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Ab initio structure determination of new compound $Ba_3(BO_3)(PO_4)$

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Abstract

The crystal structure of new compound Ba₃BPO₇ was determined by ab initio method from high-resolution conventional X-ray powder diffraction data. The Rietveld refinement converged to $R_p = 5.92\%$, $R_{wp} = 8.87\%$, $R_{exp} = 5.00\%$ with the following details: Hexagonal, space group $P6_{3}mc$, a = 5.4898 (1)Å, c = 14.7551(1)Å, Z = 2. The basic unit of the structure is the [BaO₁₀]–[BO₃]–[PO₄] polar polyhedra-chain composed of Ba1–B–P–O cluster. These chains, running along *c*-axis, stack in a HCP mode to build the whole structure with triangular prism channels. The channels are parallel to *c*-axis too, in which Ba2 and Ba3 are located. \bigcirc 2004 Elsevier Inc. All rights reserved.

Keywords: Ba3BPO7; Borophosphate; Structure determination; X-ray powder diffraction

1. Introduction

Alkaline or alkaline earth borates and phosphates such as β -Ba₂B₂O₄, LiB₃O₅ [1,2] and KTiOPO₄ [3] have been in research focus for decades for their wide applications in laser and nonlinear optic technologies. Recently, in the search for new functional materials, attention has been partly switched to borophosphates, which contain both the borate group and the phosphate group as basic structural units. Borophosphates, such as $MBPO_5$ (M = Ca, Sr, Ba) [4,5], Zn_3BPO_7, Mg_3BPO_7 [6], Na₅B₂P₃O₁₃ [7], Co₅BP₃O₁₄ [8], Ba₃[(PO₄)₂BPO₄] [9], and $Pb_3[(PO_4)_2BPO_4]$ [10] have been synthesized and characterized. The phase relations, crystal structures and vibrational spectroscopy of some alkaline earth borophosphates have been systematically investigated [11–13]. A new series of borophosphates A_3 BPO₇ (A = Ca, Sr, Ba) have been obtained and characterized [14]. Here we report the structure determination of Ba₃BPO₇ by conventional high-resolution powder X-ray diffraction.

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2. Experimental

Polycrystalline Ba_3BPO_7 was prepared by hightemperature solid state reactions. Analytically pure $BaCO_3$, H_3BO_3 and $NH_4H_2PO_4$ were mixed in a molar ratio of 3:1:1 and grounded, heated at 1200°C in a Pt crucible for 48 h and then cooled to room temperature.

X-ray powder diffraction data of Ba₃BPO₇ were measured on an automated Philips PW3040/60 highresolution diffractometer equipped with Ge(111) monochromator. Only CuK α_1 was used for diffraction. Experimental details are listed in Table 1. The data were collected in a continuous mode and then transferred into a stepscan-like format of stepsize $2\theta = 0.008^{\circ}$.

Powder second-harmonic generation measurement was performed with laser powder frequency doubler.

3. Results and discussion

3.1. Indexing

The powder diffraction pattern was indexed using the program TREOR [15], on the basis of a hexagonal cell: a = 5.4888(1) Å, b = 5.4888(1) Å, c = 14.7513(1) Å, $\alpha = 90^{\circ}$, $\beta = 90^{\circ}$, $\gamma = 120^{\circ}$ (with figure of merit $M_{33} = 153$, Table 1

Experimental details of X-ray powder diffraction and Rietveld refinement for Ba_3BPO_7

Diffractometer	Philips PW3040/60
Radiation type	$CuK\alpha_1$
Wavelength (Å)	1.54060
Source to sample distance (mm)	240
Detector to sample distance (mm)	240
Scan mode	Continuous
Scan time (min)	570.86
Profile range $(2\theta^{\circ})$	9.00-120.00
Number of observations (N)	13750
Number of contributing reflections	140
Number of structure parameters (P_1)	14
Number of profile parameters (P_2)	12
<i>R</i> _B (%)	4.99
$R_{\rm p}$ (%)	5.92
$R_{\rm wp}$ (%)	8.87
$R_{\rm exp}$ (%)	5.00
S	1.78

