# organic papers

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#### Key indicators

Single-crystal X-ray study T = 298 KMean  $\sigma$ (C–C) = 0.004 Å R factor = 0.036 wR factor = 0.054 Data-to-parameter ratio = 12.6

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

# Diphenic acid-4,4'-bipyridine (2/1)

In the title co-crystal,  $2C_{14}H_{10}O_4 \cdot C_{10}H_8N_2$ , the diphenic acid molecules form a centrosymmetric dimeric pair, through strong  $O-H\cdots O$  hydrogen bonds involving a carboxylic acid group from each molecule. In the crystal structure, the dimers are linked by  $C-H\cdots O$  interactions to form layers parallel to the (001) plane. The adjacent layers are interlinked by bipyridine molecules,  $C_{10}H_8N_2$ , through  $O-H\cdots N$  hydrogen bonds. The bipyridine molecule lies on a centre of symmetry.

#### Comment

The binding properties of the carboxyl groups of benzoic acid and its derivatives are interesting themes of structural chemistry (Lam *et al.*, 2003). Diphenic acid (biphenyl-2,2'-dicarboxylic acid) is an extremely versatile building block for the purposes of crystal engineering. In the crystal structure of this compound, molecules form infinite chains, *via*  $R_2^2(8)$ carboxylic acid pairs (Fronczek *et al.*, 1987). A co-crystal structure of diphenic acid, diphenic acid–acridine (1/1), has been reported (Shaameri *et al.*, 2001). Here we report the cocrystal structure of diphenic acid–4,4'-bipyridine (2/1), (I), which comprises two acid and one base molecule.



The asymmetric unit of (I) consists of one molecule of diphenic acid and a half molecule of 4,4'-bipyridine. The other diphenic acid molecule and the other half of the bipyridine molecule are generated by the inversion symmetry (2 - x, 1 - y, -z) and (1 - x, 2 - y, 1 - z), respectively. In the diphenic acid, one of the carboxylic acid groups shows normal C-O distances [O1-C1 = 1.321 (3) and O2-C1 = 1.197 (3) Å], while in the other these distances are nearly equal [C14-O3 = 1.272 (2) and C14-O4 = 1.256 (2) Å], indicating that it exists as a COO<sup>-</sup> ion. However, no proton transfer from this carboxylic acid group to an N atom of the

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The structure of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme for one asymmetric unit. The other diphenic acid molecule and the other half of the bipyridine molecule are generated by the crystallographic inversion centres.

bipyridine is observed. Instead, the H<sup>+</sup> ion is located at a distance of 1.33 (4) Å from O3 and 1.29 (4) Å from O4<sup>i</sup>, indicating the formation of a centrosymmetric diphenic acid dimer through strong  $O-H \cdots O$  hydrogen bonds (Table 2). The dimers are linked through C4-H4 $\cdots$ O2<sup>iii</sup> and C6-H6 $\cdots$ O3<sup>iv</sup> interactions to form molecular layers parallel to the (001) plane. The adjacent layers are interlinked by the bipyridine molecules through O1-H1 $\cdots$ N1<sup>ii</sup> hydrogen bonds. (Symmetry codes are given in Table 2.)

# **Experimental**

Diphenic acid and 4,4'-bipyridine were obtained from Aldrich. Equimolar quantities of diphenic acid (27 mg) and 4,4'-bipyridine (15 mg) were dissolved in 15 ml of ethanol. The resulting solution was then set aside to crystallize at room temperature.

#### Crystal data

$2C_{14}H_{10}O_4 \cdot C_{10}H_8N_2$
$M_r = 640.62$
Triclinic, $P\overline{1}$
a = 6.498 (2)  Å
b = 8.002 (7)  Å
c = 16.208 (6)  Å
$\alpha = 99.45 \ (5)^{\circ}$
$\beta = 100.93 \ (3)^{\circ}$
$\gamma = 98.42 \ (5)^{\circ}$
$V = 802.4 (8) \text{ Å}^3$

#### Data collection

AFC-7*R* diffractometer  $\omega/2-\theta$  scans Absorption correction:  $\psi$  scan (North *et al.*, 1968)  $T_{min} = 0.957$ ,  $T_{max} = 0.996$ 3049 measured reflections 2830 independent reflections 1448 reflections with  $I > 2\sigma(I)$ 

#### Refinement

Refinement on FR = 0.036wR = 0.054S = 1.312830 reflections 225 parameters Z = 1  $D_x = 1.326 \text{ Mg m}^{-3}$ Mo K\alpha radiation Cell parameters from 25 reflections  $\theta = 13.2-16.3^{\circ}$   $\mu = 0.09 \text{ mm}^{-1}$  T = 298 (2) KBlock, colourless  $0.44 \times 0.32 \times 0.28 \text{ mm}$ 

 $\begin{aligned} R_{\text{int}} &= 0.016 \\ \theta_{\text{max}} &= 25.0^{\circ} \\ h &= -7 \rightarrow 7 \\ k &= 0 \rightarrow 9 \\ l &= -19 \rightarrow 18 \\ 3 \text{ standard reflections} \\ \text{every } 250 \text{ reflections} \\ \text{intensity decay: } 0.9\% \end{aligned}$ 

H atoms treated by a mixture of independent and constrained refinement  $w = 1/[(\sigma)^2(F_o) + 0.00016(F_o)^2]$  $(\Delta/\sigma)_{max} < 0.001$  $\Delta\rho_{max} = 0.30 \text{ e } \text{\AA}^{-3}$  $\Delta\rho_{min} = -0.27 \text{ e } \text{\AA}^{-3}$ 



### Figure 2

Molecular packing of (I), viewed down the a axis.

#### Table 1

Selected geometric parameters (Å).

8 (3)
6 (3)
5 (4)
0(3)
7 (4)
8 (3)
2 (4)

Symmetry code: (v) 1 - x, 2 - y, 1 - z.

#### Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
O3−H2···O4 <sup>i</sup>	1.33 (4)	1.29 (4)	2.618 (3)	171 (3)
$O1-H1\cdots N1^{ii}$	1.04 (3)	1.65 (3)	2.691 (4)	172 (3)
C4−H4···O2 <sup>iii</sup>	0.95	2.56	3.235 (4)	128
C6−H6···O3 <sup>iv</sup>	0.95	2.57	3.472 (4)	158
C3−H3···O1	0.95	2.33	2.683 (4)	101
C12−H10···O3	0.95	2.39	2.730 (4)	100

Symmetry codes: (i) 2 - x, 1 - y, -z; (ii) 2 - x, 1 - y, 1 - z; (iii) x, 1 + y, z; (iv) 1 - x, 1 - y, -z.

The C-bound H atoms were placed at their geometrically calculated positions, with C-H = 0.95 Å and refined in the riding model with  $U_{iso}(H) = 1.2U_{eq}(C)$ . Attempts to place H2 at the calculated position resulted in a non-planar  $R_2^2(8)$  ring formation, with an  $O-H \cdots O$  angle of 133°. Hence, H1 and H2 were located in a difference Fourier map and both positional and isotropic displacement para-

meters were refined. H2 is involved in nearly symmetrical O-H···O hydrogen bonds with O···H distances of 1.33 (4) and 1.29 (4) Å, respectively.

Data collection: AFC Diffractometer Control Software (Molecular Structure Corporation, 1992); cell refinement: AFC Diffractometer Control Software; data reduction: TEXSAN (Molecular Structure Corporation, 1992); program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: TEXSAN; molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: TEXSAN.

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