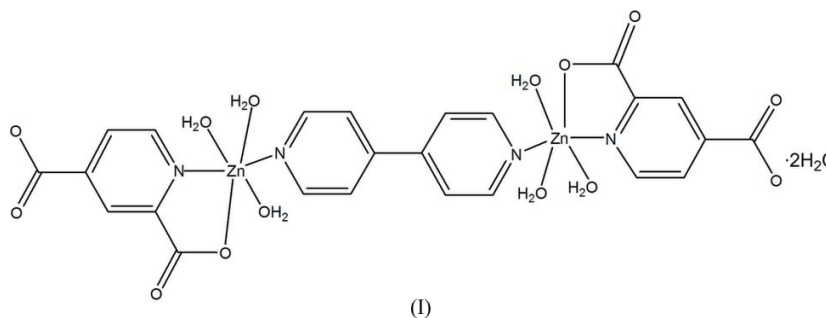


μ -4,4'-Bipyridine- $\kappa^2N:N'$ -bis[triaqua(pyridine-2,4-dicarboxylato- κ^2N,O^2)zinc(II)] dihydrateXiu-Mei Li,^{a*} Yan-Ling Niu,^a
Qing-Wei Wang^b and Bo Liu^b^aDepartment of Chemistry, Tonghua Teachers College, Tonghua 134002, People's Republic of China, and ^bDepartment of Chemistry, Jilin Normal University, Siping 136000, People's Republic of ChinaCorrespondence e-mail:
lixm20032006@yahoo.com.cn**Key indicators**Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(C-C) = 0.005$ Å
 R factor = 0.036
 wR factor = 0.061
Data-to-parameter ratio = 13.8For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

In the centrosymmetric title compound, $[Zn_2(C_7H_3NO_4)_2(C_{10}H_8N_2)(H_2O)_6] \cdot 2H_2O$, each Zn^{II} atom is six-coordinated in an octahedral geometry by one N atom of 4,4'-bipyridine (bpy), one N atom and one O atom of pyridine-2,4-dicarboxylate (pydc) and three water molecules. The crystal structure features O—H...O hydrogen bonds between carboxylate groups and water molecules, and π – π stacking interactions between the bpy and pydc molecules.

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Metal–organic complexes show a variety of supramolecular architectures (Eddaoudi *et al.*, 2001). These metal–organic network structures depend on judicious selections of metal centers and organic ligands that may provide desirable coordination covalent bonds, hydrogen-bonding linkages or π – π interactions for the construction of new supramolecular complexes. The 4,4'-bipyridine (bpy) ligand is a common linker in the construction of metal–organic frameworks, and a wide range of coordination polymers have been generated with it. Pyridinedicarboxylic acids, which can serve as bifunctional ligands, have been utilized to build coordination networks (Zhang & Chen, 2003; Wang *et al.*, 2004; Gu *et al.*, 2004). In this work, we employed pyridine-2,4-dicarboxylate (pydc) and bpy ligands, generating a new dinuclear compound, $[Zn_2(pydc)_2(bpy)(H_2O)_6] \cdot 2H_2O$, (I), under hydrothermal conditions.



In (I), each Zn^{II} atom is six-coordinated in an octahedral geometry by one N atom from the bpy ligand, one N atom and one O atom from the pydc ligand, and three water molecules (Fig. 1). The compound is binuclear, bridged by a bpy ligand across a center of inversion. π – π Interactions between the bpy and pydc molecules, with a shortest atom-to-atom distance of 3.621 (4) Å between C3 and C10(1 + x, y, z), lead to the formation of a one-dimensional chain structure. The supramolecular bands are linked into a three-dimensional network by O—H...O hydrogen bonds (Fig. 2 and Table 1).

Experimental

Compound (I) was prepared from a mixture of Zn(NO₃)₂·6H₂O (0.149 g, 0.5 mmol), H₂pydc (0.083 g, 0.5 mmol), bpy (0.078 g, 0.5 mmol) and H₂O (18 ml) in a 30 ml Teflon-lined autoclave under autogenous pressure at 433 K for 5 d. After cooling to room temperature, colorless needle-shaped crystals suitable for X-ray structure analysis were obtained. Analysis calculated for C₂₄H₃₀N₄O₁₆Zn₂: C 37.9, H 4.0, N 7.4%; found: C 37.7, H 3.8, N 7.3%.

