The influence of the $6s^2$ configuration of Bi^{3+} on the structures of A'BiNb₂O₇ (A' = Rb, Na, Li) layered perovskite oxides.

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Figure S2. Observed, calculated and difference plots from the structural refinement of RbBiNb₂O₇ against synchrotron X-ray diffraction data collected at room temperature.

2. Structural Characterisation of LiBiNb₂O₇.

SXRD and PND data collected from LiBiNb₂O₇ at room temperature can be indexed using an orthorhombic unit cell (a = 5.456 Å, b = 5.341 Å, c = 20.819 Å) consistent with an $a' \approx \sqrt{2} \times a$, $b' \approx \sqrt{2} \times b$, $c' \approx c$ geometric expansion of the undistorted n = 2 Ruddlesden-Popper aristotype unit cell, directly analogous to the reported structure of LiNdNb₂O₇. Powder SHG measurements indicate LiBiNb₂O₇ exhibits an SHG activity ~ 2 times that of KDP (Figure 1 in the main text) and thus adopts non-centrosymmetric crystal structure. Symmetry analysis of the distortions of the n = 2 Ruddlesden-Popper framework indicates that there are four non-centrosymmetric candidate structures described in space groups $Bb2_1m$, $P2_1nm$, B2cm and P2cm respectively.

Further examination of diffraction data allowed us to eliminate the $Bb2_1m$ structure based on extinction conditions, as shown in Figure S3. Thus, structural models were constructed for LiBiNb₂O₇ based on the distortions described in space groups $P2_1nm$, B2cm and P2cm and these models were refined against the NPD data collected at room temperature. Refinement of $P2_1nm$ model results in a significantly worse fit to the data than either B2cm or P2cm models as detailed in Table S1, and so this model was eliminated.

The Li⁺ cations in cation-exchanged LiAB₂O₇ Ruddlesden-Popper phases occupy only half of the available pseudo-tetrahedral coordination sites in the framework, so can adopt either ordered or disordered arrangements. Refinement of lithium ordered and disordered models against the PND data revealed that ordered models give superior fits compared to disordered models for both the B2cm and P2cm distorted structures. The refinement parameters shown in Table S1 indicate a slightly better fit for the unconstrained P2cm model compared to the B2cm model. However, examination of the unconstrained P2cm model reveals that that it has twice as many atoms in the asymmetric unit as the B2cm model and that when refined against the NPD data, chemically equivalent atoms in the P2cm model move to very different local environments. As a result, bond valence sums (BVS) calculated for pairs of chemically equivalent cations in the BiNb₂O₇ perovskite blocks are very different (Table S2) making the resulting structure chemically implausible. To address this situation, we added constrains to the P2cm model to make the environments of the two niobium cations more similar, following the procedure adopted during the refinement of LiNdNb₂O₇.¹ This more symmetric P2cm_{constrained} model fits the data more poorly than the B2cm model as shown in Table S1, confirming that LiBiNb₂O₇ adopts and $a^-a^-c^+/-(a^-a^-c^+)$ distorted n = 2 Ruddlesden-Popper structure, described in space group B2cm analogous to that reported for LiNdNb₂O₇.



Figure S3. Synchrotron X-ray diffraction data collected from LiBiNb₂O₇. Presence of [012] and [016] peaks rules out the $Bb2_1m$ structural model. Peak marked with * is from a (Li/Rb)NbO₃ impurity.

Space group	Tilt system	Glazer tilt	#parameters	R _p (%)	_w R _p (%)
P2₁nm	$\Phi_1 \Phi_2 \Psi_z / \Phi_2 \Phi_1 \Psi_z$	a [_] b [_] c ⁺ /b [_] a [_] c ⁺	128	7.645	9.081
B2cm	ΦΦΨ _z /–(ΦΦΨ _z)	a ⁻ a ⁻ c ⁺ /–(a ⁻ a ⁻ c ⁺)	96	5.368	6.924
P2cm	$\Phi_1\Phi_2\Psi_z/-(\Phi_1\Phi_2)\Psi_z$	a ⁻ b ⁻ c ⁺ /–(a ⁻ b ⁻)c ⁺	120	4.870	5.773
P2cm _{constrained}	$\Phi_1\Phi_2\Psi_z/-(\Phi_1\Phi_2)\Psi_z$	<i>a</i> -b- <i>c</i> +/–(<i>a</i> - <i>b</i> -) <i>c</i> +	117	6.464	7.776

Table S1. Fitting statistics from the refinement of a series of structural models against the powder neutron diffraction data collected from LiBiNb₂O₇.

