

**Electronic Supplementary Information**

**Phase Behaviour and Composition in the Formamidinium-Methylammonium Hybrid Lead Iodide  
Perovskite Solid Solution**

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**Table S1** Reactant quantities used for crystal synthesis via precipitation from hydroiodic acid (method HI)

Sample	Ratios		Weights (g)			Volume HI (ml)
	MA	FA	MAI	FAI	Pb(CH <sub>3</sub> CO <sub>2</sub> ) <sub>2</sub> · 3H <sub>2</sub> O	
<b>1</b>	1	0	0.2564	0	0.6119	1.6130
<b>2</b>	0.9	0.1	0.2303	0.0277	0.6106	1.6096
<b>3</b>	0.8	0.2	0.2043	0.0552	0.6093	1.6062
<b>4</b>	0.7	0.3	0.1784	0.0827	0.6080	1.6029
<b>5</b>	0.6	0.4	0.1526	0.1100	0.6068	1.5996
<b>6</b>	0.5	0.5	0.1269	0.1373	0.6055	1.5962
<b>7</b>	0.4	0.6	0.1013	0.1644	0.6043	1.5929
<b>8</b>	0.3	0.7	0.0758	0.1914	0.6030	1.5896
<b>9</b>	0.2	0.8	0.0504	0.2182	0.6018	1.5864
<b>10</b>	0.1	0.9	0.0252	0.2450	0.6005	1.5831
<b>11</b>	0	1	0	0.2717	0.5993	1.5798

**Table S2** Reactant quantities used for crystal synthesis via Inverse solubility (method IS)

0.001 mol		1 M solutions in 1 mL $\gamma$ -butyrolactone								
Ratio MA/FA	9:1	8:2	7:3	6:4	5:5	4:6	3:7	2:8	1:9	
	wgt / g									
<b>MAI</b>	0.1431	0.1272	0.1113	0.0954	0.0795	0.0636	0.0477	0.0318	0.0159	
<b>FAI</b>	0.0172	0.0344	0.0516	0.0688	0.0860	0.1032	0.1204	0.1376	0.1548	
<b>PbI<sub>2</sub></b>	0.4610	0.4610	0.4610	0.4610	0.4610	0.4610	0.4610	0.4610	0.4610	

**Table S3** Lattice Parameters and Cell Volumes at 300K of FA<sub>x</sub>MA<sub>1-x</sub>PbI<sub>3</sub> crystals grown via inverse solubility (IS) method.

x in FA <sub>x</sub> MA <sub>1-x</sub> PbI <sub>3</sub>	Reduced lattice parameter / Å	Cell Volume / Å <sup>3</sup>
0.1	6.3047(15)	249.946(14)
0.2	6.3102(15)	251.267(1)
0.3	6.3122(2)	251.501(17)
0.4	6.3199(3)	252.42(2)
0.5	6.3319(15)	253.859(1)
0.6	6.3333(3)	254.03(2)
0.7	6.3479(2)	255.795(18)
0.8	6.3543(2)	256.573(2)
0.9	6.3486(2)	255.878(15)

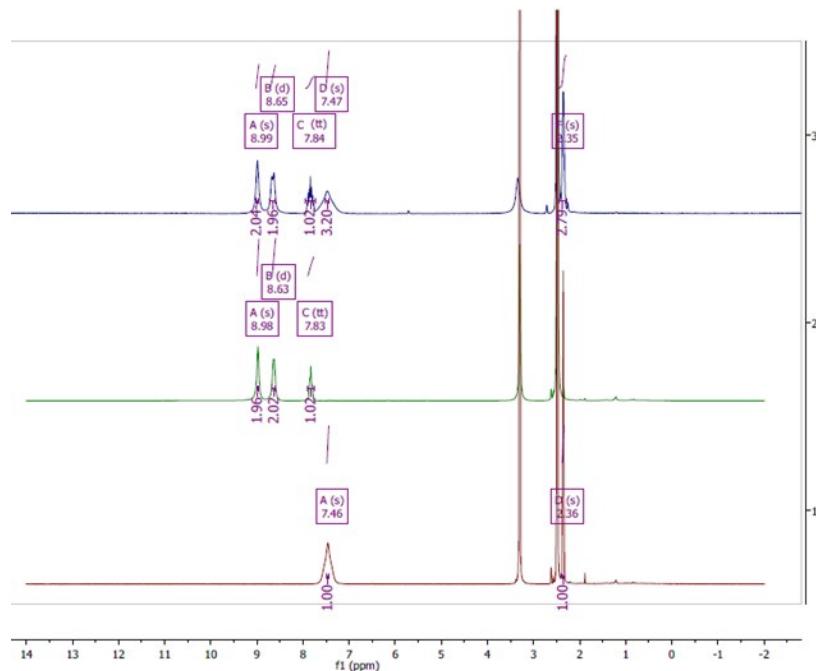


Figure 1  $^1\text{H}$  NMR in  $\text{d}_6\text{-DMSO}$  of 1)  $\text{MAPbI}_3$  (HI), 2)  $\text{FAPbI}_3$  (HI) and 3)  $\text{FA}_{0.5}\text{MA}_{0.5}\text{PbI}_3$  (IS).

