

Article

Charge-Assisted Hydrogen-Bonded Networks of NH_4^+ and $[\text{Co}(\text{NH}_3)_6]^{3+}$ with the New Linker Anion of 4-Phosphono-Biphenyl-4'-Carboxylic Acid

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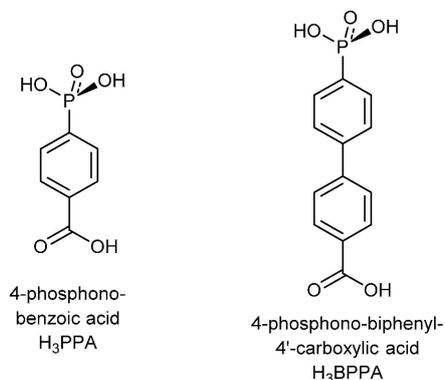
Abstract: The new linker molecule 4-phosphono-biphenyl-4'-carboxylic acid ($\text{H}_2\text{O}_3\text{P}(\text{C}_6\text{H}_4)_2\text{-COOH}$, H_3BPPA) has been structurally elucidated in hydrogen-bonded networks with the ammonium cation $\text{NH}_4(\text{H}_2\text{BPPA})(\text{H}_3\text{BPPA})$ (**1**) and the hexaamminecobalt(III) cation $[\text{Co}(\text{NH}_3)_6](\text{BPPA}) \cdot 4\text{H}_2\text{O}$ (**2**). The protic O-H and N-H hydrogen atoms were found and refined in the low-temperature single-crystal X-ray structures. The hydrogen bonds in both structures are so-called charge-assisted; that is, the H-bond donor and/or acceptor carry positive and/or negative ionic charges, respectively. The H-bonded network in **1** consists of one formally mono-deprotonated 4-phosphonato-biphenyl-4'-carboxylic acid group; that is, a H_2BPPA^- anion and a neutral H_3BPPA molecule, which together form a 3D hydrogen-bonded network. However, an almost symmetric resonance-assisted hydrogen bond (RAHB) bond [$\text{O} \cdots \text{H} = 1.17$ (**3**) and 1.26 (**3**) Å, $\text{O} \cdots \text{H} \cdots \text{O} = 180$ (**3**)°] signals charge delocalization between the formal H_2BPPA^- anion and the formally neutral H_3BPPA molecule. Hence, the anion in **1** is better formulated as $[\text{H}_2\text{BPPA} \cdots \text{H} \cdots \text{H}_2\text{BPPA}]^-$. In the H-bonded network of **2** the 4-phosphonato-biphenyl-4'-carboxylic acid is triply deprotonated, BPPA^{3-} . The $[\text{Co}(\text{NH}_3)_6]^{3+}$ cation is embedded between H-bond acceptor groups, $-\text{COO}^-$ and $-\text{PO}_3^-$ and H_2O molecules. The incorporation of sixteen H_2O molecules per unit cell makes **2** an analogue of the well-studied guanidinium sulfonate frameworks.

Keywords: hydrogen-bonded network; phosphonate-carboxylate; symmetric hydrogen bond; crystal engineering; charge-assisted H-bonds

1. Introduction

The organophosphonic acid function, which has a $\text{pK}_{\text{a}1}$ of 2.0 for the first and a $\text{pK}_{\text{a}2}$ of 6.59 for the second proton, is capable of forming strong metal-to-ligand coordinative bonds in thermodynamically stable complexes with high stability constants [1]. Metal organophosphonate compounds are multifunctional organic-inorganic hybrid materials and as open frameworks can be regarded in between zeolite-like [1,2] and metal-organic framework materials [3,4], whereas phosphonate metal-organic frameworks (MOFs) are considerably rarer than MOFs with carboxylate linkers, with phosphonates forming stronger bonds to metals than carboxylate groups [3]. Metal organophosphonates are stable in water or aqueous environment [5]. The use of metal phosphonates in catalysis, luminescence [6], ion or proton exchange or conductivity [7,8] and in separation is discussed and investigated [9]. Further, cobalt and iron organophosphonates are investigated for their magnetic properties [10–13]. Metal phosphonates are also promising porous materials [14,15], and can be reversibly hydrated and dehydrated [16].

Organophosphonates can contain additional functional groups such as carboxylate, hydroxyl or amino in the organo-moiety which presents a tunable functionality with a wide variety of structural motifs and properties [1,3,17]. Carboxy-phosphonates (Scheme 1) can be seen as intermediates between pure carboxylates and pure phosphonates, sharing synergies of both ligand classes. Carboxy-phosphonates can form porous or 3D metal-ligand networks [18–20]. Weng *et al.* described a 3D zinc carboxy-phosphonate, ZnPC-2, as a material for CO₂ adsorption [21].



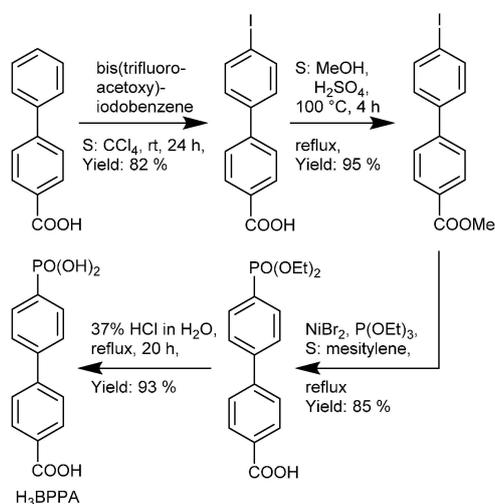
Scheme 1. Examples of phosphono-carboxylic acids.