	P2cm	P2cm _{constrained}	B2cm
Li1	0.98	0.99	0.97
Li2	1.02	1.05	
Bi1	2.42	2.40	3.00
Bi2	3.53	3.18	
Nb1	4.55	5.03	5.03
Nb2	1.52	5.12	

Table S2. Cation bond valence sums from the *P*2*cm*, constrained *P*2*cm* and *B*2*cm* structural models of LiBiNb₂O₇.



Figure S4. Observed, calculated and difference plots from the structural refinement of LiBiNb₂O₇ against synchrotron X-ray diffraction data.

3. Structural Characterisation of NaBiNb₂O₇.

High-resolution SXRD and PND data collected from NaBiNb₂O₇ can be indexed using an orthorhombic unit cell (a = 5.47 Å, b = 5.38 Å, c = 21.67 Å) which is consistent with an $a' \approx \sqrt{2} \times a$, $b' \approx \sqrt{2} \times b$, $c' \approx c$ geometric expansion of the undistorted aristotype unit cell of an n = 2 Ruddlesden-Popper structure. The extinction conditions observed in the SXRD and NPD data allow us to eliminate the majority of n = 2 Ruddlesden-Popper phases distorted by cooperative tilting on the NbO₆ units ¹ leaving 5 candidate structures described in space groups *Pbcm*, *Pnnm*, *Pnam*, *P*₂₁*nm* and *P*₂*cm*.

Models of these distorted structures were constructed and refined against the NPD data. The model described in space group $P2_1nm$ gave the best fit to the data (Table S3). However, close inspection of the refined $P2_1nm$ model reveals that the resulting structure is very irregular, with chemically equivalent cations having very different local bonding environments, as evident from the bond valence sums of the metal cations (Table S4). To address this problem, constraints were added to the model to make the bond valence sums of chemically equivalent cations the same, resulting in a poor fit to the NPD data as noted in Table S3, so this model was discarded.

The model described in space group *Pnam* gives the best fit to the data of the remaining models as shown in Table S3, and yields a chemically reasonable structure. However, this model is centrosymmetric (incompatible with observed SHG activity), suggesting that it describes the tilting distortion of the NbO₆ units accurately, but the 'true' structure of NaBiNb₂O₇ is subject to a further distortion which breaks the inversion symmetry of this *Pnam* model. Thus we considered the inclusion of Γ -point distortion modes in the *Pnam* model with the help of the ISODISTORT software.² This symmetry analysis yields four candidate non-centrosymmetric models described in space groups *P*2₁2₁2₁, *Pn*2₁*m*, *P*2₁*am* and *Pna*2₁. Fitting statistics from the refinement of these four non-centrosymmetric models (Table S5) are comparable to one another. We therefore examined the refined structures to see if they were chemically reasonable.

Pn2₁*m* model: There are 2 sites for Na, Bi and Nb in the $Pn2_1m$ model. The Na site occupancies were refined within the constraint of total composition being NaBiNb₂O₇, to establish if the Na⁺ cations were ordered over the two sites. After refinement, the occupancy of one site reached 0.68 and the other 0.32. As the Na ordering is not significant we continued further analysis with this level of ordering. Examining the refined models, it can be seen in Table S6 that the two Bi sites have very different bond valence sum values, and the fitting statistics for this model are the poorest of the 4 considered, so this structure was discarded.

Pna2₁ model: There are 2 sites for Na and Nb and one site for Bi in this model. The Na site occupancies were refined within the constraint of total composition being NaBiNb₂O₇, to establish if the Na⁺ cations were ordered over the two sites. The occupancy of one site moves to 0.81 and the other site to 0.19, indicating the Na⁺ cations are ordered. The site with 0.81 occupancy was fixed as fully occupied which did not alter the fitting statistics significantly. In the final refined structure, we observe that the two chemically equivalent Nb sites have very different local bonding environments which result in significantly different bond valence sum values (Table S6) which make the structure chemically implausible and thus so model was discarded.

