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Electronic supplementary information

Crystal structure, lattice dynamics and superexchange in MAgF₃ 1D antiferromagnets (M = K, Rb, Cs) and Rb₃Ag₂F₇ Ruddlesden-Popper phase

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Figure S1. Deconvoluted Raman spectrum of CsAgF₃.



Figure S2. Deconvoluted Raman spectrum of RbAgF₃.



Figure S3. Different Raman spectra collected for of RbAgF₃.



Figure S4. Full set of Raman spectra at low temperature for RbAgF3.



Figure S5. X-ray diffraction pattern of $RbAgF_3$ sample containing novel $Rb_3Ag_2F_7$ phase and the results of the Rietveld refinement. Bottom part shows simulated Bragg positions for each present phase (from top to bottom: $RbAgF_3$, AgF and $Rb_3Ag_2F_7$).

| Formula | RbAgF ₃ | | |
|---------------------|--|--|--|
| Colour | brown | | |
| Space group | <i>I4/mcm</i> (No. 140) | | |
| Z | 4 | | |
| V /Å ³ | 338.991(6) | | |
| a /Å | 6.33855(6) | | |
| b /Å | 6.33855(6) | | |
| c /Å | 8.43740(11) | | |
| α /° | 90 | | |
| β /° | 90 | | |
| γ /° | 90 | | |
| Positions | Rb1 0.00000 0.00000 0.25000 Ag1 0.00000 0.50000 0.00000 F1 0.00000 0.50000 0.25000 F2 0.240(4) 0.740(4) 0.00000 | | |
| Reliability factors | Not corrected for background (all data) $R_p = 0.84 \%$ $R_{wp} = 1.78 \%$ Corrected for background (all data) $cR_p = 27.86 \%$ $cR_{wp} = 15.16 \%$ R Bragg factor (this phase) $R_{Bragg} = 9.53 \%$ | | |
| Atomic distances /Å | Ag-F (apical) 2.10935(3) Ag-F (equatorial) 2.33(3) 2.15(3) | | |

Table S1. Crystallographic data for RbAgF₃.

| RbAgF ₃ Pnma (HSE06) | | RbAgF ₃ <i>I</i> 4/ <i>mcm</i> (HSE06) | | CsAgF ₃ <i>I</i> 4/ <i>mcm</i> (HSE06) | |
|---------------------------------|-----|---|-----|---|-----|
| B _{2u} | 471 | A _{2u} | 484 | Eu | 450 |
| Au | 470 | B _{2g} | 472 | B ₂ | 449 |
| B ₃ g | 467 | E _u | 467 | A20 | 449 |
| B ₁ , | 464 | Ala | 381 | Ala | 370 |
| Ba | 464 | B ₂ | 312 | R ₂ | 302 |
| | 467 | B _{2g} | 233 | B _{2g} | 227 |
| A _u | 402 | | 184 | D _{lu} | 102 |
| D _{2u} | 401 | L _g | 104 | A _{2u} | 193 |
| B _{2g} | 449 | A _{2u} | 105 | E _u | 186 |
| B _{1g} | 414 | E _u | 181 | E _g | 186 |
| Ag | 380 | Eu | 159 | Eu | 162 |
| B _{1g} | 365 | A _{2g} | 144 | B _{1u} | 155 |
| B _{3g} | 303 | B _{1u} | 139 | A _{2g} | 150 |
| B _{3u} | 271 | Eu | 124 | Eu | 118 |
| B _{1u} | 240 | A _{2g} | 69 | B _{1g} | 95 |
| B_{3u} | 205 | A _{2u} | 67 | Eg | 88 |
| A_{g} | 201 | Eu | 66 | A_{2u} | 72 |
| B_{2g} | 200 | Eg | 65 | Eu | 72 |
| B _{1u} | 199 | Eg | 26 | A _{2g} | 65 |
| B _{2u} | 186 | A _{2u} | -1 | Eg | 56 |
| B _{2g} | 182 | Eu | -2 | A _{2v} | -5 |
| B _{1u} | 175 | B _{1g} | -39 | En | -5 |
| B ₃ , | 174 | | | ч | - |
| A | 174 | | | | |
| A | 164 | | | | |
| Ba | 167 | | | | |
| Ba | 161 | | | | |
| B. | 157 | | | | |
| D _{1u} | 157 | | | | |
| D _{3g} | 133 | | | | |
| | 140 | | | | |
| Au | 140 | | | | |
| B _{1g} | 139 | | | | |
| B _{1u} | 139 | | | | |
| B _{3g} | 134 | | | | |
| B _{3u} | 134 | | | | |
| B _{3u} | 129 | | | | |
| B _{2g} | 125 | | | | |
| B _{1u} | 118 | | | | |
| Ag | 98 | | | | |
| Ag | 87 | | | | |
| B _{1g} | 86 | | | | |
| B _{1u} | 82 | | | | |
| Ag | 78 | | | | |
| B _{3g} | 77 | | | | |
| A _u | 76 | | | | |
| B _{3u} | 76 | | | | |
| B ₂₀ | 74 | | | | |
| B ₂ _σ | 73 | | | | |
| B ₂₁₁ | 71 | | | | |
| Ag | 69 | | | | |
| B ₂ | 68 | | | | |
| B ₁ | 65 | | | | |
| Δ | 64 | | | | |
| R. | 60 | | | | |
| D _{2g} | 50 | | | | |
| D _{1g} | 30 | | | | |
| A _u | 49 | | | | |
| B _{2g} | 34 | | | | |
| | 33 | | | | |
| B _{2u} | -1 | | | | |
| $ $ B_{1u} | -1 | | | | 1 |