#### Assignments:

$\text{MAPbI}_3$   $^1\text{H}$  NMR (300 MHz,  $\text{d}_6\text{-DMSO}$ )  $\delta$  2.36 (3H, s,  $\text{CH}_3$ ), 7.46 (3H,  $\text{NH}_3$ ).

$\text{FAPbI}_3$   $^1\text{H}$  NMR (300 MHz,  $\text{d}_6\text{-DMSO}$ )  $\delta$  7.83 (1H, tt, CH), 8.63 (2H, d,  $\text{NH}_2$ ), 8.98 (2H, s,  $\text{NH}_2$ ).

$\text{FA}_{0.5}\text{MA}_{0.5}\text{PbI}_3$   $^1\text{H}$  NMR (300 MHz,  $\text{d}_6\text{-DMSO}$ )  $\delta$  2.35 (3H, s,  $\text{CH}_3$ ), 7.46 (3H,  $\text{NH}_3$ ), 7.83 (1H, tt, CH), 8.63 (2H, d,  $\text{NH}_2$ ), 8.97 (2H, s,  $\text{NH}_2$ ).

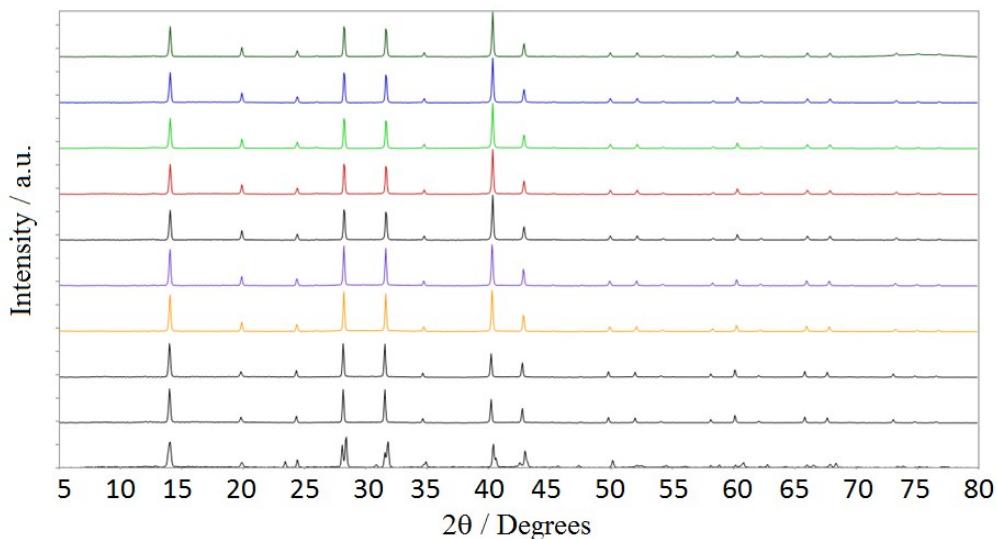


Figure 2 Powder X-ray diffraction patterns of HI method  $\text{MAPbI}_3$  (bottom)  $\text{FA}_x\text{MA}_{1-x}\text{PbI}_3$   $0 \leq x \leq 1$  in  $x = 0.1$  increments to the top.