Note: $R_{\rm B} = \sum |I_o - I_c| / \sum |I_o|, \quad R_{\rm p} = \sum |Y_{io} - Y_{ic}| / \sum |Y_{io}|, \quad R_{\rm wp} = \sum |w_i (Y_{io} - Y_{ic})^2 / \sum (w_i Y_{io})^2]^{1/2}, \quad R_{\rm exp} = [(N - P_1 - P_2) / \sum (w_i Y_{io})^2]^{1/2}, \quad S = \sum [w_i (Y_{io} - Y_{ic})^2 / (N - P_1 - P_2)]^{1/2}.$

 $F_{33} = 144$ (0.002953, 78)). The crystal system was assigned as hexagonal or triagonal with systematic absences: 0 0 l l = 2n + 1; h h l l = 2n + 1.

3.2. Cell contents determination

The chemical formula, Ba_3BPO_7 , was derived from the molar ratio and reaction properties of the starting materials. The unit cell should contain 2 formula, i.e., 6 barium, 2 phosphorus, 2 boron, and 14 oxygen atoms since the calculated density (4.883 g/cm³) is comparable with the measured density of known Ba-borophosphates [6]. Room temperature Raman scattering spectra and infrared transmission spectra measurements show that the phosphorous atoms are tetrahedrally coordinated while the boron atoms are triangly coordinated [14] by oxygen atoms.

3.3. Space group determination

According to the cell dimensions and systematic absences, the possible space groups are $P3_1c$, $P\overline{3}_1c$, $P6_3mc$, $P\overline{6}2c$ and $P6_3/mmc$. Of these five space groups, two are centrosymmetric and the other three are noncentrosymmetric. The SHG measurement shows minor second harmonic effects, which indicates Ba_3BPO_7 is either intrinsically noncentrosymmetric or centrosymmetric but with a little deviation. The noncentrosymmetric space groups are preferred because the center of symmetry can be easily added to the structure if it is centrosymmetric. Of the three noncentrosymmetric space groups, $P6_3mc$ was first chosen for its higher symmetry. In fact, $P3_1c$ is a subgroup of $P6_3mc$. All the symmetry elements of $P3_1c$ are included in $P6_3mc$. The noncentrosymmetric space group $P\overline{6}_2c$ was also tried, but the structure could not be solved. After the structure was solved, symmetries among the unique atoms were carefully checked. The results show that there is no additional symmetry element(s).

3.4. Pattern decomposition

Integrated intensities for solving the structure through the techniques for single crystal analysis were extracted from the powder diffractogram in the range $9^{\circ} < 2\theta < 120^{\circ}$ using the program Fullprof [16] on the basis of Le Bail profile-fitting procedure. Pattern decomposition with the space group $P6_3mc$ converged to $R_p = 3.98\%$, $R_{wp} = 3.88\%$. A set of 140 reflections was extracted from the powder diffraction pattern. For this space group, the intensities of exact overlapping reflections are equi-partitionable among the exact overlapping reflections [17]. The intensities of these reflections are thus accurate enough for the following work.

3.5. Structure determination

The direct methods program SHELX97 [18] was used to build the structure model from the extracted intensities. Although no identifiable molecular fragments showed in the E-map, the 3 independent Ba atoms are very obvious. The highest peak in the sequential differential Fourier electron density map was located at (0, 0, 0), which was assigned as phosphorus. Several more cycles of differential Fourier synthesis revealed the position of the rest independent atoms in the unit cell (Table 3).

3.6. Rietveld refinement

The trial structure built through direct methods and differential Fourier synthesis was then used as the starting model for the sequential Rietveld refinement. At the beginning, all the refinable atomic coordinates were attempted to be refined, but the refinement would not converge. So, the atomic coordinate x and y of O1 were adjusted to an ideal position, i.e., the [BO₃] cluster was restrained to be a equilateral triangle with B at the center and O1 (including two symmetry equivalent atoms) on the vertexes of the triangle. The B-O bond length was determined in such a way that the bond valence summation of B should be 3^+ , the formal oxidation state of boron, in accordance with Brown's bond valence theory [19]. Similarly, the positions of O2 and O3 were also fixed to restrain the $[PO_4]$ cluster to be a regular tetrahedra. Then, the z coordinates of Bal, Ba2, Ba3, O1, B and the profile parameters were refined. In the following cycles, the thermal parameters of Ba1, Ba2, Ba3 and O1, O2, O3 were set to the same values, respectively. And then, all the atoms were refined



Fig. 1. The final Rietveld refinement plot for Ba_3BPO_7 . Continuous line corresponds to observed pattern, the small crosses (+) the calculated values; vertical bars (|) indicate the Bragg positions. The bottom traces depicts the differential plot of experimental and calculated intensities.