Table S2. Theoretically calculated modes.

Cif files of structures used in lattice dynamics calculations

RbAgF₃ Pnma (HSE06) data findsym-output audit creation method FINDSYM _cell_length_a 6.2633870000 _cell_length_b 8.3574660000 cell length c 6.3444310000 cell angle alpha 90.000000000 cell angle beta 90.0000000000 cell angle gamma 90.0000000000 _cell_volume 332.1058636297 _symmetry_space_group_name_H-M "P 21/n 21/m 21/a" _space_group.reference_setting '062:-P 2ac 2n' space group.transform Pp abc a,b,c;0,0,0 loop _space_group_symop_id _space_group_symop_operation_xyz 1 x,y,z 2 x+1/2,-y+1/2,-z+1/2 3 -x,y+1/2,-z 4 -x+1/2,-y,z+1/2 5 -x,-y,-z 6 -x+1/2,y+1/2,z+1/2 7 x,-y+1/2,z 8 x+1/2,y,-z+1/2 loop _atom_site_label _atom_site_type_symbol atom_site_symmetry_multiplicity atom_site_Wyckoff_label atom site fract x _atom_site_fract_y _atom_site_fract_z _atom_site_occupancy atom_site_fract_symmform Ag1 Ag 4 a 0.0000 0.00000 0.00000 1.00000 0.0.0 Rb1 Rb 4 c 0.00381 0.25000 0.49913 1.00000 Dx,0,Dz F1 F 4 c 0.49600 0.25000 0.45836 1.00000 Dx,0,Dz F2 F 8 d 0.26645 -0.02118 0.27814 1.00000 Dx,Dy,Dz

RbAgF₃ I4/mcm (HSE06)

data_findsym-output _audit_creation_method FINDSYM

_cell_length_a 6.2963000000 _cell_length_b 6.2963000000 _cell_length_c 8.4031760000 _cell_angle_alpha 90.0000000000 _cell_angle_beta 90.0000000000 _cell_angle_gamma 90.0000000000 _cell_volume 333.1304144144

_symmetry_space_group_name_H-M "I 4/m 2/c 2/m" _symmetry_Int_Tables_number 140 _space_group.reference_setting '140:-I 4 2c' _space_group.transform_Pp_abc a,b,c;0,0,0

loop_ _space_group_symop_id _space_group_symop_operation_xyz 1 x,y,z 2 x,-y,-z+1/2 3 -x,y,-z+1/2 4 -x,-y,z 5 -y,-x,-z+1/2 6 -y,x,z 7 y,-x,z 8 y,x,-z+1/2 9 -x,-y,-z 10 - x, y, z + 1/211 x,-y,z+1/2 12 x,y,-z 13 y,x,z+1/2 14 y,-x,-z 15 -y,x,-z 16 - y, -x, z + 1/217 x+1/2,y+1/2,z+1/2 18 x+1/2,-y+1/2,-z 19 -x+1/2,y+1/2,-z 20 -x+1/2,-y+1/2,z+1/2 21 -y+1/2,-x+1/2,-z 22 -y+1/2,x+1/2,z+1/2 23 y+1/2,-x+1/2,z+1/2 24 y+1/2,x+1/2,-z 25 -x+1/2,-y+1/2,-z+1/2 26 -x+1/2,y+1/2,z 27 x+1/2,-y+1/2,z 28 x+1/2,y+1/2,-z+1/2 29 y+1/2,x+1/2,z 30 y+1/2,-x+1/2,-z+1/2 31 -y+1/2,x+1/2,-z+1/2 32 -y+1/2,-x+1/2,z

loop_ _atom_site_label _atom_site_type_symbol _atom_site_symmetry_multiplicity _atom_site_symmetry_multiplicity _atom_site_fract_x _atom_site_fract_y _atom_site_fract_z _atom_site_occupancy _atom_site_fract_symmform Ag1 Ag 4 d 0.00000 0.50000 0.00000 1.00000 0,0,0 Rb1 Rb 4 a 0.00000 0.50000 0.25000 1.00000 0,0,0 F1 F 4 b 0.00000 0.50000 0.25000 1.00000 0,0,0 F2 F 8 h 0.22870 0.72870 0.00000 1.00000 Dx,Dx,0