P21am model: There are two sites for Na, Nb and Bi in this model. The Na site occupancies were refined within the constraint of total composition being NaBiNb₂O₇, to establish if the Na⁺ cations were ordered over the two sites. The occupancy of one site moves to 0.88 and the other site to 0.12, indicating the Na⁺ cations are ordered. The site with 0.88 occupancy was fixed to be fully occupied which did not alter the fitting statistics significantly. Table S6 reveals that the bond valence sums of chemically equivalent Bi and Nb cations are significantly different from one another in the final refined structure. Close inspection reveals that the Bi1 and Nb1 sites belong to one Nb2O7 layer and Bi2 and Nb2 belong to another Nb2O7 layer. This allows the structure to have a very irregular local bonding environment for these chemically equivalent cations. From the Cartesian coordinates and bond valence sum values of the two Bi sites, it is clear that they are uneven and if this were the true structure we would be able to distinguish them in high-resolution X-ray powder diffraction given the strong X-ray scattering power of bismuth. Indeed, we observe that the a calculated X-ray powder diffraction pattern of this P21am structure yields calculated intensities that are not observed in the experimental X-ray powder diffraction pattern as shown in Figure S5. For these reasons, the P21am model was discarded.

*P*2₁2₁2₁ model: There are two independent sites for Na and Nb and one site for Bi in this model. The Na site occupancies were refined within the constraint of total composition being NaBiNb₂O₇, to establish if the Na⁺ cations were ordered over the two sites. The occupancy of one site moves to 0.88 and the other site to 0.12, indicating the Na⁺ cations are ordered. The site with occupancy 0.88 was fixed to be completely occupied which did not alter the fitting statistics significantly Bond valence sums of the chemically equivalent cations in the refined structure (Table S6) are similar to one another and close to their ideal value. Furthermore, the *P*2₁2₁2₁ model allows the 'checkerboard' ordering of Na, which minimizes the Na-Na repulsion as evident from the large Na-Na distance (3.77 Å). For these reasons the *P*2₁2₁2₁ model is thought to be the best description of the structure of NaBiNb₂O₇. In the final cycles of

refinement the occupancy of the Na cation site was allowed to deviate from full occupancy to yield a value of 0.81(4), without any significant changes to any other structural parameters.

It was observed that the both the SXRD and NPD data collected from NaBiNb₂O₇ exhibited strong (00*I*) preferred orientation and *hkl*-dependent peak broadening. The preferred orientation was modelled using an 4th order spherical harmonic expression,³ and the anisotropic peak broadening was modelled using an anisotropic line-shape broadening function described by Stephens ⁴. Figure S7 shows the effect of these contributions.

Space group	Tilt system	Glazer tilt	#parameters	R _p (%)	_w R _p (%)
Pbcm(#57)	$\Phi \Phi \Psi_z / - (\Phi \Phi) \Psi_z$	a ⁻ a ⁻ c ⁺ /–(a ⁻ a ⁻)c ⁺	96	7.556	8.019
Pnnm(#58)	$\Phi_1 \Phi_2 0 / \Phi_2 \Phi_1 0$	a [_] b [_] c ⁰ /b [_] a [_] c ⁰	98	7.848	8.114
<i>Pnam</i> (#62)	$ΦΦΨ_z/ΦΦ-Ψ_z$	a [_] a [_] c ⁺ /a [_] a [_] -c ⁺	98	6.366	6.635
<i>P</i> 2₁ <i>nm</i> (#31)	$\Phi_1 \Phi_2 \Psi_z / \Phi_2 \Phi_1 \Psi_z$	a [_] b [_] c ⁺ /b [_] a [_] c ⁺	118	5.840	6.127
P2 ₁ nm _{constrained}	$\Phi_1 \Phi_2 \Psi_z / \Phi_2 \Phi_1 \Psi_z$	a [_] b [_] c ⁺ /b [_] a [_] c ⁺	105	7.405	7.700
P2cm(#28)	$\Phi_1\Phi_2\Psi_z/-(\Phi_1\Phi_2)\Psi_z$	a ⁻ b ⁻ c ⁺ /-(a ⁻ b ⁻ -c ⁺	116	7.633	8.028

Table S3. Fitting statistics from the refinement of different models against the powder neutron diffraction data collected from NaBiNb₂O₇.

	P2₁nm	P2 ₁ nm _{constrained}	Pnam
Bi1	2.138	2.671	2.948
Bi2	3.722	3.184	
Nb1	4.697	5.033	4.908
Nb2	5.513	4.900	

Table S4. Cation bond valence sum calculated for the $P2_1nm$, constrained $P2_1nm$ and *Pnam* models of NaBiNb₂O₇.

