## inorganic papers

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#### **Key indicators**

Powder X-ray study T = 295 KMean  $\sigma$ (P–O) = 0.015 Å R factor = 0.076 wR factor = 0.106 Data-to-parameter ratio = 19.9

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

# TIHP<sub>2</sub>O<sub>7</sub>: structure determination from X-ray powder diffraction data

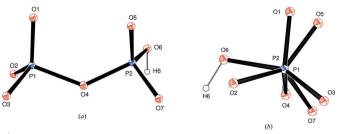
The crystal structure of monoclinic thallium(III) hydrogendiphosphate, TlHP<sub>2</sub>O<sub>7</sub>, has been solved from laboratory X-ray powder diffraction data and refined by the Rietveld method. All atoms occupy general positions. The Tl<sup>III</sup> ions have a distorted octahedral coordination environment of O atoms from four diphosphate groups. Adjacent TlO<sub>6</sub> octahedra form edge-shared Tl<sub>2</sub>O<sub>10</sub> dimers which share corners with PO<sub>4</sub> tetrahedra to build a three-dimensional polymeric structure.

#### Comment

Hydrogen diphosphates  $M^{\rm III}$ HP<sub>2</sub>O<sub>7</sub> have the ability to incorporate monovalent cations in place of hydrogen to give double salts  $M^{\rm III}M^{\rm I}$ P<sub>2</sub>O<sub>7</sub> (Lyutsko & Prisedskaya, 1986), which are of interest because of their ionic conductivity at high temperatures (Vītiņš *et al.*, 2000).

Structural investigations of  $M^{\rm III}\rm HP_2O_7$  have been performed only for  $M^{\rm III} = \rm Mn$  and Fe, including an  $\alpha$  form for  $M^{\rm III} = \rm Mn$  (Ivashkevich *et al.*, 2006), a  $\beta$  form for  $M^{\rm III} = \rm Mn$ (Durif & Averbuch-Pouchot, 1982) and Fe (Selevich *et al.*, 2006), and an unnamed form for  $M^{\rm III} = \rm Fe$  (Genkina *et al.*, 1988). All these structures are monoclinic and based on threedimensional networks of metal atoms and diphosphate groups. Because the  $\rm Mn^{\rm III}$  and Fe<sup>III</sup> cations are rather small in size, it is of interest to know whether the above crystal forms are also typical for larger trivalent cations. Two such compounds have been synthesized to date, including TIHP<sub>2</sub>O<sub>7</sub> (Selevich & Lesnikovich, 1996) and YbHP<sub>2</sub>O<sub>7</sub> (Selevich *et al.*, 2005). However, their crystal structures have not been determined. In the present paper, we report the crystal structure of TIHP<sub>2</sub>O<sub>7</sub> determined from X-ray powder diffraction data.

The unit-cell dimensions of TlHP<sub>2</sub>O<sub>7</sub> unambiguously show that the compound does not belong to the  $\alpha$  or  $\beta$  crystal forms mentioned above. However, TlHP<sub>2</sub>O<sub>7</sub> shows similarities with the second form of FeHP<sub>2</sub>O<sub>7</sub> investigated previously (Genkina *et al.*, 1988). The two compounds crystallize in the same space



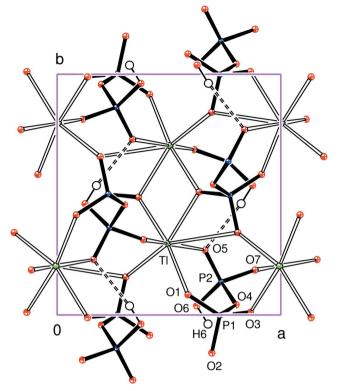
#### Figure 1

The conformation of the diphosphate anion in TlHP<sub>2</sub>O<sub>7</sub>. (*a*) Viewed approximately perpendicular to the plane P1/O4/P2. (*b*) Viewed along the P1 $\cdots$ P2 vector. Displacement parameters are drawn at the 50% probability level. The H atom is shown as a sphere of arbitrary radius.

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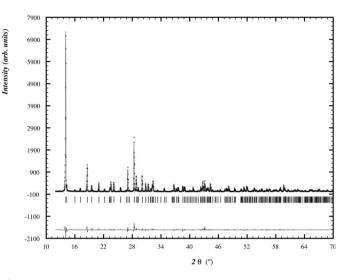
#### Figure 2

The crystal structure of  $\text{TlHP}_2\text{O}_7$ , viewed along the *c* axis. Dashed lines indicate hydrogen bonds.

group and show similar atomic arrangements. The unit-cell setting for FeHP<sub>2</sub>O<sub>7</sub> (Genkina *et al.*, 1988) differs from that for TlHP<sub>2</sub>O<sub>7</sub> and the latter is used for the purpose of comparing the two structures.

