

Dimorphism of ethyl 1,4-dihydro-2,4,6-triphenylpyridine-3-carboxylate

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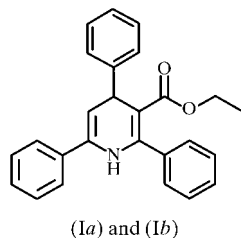
Received 21 March 2001

Accepted 29 March 2001

The title compound, $C_{26}H_{23}NO_2$, (*Ia*) and (*Ib*), shows polymorphism with crystals obtained from different solvents displaying different crystal structures. However, it is not the geometry of the single molecules nor the hydrogen-bond pattern that is different in (*Ia*) and (*Ib*), but the way in which the hydrogen-bonded chains, running along the *a*-axis direction, are arranged with respect to each other.

Comment

The 1,4-dihydropyridine ring system is of biological importance because it occurs in the reduced forms of nicotinamide adenine dinucleotide (NADH) and NADPH (Kosower & Sorensen, 1962). Since very little information is available on the geometry of aryl-substituted 1,4-dihydropyridines, the title compound, (*I*), was synthesized and its conformation was determined by means of an X-ray structure analysis. Surprisingly, crystals of (*I*) obtained from two different solvents showed two different crystal structures, (*Ia*) and (*Ib*). Thus, compound (*I*) shows polymorphism. However, since only two polymorphs have been characterized to date, one might speak of dimorphism.



Polymorphism is a well known phenomenon in crystallography (Mitscherlich, 1822) and it is extensively described in crystallographic textbooks (e.g. Dunitz, 1979; Glusker, 1994). In the case of (*I*), two polymorphs, (*Ia*) (Fig. 1) and (*Ib*) (Fig. 2), were obtained *via* crystallization from different solvents. Although both polymorphs crystallize in space group $P\bar{1}$ (No. 2) with $Z = 2$, they have slightly different unit-cell volumes and their unit-cell dimensions, apart from the *a* axis, are significantly different.

The Niggli values (Spek, 1990; Table 5) demonstrate clearly that the reduced cells are definitely different. The molecular structures of (*I*) found in the two polymorphs are very similar, indeed nearly identical. A least-squares fit of all non-H atoms in (*Ia*) and (*Ib*) gives an r.m.s. deviation of 0.263 Å. Bond lengths and angles are typical of their types (Allen *et al.*, 1987). The 1,4-dihydropyridine ring system is essentially planar [r.m.s. deviations 0.082 and 0.098 Å for (*Ia*) and (*Ib*), respectively]. The phenyl ring in the *para* position to the NH group is almost perpendicular to the heterocycle [80.54 (5)° in (*Ia*) and

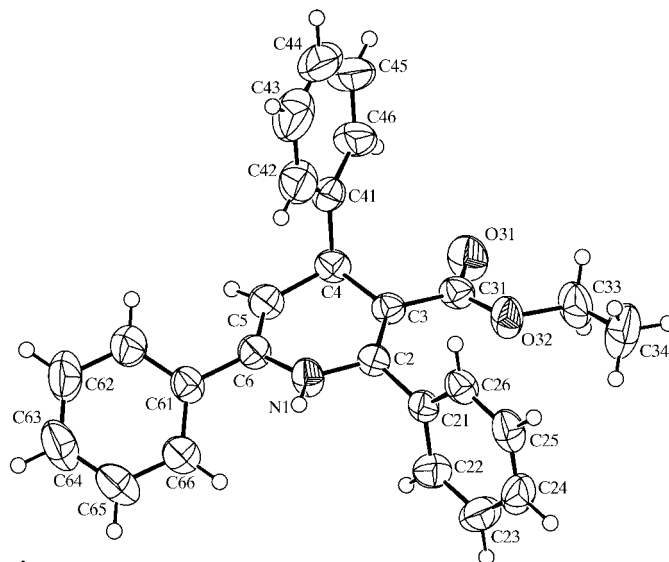


Figure 1
The molecular structure of (*Ia*) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