Space group	Distortion	#parameters	R _p (%)	_w R _p (%)	Na ordering
	mode				pattern
<i>Pnam</i> (#62)		98	6.37	6.63	Disordered
<i>Pn</i> 2₁ <i>m</i> (#31)	Γ2 ⁻	119	6.50	5.56	Checkerboard
Pna21(#33)	Γ ₄ -	112	6.05	6.29	Stripes
<i>P</i> 2₁ <i>am</i> (#26)	Γ ₃ -	114	5.23	5.37	Stripes
<i>P</i> 2 ₁ 2 ₁ 2 ₁ (#19)	Γ ₁ -	111	6.13	6.35	Checkerboard

Table S5. Fitting statistics from the structural refinement, against NPD data, of a series of different models obtained by the addition of Γ -point distortion modes to the *Pnam* structural model of NaBiNb₂O₇.

	Pnam					
	x	У	Z	BVS		
Na	0.9984(22)	0.2568(27)	0.4908(8)	+1.037		
Bi	0.6917(6)	0.0237(7)	0.25	+2.948		
Nb	0.7538(5)	0.0139(5)	0.8583(1)	+4.908		
	•	P212121	1			
Na	0.991(2)	0.995(3)	0.5047(9)	+0.989		
Bi	0.6936(6)	0.2716(7)	0.2510(8)	+2.960		
Nb1	0.7481(12)	0.2610(20)	0.8630(4)	+4.863		
Nb2	0.2420(11)	0.2330(20)	0.1439(4)	+5.092		
	l	Pn2₁m	1	I		
Na1	0.7197(23)	0.6986(25)	0.2603(7)	+1.183		
Na2	0.4945(65)	0.5650(37)	0.6947(15)	+1.105		
Bi1	0.4891(21)	0.9247(15)	0.0000	+2.723		
Bi2	0.9216(13)	0.4949(16)	0.0000	+3.196		
Nb1	0.5152(13)	0.9474(10)	0.6117(3)	+5.112		
Nb2	0.0007(13)	0.4839(8)	0.3953(3)	+5.081		
	1	P2₁am		l		
Na	0.9761(21)	0.4855(20)	0.2502(6)	+1.010		
Bi1	0.3110(14)	0.7135(13)	0.0000	+3.650		
Bi2	0.8075(14)	0.7737(13)	0.0000	+2.481		
Nb1	0.2328(10)	0.7412(11)	0.6135(2)	+4.871		
Nb2	0.7486(9)	0.7655(11)	0.1033(2)	+5.183		
		Pna2₁	•			
Na	0.4957(25)	0.7522(31)	0.0026(10)	+0.859		
Bi	0.6870(6)	0.9816(7)	0.7621(6)	+2.950		
Nb1	0.7553(17)	0.9794(10)	0.1470(2)	+5.472		
Nb2	0.2549(18)	0.5168(14)	0.3651(2)	+4.683		

Table S6. Fractional coordinates and cation bond valence sums of from the structural refinement of a series of distorted non-centrosymmetric models against the powder neutron diffraction data collected from NaBiNb₂O₇.



Figure S5. Observed, calculated and difference plots from the structural refinement of *Pnam*, *P*2₁*am* and *P*2₁2₁2₁ symmetry models against synchrotron X-ray diffraction data collected from NaBiNb₂O₇, showing the poor fits obtained by the *Pnam*, *P*2₁*am* models.



Figure S6. Observed, calculated and difference plots from the structural refinement of a *P*2₁2₁2₁ model against synchrotron X-ray diffraction data collected from NaBiNb₂O₇.



Figure S7. Fit to NPD data collected from NaBiNb₂O₇ (top) without accounting for preferred orientation or anisotropic peak broadening; (middle) accounting for anisotropic broadening only; (bottom) accounting for both anisotropic peak broadening and preferred orientation.

4. Microstructural Characterisation of NaBiNb₂O₇.



Figure S8. HAADF-STEM image collected from NaBiNb₂O₇. Green brackets outline the thin slabs of a Dion-Jacobson phase with [010] orientation. Pink bracket [001] or [101] zone of BiNbO₄ (Pnna, a = 5.68 Å, b = 11.72 Å, c = 4.98 Å). The models are inserted: green – Bi, orange – Nb, blue – Na.

5. Distortion modes of A'ANb₂O₇ phases.

The structures of the *A*'*A*Nb₂O₇ phases can be analysed in terms the of distortion modes which need to be applied to the undistorted aristotype structures to obtain the observed experimental structures. The atom displacements, which occur on going from the aristotype structure to the distorted structure, can be deconvoluted into contributions from the individual distortion modes.² The data tabulated below reports the amplitudes of the displacements due to the distortion modes in three ways:

A_s : the supercell normalised amplitude, which is the square root of the sum of the squares of the mode-induced displacements.

