metal-organic papers

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Key indicators

Single-crystal X-ray study T = 298 KMean $\sigma(C-C) = 0.073 \text{ Å}$ R factor = 0.106 wR factor = 0.262 Data-to-parameter ratio = 16.4

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

Tris(1,4-dimethylpyridinium) aquaheptakis-(dimethylformamide-κΟ)lanthanum(III) diphosphatooctadecatungsten dimethylformamide disolvate

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In the title compound, $(C_7H_{10}N)_3[La(P_2W_{18}O_{62})(C_3H_7NO)_7(H_2O)]\cdot 2C_3H_7NO$, 1,4-dimethylpyridinium cations, [La-(DMF)₇(H₂O)]³⁺ (DMF = dimethylformamide) complex cations and octadecatungstodiphosphate ($P_2W_{18}O_{62}^{6-}$) anions form an infinite three-dimensional assembly *via* La–O–W coordination bonds, electrostatic forces, and C–H···O and O–H···O hydrogen bonds.

Comment

In crystal engineering and polyoxometalate chemistry, one of the major challenges is to establish reliable connections between new molecular and supramolecular structures on the basis of intermolecular interactions (Desiraju 1995; Niu *et al.*, 2003). In this paper we report the preparation and crystal structure of the title compound, (I).



The asymmetric unit of (I) consists of three 1,4-dimethylpyridinium (DMAP) cations, one $[La(DMF)_7(H_2O)]^{3+}$ complex cation (DMF = dimethylformamide), one octadecatungstodiphosphate anion (P₂W₁₈O₆₂⁶⁻; abbreviated as POM) and two DMF solvent molecules. The La atom is nine-coordinated by seven O atoms from DMF, one O atom from water and one O atom from POM (Figs. 1 and 2), exhibiting a distorted tricapped trigonal–prismatic coordination geometry. The La1–O58 (from POM) bond distance is 2.70 (3) Å, which is longer than the other La–O coordinate bonds, 2.46 (4)– 2.65 (4) Å. In the Dawson structure P₂W₁₈O₆₂⁶⁻, there are 18 terminal O atoms, which are only bonded to one W atom, and the W–O distances range from 1.62 (3) to 1.74 (3) Å. The bond distance of W7–O58 is 1.67 (3) Å, a median, indicating

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The La(DMF)₇H₂O³⁺ cation and octadecatungstodiphosphate anion in the asymmetric unit of (I), showing 20% probability displacement ellipsoids.





Chain of $[La(DMF)_7(H_2O)]^{3+}$ cations and $P_2W_{18}O_{62}^{6-}$ anions along the c axis; displacement ellipsoids are depicted at the 10% probability level. Dashed lines indicate hydrogen bonds.



Figure 2

1,4-Dimethylpyridinium cations and DMF solvent molecules in the asymmetric unit of (I), showing 20% probability displacement ellipsoids.

that the formation of the O58-La1 coordinate bond has little effect on the W7-O58 bond.

In the crystal structure there are $C-H \cdots O$ and $O-H \cdots O$ hydrogen bonds (Table 2). $[La(DMF)_7(H_2O)]^{3+}$ cations and POM anions form infinite chains along the *c* axis *via* La–O– W bonds and hydrogen bonds (Fig. 3), while DMP cations and POM anions form infinite chains along the b axis via hydrogen bonds (Fig. 4). These chains crosslink to form an infinite three-





Figure 4

Chain of DMP cations and $P_2W_{18}O_{62}^{6-}$ anions along the *b* axis; displacement ellipsoids are depicted at the 10% probability level. Dashed lines indicate hydrogen bonds.

dimensional assembly through hydrogen bonds and electrostatic forces. DMP cations are located around POM anions and DMF solvent molecules are dispersed throughout the framework.

Experimental

 $K_6P_2W_{18}O_{62}$ was prepared according to the literature method (Contant, 1990). 1,4-Dimethylpyridinium iodide (DMPI) was prepared following Ren et al. (2000). Compound (I) was prepared by mixing a 0.01 mol l⁻¹ LaCl₃/DMF solution, a 0.01 mol l⁻¹ DMPI/ DMF solution and a 0.01 mol l^{-1} K₆P₂W₁₈O₆₂/DMF solution in the molar ratio 1:3:1. Bright-blue crystals of (I) were obtained from the reaction mixture by slow diffusion of water at room temperature. Yield ca 10% (evaluated from LaCl₃). Elemental analysis calculated: C 10.47, H 1.74, N 3.05%; found: C 9.83, H 1.67, N 2.57%

m2253

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Z = 4

 $D_x = 3.431 \text{ Mg m}^{-3}$

 $0.58 \times 0.25 \times 0.05 \; \text{mm}$

61983 measured reflections 21381 independent reflections 15566 reflections with $I > 2\sigma(I)$

Mo $K\alpha$ radiation

 $\mu = 19.90 \text{ mm}^{-1}$

T = 298 (2) K

Plate, blue

 $R_{\rm int}=0.091$

 $\theta_{\rm max} = 26.8^\circ$

Crystal data

(C7H10N)3[La(P2W18O62)-(C₃H₇NO)₇(H₂O)]·2C₃H₇NO $M_r = 5502.51$ Orthorhombic, Pna2, a = 38.970 (19) Åb = 14.804 (7) Å c = 18.441 (9) Å V = 10639 (9) Å³