Various metal complexes have been synthesized with a ligand from deprotonated 4-phosphono-benzoic acid, including the metals barium [22], cobalt [23], copper [23], europium [24], lead [20], lithium [25], silver [26], strontium [27], thorium [28], titanium [29], uranium [28] and zinc [30,31]. However, the extended biphenyl-based variant 4-phosphono-biphenyl-4'-carboxylic acid (H₃BPPA) was unknown so far (Scheme 1).

Herein, we present the new linker 4-phosphono-biphenyl-4'-carboxylic acid, H₃BPPA, and its deprotonated structure in hydrogen-bonded networks with NH₄⁺ and [Co(NH₃)₆]³⁺ cations.

2. Results and Discussion

4-Phosphono-biphenyl-4'-carboxylic acid H₃BPPA has been synthesized, following a known procedure by Merkushev *et al.* from 4-biphenyl carboxylic acid through the intermediates 4'-iodo-biphenyl-4-carboxylic acid [32] and its methyl ester, followed by the nickel(II)-catalyzed conversion to a phosphonate ester, which after hydrolysis gave H₃BPPA (Scheme 2).



Scheme 2. Reaction sequence for the synthesis of H₃BPPA from 4-biphenyl carboxylic acid.

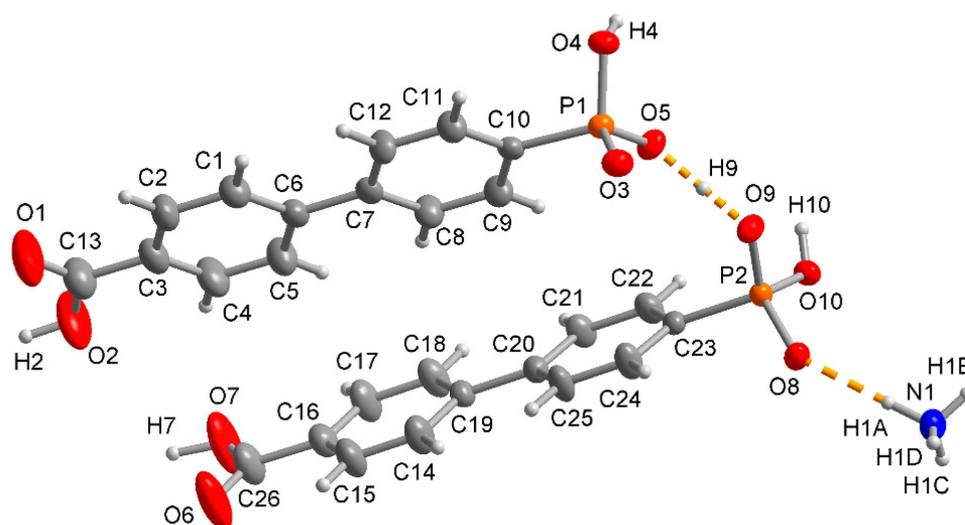
Neutralization of H₃BPPA with one equivalent of ammonium acetate yielded colorless crystals of formula NH₄(HO₃P-(C₆H₄)₂-COOH)(H₂O₃P-C₆H₄-C₆H₄-COOH) (**1**). The ammonium monohydrogenphosphonato-biphenyl-carboxylic acid crystallized with one molecule of the free H₃BPPA acid. The best results were obtained using a 1:1 ratio, though less ammonium acetate also led to product formation of lower quality. When the neutralization of H₃BPPA was carried out with excess conc. aqueous NH₃ instead of stoichiometric ammonium acetate, the same product, **1**, was formed, albeit of lower purity. Importantly, no complete or even twofold deprotonation of H₃BPPA was achieved in that way.

The asymmetric unit of **1** consists of the ammonium-cation, and formally a H₂BPPA[−] anion and a neutral H₃BPPA molecule (see below) (Figure 1a). The H₂BPPA[−] anion is derived by mono-deprotonation of the phosphonic acid group. The protic O–H and N–H hydrogen atoms were found and refined with $U_{eq} = 1.5 U_{eq}(O,N)$. The three building blocks form a three-dimensional (3D) hydrogen bonded network.

The carboxylic acid groups are oriented towards each other with the typical tail-to-tail arrangement, also known as R₂²(8)-motif in the Etter-notation (Figure 1b) [33].

The biphenyl systems of the BPPA molecules are in nearly planar geometry with 0.31 (14) and 2.79 (14)° for the dihedral angles between the aryl ring planes, and 1.7 (2)° and 3.3 (2)° for the dihedral angle between –COOH and its aryl ring in the P1 and P2 molecule, respectively. The shape of the thermal ellipsoids of the carboxyl oxygen atoms O1, O2, O6, and O7 is indicative of some rotational movement (vibration) around the (carboxyl)C–C(aryl) bond (yet, no split refinement was suggested by SHELX from the principal mean square atomic displacements). Despite the presence of the biphenyl π-systems in **1**, there are no π–π interactions [34] and only few intermolecular C–H ··· π [35] are evident. The angle is 57° for the plane formed by one biphenyl system to its neighbor.

