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2,6-Diphenylpyridine-4-carboxylic acid

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The distinctive feature of the crystal structure of 2,6diphenylpyridine-4-carboxylic acid, $C_{18}H_{13}NO_2$, is the formation of intermolecular $O-H \cdot \cdot \cdot O$ hydrogen bonds that lead to the formation of centrosymmetric cyclic dimers with $R_2^2(8)$ topology. Molecules related by translation along the *b* axis exhibit strong $\pi-\pi$ stacking of aromatic rings, with an average interplanar distance of 3.3 Å.

Comment

The most common packing motif of carboxylic acids is based on the formation of cyclic dimers through a pair of $O-H\cdots O$ bonds (Leiserowitz, 1976). By contrast, the solid-state chemistry of pyridinecarboxylic acids is dominated by the formation of an intermolecular $O-H\cdots N$ hydrogen bond between the carboxylic acid group and the pyridine N atom of a second molecule (Takusagawa & Shimada, 1976; Wright & King, 1953). The latter hydrogen-bonding interaction can be described more accurately as occurring *via* $O-H\cdots N$ or $^+N H\cdots O^-$, bonds depending on the tautomeric form present. Similar arguments also apply to other heterocyclic aromatic carboxylic acids, such as pyrazinic acid (Takusagawa *et al.*, 1974) and quinoline-4-carboxylic acid (Dobson & Gerkin, 1998). The directional nature of the $O-H\cdots N$ bonding interaction for the isonicotinic acid has been exploited in the



promotion of highly oriented structures (O'Shea *et al.*, 2001). On the other hand, when carboxylic acid groups occupy both α and α' positions of the pyridine ring, as in dipicolinic acid, the N atom is not involved in head-to-tail interactions of the type described, although O-H···N hydrogen bonds with a



Figure 1

The molecular structure of (I) showing the atom-labelling scheme and displacement ellipsoids at the 50% probability level.



Figure 2 The crystal packing of (I) viewed down the *b* axis.

cocrystallized water molecule are still present (Takusagawa et al., 1973).

The molecular structure of (I) is shown in Fig. 1. Geometric parameters are normal (Table 1) and compare well with those of other 2,6-diphenylpyridines (Krygowski *et al.*, 1994; Silva *et al.*, 1997). The whole molecule is nearly planar, as clearly shown by the values of the torsion angles reported in Table 1. Dihedral angles between the best-fit plane of the pyridine and those of the phenyl substituents are 8.7 (1) (ring atoms C8–C13) and 9.8 (1)° (ring atoms C14–C19).

The supramolecular structure of (I) can be described in terms of π - π interactions and intermolecular hydrogen bonding (Table 2). In the crystal packing, the molecules are stacked along the *b* axis (Fig. 2). The shortest interplanar distance of 2.919 (2) Å is found between the C14-C19 phenyl

ring and its translation-related ring. The stacks are organized in a pseudo-herring-bone fashion since the π ···H interactions do not seem to be dominant like in a classical herring-bone stacking mode (Gavezzotti & Desiraju, 1988; André et al., 1997). The dominant supramolecular interaction is the O-H···O hydrogen bonding between centrosymmetric carboxylic acid groups (Fig. 3). The corresponding graph-set is



Figure 3

A partial view of the crystal structure of (I) showing the formation of ring motifs with graph-set $R_2^2(8)$ that further interact through lateral C-H···O interactions to afford larger rings with graph-set $R_6^4(44)$.

therefore $R_2^2(8)$ (Etter *et al.*, 1990), which is typical for carboxyl dimers. In addition, a second motif with graph-set $R_6^4(44)$ can be identified in the crystal packing when weaker hydrogen bonds of the $C-H \cdots O$ type are considered (Fig. 3).

Experimental

The title compound was prepared in one step from 3-benzoylacrylic acid and N-phenacylpyridinium bromide in the presence of excess ammonium acetate (Blumbergs et al., 1972). The compound was recrystallized from hot acetic acid as pale-yellow crystals.

Crystal data

 $\theta_{\rm max} = 25.1^\circ$

$C_{18}H_{13}NO_2$	$D_x = 1.32 \text{ Mg m}^{-3}$
$M_r = 275.29$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 25
a = 16.524 (4) Å	reflections
b = 5.2898 (9) Å	$\theta = 8.0 - 12.9^{\circ}$
c = 17.021 (3) Å	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 111.229 \ (17)^{\circ}$	T = 293 (2) K
$V = 1386.8 (5) \text{ Å}^3$	Block, pale yellow
Z = 4	$0.23 \times 0.20 \times 0.17 \text{ mm}$
Data collection	
Siemens R3m/V diffractometer	$h = 0 \rightarrow 19$
$\theta/2\theta$ scans	$k = 0 \rightarrow 6$
2559 measured reflections	$l = -20 \rightarrow 18$
2469 independent reflections	2 standard reflections
980 reflections with $I > 2\sigma(I)$	frequency: 48 min
$R_{\rm int} = 0.069$	intensity decay: none

Table 1

Selected geometric parameters (Å, °).

N1-C2	1.343 (3)	C3-C4	1.378 (3)
N1-C6	1.350 (3)	C4-C5	1.365 (3)
O1-C7	1.297 (3)	C4-C7	1.478 (3)
O2-C7	1.243 (3)	C5-C6	1.394 (3)
C2-C3	1.393 (3)		
O2-C7-O1	122.8 (3)	O1-C7-C4	115.6 (3)
O2-C7-C4	121.6 (2)		
C3-C4-C7-O2	0.7 (4)	N1-C6-C8-C13	-7.6(4)
C5-C4-C7-O1	4.1 (4)	N1-C2-C14-C15	-8.6 (3)

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	$D-{\rm H}$	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$D1 - H1 \cdots O2^{i}$	0.82	1.83	2.640 (3)	172
$D1 - H11 \cdots O2^{ii}$	0.93	2.70	3.406 (3)	134

Symmetry codes: (i) 2 - x, 2 - y, 1 - z; (ii) $\frac{1}{2} + x$, $\frac{1}{2} - y$, $\frac{1}{2} + z$.

Refinement

Refinement on F^2	H-atom parameters constrained
R(F) = 0.045	$w = 1/[\sigma^2(F_o^2) + (0.0250P)^2]$
$wR(F^2) = 0.089$	where $P = (F_o^2 + 2F_c^2)/3$
S = 0.87	$(\Delta/\sigma)_{\rm max} < 0.001$
2469 reflections	$\Delta \rho_{\rm max} = 0.13 \ {\rm e} \ {\rm \AA}^{-3}$
190 parameters	$\Delta \rho_{\rm min} = -0.13 \text{ e } \text{\AA}^{-3}$

Data collection: XPREP (Bruker, 1997); cell refinement: XPREP; data reduction: XPREP; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: XP (Bruker, 1997).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: NA1541). Services for accessing these data are described at the back of the journal.

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