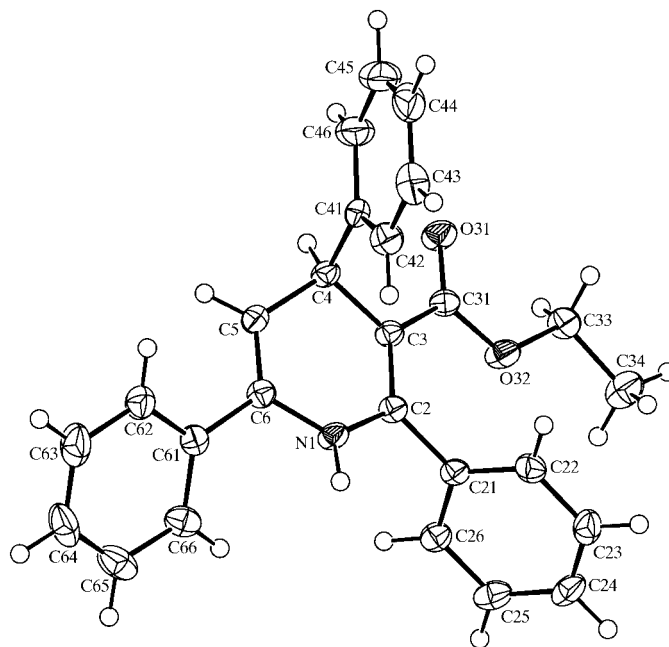


Figure 2
The molecular structure of (*Ib*) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

87.13 (3)° in (Ib)]. The phenyl rings in the *ortho* positions with respect to the NH group form smaller dihedral angles with the central ring: C21–C26 71.37 (6)° in (Ia) and 68.07 (4)° in (Ib), and C61–C66 36.96 (9)° in (Ia) and 35.58 (5)° in (Ib). Since the C21–C26 aryl ring is *ortho* with respect to the ester group, it is obvious that its dihedral angle with the heterocycle is increased due to steric repulsion with the ester group. This ester group is also planar [r.m.s. deviation 0.040 Å in (Ia) and 0.021 Å in (Ib)] and more or less coplanar with the heterocycle [17.6 (1)° in (Ia) and 7.57 (6)° in (Ib)].

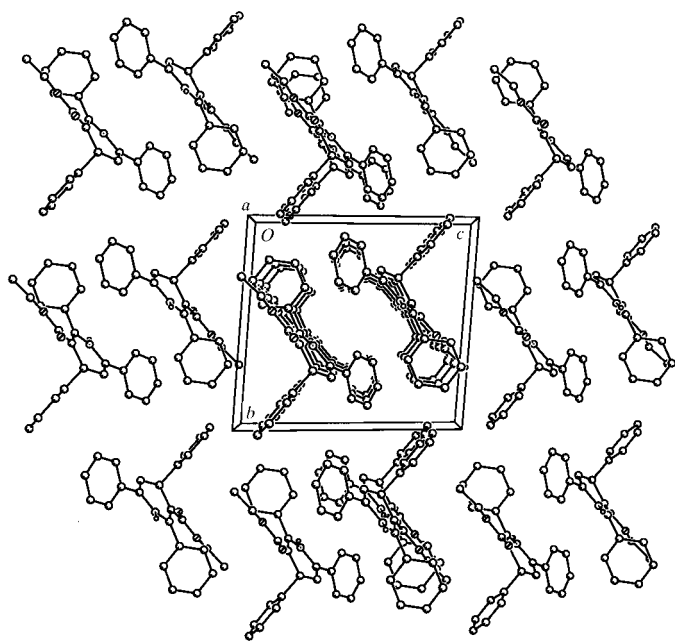


Figure 3
The packing diagram of (Ia) viewed down the *a* axis.