The ammonium cation engages all of its four (found and refined) N–H bonds in the hydrogen network to four different phosphono groups. The ammonium cations and phosphono groups form hydrogen-bonded layers parallel to the *ab*-plane, separated by the biphenyl-carboxylic acid parts (Figure 1c). Ammonium benzenephosphonate, NH₄(HO₃PC₆H₅) [36], consists of a layered structure due to hydrogen bonds, with a similar motif to that of **1**.



(a)

Figure 1. Cont.

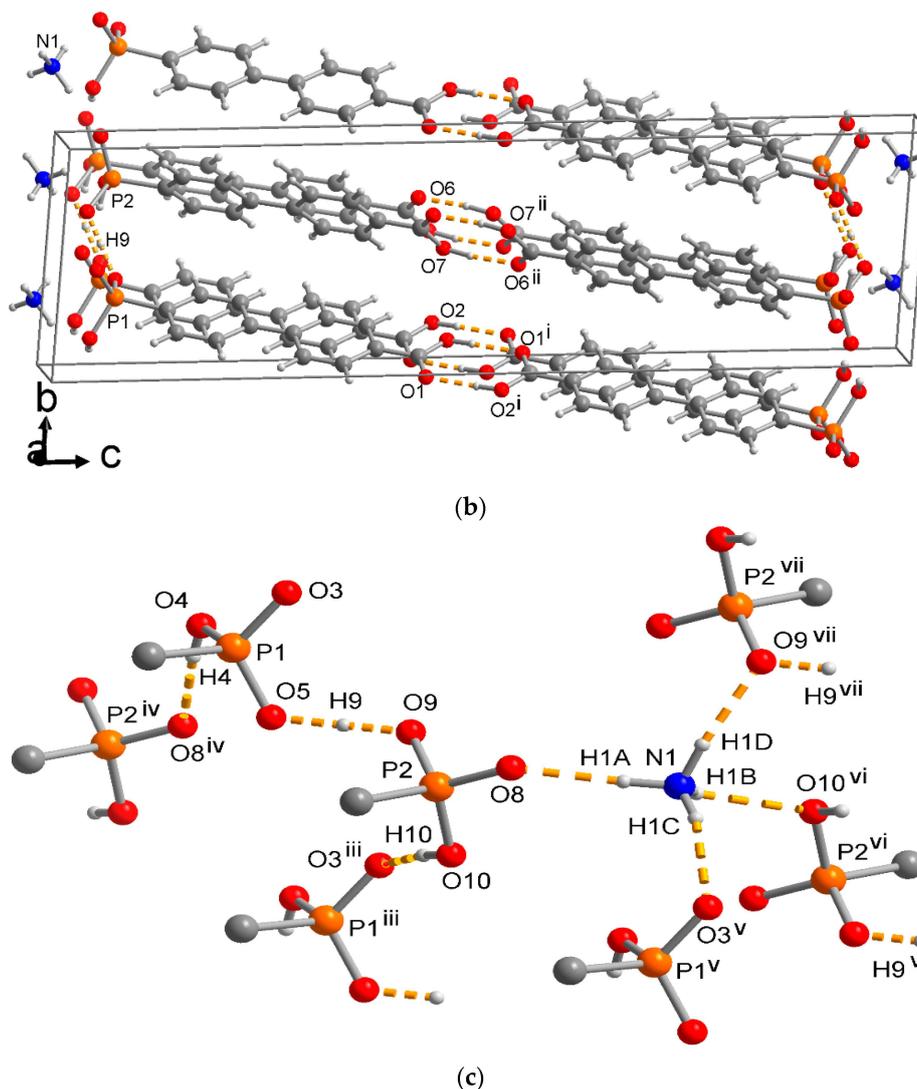


Figure 1. (a) Asymmetric unit of **1** (50% thermal ellipsoids); (b) unit-cell packing diagram with tail-to-tail arrangement of the carboxylic acid groups (showing only the carboxyl and the O9-H-O5 H bonds for clarity); and (c) full hydrogen-bonding arrangement around the NH_4^+ cation and the phosphonate and phosphonic acid groups. Details of the H-bonding interactions (orange dashed lines) are given in Table 1, selected non-hydrogen bonds and angles in Table 2. Symmetry transformations: $i = -1 - x, -y, 1 - z$; $ii = -x, 1 - y, 1 - z$; $iii = 1 + x, y, z$; $iv = x, -1 + y, z$; $v = 1 + x, 1 + y, z$; $vi = 2 - x, 2 - y, -z$; $vii = 1 - x, 2 - y, -z$.

Table 1. Details of the hydrogen bonding interactions in **1**^a.