Crystal data

[Zn₂(C₇H₃NO₄)₂(C₁₀H₈N₂)·(H₂O)₆]·2H₂O
M_r = 761.26
 Monoclinic, *P*2₁/*c*
a = 10.023 (2) Å
b = 13.941 (3) Å
c = 11.239 (2) Å
 β = 106.478 (3)°
V = 1505.9 (5) Å³
Z = 2
D_x = 1.679 Mg m⁻³
 Mo *K*α radiation
 μ = 1.68 mm⁻¹
T = 292 (2) K
 Needle, colorless
 0.43 × 0.07 × 0.06 mm

Data collection

Bruker SMART APEX CCD diffractometer
 φ and ω scans
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
T_{min} = 0.860, *T_{max}* = 0.899
 8056 measured reflections
 2985 independent reflections
 1979 reflections with *I* > 2σ(*I*)
R_{int} = 0.069
 θ_{max} = 26.1°

Refinement

Refinement on *F*²
R [*F*² > 2σ(*F*²)] = 0.036
wR (*F*²) = 0.061
S = 0.83
 2985 reflections
 216 parameters
 H-atom parameters constrained
w = 1/[σ²(*F_o*²)]
 where *P* = (*F_o*² + 2*F_c*²)/3
 (Δ/σ)_{max} = 0.001
 Δρ_{max} = 0.33 e Å⁻³
 Δρ_{min} = -0.53 e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O5—H12···O2 ⁱ	0.80	1.81	2.599 (3)	169
O5—H13···O4 ⁱⁱ	0.83	1.94	2.744 (3)	165
O6—H11···O1 ⁱⁱⁱ	0.78	2.06	2.827 (3)	168
O6—H10···O3 ^{iv}	0.99	1.73	2.719 (3)	175
O7—H8···O8 ⁱ	0.92	1.94	2.808 (3)	157
O7—H9···O8 ^v	1.00	1.74	2.735 (3)	171
O8—H1W···O5	0.73	2.17	2.889 (4)	173
O8—H2W···O4 ^{vi}	0.94	1.75	2.689 (4)	174

Symmetry codes: (i) *x*, -*y* + ½, *z* + ½; (ii) *x* - 1, *y*, *z*; (iii) -*x* + 1, -*y* + 2, -*z*; (iv) -*x* + 2, *y* + ½, -*z* + ½; (v) -*x* + 1, *y* + ½, -*z* + ½; (vi) *x* - 1, -*y* + ½, *z* - ½.

All H atoms on C atoms were positioned geometrically and refined as riding atoms, with C—H = 0.93 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C). H atoms of water molecules were located in a difference Fourier map and refined as riding in their as-found relative positions, with individual isotropic displacement parameters.

Data collection: SMART (Bruker, 2001); cell refinement: SAINT (Bruker, 2001); 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, 2001) and Mercury (Macrae *et al.*, 2006); software used to prepare material for publication: SHELXTL.

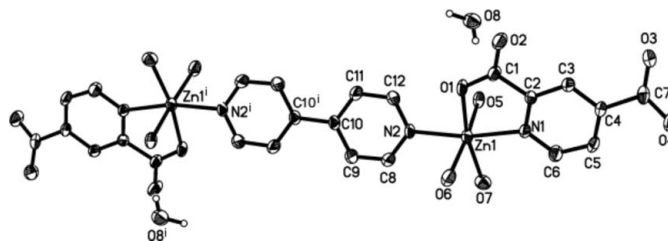


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 50% probability level. H atoms have been omitted except for those of the uncoordinated water molecules. [Symmetry code: (i) -*x*, 2 - *y*, -*z*.]

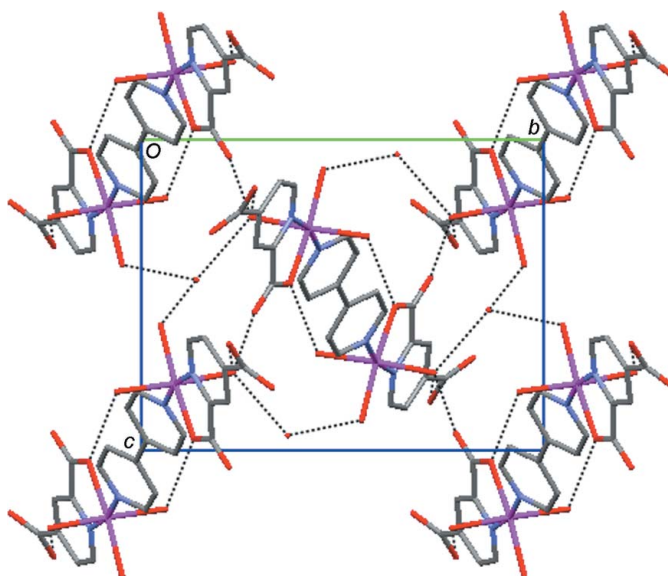


Figure 2

A view of crystal packing along the *a* axis. Dashed lines indicate hydrogen bonds. H atoms have been omitted for clarity.

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