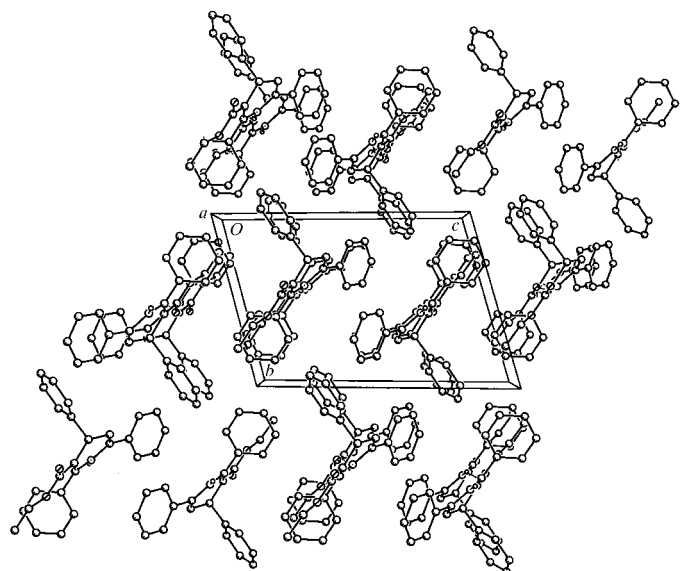


Figure 4
The packing diagram of (Ib) viewed down the *a* axis.

In the crystal, the molecules form infinite chains which are stabilized by a hydrogen bond from the NH group to the carbonyl O atom. The remarkable feature of (Ia) and (Ib), however, is their different crystal packing (Figs. 3 and 4).

Experimental

Compound (I) was synthesized according to the reaction described by Weiss (1952). Crystals of (Ia) were obtained from ethyl acetate and those of (Ib) were obtained from acetone. Having discovered that (Ib) was a polymorph of (Ia), the cell parameters of several further crystals of (Ib) were checked at room temperature and at 163 K. They all turned out to be identical with those found for the first crystal of (Ib).

Compound (Ia)

Crystal data

$C_{26}H_{23}NO_2$
 $M_r = 381.45$
Triclinic, $P\bar{1}$
 $a = 7.5083$ (2) Å
 $b = 11.6140$ (3) Å
 $c = 12.6644$ (3) Å
 $\alpha = 93.862$ (1)°
 $\beta = 98.759$ (1)°
 $\gamma = 94.527$ (1)°
 $V = 1084.53$ (5) Å³

$Z = 2$
 $D_x = 1.168$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 6287 reflections
 $\theta = 1-26^\circ$
 $\mu = 0.07$ mm⁻¹
 $T = 293$ K
Block, colourless
0.5 × 0.4 × 0.3 mm

Data collection

Siemens CCD three-circle diffractometer
 ω scans
9841 measured reflections
3961 independent reflections
3332 reflections with $I > 2\sigma(I)$

$R_{int} = 0.038$
 $\theta_{max} = 26.1^\circ$
 $h = -9 \rightarrow 9$
 $k = -14 \rightarrow 14$
 $l = -15 \rightarrow 15$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.141$
 $S = 1.10$
3961 reflections
262 parameters
H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0510P)^2 + 0.3729P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{max} < 0.001$
 $\Delta\rho_{max} = 0.22$ e Å⁻³
 $\Delta\rho_{min} = -0.28$ e Å⁻³

Table 1

Selected geometric parameters (Å, °) for (Ia).

N1–C2	1.388 (2)	C3–C4	1.541 (2)
N1–C6	1.411 (2)	C4–C5	1.514 (3)
C2–C3	1.378 (2)	C5–C6	1.349 (2)
C2–N1–C6	122.03 (14)	C5–C4–C3	110.26 (13)
C3–C2–N1	120.25 (15)	C6–C5–C4	123.29 (16)
C2–C3–C4	121.10 (15)	C5–C6–N1	119.41 (16)

Table 2

Hydrogen-bonding geometry (Å, °) for (Ia).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N1–H1 \cdots O31 ⁱ	0.86	2.17	3.0103 (19)	164

Symmetry code: (i) 1 + *x*, *y*, *z*.