D-H...A	D-H [Å]	H...A [Å]	D...A [Å]	D-H...A [°]	Symmetry Transformations
N1-H1A...O8	0.94 (3)	1.90 (3)	2.840 (3)	177 (3)	
N1-H1B...O10 ^{vi}	0.86 (3)	2.25 (3)	2.945 (3)	138 (3)	$vi = 2 - x, 2 - y, -z$
N1-H1C...O3 ^v	0.84 (3)	2.00 (3)	2.817 (3)	165 (3)	$v = 1 + x, 1 + y, z$
N1-H1D...O9 ^{vii}	0.95 (3)	1.92 (3)	2.845 (3)	164 (2)	$vii = 1 - x, 2 - y, -z$
O2-H2...O1 ⁱ	0.94 (5)	1.71 (5)	2.623 (3)	164 (4)	$i = -1 - x, -y, 1 - z$
O4-H4...O8 ^{iv}	0.78 (3)	1.78 (3)	2.563 (2)	175 (3)	$iv = x, -1 + y, z$
O7-H7...O6 ⁱⁱ	0.99 (5)	1.66 (5)	2.642 (3)	172 (4)	$ii = -x, 1 - y, 1 - z$
O9-H9...O5	1.17 (3)	1.26 (3)	2.428 (2)	180 (3)	
O10-H10...O3 ⁱⁱⁱ	0.83 (3)	1.72 (3)	2.537 (2)	170 (3)	$iii = 1 + x, y, z$

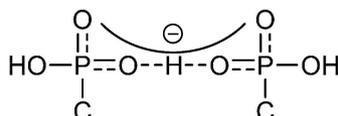
Notes: ^a D = donor, A = acceptor.

Table 2. Selected bond lengths [Å] and angles [°] in **1**.

P1–O3	1.4975 (18)	P2–O8	1.5045 (18)
P1–O5	1.5174 (18)	P2–O9	1.5190 (18)
P1–O4	1.5583 (19)	P2–O10	1.5553 (19)
P1–C10	1.787 (3)	P2–C23	1.796 (3)
C13–O1	1.235 (4)	C26–O6	1.230 (4)
C13–O2	1.283 (4)	C26–O7	1.275 (4)
O3–P1–O5	115.63 (11)	O8–P2–O9	112.38 (11)
O3–P1–O4	107.83 (10)	O8–P2–O10	109.16 (10)
O5–P1–O4	108.78 (11)	O9–P2–O10	110.04 (10)
O3–P1–C10	109.35 (11)	O8–P2–C23	109.32 (11)
O5–P1–C10	107.54 (11)	O9–P2–C23	108.76 (11)
O4–P1–C10	107.44 (11)	O10–P2–C23	107.04 (11)

Noteworthy, the H-bond O9–H9···O5 refined to an almost symmetric O9···H9···O5 hydrogen bridge with very similar distances of the H atom to both oxygen neighbors (1.17 (3) and 1.26 (3) Å) and a 180 (3)° O–H···O bond angle. This symmetric resonance-assisted hydrogen bond (RAHB) [37–41] O···H···O signals charge delocalization between the formal H₂BPPA[−] anion (of P2) and the formally neutral H₃BPPA molecule (of P1). Hence, the anion is better formulated as [H₂BPPA···H···H₂BPPA][−].

The interpretation of delocalized anion charge over the two phosphono groups is in agreement with the P–O bond lengths (Scheme 2). In each phosphonate group, there is a longer P–O bond of ~1.56 Å and two shorter P–O bonds between 1.50–1.52 Å. The P–O(H) bonds are 1.5583 (19) Å and 1.5553 (19) Å. One cannot clearly distinguish between a formally P=O double bond and a formally deprotonated P–O[−] bond. The P–O bond lengths of the symmetric O9···H9···O5 hydrogen bridge are only slightly longer (~1.52 Å) than what should be P=O double bonds (~1.50 Å). The negative charge is delocalized over the P–O[−] and P=O bonds, giving both of them a partial double bond character with P–O bond lengths between 1.50–1.52 Å (Scheme 3).

**Scheme 3.** Lewis valence structure for the bond order and charge-delocalization in the phosphonate groups in **1**.

Thermogravimetric analysis (TGA) of **1** shows a first a mass loss of ~4% up to 240 °C (Figure 2), which can be assigned to one molecule of ammonia (~3%), which is in agreement with literature values [42]. In a second step decarboxylation of one mol CO₂ (44 g/mol) leads to a mass loss of ~8%. With a third step of ~24% rapid dephosphonation of one mol PO(OH)₂ and final decarboxylation of another mol CO₂ takes place, which is followed by steady decomposition up to 700 °C.

The hydrated salt [Co(NH₃)₆](O₃P–(C₆H₄)₂–COO)·4H₂O, **2** could be crystallized from an aqueous ammonia solution 4-phosphono-biphenyl-4'-carboxylic acid. In a similar approach, 4-phosphono benzoic acid was crystallized with hexaaquacobalt(II) [43], and sulfonate ligands were crystallized with [Co(NH₃)₆]³⁺ cations, resulting in hydrogen-bonded networks [44].

In the asymmetric unit of **2** there is one trivalent hexaamminecobalt cation, one completely deprotonated BPPA^{3−} trianion and four water molecules (Figure 3a). The coordination sphere of Co³⁺ with six crystallographically different ammine ligands results in the well-known [Co(NH₃)₆]³⁺ octahedron [45]. Despite its high symmetry, the Co(NH₃)₆³⁺ octahedron does not reside on a special

position. The Co-N distances (Table 3) are comparable with that of $[\text{Co}(\text{NH}_3)_6]^{3+}$ in related complexes (Co-N = 1.951 (2) – 1.976 (2) Å, av. 1.956 (2) Å) [43,44,46].