 A_p : the parent cell normalised amplitude. $A_p = A_s * \sqrt{(V_p/V_s)}$, where V_p and V_s are the volumes of the parent cell and supercell respectively.

d_{max} is the maximum displacement experienced by an atom due to this distortion mode.

RbNdNb2O7						
Distortion mode	As	Ap	d _{max} (Å)			
[0,0,0] Γ ₁ + [Nb1:h:dsp] A1(a)	0.06720	0.04752	0.03360			
[0,0,0] Γ ₁ + [O1:i:dsp] A1(a)	0.24133	0.17065	0.08532			
[0,0,0] Γ ₁ + [O2:h:dsp] A1(a)	0.11508	0.08137	0.05754			
[0,0,0] Г₁⁺ all	0.27568	0.19494				
[0,0,0] Г₅⁻ [Rb1:b:dsp] Eu(a)	0.0000	0.0000	0.0000			
[0,0,0] Г₅⁻ [Nd1:a:dsp] Eu(a)	0.01537	0.01087	0.01087			
[0,0,0] Г₅⁻ [Nb1:h:dsp] E(a)	-0.07282	-0.05149	0.03641			
[0,0,0] Г₅⁻ [O1:i:dsp] B2(a)	0.53204	0.37621	0.18810			
[0,0,0] Г₅⁻ [O1:i:dsp] B1(a)	0.12119	0.08569	0.04285			
[0,0,0] Г₅⁻ [O2:h:dsp] E(a)	-0.08043	-0.05687	0.04022			
[0,0,0] Г₅ [–] [O3:c:dsp] Eu(a)	-0.25286	-0.17880	0.17880			
[0,0,0] Г₅⁻all	0.61131	0.43226				
[1/2,1/2,1/2] A2+ [O1:i:dsp] B2(a)	-0.72225	-0.51071	0.25535			
$[\frac{1}{2},\frac{1}{2},\frac{1}{2}]$ A ₂ + all	0.72225	0.51071				
[1/2,1/2,1/2] A3+ [O1:i:dsp] B1(a)	-0.10054	-0.07109	0.03555			
$[\frac{1}{2},\frac{1}{2},\frac{1}{2}]$ A ₃ ⁺ all	0.10054	0.07109				
[1/2,1/2,1/2] A5 ⁻ [Nd1:a:dsp] Eu(a)	-0.00769	-0.00544	0.00544			
[½,½,½] A ₅ - [Nb1:h:dsp] E(a)	0.00978	0.00692	0.00489			
[1/2,1/2,1/2] A5- [O1:i:dsp] A1(a)	-0.76748	-0.54269	0.27135			
[½,½,½] A5 ⁻ [O2:h:dsp] E(a)	-0.57280	-0.40503	0.28640			
[1/2,1/2,1/2] A5- [O3:c:dsp] Eu(a)	0.56182	0.39727	0.39727			
$[\frac{1}{2},\frac{1}{2},\frac{1}{2}]$ A ₅ ⁻ all	1.1104	0.78515				

Table S7. Distortion mode amplitudes extracted from the refined structure of RbNdNb₂O₇ (space group *I*2*cm*)⁵ compared to the *P*4/*mmm* symmetry aristotype structure.

RbBiNb ₂ O ₇						
Distortion mode	As	Ap	d _{max} (Å)			
[0,0,0] Γ ₁ + [Nb1:h:dsp] A1(a)	0.25562	0.18075	0.12781			
[0,0,0] Γ ₁ + [O1:i:dsp] A1(a)	0.31711	0.22423	0.11212			
[0,0,0] Γ ₁ + [O2:h:dsp] A1(a)	0.19732	0.13953	0.09866			
[0,0,0] Γ ₁ + all	0.45259	0.32003				
[0,0,0] Г₅⁻ [Rb1:b:dsp] Eu(a)	0.0000	0.0000	0.0000			
[0,0,0] Γ₅⁻ [Bi1:a:dsp] Eu(a)	-0.29270	-0.20697	0.20697			
[0,0,0] Γ₅⁻ [Nb1:h:dsp] E(a)	0.11920	0.08429	0.05960			
[0,0,0] Г₅⁻ [O1:i:dsp] B2(a)	0.62795	0.44403	0.22201			
[0,0,0] Г₅⁻ [O1:i:dsp] B1(a)	0.22485	0.15899	0.07950			
[0,0,0] Г₅⁻ [O2:h:dsp] E(a)	0.12570	0.08888	0.06285			
[0,0,0] Г₅⁻ [O3:c:dsp] Eu(a)	0.34480	0.24381	0.24381			
[0,0,0] Г₅⁻all	0.82429	0.58286				
[½,½,½] A ₂ + [O1:i:dsp] B2(a)	-0.73305	-0.51834	0.25917			
$[\frac{1}{2},\frac{1}{2},\frac{1}{2}]$ A ₂ + all	0.73305	0.51834				
[½,½,½] A ₃ + [O1:i:dsp] B1(a)	-0.04605	-0.03256	0.01628			
$[\frac{1}{2},\frac{1}{2},\frac{1}{2}]$ A ₃ + all	0.04605	0.03256				
[½,½,½] A5 ⁻ [Bi1:a:dsp] Eu(a)	0.12872	0.09102	0.09102			
[½,½,½] A ₅ - [Nb1:h:dsp] E(a)	-0.05201	-0.03678	0.02601			
[½,½,½] A5 ⁻ [O1:i:dsp] A1(a)	-0.45980	-0.32513	0.16256			
[½,½,½] A5 ⁻ [O2:h:dsp] E(a)	-0.33483	-0.23676	0.16742			
[½,½,½] A ₅ - [O3:c:dsp] Eu(a)	0.53406	0.37764	0.37764			
[½,½,½] A₅- all	0.79248	0.56037				