Compound (Ib)

Crystal data

C₂₆H₂₃NO₂
M_r = 381.45
 Triclinic, P1̄
a = 7.5178 (1) Å
b = 10.0669 (2) Å
c = 14.6364 (2) Å
 α = 73.534 (1)°
 β = 80.826 (1)°
 γ = 81.478 (1)°
V = 1042.46 (3) Å³

Data collection

Siemens CCD three-circle diffractometer
 ω scans
 15 858 measured reflections
 4188 independent reflections
 3545 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.022

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.036
wR(*F*²) = 0.095
S = 1.03
 4188 reflections
 263 parameters
 H-atom parameters constrained

Z = 2
D_x = 1.215 Mg m⁻³
 Mo *K*α radiation
 Cell parameters from 5616 reflections
 θ = 1–25°
 μ = 0.08 mm⁻¹
T = 163 K
 Block, colourless
 0.6 × 0.4 × 0.2 mm

θ_{max} = 26.5°
h = -9 → 9
k = -12 → 12
l = -18 → 17
 260 standard reflections
 frequency: 720 min
 intensity decay: none

w = 1/[σ²(*F_o*²) + (0.0462*P*)² + 0.2564*P*]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} < 0.001
 Δρ_{max} = 0.27 e Å⁻³
 Δρ_{min} = -0.15 e Å⁻³
 Extinction correction: *SHELXL97* (Sheldrick, 1997)
 Extinction coefficient: 0.017 (2)

Table 3

Selected geometric parameters (Å, °) for (Ib).

N1—C2	1.3773 (14)	C3—C4	1.5310 (15)
N1—C6	1.4042 (14)	C4—C5	1.5186 (15)
C2—C3	1.3696 (15)	C5—C6	1.3401 (15)
C2—N1—C6	121.81 (9)	C5—C4—C3	109.82 (9)
C3—C2—N1	120.25 (10)	C6—C5—C4	122.36 (10)
C2—C3—C4	120.69 (9)	C5—C6—N1	119.72 (10)

Table 4

Hydrogen-bonding geometry (Å, °) for (Ib).

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...O31 ⁱⁱ	0.88	2.17	2.9912 (12)	155

Symmetry code: (ii) *x* - 1, *y*, *z*.

Table 5

Comparative table of reduced (Niggli) cell parameters of the two polymorphs, (Ia) and (Ib) (Å²).

Polymorph	<i>a.a</i>	<i>b.b</i>	<i>c.c</i>	<i>b.c</i>	<i>a.c</i>	<i>a.b</i>
(Ia)	56.370	134.890	160.390	-9.905	-14.480	-6.885
(Ib)	56.520	101.340	214.220	41.765	17.545	11.215

All H atoms were located by difference Fourier synthesis and refined with fixed individual displacement parameters [*U*(H) = 1.5*U*_{eq}(C_{methyl}), 1.2*U*_{eq}(C) or 1.2*U*_{eq}(N)], using a riding model with N—H = 0.86 and 0.88 Å, C—H(aromatic) = 0.93 and 0.95 Å, C—H(methyl) = 0.96 and 0.98 Å, C—H(secondary) = 0.97 and 0.99 Å, and C—H(tertiary) = 0.98 and 1.00 Å, for (Ia) and (Ib), respectively.

For both compounds, data collection: *SMART* (Siemens, 1995); cell refinement: *SMART*; data reduction: *SAINT* (Siemens, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 1990) and *XP* in *SHELXTL-Plus* (Siemens, 1991).

The author thanks the late Dr Schütz (University of Frankfurt) for providing the samples.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1471). Services for accessing these data are described at the back of the journal.

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