Table 2 Crystallographic data for Ba₃BPO₇

Identification code	Ba ₃ BPO ₇
Empirical formula	Ba ₃ (BO ₃)(PO ₄)
Formula weight	565.80
Temperature	293(2) K
Wavelength	1.54060 Å
Crystal system, space group	Hexagonal, P63mc
Unit cell dimensions	$a = 5.4898(1) \text{ Å}, \ \alpha = 90^{\circ}$
	$b = 5.4898(1) \text{ Å}, \ \beta = 90^{\circ}$
	$c = 14.7551(1)$ Å, $\gamma = 120^{\circ}$
Volume	385.141(1)Å ³
Z, Calculated density	2, 4.883 g/cm^3
Absorption coefficient	$120.079 \mathrm{mm}^{-1}$
F(000)	488

Note: Cell parameters listed here are the final refined results, which are slightly different from the results of indexing procedure.

isotropically. The refinement procedure was very stable and converged to $R_p = 5.92\%$, $R_{wp} = 8.87\%$, $R_{exp} = 5.00\%$. The experimental pattern, the calculated pattern and the corresponding differential profile are shown in Fig. 1. The crystallographic details, final refined fractional coordinates and isotropic atomic thermal parameters for Ba₃BPO₇ are listed in Tables 2 and 3, respectively. The observed and calculated intensities are given in Table 4.

3.7. Structure description and evaluation

Selected bond lengths and angles are given in Table 5. Fig. 2 shows the distribution of $[PO_4]$ tetrahedron, $[BO_3]$ triangles, and the barium atoms in the unit cell. As shown in Fig. 3(a), Ba1 is coordinated by 10 oxygen atoms, of which 6 from 3 $[BO_3]$ triangles and 4 from 2 $[PO_4]$ tetrahedron, to form the $[BaO_{10}]$ polyhedron. The clusters are then linked together by oxygen atom O3

Table 3 Atomic coordinates and equivalent isotropic thermal parameters $(Å^2)$ for Ba₃BPO₇

Atom		r	v	7	R
		24	y	2	Deq
Ba1	2(a)	0.0000	0.0000	0.2186(7)	/
		0.0000	0.0000	0.2165(1)	1.030(17)
Ba2	2(b)	0.3333	0.6667	0.0321(6)	/
		0.3333	0.6667	0.0338(2)	1.030(17)
Ba3	2(b)	0.3333	0.6667	0.3877(5)	/
		0.3333	0.6667	0.3875(2)	1.030(17)
Р	2(a)	0.0000	0.0000	0.0000	/
		0.0000	0.0000	0.0000	0.971(67)
В	2(b)	0.3333	0.6667	0.749(9)	/
		0.3333	0.6667	0.7540(3)	1.405(300)
01	6(c)	0.524(10)	0.476(10)	0.259(6)	/
		0.5230(0)	0.4770(0)	0.2540(3)	1.570(64)
O2	6(c)	0.849(7)	0.151(7)	0.045(4)	/
		0.8478(0)	0.1522(0)	0.0345(0)	1.570(64)
O3	2(a)	0.0000	0.0000	0.407(18)	/
		0.0000	0.0000	0.3960(0)	1.570(64)

Note: 1st line, coordinates from SHELXL97; 2nd line, coordinates from Rietveld refinement B_{eq} is defined as one-third of the trace of the orthogonalized B_{ij} tensor. *B* for all atoms during the structure solution are set to be the default value by the program SHELEX97 and not given here.