Table S8. Distortion mode amplitudes extracted from the refined structure of RbBiNb₂O₇ (space group *l*2*cm*) compared to the *P*4/*mmm* symmetry aristotype structure.

LiNdNb2O7						
Distortion mode	As	Ap	d _{max} (Å)			
[0,0,0] Γ1+ [Nb1:e:dsp] A1(a)	-0.44990	-0.31813	0.22495			
[0,0,0] Γ ₁ + [O2:g:dsp] A1(a)	-0.46851	-0.33129	0.16564			
[0,0,0] Γ ₁ + [O3:e:dsp] A1(a)	0.67894	0.48008	0.33947			
[0,0,0] Г₁⁺ all	0.93961	0.66441				
[0,0,0] Г₅⁻ [Li1:d:dsp] E(a)	0	0	0			
[0,0,0] Г₅⁻ [Nd1:a:dsp] Eu(a)	0.03605	0.02549	0.02549			
[0,0,0] Г₅⁻ [Nb1:e:dsp] E(a)	0.24514	0.17334	0.12257			
[0,0,0] Г₅⁻ [O1:b:dsp] Eu(a)	0.21859	0.15457	0.15457			
[0,0,0] Г₅ [–] [O2:g:dsp] B2(a)	0.70126	0.49587	0.24793			
[0,0,0] Г₅ [–] [O2:g:dsp] B1(a)	0.38670	0.27344	0.13672			
[0,0,0] Γ ₅ - [O3:e:dsp] E(a)	0.20718	0.14650	0.10359			
[0,0,0] Г₅ [–] all	0.89073	0.62984				
[½,½,0] X ₂ + [Li1:d:dsp] E(a)	0.44473	0.31447	0.22236			
[½,½,0] X ₂ + [O2:g:dsp] B2(a)	-0.66438	-0.46979	0.23489			
[½,½,0] X ₂ + [O2:g:dsp] B1(a)	0.01573	0.01112	0.00556			
[½,½,0] X ₂ + all	0.79965	0.56543				
[½,½,0] X ₄ - [Nd1:a:dsp] Eu(a)	0.01534	0.01085	0.01085			
[½,½,0] X ₄ - [Nb1:e:dsp] E(a)	-0.01952	-0.01380	0.00976			
[½,½,0] X ₄ - [O1:b:dsp] Eu(a)	0.52233	0.36934	0.36934			
[½,½,0] X ₄ - [O2:g:dsp] A1(a)	0.79821	0.56442	0.28221			
[½,½,0] X ₄ - [O3:e:dsp] E(a)	-0.96105	-0.67956	0.48052			
[½,½,0] X₄⁻ all	1.3543	0.95765				

Table S9. Distortion mode amplitudes extracted from the refined structure of LiNdNb₂O₇ (space group B2cm)¹ compared to the *I*4/*mmm* symmetry aristotype structure.