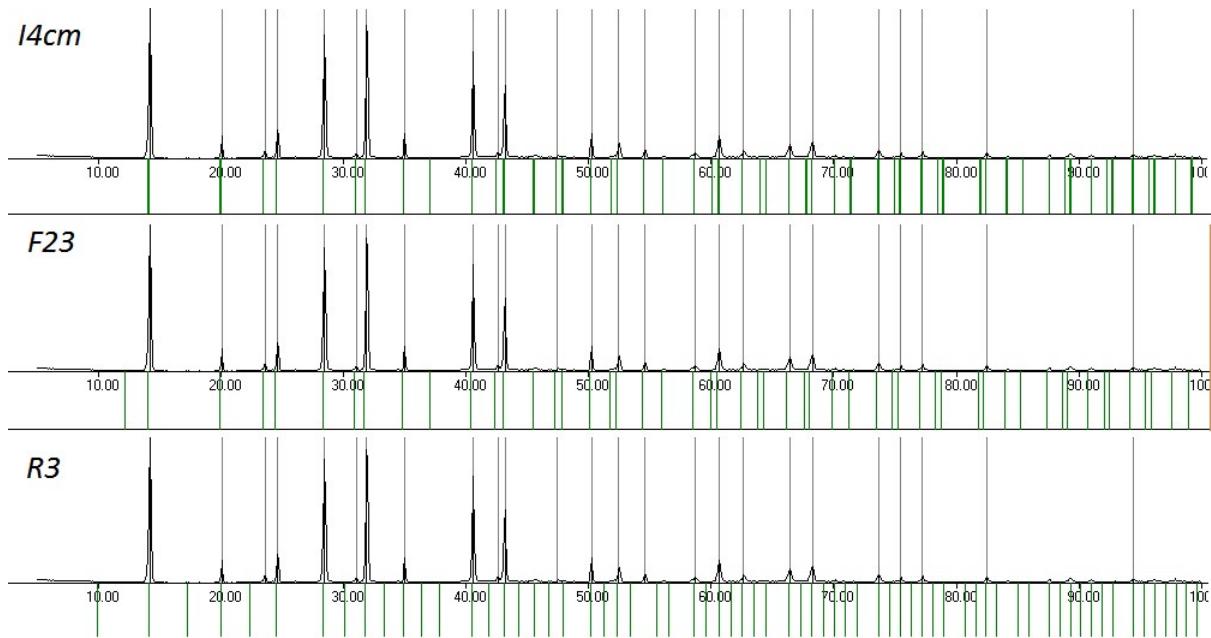


Figure 3  $\text{FA}_{0.1}\text{MA}_{0.9}\text{PbI}_3$  (HI) PXD pattern (black lines), peaks used for unit cell refinement (grey lines) and calculated peak positions (green lines) from CELREF for *I*<sub>4</sub><sub>*c*</sub>  $a = 8.89$   $c = 12.61$ , *F*2<sub>3</sub>  $a = 12.6$  and *R*3  $a = 8.91$ .