The asymmetric unit of TlHP<sub>2</sub>O<sub>7</sub> contains one Tl<sup>III</sup> ion and one HP<sub>2</sub>O<sub>7</sub><sup>3-</sup> group with a conformation intermediate between eclipsed and staggered (Fig. 1). The average of the pseudotorsion angles O1–P1···P2–O5, O2–P1···P2–O6 and O3–P1···P2–O7 is *ca* 24°. The Tl<sup>III</sup> ions have a distorted octahedral environment of O atoms from four diphosphate groups. By sharing one edge, two adjacent TlO<sub>6</sub> octahedra form a Tl<sub>2</sub>O<sub>10</sub> dimer [with a Tl···Tl distance of 3.742 (2) Å] located near the (101) plane and aligned approximately along the *b* axis (Fig. 2). The dimers and the diphosphate groups share O atoms to build a three-dimensional polymeric structure. Atom O6 is not bonded to Tl and most likely carries the H atom of the HP<sub>2</sub>O<sub>7</sub> group. The probable hydrogen bonds (Table 2) connect neighbouring diphosphate anions to form polymeric macroanions extending along the *b* axis.

Although the structures of TlHP<sub>2</sub>O<sub>7</sub> and FeHP<sub>2</sub>O<sub>7</sub> show a strong similarity, a number of differences should be mentioned. Firstly, the elongation of the unit cell in TlHP<sub>2</sub>O<sub>7</sub> [b = 9.09283 (17) Å] relative to FeHP<sub>2</sub>O<sub>7</sub> [b = 8.270 (4) Å] may be caused by the presence of the rather rigid  $M^{\rm III}_2O_{10}$  dimers aligned along the b axis and the elongation of the  $M^{\rm III}$ –O bonds (mean Tl–O = 2.250 Å versus mean Fe–O = 2.004 Å). Secondly, the P<sub>2</sub>O<sub>7</sub> groups show opposite orientations of the terminal PO<sub>3</sub> groups relative to each other (pseudo-torsion angle of *ca* 24° in TlHP<sub>2</sub>O<sub>7</sub> *versus ca* –22° in



#### Figure 3

Rietveld refinement plots of TlHP<sub>2</sub>O<sub>7</sub>, showing the experimental diffraction pattern (circles), the calculated pattern (full line), their difference (bottom) and the positions of the Bragg reflections (vertical lines). Only the low-angle region is shown  $(10-70^{\circ} 2\theta)$ .

FeHP<sub>2</sub>O<sub>7</sub>). Nevertheless, there is one-to-one correspondence between the O atoms in the two structures, so that the OH group in FeHP<sub>2</sub>O<sub>7</sub> corresponds to atom O7 in TlHP<sub>2</sub>O<sub>7</sub>. Finally, the hydrogen bonding also differs in the two structures. In FeHP<sub>2</sub>O<sub>7</sub>, the hydrogen-bonded diphosphate anions form macroanions extending along the *a* axis, whereas they are parallel to the *b* axis in TlHP<sub>2</sub>O<sub>7</sub>.

#### Experimental

The title compound was synthesized by the thin-layer method described by Selevich & Lesnikovich (1996). A mixture of  $Tl(OH)_3$  (3.83 g) and  $H_3PO_4$  (5 ml, D = 1.7 Mg m<sup>-3</sup>), in the molar ratio 1:5, was placed in a quartz crucible and heated in steps up to 423 K. The mixture was kept at this temperature until crystallization of  $TlHP_2O_7$  occurred after 2–3 d. The crystalline product was separated from the unreacted mixture by washing with water, and was dried at room temperature. The product was identified by chemical analysis, X-ray diffraction, quantitative thin-layer chromatography and IR spectroscopy.

