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## Key indicators

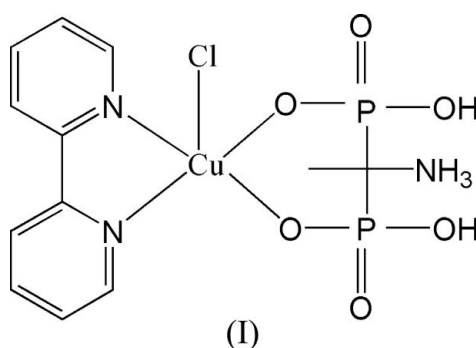
Single-crystal X-ray study  
 $T = 293\text{ K}$   
Mean  $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$   
 $R$  factor = 0.029  
 $wR$  factor = 0.083  
Data-to-parameter ratio = 17.0For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.(1-Ammonioethylidenediphosphonato- $\kappa^2\text{O},\text{O}$ )-  
(2,2'-bipyridyl- $\kappa^2\text{N},\text{N}'$ )chloridocopper(II)

In the title complex,  $[\text{Cu}(\text{C}_2\text{H}_8\text{NO}_6\text{P}_2)\text{Cl}(\text{C}_{10}\text{H}_8\text{N}_2)]$ , the  $\text{Cu}^{\text{II}}$  atom has a slightly distorted square-pyramidal geometry.  $\text{O}-\text{H}\cdots\text{O}$ ,  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{O}-\text{H}\cdots\text{Cl}$  interactions form a two-dimensional layer structure. Adjacent layers are linked to form a three-dimensional structure through  $\pi-\pi$  interactions.

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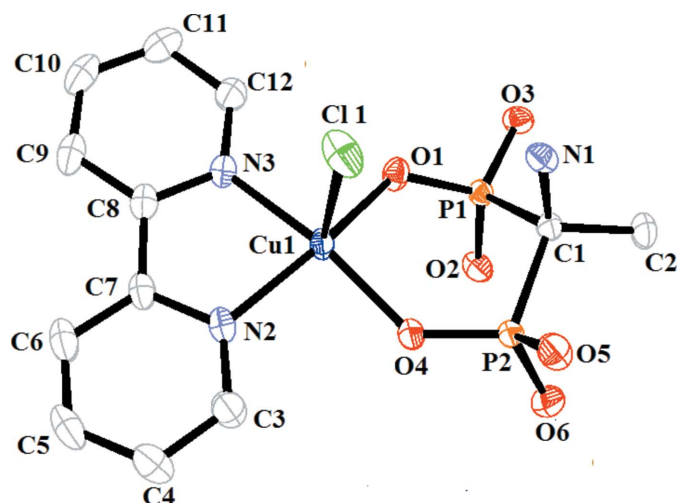
## Comment

Metal organophosphonates have been widely investigated in the past two decades due to their potential applications as catalysts, sensors, sorbents, and magnetic and luminescent materials (Clearfield, 1996; Finn *et al.*, 2003). Among the diverse organophosphonic acids, phosphonic acids and diphosphonic acids containing functional groups, such as crown ether,  $-\text{COOH}$ ,  $-\text{OH}$ ,  $-\text{NR}_2$  or mixed groups, are excellent precursors for the preparation of hybrid materials (Clearfield, 2002, and references therein; Yin *et al.*, 2003). 1-Aminoethylidenediphosphonic acid (AEDPH<sub>4</sub>) contains an  $-\text{NH}_2$  group, which exists as a zwitterion by transfer of one H atom to the amino group. Deprotonation of this group will result in predictable hydrogen-bonded aggregates from stronger  $\text{P}-\text{O}-\text{H}\cdots\text{O}-\text{P}$  to weaker  $\text{C}-\text{H}\cdots\text{X}$  hydrogen bonds. However, little attention has been paid to the structural study of metal-AEDP compounds (Yin *et al.*, 2005; Ding *et al.*, 2006; Li *et al.*, 2006). We present here the title new AEDP-containing complex, (I).

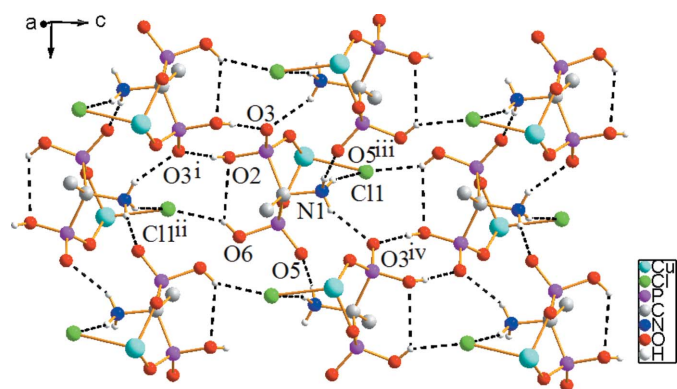


The  $\text{Cu}^{\text{II}}$  ion of (I) is coordinated by two phosphonate O atoms from the 1-aminoethylidenediphosphonate ligand, two N atoms from the 2,2'-bipyridyl ligand and one  $\text{Cl}^-$  ion in a distorted square-pyramidal geometry (Fig. 1).

