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Key indicators

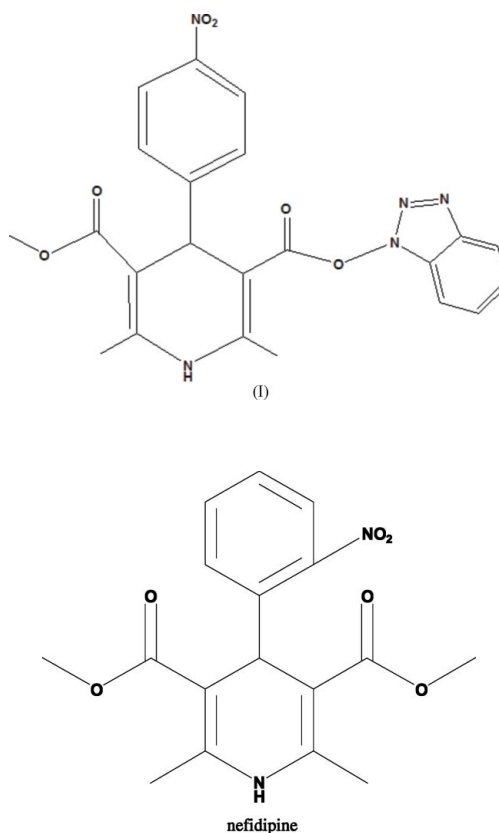
Single-crystal X-ray study
 $T = 293\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.004\text{ \AA}$
Disorder in main residue
 R factor = 0.058
 wR factor = 0.147
Data-to-parameter ratio = 11.5For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.3-Benzotriazol-1-yl 5-methyl 2,6-dimethyl-
4-(*p*-nitrophenyl)-1,4-dihydropyridine-
3,5-dicarboxylate

The title compound, $\text{C}_{22}\text{H}_{19}\text{N}_5\text{O}_6$, is an important intermediate in the synthesis of analogs of the dihydropyridine calcium channel blocker nefidipine. The crystal packing is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

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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 & Knaus, 1999; Goldmann & Stoltefuss, 1991).



The title compound, (I), is a key intermediate for preparing nefidipine analogs. In (I) (Fig. 1), the dihydropyridine ring has a flattened boat conformation, which compares well with the structure of a nefidipine analog reported earlier and with nefidipine itself (Sun *et al.*, 2006; Hofmann & Cimraglia, 1990). The crystal packing is stabilized by intermolecular $\text{N}-\text{H}\cdots\text{N}$ and $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds (Table 1), which link the molecules into chains running parallel to the *a* axis.

Experimental

The title compound was prepared by dissolving 2,6-dimethyl-4-(*p*-nitrophenyl)-1,4-dihydropyridine-3,5-dicarboxylic acid monomethyl ester (332 mg, 1 mmol) and benzotriazol-1-ol (135 mg, 1 mmol) in CH₂Cl₂ (25 ml). Dicyclohexylcarbodiimide (206 mg, 1 mmol) in CH₂Cl₂ (10 ml) was added dropwise to this solution at 278 K. The reaction mixture was stirred at 276–279 K for a further 5 h. The solvent, CH₂Cl₂, was removed by vacuum evaporation at 293 K. The desired compound, obtained in almost quantitative yield, was purified by chromatography on a silica gel column (eluted by ethyl acetate and petroleum, 1:6) at room temperature. Suitable crystals were obtained by slow evaporation of a solution in ethyl acetate.

Crystal data

C ₂₂ H ₁₉ N ₅ O ₆	$V = 1065.9 (4) \text{ \AA}^3$
$M_r = 449.42$	$Z = 2$
Triclinic, $P\bar{1}$	$D_x = 1.400 \text{ Mg m}^{-3}$
$a = 10.318 (2) \text{ \AA}$	Mo $K\alpha$ radiation
$b = 10.751 (2) \text{ \AA}$	$\mu = 0.11 \text{ mm}^{-1}$
$c = 11.491 (2) \text{ \AA}$	$T = 293 (2) \text{ K}$
$\alpha = 64.37 (3)^\circ$	Block, yellow
$\beta = 76.09 (3)^\circ$	$0.10 \times 0.08 \times 0.06 \text{ mm}$
$\gamma = 68.77 (3)^\circ$	