Table 4

List of calculated and observed intensities and d-spacing of selected reflections for Ba_3BPO_7

hkl	$I_{\rm calc}$	$I_{\rm obs}$	$d_{\rm calc}$	$d_{\rm obs}$	hkl	$I_{\rm calc}$	$I_{\rm obs}$	d_{calc}	$d_{\rm obs}$
103	100.00	100.00	3.418	3.417	205	12.32	11.60	1.851	1.851
110	97.51	96.93	2.745	2.744	108	12.05	11.22	1.719	1.719
102	60.10	57.22	3.996	3.994	226	10.95	10.88	1.198	1.198
104	37.01	40.20	2.914	2.913	201	9.70	10.10	2.347	2.346
116	42.60	40.00	1.832	1.831	416	10.14	10.10	0.956	0.956
203	36.60	35.99	2.140	2.140	218	9.31	8.82	1.287	1.287
202	30.89	31.52	2.263	2.262	215	8.58	8.60	1.535	1.535
101	23.78	27.37	4.525	4.523	313	8.50	8.60	1.274	1.273
006	24.17	23.98	2.459	2.458	004	9.90	8.50	3.689	3.687
105	21.69	20.40	2.507	2.506	410	7.87	8.21	1.037	1.037
204	17.83	19.54	1.998	1.998	208	7.28	7.34	1.457	1.457
213	19.01	19.00	1.688	1.688	211	6.08	6.69	1.784	1.783
300	16.01	15.86	1.585	1.585	002	5.06	5.66	7.377	7.370
212	13.98	14.33	1.746	1.746	2114	6.02	5.61	0.909	0.909
220	13.71	14.17	1.372	1.372	312	5.02	5.46	1.298	1.298
214	12.27	12.92	1.615	1.615	323	5.20	5.43	1.065	1.065
306	13.45	12.80	1.332	1.332					

Note: Only reflections with relative intensity greater than 5 are listed.

to form a $[BaO_{10}]$ – $[BO_3]$ – $[PO_4]$ polar polyhedra-chain running along the *c*-axis, which is the basic unit of the crystal structure. These parallel chains, connected to each other by sharing oxygen atom O1, stack in the hexagonal close packing (HCP) mode to build a structure with triangular prism channels in which Ba2, Ba3 and their symmetry equivalents are located (Fig. 3(b)). The coordination geometries of Ba2 and Ba3 are shown in Figs. 4(a) and (b), respectively.

Table 5 Selected Bond lengths [Å] and angles [°] for Ba₃BPO₇

Ba1–O1	2.820(1)	Ba3–O2#15	2.769(2)
Ba1–O1#1	2.820(1)	Ba3-O2#16	2.769(2)
Ba1-O1#2	2.820(1)	Ba3-O2#17	2.769(2)
Ba1–O1#3	2.820(1)	Ba3-O3#11	3.172(1)
Ba1–O1#4	2.820(1)	Ba3-O3#12	3.172(1)
Ba1–O1#5	2.820(1)	Ba3–O3	3.172(1)
Ba1-O2#6	2.998(2)		
Ba1–O2#7	2.998(2)	PO2#8	1.534(0)
Ba1-O2#8	2.998(2)	PO2#6	1.534(0)
Ba1–O3	2.706(2)	PO2#7	1.534(0)
	~ /	P-O3#18	1.534(0)
Ba2–O1	3.716(4)		
Ba2-O1#14	3.716(4)	B-O1#17	1.366(0)
Ba2–O1#3	3.716(4)	B-O1#15	1.366(0)
Ba2–O2#5	2.748(0)	B-O1#16	1.366(0)
Ba2-O2#10	2.748(0)		
Ba2–O2#7	2.748(0)	O2#8-P-O2#6	109.537(0)
Ba2–O2#11	2.748(0)	O2#8-P-O2#7	109.537(0)
Ba2–O2#8	2.748(0)	O2#6-P-O2#7	109.537(0)
Ba2–O2#2	2.748(0)	O2#8-P-O3#18	109.405(0)
Ba2–O3#18	3.765(1)	O2#6-P-O3#18	109.405(0)
Ba2–O3#9	3.765(1)	O2#7-P-O3#18	109.405(0)
Ba2–O3#13	3.765(1)		
		O1#17-B-O1#17	120.000(0)
Ba3–O1#3	2.670(3)	O1#17-B-O1#15	120.000(0)
Ba3-O1#14	2.670(3)	O1#18-B-O1#15	120.000(0)
Ba3–O1	2.670(3)		