CsAgF₃ I4/mcm (HSE06)

data_findsym-output _audit_creation_method FINDSYM

_cell_length_a 6.4411370000 _cell_length_b 6.4411370000 _cell_length_c 8.4690630000 _cell_angle_alpha 90.0000000000 _cell_angle_beta 90.0000000000 _cell_angle_gamma 90.0000000000 _cell_volume 351.3665678866

_symmetry_space_group_name_H-M "I 4/m 2/c 2/m" _symmetry_Int_Tables_number 140 _space_group.reference_setting '140:-I 4 2c' _space_group.transform_Pp_abc a,b,c;0,0,0

loop_ _space_group_symop_id _space_group_symop_operation_xyz 1 x,y,z 2 x,-y,-z+1/2 3 -x,y,-z+1/2 4 -x,-y,z 5 -y,-x,-z+1/2 6 -y,x,z 7 y,-x,z 8 y,x,-z+1/2 9 -x,-y,-z 10 - x, y, z + 1/211 x,-y,z+1/2 12 x,y,-z 13 y,x,z+1/2 14 y,-x,-z 15 -y,x,-z 16 -y,-x,z+1/2 17 x+1/2,y+1/2,z+1/2 18 x+1/2,-y+1/2,-z 19 -x+1/2,y+1/2,-z 20 -x+1/2,-y+1/2,z+1/2 21 -y+1/2,-x+1/2,-z 22 -y+1/2,x+1/2,z+1/2 23 y+1/2,-x+1/2,z+1/2 24 y+1/2,x+1/2,-z 25 -x+1/2,-y+1/2,-z+1/2 26 -x+1/2,y+1/2,z 27 x+1/2,-y+1/2,z 28 x+1/2,y+1/2,-z+1/2 29 y+1/2,x+1/2,z 30 y+1/2,-x+1/2,-z+1/2 31 -y+1/2,x+1/2,-z+1/2 32 -y+1/2,-x+1/2,z

loop_

_atom_site_label _atom_site_type_symbol _atom_site_symmetry_multiplicity _atom_site_fract_x _atom_site_fract_y _atom_site_fract_z _atom_site_fract_z _atom_site_fract_symmform Ag1 Ag 4 d 0.00000 0.50000 0.00000 1.00000 0,0,0 Cs1 Cs 4 a 0.00000 0.50000 0.25000 1.00000 0,0,0 F1 F 4 b 0.00000 0.50000 0.25000 1.00000 0,0,0 F2 F 8 h 0.22363 0.72363 0.00000 1.00000 Dx,Dx,0

Rb₃Ag₂F₇ Cmca (GGA+U)

data_findsym-output _audit_creation_method FINDSYM

_cell_length_a 21.6756500000 _cell_length_b 6.2943600000 _cell_length_c 6.2937100000 _cell_angle_alpha 90.00000000000 _cell_angle_beta 90.0000000000 _cell_angle_gamma 90.0000000000 _cell_volume 858.6781972783

_symmetry_space_group_name_H-M "C 2/m 2/c 21/a" _symmetry_Int_Tables_number 64 _space_group.reference_setting '064:-C 2ac 2' _space_group.transform_Pp_abc a,b,c;0,0,0

loop_

_space_group_symop_id _space_group_symop_operation_xyz 1 x,y,z 2 x,-y,-z 3 -x+1/2,y,-z+1/2 4 -x+1/2,-y,z+1/2 5 -x,-y,-z 6 -x,y,z 7 x+1/2,-y,z+1/2 8 x+1/2,y,-z+1/2 9 x+1/2,y+1/2,z 10 x+1/2,-y+1/2,-z 11 -x,y+1/2,-z+1/2 12 -x,-y+1/2,z+1/2 13 -x+1/2,-y+1/2,-z 14 -x+1/2,y+1/2,z 15 x,-y+1/2,z+1/2 16 x,y+1/2,-z+1/2

loop_

_atom_site_label _atom_site_type_symbol _atom_site_symmetry_multiplicity _atom_site_Wyckoff_label _atom_site_fract_x _atom_site_fract_y _atom_site_fract_z _atom_site_occupancy _atom_site_fract_symmform Ag1 Ag 8 d 0.40284 0.00000 0.00000 1.00000 Dx,0,0 Rb1 Rb 8 d 0.18085 0.00000 0.00000 1.00000 Dx,0,0 Rb2 Rb 4 a 0.00000 0.00000 0.00000 1.00000 Dx,0,0 F1 F 8 d 0.30625 0.00000 0.00000 1.00000 Dx,0,0 F2 F 4 b 0.50000 0.00000 0.00000 1.00000 Dx,0,0 F3 F 16 g 0.09527 0.26940 0.23077 1.00000 Dx,Dy,Dz