Data collection

Bruker SMART CCD area detector diffractometer	6615 measured reflections
ω scans	3740 independent reflections
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)	2226 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.990$, $T_{\max} = 0.994$	$R_{\text{int}} = 0.051$
	$\theta_{\max} = 25.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0742P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.058$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.147$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 0.99$	$\Delta\rho_{\max} = 0.28 \text{ e \AA}^{-3}$
3740 reflections	$\Delta\rho_{\min} = -0.22 \text{ e \AA}^{-3}$
324 parameters	Extinction correction: SHELXL97
H atoms treated by a mixture of independent and constrained refinement	Extinction coefficient: 0.041 (5)

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots N5^i$	0.86 (3)	2.35 (3)	3.156 (3)	155 (2)
$C15-H15\cdots O2^{ii}$	0.93	2.45	3.184 (4)	136
$C18-H18\cdots O3^{iii}$	0.93	2.42	3.333 (8)	167
$C21-H21\cdots O4^{iv}$	0.93	2.60	3.230 (9)	126

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

The H atom bonded to N1 was located in a difference map and its positional parameters were refined freely [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{N})$]. All other H atoms were positioned geometrically and refined using a riding model, with $C-H = 0.93\text{--}0.98 \text{ \AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ or $1.5U_{\text{eq}}(\text{C})$. The nitro group is disordered; each O atom was refined on two alternative sites with equal occupancy, and restraints were applied for geometrical similarity, planarity and approximately isotropic displacement parameters.

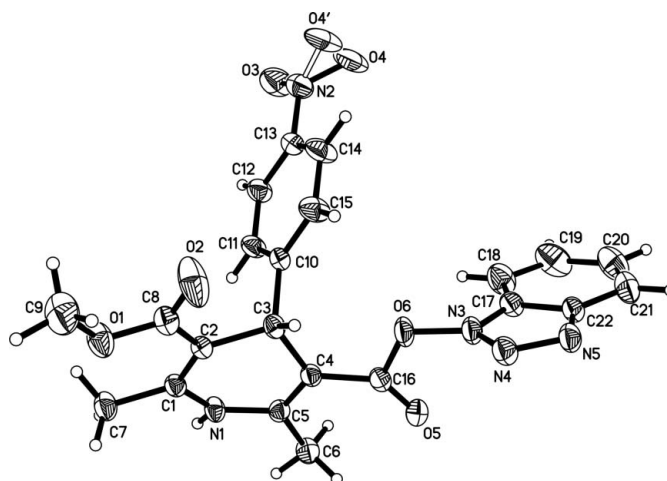


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii. The nitro group is disordered with each O atom refined on two alternative sites with equal occupancy.

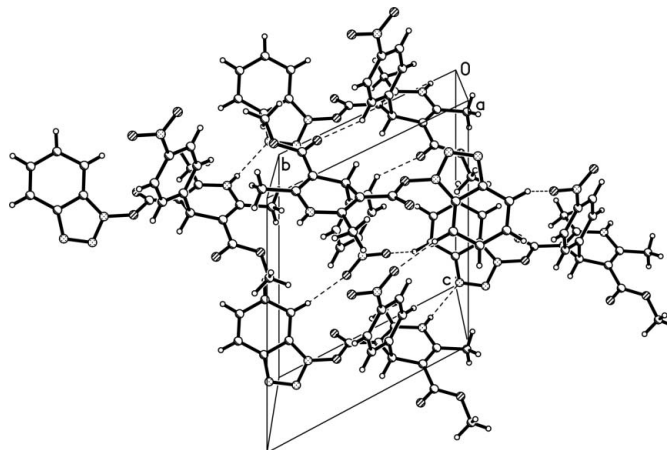


Figure 2

A packing diagram of (I). Dashed lines indicate hydrogen bonds.

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); software used to prepare material for publication: *SHELXTL*.

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