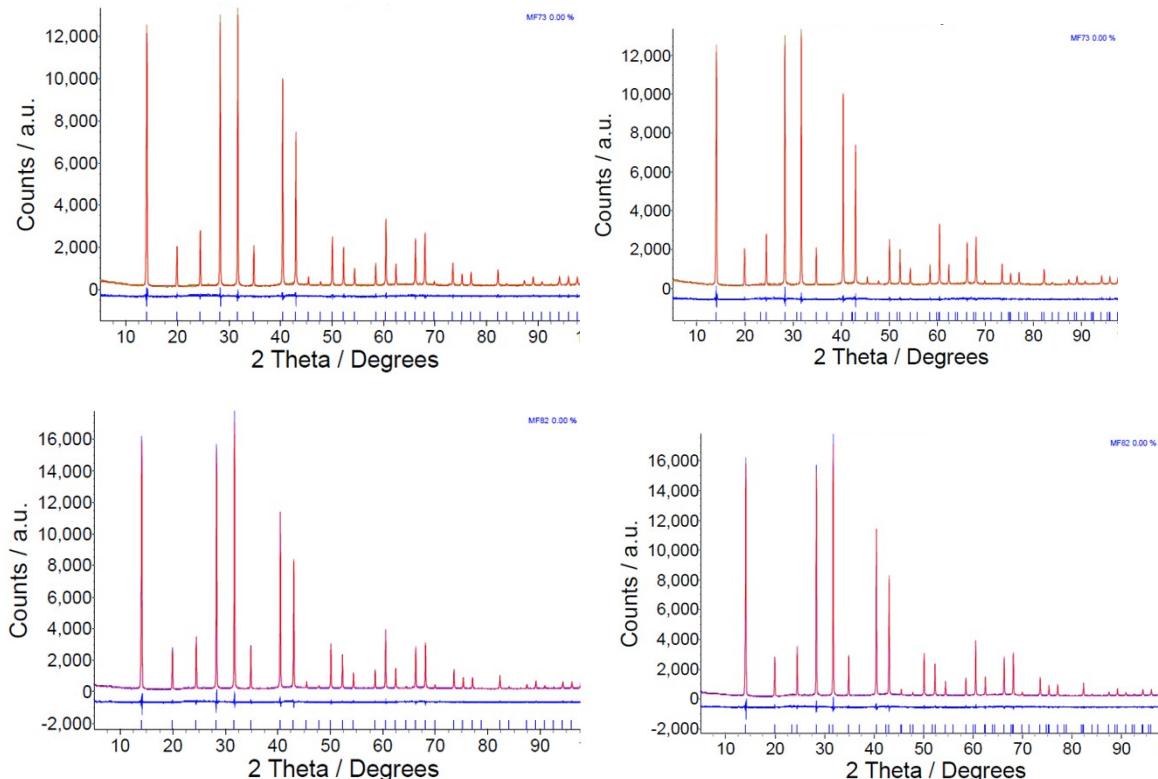
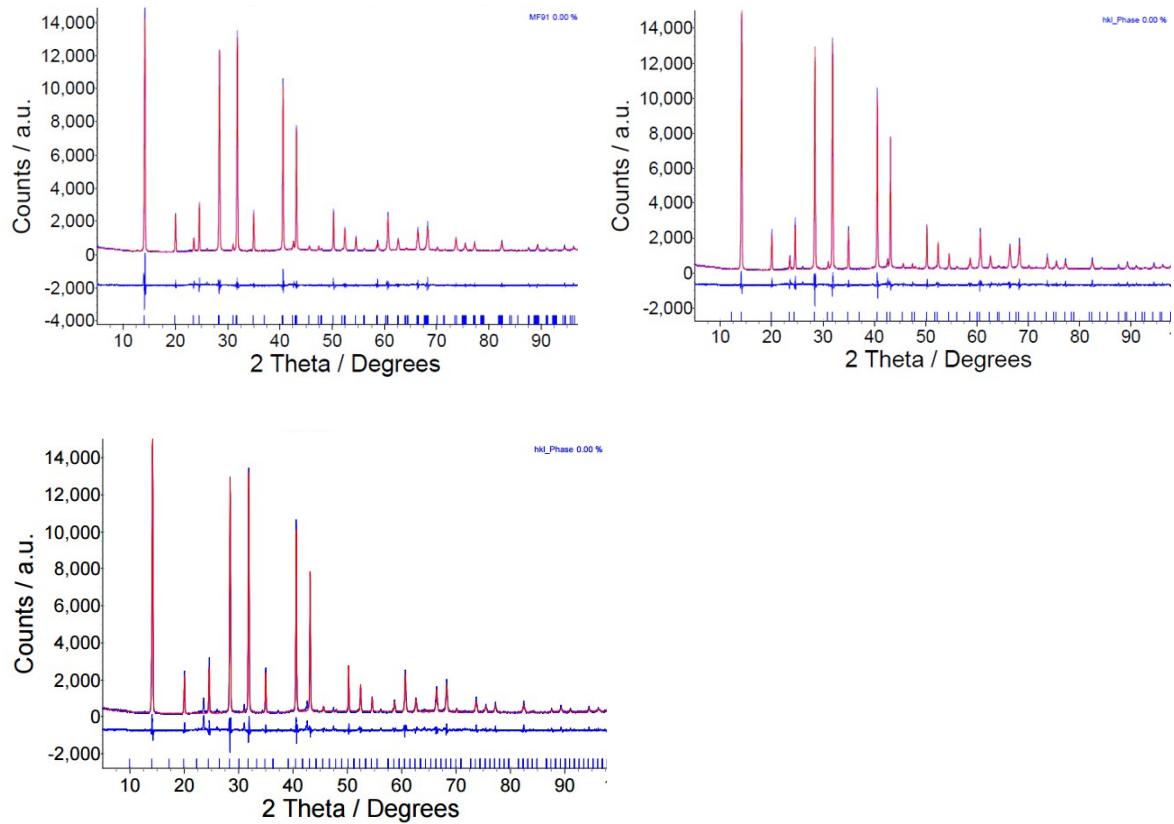


Figure 4 Pawley fits to PXD patterns of  $\text{FA}_{0.3}\text{MA}_{0.7}\text{PbI}_3$ , cubic *Pm*-3*m*,  $a \approx 6.31$  Å (top left) and tetragonal *I*<sub>4</sub><sub>*c*</sub>  $a \approx 8.90$  Å,  $c \approx 12.59$  Å (top right).  $\text{FA}_{0.2}\text{MA}_{0.8}\text{PbI}_3$  *Pm*-3*m* (bottom left) and *I*<sub>4</sub><sub>*c*</sub> (bottom right). Both samples were synthesised by the HI method. Difference profiles are plotted in blue and calculated peak positions are designated by pink ticks.



**Figure 5** Pawley fits to the room temperature PXD pattern of  $\text{FA}_{0.1}\text{MA}_{0.9}\text{PbI}_3$  for  $I4\text{cm}$ ,  $a \approx 8.91 \text{ \AA}$   $c \approx 12.59 \text{ \AA}$  (top left),  $F23$   $a \approx 12.6 \text{ \AA}$  (top right) and  $Im-3$   $a \approx 12.6 \text{ \AA}$  (bottom left).