

## 4,4'-Bipyridine–2,4-dihydroxybenzoic acid (1/1)

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## Key indicators

Single-crystal X-ray study  
 $T = 292\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$   
 $R$  factor = 0.056  
 $wR$  factor = 0.155  
Data-to-parameter ratio = 16.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.The title compound,  $\text{C}_{10}\text{H}_8\text{N}_2 \cdot \text{C}_7\text{H}_6\text{O}_4$ , consists of 4,4'-bipyridine and 2,4-dihydroxybenzoic acid molecules, which are linked *via*  $\text{O}-\text{H} \cdots \text{N}$  hydrogen bonds, forming infinite one-dimensional chains. Adjacent chains are further linked into a two-dimensional structure by  $\text{C}-\text{H} \cdots \pi$  interactions.

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## Comment

The reliability of hydrogen bonds has been widely applied to organize one-, two- and three-dimensional networks. Moreover, hydrogen-bonded networks are organized according to their dimensionality and shape (Beatty, 2003). We report here the structure of the title compound, (I).

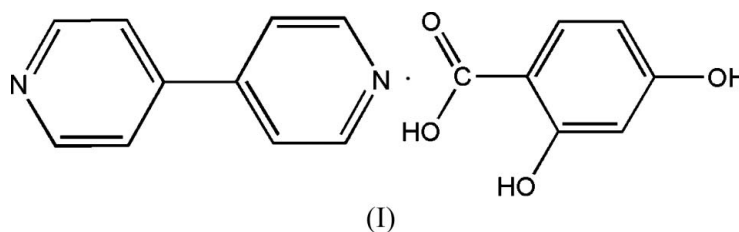
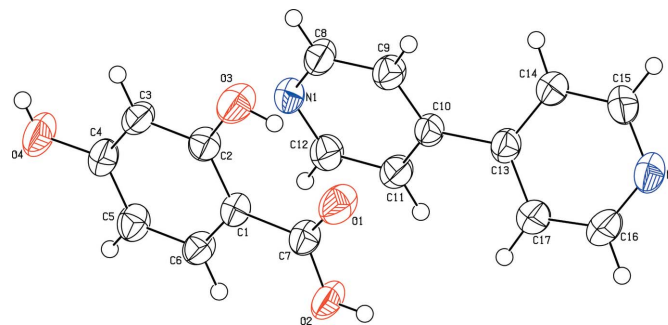
Compound (I) consists of 4,4'-bipyridine and 2,4-dihydroxybenzoic acid molecules (Fig. 1). Atoms O2 and O4 of 2,4-dihydroxybenzoic acid acts as hydrogen-bond donors to atoms N1 and N2 of 4,4'-bipyridine (Table 1), generating an infinite one-dimensional chain along the  $[40\bar{1}]$  direction (Fig. 2). There is also an intramolecular  $\text{O}-\text{H} \cdots \text{O}$  hydrogen bond in 2,4-dihydroxybenzoic acid, leading to an  $S(6)$  ring. In addition, adjacent chains are linked into a two-dimensional framework by  $\text{C}-\text{H} \cdots \pi$  interactions with an  $\text{H}15 \cdots \text{Cg}1(x, \frac{3}{2} - y, z + \frac{1}{2})$  distance of  $2.73\text{ \AA}$  (Fig. 3; Cg1 is the centroid of the C1–C6 ring).

Figure 1

The asymmetric unit of (I), showing the atom-labeling scheme. Displacement ellipsoids are drawn at the 30% probability level.

## Experimental

All reagents were commercially available and of analytical grade. An ethanol solution (3 ml) of 4,4'-bipyridine (0.156 g, 1 mmol) was added dropwise to a vigorously stirred solution of 2,4-dihydroxybenzoic acid (0.31 g, 2.0 mmol) in 10 ml distilled water. The solution was then stirred for 15 min at 343 K and filtered. On slow evaporation of the filtrate for 3 d, crystals of (I) appeared and were selected. The crystal shape and IR spectrum confirmed that they were not the starting materials. We expected to prepare a bipyridinium dihydroxybenzoate salt. However, the obtained compound was, in fact, a co-crystal of the neutral molecules.