Data collection

Bruker SMART CCD area-detector
diffractometer
φ and ω scans
Absorption correction: part of the
refinement model (ΔF)
(Parkin et al., 1995)
$T_{\min} = 0.003, T_{\max} = 0.366$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.1368P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.106$	where $P = (F_0^2 + 2F_c^2)/3$
$wR(F^2) = 0.262$	$(\Delta/\sigma)_{\rm max} = 0.001$
S = 1.00	$\Delta \rho_{\rm max} = 4.97 \text{ e } \text{\AA}^{-3}$
21381 reflections	$\Delta \rho_{\rm min} = -4.08 \text{ e } \text{\AA}^{-3}$
1300 parameters	Absolute structure: Flack (1983),
H-atom parameters constrained	9699 Friedel pairs
	Flack parameter: 0.02 (1)

Table 1

Selected geometric parameters (Å, °).

La1-O58	2.70 (3)	La1-074	2.53 (4)
La1-O70	2.53 (4)	La1-075	2.65 (4)
La1-071	2.58 (4)	La1-O76	2.49 (3)
La1-072	2.46 (4)	La1-077	2.46 (4)
La1-073	2.57 (4)		
W7-O58-La1	172.9 (17)		

Table	2
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Hydrogen-bond geometry (Å, °).

$\overline{D-\mathrm{H}\cdots A}$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
070−H702···O41 ⁱ	0.85	2.41	2.82 (5)	111
$C1-H1\cdots O46^{ii}$	0.93	2.49	3.33 (5)	150
C15-H15···O78 ⁱⁱ	0.93	2.39	3.21 (10)	148
C16−H16B····O49 ⁱⁱⁱ	0.96	2.45	3.30 (6)	148
C21-H21···O33	0.93	2.34	3.26 (7)	171
C22-H22···O49	0.93	2.45	3.33 (7)	158
C24-H24···O31 ^{iv}	0.93	2.45	3.34 (7)	161
$C25-H25\cdots O2^{iv}$	0.93	2.53	3.35 (6)	148
$C26-H26C\cdots O60^{iv}$	0.96	2.30	3.25 (11)	171
C31-H31A···O71	0.96	2.31	2.74 (7)	106
C31−H31C···O31 ^v	0.96	2.49	3.25 (6)	135
C32−H32C···O32 ⁱⁱ	0.96	2.53	3.17 (6)	124
C40-H40···O58	0.93	2.54	3.11 (5)	120
C41-H41A···O72	0.96	2.42	2.80 (6)	103
$C41 - H41C \cdot \cdot \cdot O29^{v}$	0.96	2.51	3.44 (7)	163
C42-H42A···O31	0.96	2.57	3.32 (9)	135
$C51 - H51A \cdots O50^{v}$	0.96	2.23	3.12 (6)	154

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$C51-H51B\cdots O54^{i}$	0.96	2.58	3.35 (6)	138
$C51-H51C\cdots O31^{v}$	0.96	2.55	3.42 (6)	150
C52-H52A···O73	0.96	2.24	2.66 (8)	105
$C52-H52C\cdots O54^{i}$	0.96	2.24	3.12 (9)	152
$C61 - H61B \cdot \cdot \cdot O61$	0.96	2.48	3.14 (9)	126
C62-H62A···O74	0.96	2.33	2.73 (6)	104
$C62 - H62B \cdot \cdot \cdot O26^{iii}$	0.96	2.57	3.11 (5)	116
C71−H71A···O75	0.96	2.39	2.80 (7)	105
$C71 - H71B \cdots O52^{i}$	0.96	2.47	3.08 (5)	121
C82−H82C···O49	0.96	2.59	3.52 (7)	164
C100-H100···O51 ⁱⁱ	0.93	2.59	3.37 (6)	142
C102-H97A···O78	0.96	2.25	2.66 (8)	104
$C102 - H97C \cdot \cdot \cdot O29^{vi}$	0.96	2.50	3.30 (6)	141
$C111-H11A\cdots O16^{vii}$	0.96	2.49	3.14 (9)	125

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

Because a regularly shaped crystal could not be obtained, and the faces could not be indexed adequately, our attempts to apply a numerical absorption correction were unsuccessful. Multi-scan and refdelf absorption corrections were used as comparison, and the latter can lead to a better result, though the R value (0.1057) was still high. Restraints and constraints (ISOR, FLAT, DFIX, SADI and DELU instructions in SHELXL97; Sheldrick, 1997) were used in order to maintain a reasonable geometry and atomic displacement parameters for DMP and DMF molecules. The H atoms of the water molecule were positioned with the CALCOH instruction (Nardelli, 1999) in WinGX (Farrugia, 1999), the O-H distance being 0.82-0.85 Å; $U_{iso}(H) = 1.5U_{eq}(O)$. All other H atoms were positioned geometrically and allowed to ride on their parent atoms, with C-H distances 0.93–0.96 Å, and with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C)$ for methyl groups.

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

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