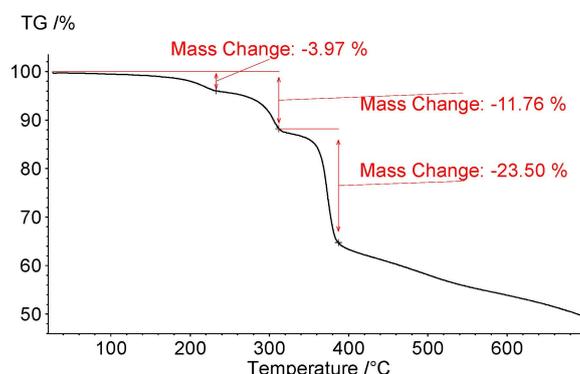


Figure 2. Thermogravimetric analysis (TGA) of **1** in the temperature range 20–700 °C.

Again, the hydrophilic groups, $[\text{Co}(\text{NH}_3)_6]^{3+}$, $-\text{COO}^-$, and $-\text{PO}_3^{2-}$, and the crystal water molecules are arranged in slabs (parallel to the *ac* plane) with slabs of the hydrophobic biphenyl part in-between (Figure 3b,c). Such a separation of hydrophilic and hydrophobic parts of molecules is a common packing motif [47–49]. Here the hydrophilic region is organized by hydrogen bonding, the biphenyl rings are arranged by singular N–H \cdots π , O–H \cdots π , C–H \cdots π or van-der-Waals interactions (see Supplementary Information). The dihedral angles within the biphenyl-carboxylate are 26.8 (4)° (ring to ring) and 42.0 (4)° ($-\text{CCOO}^-$ to aryl ring).

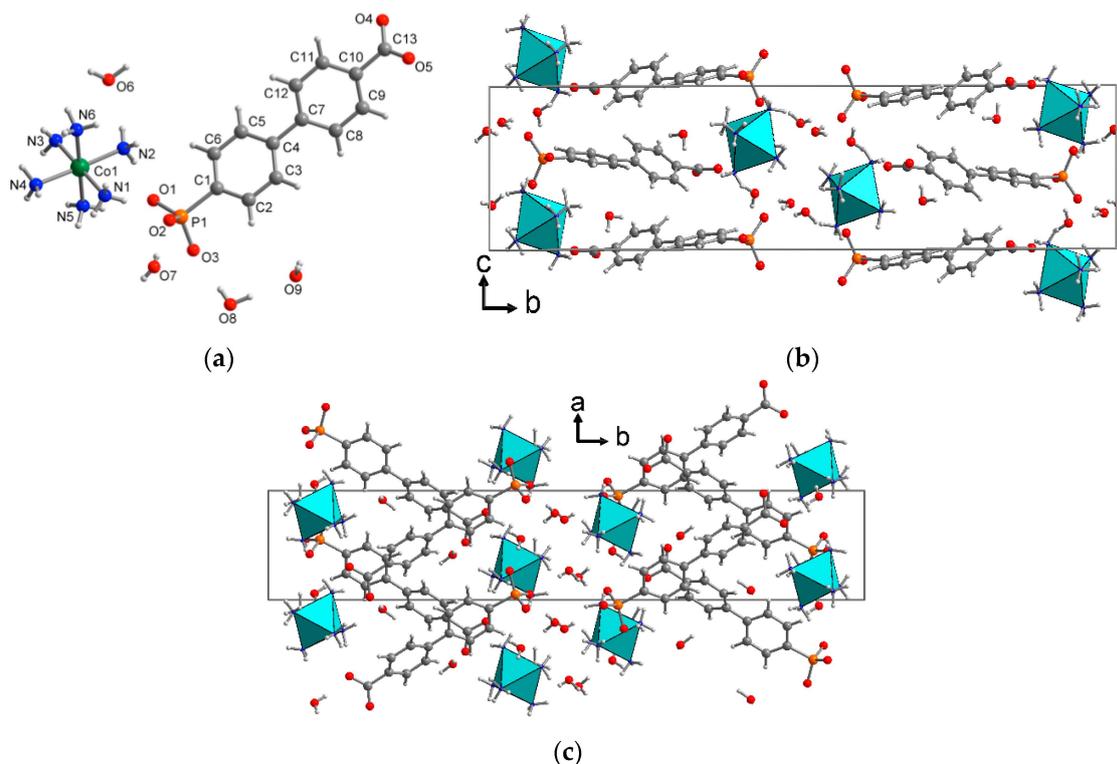


Figure 3. (a) Asymmetric unit of **2**; and (b,c) projections of the unit-cell packing on different planes. The $[\text{Co}(\text{NH}_3)_6]^{3+}$ cations are illustrated as octahedra; hydrogen bonds are not shown in (a–c) for clarity. Selected bond distances and angles are given in Table 3.

Table 3. Selected bond lengths and angles (Å, °) in **2**.

