrm{C} 8-\mathrm{N} 7$ | $-59.78(17)$ |
| N1-C2-C3-N31 | $173.52(12)$ | $\mathrm{C} 5-\mathrm{C} 4-\mathrm{C} 8-\mathrm{C} 9$ | $120.03(15)$ |

graphics: XtalView (McRee, 1998); software used to prepare material for publication: XtalView and ZORTEP (Zsolnai \& Huttner, 1994).

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD1058). Services for accessing these data are described at the back of the journal.

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[^0]:    $\dagger$ Medical Research Council of Canada Group in Protein Structure and Function.