Crystal data	
TIHP <sub>2</sub> O <sub>7</sub>	$D_x = 4.527 \text{ Mg m}^{-3}$
$M_r = 379.33$	Cu Kα radiation
Monoclinic, $P2_1/n$	$\lambda = 1.54180 \text{ Å}$
a = 8.78516 (15)  Å	T = 295  K
b = 9.09283 (17)  Å	Specimen shape: flat sheet
c = 7.24420 (12)  Å	$30 \times 30 \times 1 \text{ mm}$
$\beta = 105.8725 \ (9)^{\circ}$	Specimen prepared at 423 K
$V = 556.62 (2) \text{ Å}^3$	Particle morphology: plate-like,
Z = 4	colourless

#### Data collection

- HZG-4A (Carl Zeiss, Jena) diffractometer
- Specimen mounting: packed powder pellet

Specimen mounted in reflection mode Scan method: step  $2\theta_{\min} = 12.0, 2\theta_{\max} = 137.0^{\circ}$ Increment in  $2\theta = 0.02^{\circ}$ 

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Refinement

-	
Refinement on $I_{\rm net}$	H-atom parameters constrained
$R_{\rm p} = 0.076$	Weighting scheme based on $1/\sigma^2$
$R_{\rm wp} = 0.106$	$(\Delta/\sigma)_{\rm max} = 0.03$
$R_{\rm exp} = 0.135$	Preferred orientation correction: A
$R_{\rm B} = 0.055$	Marsh-Dollase correction of
S = 0.79	intensities (Marsh, 1932; Dollase,
Excluded region(s): none	1986) for the $(10\overline{1})$ preferred
Profile function: pseudo-Voigt	orientation of plate-like grains
profile function; $\eta = 0.62$ .	was applied. The refined values of
1036 reflections	coefficients are $G_1 = 0.668(6)$ ,
52 parameters	$G_2 = 0.316(19).$

#### Table 1

Selected geometric parameters (Å, °).

Tl-O1	2.195 (19)	P1-O2	1.448 (15)
Tl-O5	2.250 (19)	P1-O3	1.487 (15)
Tl-O3 <sup>i</sup>	2.33 (2)	P1-O4	1.656 (16)
Tl-O2 <sup>ii</sup>	2.253 (18)	P2-O4	1.618 (16)
Tl-O3 <sup>iii</sup>	2.31 (2)	P2-O5	1.455 (17)
Tl-O7 <sup>iv</sup>	2.160 (19)	P2-O6	1.584 (18)
P1-O1	1.452 (19)	P2-O7	1.486 (18)
O1-P1-O2	111.0 (11)	O4-P2-O6	109.9 (10)
O1-P1-O3	120.1 (12)	O4-P2-O7	100.6 (11)
O1-P1-O4	108.7 (11)	O5-P2-O6	115.9 (11)
O2-P1-O3	105.0 (14)	O5-P2-O7	105.3 (11)
O2-P1-O4	106.9 (11)	O6-P2-O7	115.9 (11)
O3-P1-O4	104.4 (9)	P1-O4-P2	133.8 (12)
O4-P2-O5	108.0 (11)		

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

#### Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$\overline{O6-H6\cdots O5^{v}}$	0.85	2.23	3.06 (2)	165
Symmetry code: (v)	$-x + \frac{3}{2}, y - \frac{1}{2}, -x$	$z + \frac{1}{2}$		

The diffraction pattern was indexed on a monoclinic cell ( $M_{20} = 71$ ,  $F_{20} = 138$ ,  $M_{40} = 52$ ,  $F_{40} = 110$ ) using the program *TREOR90* (Werner et al., 1985). All non-H atoms were located as part of the structure solution with  $R_{\rm F} = 0.180$ . The structure was subsequently refined against the full data set using the Rietveld method, as implemented in the FULLPROF program (Rodríguez-Carvajal, 2001). A correction for profile asymmetry was made for reflections up to  $2\theta = 40^{\circ}$ . The

atomic displacement parameters were refined isotropically, with one common parameter for the O atoms. The identification of O6 as an OH group was confirmed by bond-valence analysis (Brese & O'Keeffe, 1991; Brown, 2002) performed using the program PLATON (Spek, 2003), resulting in a bond-valence sum of 1.13 for O6. The H atom was placed in a calculated position (O-H = 0.85 Å;Sheldrick, 1997) with a displacement parameter  $U_{iso}(H) = 0.05 \text{ Å}^2$ . The final structure refinement (Fig. 3) used soft restraints for the interatomic distances of the diphosphate group, based on a geometric analysis of a large number of diphosphates.

Data collection: local program; cell refinement: FULLPROF (Rodríguez-Carvajal, 2001); data reduction: local program; program(s) used to solve structure: EXPO (Altomare et al., 1999); program(s) used to refine structure: FULLPROF; molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: FULLPROF and PLATON (Spek, 2003).

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