Co1–N1	1.957 (2)	P1–O1	1.5235 (18)
Co1–N2	1.965 (2)	P1–O2	1.5283 (18)
Co1–N3	1.951 (2)	P1–O3	1.5247 (18)
Co1–N4	1.959 (2)	P1–C1	1.820 (2)
Co1–N5	1.976 (2)		
Co1–N6	1.961 (2)		
N5–Co1–N1	87.17 (9)	N3–Co1–N4	89.51 (9)
N4–Co1–N1	91.46 (9)	N3–Co1–N2	90.46 (9)
N4–Co1–N5	90.01 (9)	N6–Co1–N1	91.82 (9)
N2–Co1–N1	88.66 (9)	N6–Co1–N5	178.81 (9)
N2–Co1–N5	92.53 (9)	N6–Co1–N2	88.07 (9)
N2–Co1–N4	177.47 (9)	N6–Co1–N3	90.31 (9)
N3–Co1–N1	177.66 (9)	N6–Co1–N4	89.40 (9)
N3–Co1–N5	90.71 (9)		

The BPPA^{3−} anions and the [Co(NH₃)₆] octahedra are connected to each other by hydrogen bonding (Table 4, Figure 4). The fully deprotonated phosphonate-carboxylate is solely an H-acceptor for the N–H and water O–H bonds. The carboxylate group is acceptor to O–H from water molecules. The four water molecules are held by hydrogen bonding from N–H and O–H-donors and –COO[−] and –P(O)₂O^{2−} acceptors.

Table 4. Details of the hydrogen bonding interactions in **2**^a.

D–H···A	D–H [Å]	H···A [Å]	D···A [Å]	D–H···A [°]	Symmetry Transformations
N1–H1A···O8 ⁱ	0.84 (5)	2.71 (4)	3.327 (3)	132 (4)	i = x, y, z – 1
N1–H1B···O2	0.84 (4)	2.14 (4)	2.976 (3)	173 (3)	
N1–H1C···O8 ⁱⁱ	0.87 (4)	2.15 (4)	2.971 (3)	156 (3)	ii = –x + 1, –y, –z + 1
N2–H2A···O4 ^{vi}	0.86 (4)	2.28 (4)	3.086 (3)	157 (3)	vi = x + 1/2, –y + 1/2, z – 1/2
N2–H2B···O6	0.86 (4)	2.07 (4)	2.909 (4)	163 (3)	
N2–H2C···O1	0.89 (3)	2.00 (3)	2.864 (3)	165 (3)	
N3–H3A···O6	0.82 (5)	2.60 (5)	3.177 (4)	129 (4)	
N3–H3B···O2 ⁱⁱⁱ	0.90 (4)	1.90 (4)	2.791 (3)	174 (3)	iii = x + 1, y, z
N3–H3C···O5 ^{viii}	0.69 (5)	2.56 (4)	3.048 (3)	130 (4)	viii = x + 3/2, –y + 1/2, z – 1/2
N3–H3C···O4 ^{viii}	0.69 (5)	2.56 (4)	3.180 (3)	152 (4)	viii = x + 3/2, –y + 1/2, z – 1/2
N4–H4A···O9 ^{iv}	0.90 (4)	2.05 (4)	2.937 (3)	171 (3)	iv = x + 1, y, z – 1
N4–H4B···O7 ^v	0.83 (4)	2.77 (3)	3.270 (4)	120 (3)	v = –x + 1, –y, –z
N4–H4C···O7 ⁱⁱⁱ	0.86 (4)	2.00 (4)	2.852 (3)	171 (3)	iii = x + 1, y, z
N5–H5A···O8 ⁱⁱ	0.90 (4)	2.19 (4)	3.035 (3)	157 (3)	ii = –x + 1, –y, –z + 1
N5–H5B···O2 ⁱⁱⁱ	0.88 (4)	2.63 (4)	3.437 (3)	153 (3)	iii = x + 1, y, z
N5–H5C···O1	0.83 (4)	2.13 (4)	2.939 (3)	162 (3)	
N6–H6A···O3 ⁱ	0.82 (4)	2.09 (5)	2.904 (3)	175 (3)	i = x, y, z – 1
N6–H6B···O5 ^{viii}	0.93 (4)	2.26 (4)	3.170 (3)	167 (3)	viii = x + 3/2, –y + 1/2
N6–H6C···O4 ^{vi}	0.85 (4)	2.13 (4)	2.948 (3)	160 (3)	vi = x + 1/2, –y + 1/2, z – 1/2
O6–H6E···O5 ^{viii}	0.87	1.82 (1)	2.657 (4)	161 (4)	viii = x + 3/2, –y + 1/2, z – 1/2
O7–H7A···O3 ⁱⁱ	0.62 (5)	2.12 (5)	2.740 (3)	171 (6)	ii = –x + 1, –y, –z + 1
O7–H7B···O2	0.85 (4)	1.89 (5)	2.690 (3)	156 (4)	
O8–H8A···O9	0.82 (5)	1.99 (5)	2.802 (3)	175 (4)	
O8–H8B···O3	0.73 (5)	1.97 (5)	2.693 (3)	175 (5)	
O9–H9A···O1 ^{vii}	0.74 (4)	1.96 (4)	2.704 (3)	176 (4)	vii = x – 1, y, z
O9–H9B···O4 ^{ix}	0.78 (4)	1.95 (4)	2.728 (3)	173 (4)	ix = x + 1/2, –y + 1/2, z + 1/2

Notes: ^a D = donor, A = acceptor.