LiBiNb ₂ O ₇					
Distortion mode	As	Ap	d _{max} (Å)		
[0,0,0] Γ ₁ + [Nb1:e:dsp] A1(a)	-0.63705	-0.45046	0.31853		
[0,0,0] Γ ₁ + [O2:g:dsp] A1(a)	-0.41219	-0.29146	0.14573		
[0,0,0] Γ ₁ + [O3:e:dsp] A1(a)	0.56210	0.39746	0.28105		
[0,0,0] Γ ₁ + all	0.94429	0.66772			
[0,0,0] Γ₅⁻ [Li1:d:dsp] E(a)	0.0000	0.0000	0.0000		
[0,0,0] Γ₅⁻ [Bi1:a:dsp] Eu(a)	-0.46107	-0.32603	0.32603		
[0,0,0] Γ₅⁻ [Nb1:e:dsp] E(a)	-0.06753	-0.04775	0.03377		
[0,0,0] Γ₅⁻ [O1:b:dsp] Eu(a)	0.01795	0.01269	0.01269		
[0,0,0] Γ₅⁻ [O2:g:dsp] B2(a)	0.42192	0.29834	0.14917		
[0,0,0] Γ₅⁻ [O2:g:dsp] B1(a)	-0.00486	-0.00344	0.00172		
[0,0,0] Γ₅⁻ [O3:e:dsp] E(a)	-0.14640	-0.10352	0.07320		
[0,0,0] Γ₅⁻ all	0.64571	0.45659			
[½,½,0] X ₂ + [Li1:d:dsp] E(a)	-0.00378	-0.00267	0.00189		
[½,½,0] X ₂ + [O2:g:dsp] B2(a)	-0.67097	-0.47445	0.23722		
[1/2,1/2,0] X2+ [O2:g:dsp] B1(a)	0.00216	0.00153	0.00076		
[½,½,0] X ₂ + all	0.67098	0.47446			
[1/2,1/2,0] X4- [Bi1:a:dsp] Eu(a)	0.11613	0.08212	0.08212		
[½,½,0] X4 ⁻ [Nb1:e:dsp] E(a)	-0.04970	-0.03514	0.02485		
[1/2,1/2,0] X4- [O1:b:dsp] Eu(a)	0.54320	0.38410	0.38410		
[1/2,1/2,0] X4- [O2:g:dsp] A1(a)	0.60062	0.42470	0.21235		
[½,½,0] X ₄ - [O3:e:dsp] E(a)	-0.81899	-0.57911	0.40950		
[1/2,1/2,0] X4 ⁻ all	1.1587	0.81930			

Table S10. Distortion mode amplitudes extracted from the refined structure of LiBiNb₂O₇ (space group *B*2*cm*) compared to the *I*4/*mmm* symmetry aristotype structure.