Molecules of (I) are connected by a pair of strong  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds [ $\text{O}2-\text{H}2\cdots\text{O}3^{\ddagger}$ ; symmetry code: (i)  $-x, 1-y, 1-z$ ] to form a dimer. These dimers are then linked to form a layer structure parallel to the  $bc$  plane via  $\text{O}-\text{H}\cdots\text{Cl}$  and  $\text{N}-\text{H}\cdots\text{O}$  interactions, as shown in Fig. 2. Details of the hydrogen bonds are given in Table 1. The shortest



**Figure 1**  
The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level. All H atoms have been omitted for clarity.



**Figure 2**  
The two-dimensional layer structure in (I), formed *via* various hydrogen bonds, including O—H...O, N—H...O and O—H...Cl. The 2,2'-bipyridyl ligands have been omitted for clarity. [Symmetry codes: (i)  $-x, -y + 1, -z + 1$ ; (ii)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ ; (iii)  $-x, y + \frac{1}{2}, -z + \frac{1}{2}$ ; (iv)  $-x, y - \frac{1}{2}, -z + \frac{1}{2}$ .]

plane-to-plane (3.311 Å) and centroid-to-centroid (3.544 Å) distances between 2,2'-bipyridyl ligands in two neighbouring layers (symmetry relationship:  $1 - x, 1 - y, 1 - z$ ) suggest a  $\pi$ - $\pi$  stacking interaction between them. These weak  $\pi$ - $\pi$  interactions, together with the C—H...O and C—H...Cl hydrogen bonds, link adjacent layers to form a three-dimensional structure.

## Experimental

CuCl<sub>2</sub>·2H<sub>2</sub>O (0.0426 g, 0.25 mmol), 2,2'-bipyridyl (0.0390 g, 0.25 mmol), AEDPH<sub>4</sub> (0.1025 g, 0.5 mmol) and H<sub>2</sub>O (1.0 ml) were mixed and sealed in a Teflon-lined autoclave and heated at 353 K. Blue crystals of (I) suitable for single-crystal analysis were obtained after 6 d (0.0489 g, yield 42.5% based on 2,2'-bipyridyl).

## Crystal data

[Cu(C<sub>2</sub>H<sub>8</sub>NO<sub>6</sub>P<sub>2</sub>)Cl(C<sub>10</sub>H<sub>8</sub>N<sub>2</sub>)]  
 $M_r = 459.21$   
 Monoclinic,  $P2_1/c$   
 $a = 13.1203$  (12) Å  
 $b = 8.6490$  (8) Å  
 $c = 15.1501$  (14) Å  
 $\beta = 108.3670$  (10)°

$V = 1631.6$  (3) Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 1.74$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 $0.37 \times 0.30 \times 0.22$  mm

## Data collection

Bruker SMART CCD area-detector diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 2000)  
 $T_{\min} = 0.566, T_{\max} = 0.701$   
 10176 measured reflections  
 3913 independent reflections  
 3464 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.027$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.029$   
 $wR(F^2) = 0.083$   
 $S = 1.04$   
 3913 reflections  
 230 parameters  
 H-atom parameters constrained  
 $\Delta\rho_{\max} = 0.79$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.43$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
N1—H1A...O3 <sup>i</sup>	0.89	2.04	2.845 (2)	151
N1—H1B...Cl1	0.89	2.33	3.1899 (16)	161
N1—H1C...O5 <sup>ii</sup>	0.89	1.88	2.762 (2)	174
O2—H2...O3 <sup>iii</sup>	0.82	1.73	2.5316 (18)	167
O6—H6A...Cl1 <sup>iv</sup>	0.82	2.44	3.1357 (15)	143
O6—H6A...O2	0.82	2.55	3.120 (2)	128

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

All H atoms were positioned geometrically and treated as riding, with C—H = 0.93 Å (aromatic) or 0.96 Å (methyl), N—H = 0.89 Å and O—H = 0.82 Å, and with  $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N}, \text{O})$ , or  $1.5U_{\text{eq}}(\text{C})$  for methyl H.

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1997); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2000) and DIAMOND (Brandenburg, 1999); software used to prepare material for publication: SHELXTL (Bruker, 2000).

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