Finally, we note that in both structures, **1** and **2**, the H-bonds to the phosphonate groups are so-called charge-assisted hydrogen bonds. The hydrogen bond donor and/or acceptor carry positive and negative ionic charges, respectively, hence are usually stronger and shorter than neutral H-bonds [12,46,47,50–54]. In **1** these are bonds NH₄⁽⁺⁾···^(−)O–P and NH₄⁽⁺⁾···(H)O–P, P–OH···^(−)O–P (Figure 1c). In **2** these are bonds Co–NH₃⁽⁺⁾···^(−)O–P and HOH···^(−)O–P (Figure 4).

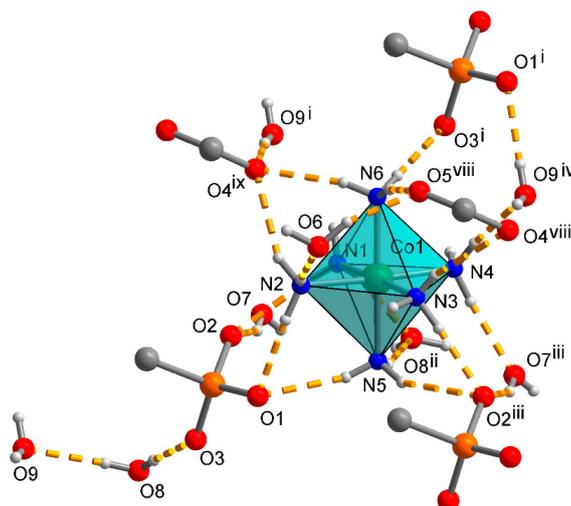


Figure 4. Most relevant H-bonding interactions (orange dashed lines) around a $[\text{Co}(\text{NH}_3)_6]^{3+}$ cation in the structure of **2**. Details of the H-bond distances and angles are listed in Table 4. Symmetry transformations: i = $x, y, z - 1$; ii = $-x + 1, -y, -z + 1$; iii = $x + 1, y, z$; iv = $x + 1, y, z - 1$; v = $-x + 1, -y, -z$; vi = $x + 1/2, -y + 1/2, z - 1/2$; vii = $x - 1, y, z$; viii = $x + 3/2, -y + 1/2, z - 1/2$; ix = $x + 1/2, -y + 1/2, z + 1/2$.

The large number of hydrogen-bonds in **2** results in a thermal stability that is higher than that of other supramolecular complexes of $[\text{Co}(\text{NH}_3)_6]^{3+}$ [55,56]. The thermal stability of BPPA^{3-} is reflected by the TGA measurement (Figure 5). The mass loss in **2** up to ($\sim 17\%$) is due to the evaporation of the four water molecules together with one ammine ligand (17.5%). The next five ammine ligands are removed along with decarboxylation of BPPA in the range from 220 °C to 500 °C, followed by decomposition of the biphenyl system ($\sim 42\%$ in total). The remaining mass of $\sim 40\%$ can be assigned to cobalt phosphonate species ($\sim 35\%$). It has been observed that metal phosphonates are stable up to 650 °C and higher [57].

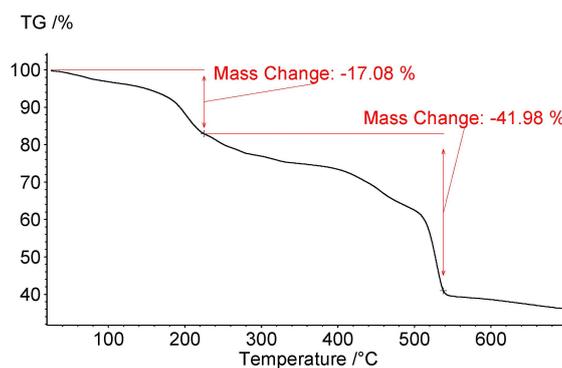


Figure 5. TGA of **2** in the temperature range 20–700 °C.

Comparison of the experimental powder X-ray diffractogram for **2** with the simulation from the single-crystal X-ray dataset (Figure 6) shows that the investigated single crystal was representative of the bulk amount when one takes into account the preferential orientation of the column- or rod-shaped crystals of **2** (Figure S1 in Supplementary Material) on the flat sample holder. Due to the preferred orientation of the rod-shaped crystals on the flat sample holder during the powder X-ray diffraction (PXRD) measurement, and their small quantity, some reflections were not present in the experimental diffraction pattern or their intensity was strongly changed. Such a behavior is discussed in detail in the literature [58–60].