### Crystal data

$C_{10}H_8N_2 \cdot C_7H_6O_4$	$Z = 4$
$M_r = 310.30$	$D_x = 1.399 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 6.6085 (8) \text{ \AA}$	$\mu = 0.10 \text{ mm}^{-1}$
$b = 10.7724 (12) \text{ \AA}$	$T = 292 (2) \text{ K}$
$c = 20.809 (2) \text{ \AA}$	Plate, colorless
$\beta = 95.942 (2)^\circ$	$0.32 \times 0.20 \times 0.08 \text{ mm}$
$V = 1473.4 (3) \text{ \AA}^3$	

### Data collection

Bruker SMART APEX CCD area-detector diffractometer	3528 independent reflections
$\omega$ scans	2537 reflections with $I > 2\sigma(I)$
Absorption correction: none	$R_{\text{int}} = 0.109$
12686 measured reflections	$\theta_{\text{max}} = 28.3^\circ$

### Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.056$	$w = 1/[\sigma^2(F_o^2) + (0.0824P)^2]$
$wR(F^2) = 0.155$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.00$	$(\Delta/\sigma)_{\text{max}} < 0.001$
3528 reflections	$\Delta\rho_{\text{max}} = 0.41 \text{ e \AA}^{-3}$
211 parameters	$\Delta\rho_{\text{min}} = -0.21 \text{ e \AA}^{-3}$

**Table 1**

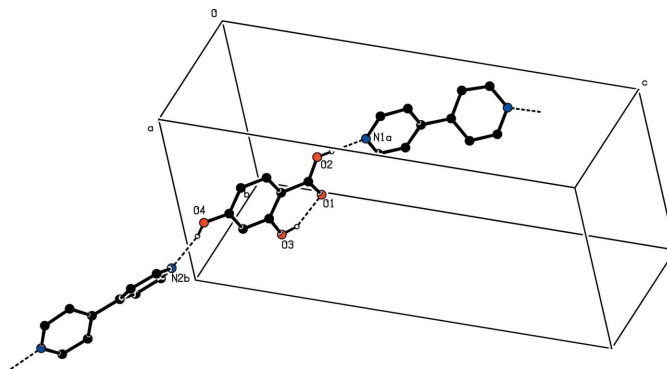
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O2-H2 \cdots N1^i$	0.82	1.82	2.6298 (15)	172
$O3-H3A \cdots O1$	0.82	1.88	2.6003 (16)	146
$O4-H4 \cdots N2^{ii}$	0.82	1.98	2.7622 (19)	160
$C15-H15 \cdots Cg1^{iii}$	0.93	2.73	3.3328 (19)	124

Symmetry codes: (i)  $x - 1, y, z$ ; (ii)  $x + 1, -y + \frac{3}{2}, z - \frac{1}{2}$ ; (iii)  $x, -y + \frac{3}{2}, z + \frac{1}{2}$ .

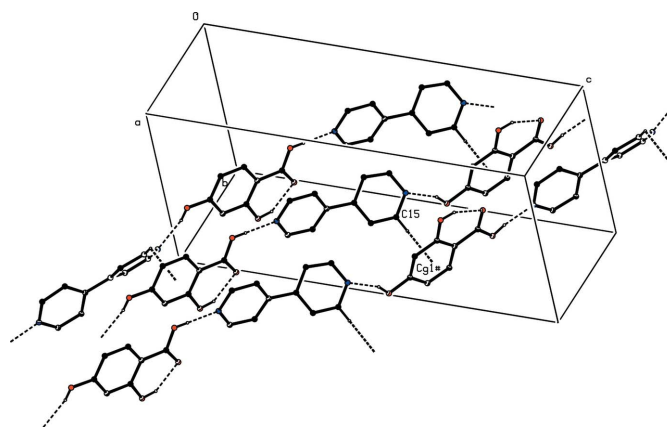
All H atoms were placed in calculated positions and refined as riding, with  $C-H = 0.93 \text{ \AA}$  and  $O-H = 0.82 \text{ \AA}$ , and  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  or  $1.5U_{\text{eq}}(\text{O})$ .

Data collection: SMART (Bruker, 2001); cell refinement: SAINT-Plus (Bruker, 2001); data reduction: SAINT-Plus; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics:



**Figure 2**

Part of the crystal structure of (I), showing the formation of a chain along the  $[40\bar{1}]$  direction. H atoms have been omitted unless involved in hydrogen bonds (dashed lines). [Symmetry codes: (a)  $x - 1, y, z$ ; (b)  $1 + x, \frac{3}{2} - y, z - \frac{1}{2}$ ]



**Figure 3**

A part of the crystal structure of (I), showing the formation of a sheet by the  $C-H \cdots \pi$  interactions between adjacent chains. H atoms have been omitted unless involved in hydrogen bonds (dashed lines). [Symmetry code: (#)  $x, \frac{3}{2} - y, \frac{1}{2} + z$ .]

PLATON (Spek, 2003); software used to prepare material for publication: PLATON.

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