NaBiNb2O7						
Distortion mode	As	Ap	d _{max} (Å)			
[0,0,0] Γ ₁ + [Nb1:e:dsp] A1(a)	-0.58470	-0.29235	0.20672			
[0,0,0] Γ ₁ + [O2:g:dsp] A1(a)	-0.83339	-0.41669	0.20835			
[0,0,0] Γ ₁ + [O3:e:dsp] A1(a)	1.1725	0.58623	0.41453			
[0.0.0] F ₁ ⁺ all	1.5528	0.77638				
[0,0,0] Γ ₄ -[O2:g:dsp] A1(a)	-0.34851	-0.17425	0.087130			
[0,0,0] Г₄⁻ all	0.34851	0.17425				
[½,½,0] X ₂ + [Na1:d:dsp] E(a)	-0.06896	-0.03448	0.02438			
$[\frac{1}{2},\frac{1}{2},0]$ X ₂ + [O2:g:dsp] B2(a)	-0.85046	-0.42523	0.21262			
[½,½,0] X ₂ + [O2:g:dsp] B1(a)	0.07662	0.03831	0.01916			
$[\frac{1}{2},\frac{1}{2},0]$ X ₂ ⁺ all	0.85668	0.42834				
[1/1/0] X ₂ + [No1:d:dop] P2(o)	0 1 1 2 9 9	0.07104	0.05097			
$[72,72,0] \land 3^{\circ}$ [Na1.0.0Sp] B2(a)	0.14300	0.07194	0.00007			
$[\frac{7}{2},\frac{7}{2},0] \times [1001.0.05p] E(a)$	0.07565	0.03793	0.02002			
$[\frac{1}{2},\frac{1}{2},0] \times [02.9,05p] \times [02.9,05p]$	0.09308	0.04034	0.02327			
[72,72,0] A3 [O3.0.050] E(a)	0.30470	0.10233	0.12094			
	0.41003	0.20502				
[½ ½ 0] X₂⁻ [Na1:d:dsp] F(a)	0.06896	0.03448	0.02438			
$[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$ [Bi1:a:dsp] A2u(a)	0.04329	0.02165	0.02165			
$[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$ [2] [2] [2] [2] [2] [2] [2] [2] [2] [2]	0 21123	0 10562	0.07468			
$[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$ $[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$ $[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$	0.02165	0.01082	0.01083			
$[\frac{1}{2}, \frac{1}{2}, \frac{1}{2}, \frac{1}{2}]$ X_2^- [O2: g:dsp] B2(a)	0.26816	0 13408	0.06704			
[%,%,0] X ₂ -[O3:e:dsp] A1(a)	0.17449	0.08725	0.06169			
[½,½,0] X ₂ - all	0.39252	0.19626	0.00100			
[1/2,1/2,0] X3 ⁻ [Na1:d:dsp] B2(a)	-0.14388	-0.07194	0.05087			
[½,½,0] X ₃ - [Bi1:a:dsp] Eu(a)	-0.23405	-0.11703	0.11703			
[½,½,0] X ₃ ⁻ [Nb1:e:dsp] E(a)	-0.21453	-0.10726	0.07585			
[½,½,0] X ₃ ⁻ [O1:b:dsp] Eu(a)	0.77148	0.38574	0.38574			
[½,½,0] X ₃ ⁻ [O2:g:dsp] A1(a)	0.47406	0.23703	0.11852			
[½,½,0] X ₃ ⁻ [O3:e:dsp] E(a)	-0.56697	-0.28348	0.20045			
[½,½,0] X ₃ - all	1.1238	0.56189				
	0.00004	0.04045	0.04054			
[1,1,1] M5 ⁺ [Na1:d:dsp] E(a)	-0.03831	-0.01915	0.01354			
[1,1,1] M ₅ ⁺ [Nb1:e:dsp] E(a)	0.04597	0.02298	0.01625			
[1,1,1] M ₅ + [O2:g:dsp] B2(a)	0.02299	0.01150	0.00575			
[1,1,1] M₅+ [O2:g:dsp] B1(a)	0.06129	0.03064	0.01532			
[1,1,1] M5 ⁺ [U3:e:dsp] E(a)	0.12259	0.06130	0.04334			
[1,1,1] M₅⁺ all	0.15131	0.07565				
[1 1 1] Mr-[Na1:d:den] F(2)	-0 03831	-0 01015	0.01354			
[1, 1, 1] [No [Nat.u.usp] $L(d)$	-0.61112	-0.30556	0.01554			
[1,1,1] Ms [D1.a.usp] Lu(d)	0.01112	0.00000	0.00000			
[1, 1, 1] Ms [[101.6.030] $L(a)$	-0 13003	-0.02007	0.06502			
[1, 1, 1] Ms [01.0.059] Ed(a)	-0.62061	-0 3103	0.15515			
[1, 1, 1] Ms [02.9.039] $D2(a)$	0.02001	0.02298	0.01149			
[1, 1, 1] M ₅ ⁻ [O3:e:dsp] E(a)	-0.05057	-0.02528	0.01788			
[1,1,1] M ₅ - all	0.88536	0.44268	0.01700			
	0.00000	0				

Table S11. Distortion mode amplitudes extracted from the refined structure of NaBiNb₂O₇ (space group $P_{2_12_12_1}$) compared to the *I*4/*mmm* symmetry aristotype structure.

	Lattice parameters			Disp	lacive symm	etry adapte	d mode mag	ntides	
	a (Å)	b (Å)	c (Å)	Vol(Å ³)	Γ ₅ -	A ₅ -	A ₂ +	A ₃ +	Γ_1^+
Parent (supercell)	22.311	5.450	5.450	662.750	-	-	-	-	-
VASP	22.490	5.360	5.432	654.836	0.81287	0.88286	1.07549	0.00921	0.32531
Experiment	22.422	5.382	5.452	657.984	0.82918	0.79429	0.73741	0.04633	0.40586

6. First-principles analysis of RbBiNb₂O₇ and LiBiNb₂O₇.

Table S12. Comparison between theoretical and experimentally synthesizedstructures of RbBiNb2O7 in the polar *I2cm* phase.

Space group	Source	a (Å)	b (Å)	<i>c</i> (Å)	Volume (ų)
Bmcm	PBEsol	5.5665	5.5620	20.0563	620.9755
B2cm	Experiment	5.4560	5.3415	20.8186	606.7211
	PBEsol	5.4456	5.3314	20.8069	604.0954

Table S13. Comparison between structures of LiBiNb₂O₇ in the listed space groups from our first-principles calculations (PBEsol) and experiment.

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