Note: [PO₄] and [BO₃] are restrained to be regular tetrahedron and triangles respectively Symmetry transformations used to generate equivalent atoms: #1 x-1,y-1,z; #2 -y+1,x-y,z; #3 -x+y,-x+1,z; #4 -x+y,-x,z; #5 -y,x-y,z; #6 -y,x-y-1,z; #7 x-1,y,z; #8 -x+y+1,-x+1,z; #9 -x,-y+1,z-1/2; #10 -x+y+1,-x+2,z; #11 x,y+1,z; #12 x+1,y+1,z; #13 -x+1,-y+1,z-1/2; #14 -y+1,x-y+1,z; #15 -x+1,-y+1,z+1/2; #16 x-y,x,z+1/2; #17 y,-x+y+1,z+1/2; #18 -x,-y,z-1/2.

Bond valence calculations according to Brown's rules are listed in Table 6. It is obvious that the bond valence sums are reasonable.

4. Discussion

To our knowledge, the other two borophosphates of the same formula type with Ba_3BPO_7 are Zn_3BPO_7 and Mg_3BPO_7 [6]. But the structure type of these two compounds is different from that of Ba_3BPO_7 . On the contrary, the atomic distribution of $BaPBO_3$ [20] and $SrPBO_3$ [21] are very similar to that of Ba_3BPO_7 although they have different formula type (Table 7 and Fig. 5). For Zn_3BPO_7 , if the phosphorus atom P is taken as the origin, and axis *a* is halved, the structure is still different from the Ba_3BPO_7 structure. These differences perhaps stem from the different coordination environment of Zn and Ba. Compared with Ba, Zn has a smaller coordination number and the Zn–O polyhedron is relatively rigid. To form a new structure, these





Fig. 3. (a) $[BaO_{10}]$ - $[BO_3]$ - $[PO_4]$ polar polyhedra-chain running along the *c*-axis and (b) stacking of $[BaO_{10}]$ - $[BO_3]$ - $[PO_4]$ polar polyhedra-chain in the unit cell.

requirements must be met. As a result, the Zn, $[PO_4]$ and $[BO_3]$ have to be adjusted to a suitable position.

In this work, compound Ba_3BPO_7 was synthesized and the structure was investigated by conventional highresolution X-ray powder diffraction. All the independent atoms in the unit cell were located through ab initio procedure and the structure was refined by Rietveld method. To solve the crystal structure from X-ray



Fig. 4. (a) Coordination geometry of Ba_2 and (b) coordination geometry of Ba_3 .

Table 6 Bond valence calculations for Ba₃BPO₇

	O1	O2	O3	$\sum v_{ij}$
Ba1	0.235	0.145	0.321	2.165
Ba2	0.021	0.286	0.018	1.834
Ba3	0.352	0.270	0.091	2.141
Р	0	1.252	1.252	5.006
В	1.014	0	0	3.041
$\sum v_{ij}$	1.857	2.240	1.900	

Table 7 Crystallographic data for Zn₃BPO₇ and Ba₃PBO₃

Identification code	Ba ₃ BPO ₇	Ba ₃ PBO ₃	Zn ₃ BPO ₇
Crystal system, space group Unit cell dimensions	Hexagonal $P6_3mc$ a = 5.4898 Å	Hexagonal $P6_3/mmc$ a = 5.502 Å	Monoclinic Cm a = 9.72 Å
unicitisions	$c = 14.7551 \text{\AA}$	c = 13.506 Å	b = 12.72 Å c = 4.87 Å $\beta = 119.8^{\circ}$
Volume Z	385.14 Å ³ 2	354.10 Å ³ 2	523.20 Å ³



Fig. 5. (a) Crystal structure of Zn_3BPO_7 and (b) crystal structure of Ba_3PBO_3 .

powder diffraction data, building the model from known structure is a simple and powerful method. But this method is not infallible. As mentioned above, Zn_3BPO_7 or Mg₃BPO₇ seems to be a very good candidate model for Ba₃BPO₇. But the fact is that Ba₃PBO₃ or Sr₃PBO₃, although has some difference with Ba₃BPO₇, can provide much more hints than Zn_3BPO_7 or Mg₃BPO₇. The ab initio method is a good approach to solve the structure when no trial structures are available.

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