Single Crystal X-ray Structures

Suitable crystals were carefully selected under a polarizing microscope, covered in protective oil and mounted on a 0.05 mm cryo loop. *Data collection.* Bruker Kappa APEX2 CCD X-ray diffractometer with microfocus tube, Mo-K α radiation ($\lambda = 0.71073 \text{ \AA}$), multi-layer mirror system, ω - and θ -scans; data collection with APEX2, cell refinement and data reduction with SAINT [63], experimental absorption correction with SADABS [64]. *Structure analysis and refinement:* All structures were solved by direct methods using SHELXL2014; refinement of **1** was done by full-matrix least squares on F^2 using the SHELX-97 program suite [65], of **2** with OLEX 2 [66,67]. Non-hydrogen atoms were refined with anisotropic displacement parameters. Hydrogen atoms were positioned geometrically (C–H = 0.95 \AA) and refined using riding models (AFIX 43 for aromatic CH with C–H = 0.93 \AA and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$). In **1** the protic hydrogen atoms (O–H, N–H) were found and freely refined with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}$ (NH and OH).

In **2**, NH₃ hydrogen atoms were found and freely refined. Water hydrogen atoms were also found and refined, except for O₆, where they were positioned geometrically (O–H = 0.870 \AA) and refined using a riding model (AFIX 6) with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

Crystal data and details on the structure refinement are given in Table 5. Graphics were drawn with DIAMOND [68]. Analyses on the supramolecular C–H \cdots O, C–H $\cdots\pi$ and π - π -stacking interactions were done with PLATON for Windows [69]. CCDC 1450889 and 1450890 contain the supplementary crystallographic data for this paper. These data can be obtained free of charge via <http://www.ccdc.cam.ac.uk/conts/retrieving.html>.

Table 5. Crystal data and refinement details.

	1	2
Chemical formula	C ₂₆ H ₂₁ O ₁₀ P ₂ ·H ₄ N	C ₁₃ H ₈ O ₅ P·CoH ₁₈ N ₆ ·4(H ₂ O)
Mr	573.41	508.36
Crystal system, space group	Triclinic, P $\bar{1}$	Monoclinic, P2 ₁ /n
Temperature (K)	150	173
a (\AA)	5.9358(5)	7.0193(5)
b (\AA)	7.5309(5)	35.454(3)
c (\AA)	27.781(2)	9.2797(7)
α ($^\circ$)	95.413(4)	90
β ($^\circ$)	90.768(5)	111.921(4)
γ ($^\circ$)	92.816(5)	90
V (\AA^3)	1234.65(16)	2142.4 (3)
Z	2	4
μ (mm^{-1})	0.24	0.93
Crystal size (mm)	0.20 \times 0.15 \times 0.01	0.33 \times 0.3 \times 0.15
Absorption correction	Multi-scan, wR2(int) was 0.0937 before and 0.0571 after correction. The Ratio of minimum to maximum transmission is 0.9165. The 1/2 correction factor is 0.0000.	Multi-scan, wR2(int) was 0.1520 before and 0.0844 after correction. The Ratio of minimum to maximum transmission is 0.6784. The 1/2 correction factor is 0.0015.
T _{min} , T _{max}	0.683, 0.746	0.507, 0.748
No. of measured, independent and observed reflections	20157, 4937, 3104 [$I > 2\sigma(I)$]	89006, 4214, 4150 [$I > 2\sigma(I)$]
R _{int}	0.066	0.073
($\sin \theta / \lambda$) _{max} (\AA^{-1})	0.617	0.617
R[F ² > 2 σ (F ²)], wR(F ²), S	0.049, 0.121, 1.02	0.038, 0.100, 1.04
No. of reflections	4937	4214
No. of parameters	379	374
$\Delta\rho_{\text{max}}$, $\Delta\rho_{\text{min}}$ ($\text{e} \cdot \text{\AA}^{-3}$)	0.30, -0.34	1.10, -0.85

4. Conclusions

We investigated the hydrogen-bonding potential of the new organo-phosphonate linker, H₃BPPA, in its (partially) deprotonated forms H₂BPPA[−] and BPPA^{3−}. As expected, the protonated phosphonic acid and deprotonated phosphonate group enters into H-bonds with all of its P–O–H donors and P–O[−] acceptors. Remarkably and unexpectedly, an almost symmetric resonance-assisted hydrogen bond (RAHB) bond was found between the formal H₂BPPA[−] anion and the formally neutral H₃BPPA molecule in **1** to give the overall anion [H₂BPPA···H···H₂BPPA][−].

Supplementary Materials: The following are available online at <http://www.mdpi.com/2073-4352/6/3/22/s001>. Figure S1: Photographs of crystals of (a) NH₄(HO₃P-(C₆H₄)₂-COOH)(H₂O₃P-(C₆H₄)₂-COOH), **1** and (b) [Co(NH₃)₆](O₃P-(C₆H₄)₂-COO)·4H₂O, **2** taken with a light microscope; Figure S2: FT-IR (ATR) spectrum of NH₄(HO₃P-(C₆H₄)₂-COOH)(H₂O₃P-(C₆H₄)₂-COOH), **1**; Figure S3: FT-IR (ATR) spectrum of [Co(NH₃)₆](O₃P-(C₆H₄)₂-COO)·4H₂O, **2**. Figure S4: Comparison of the experimental PXRD pattern of **1** (black) with the simulated pattern from the X-ray data (red). Packing analyses.

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Author Contributions: Christian Heering designed the experiments, synthesized the ligand and compound **2**. Bahareh Nateghi carried out the reaction leading to **1**. Data analysis and measurements were performed by Christian Heering, while Christoph Janiak and Christian Heering wrote the manuscript.

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