

## Growth window optimization for large-size quasi two-dimensional Dion-Jacobson type perovskite

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### METHODS

#### Materials preparation

Lead (II) bromide (PbBr<sub>2</sub>, 99.5%), hydrobromic acid (HBr, 48 wt%), 3-(aminomethyl)pyridine (3AMPY, 99 wt%), and methylamine bromide (CH<sub>6</sub>BrN, 98%) were purchased from Anegi Reagent Co. Ltd (Shanghai, China). All reagents and solvents in this experiment were of reagent grade and could be used without further purification.

#### Preparation of (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub> SCs

To prepare (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub> microcrystals, 3 mmol of PbBr<sub>2</sub> (1.101 g) was dissolved in 12 mL of HBr solution. Subsequently, 0.5 mmol of 3AMPY (0.0476 g) was added to the solution, resulting in the formation of a light yellow precipitate. After warming the solution and allowing the precipitate to dissolve, 3 mmol (0.2026 g) of methylamine bromide was added. The mixture was then heated with constant stirring on a magnetic stirrer at 120 °C until the precipitate completely dissolved, producing a clarified solution. The (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub> microcrystals were obtained by rapidly

cooling this solution. To grow larger (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub> single crystals, the saturated solution was transferred to a glass vial and placed in an electrically heated, constant-temperature drying oven at 80 °C and cooled from 80-50 °C at 2 °C per day, Then cooled to room temperature (25 °C) at a rate of 1 °C per day. The ambient temperature was 21 °C, and the humidity was 48%.

### **Materials characterization**

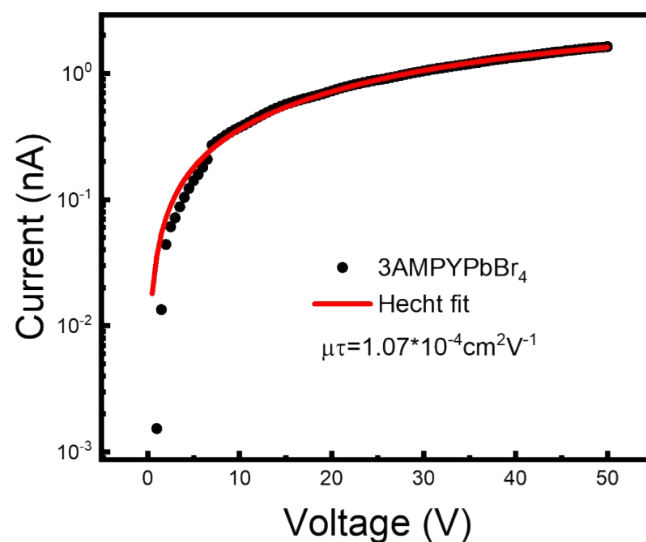
XRD patterns were collected by a Bruker D8 Advance diffractometer in 1° steps. The photoluminescence spectra were tested by a steady-state/transient fluorescence spectrometer (FLS 1000). The absorption spectra were obtained by a UV–vis–NIR spectrophotometer (Shimadzu UV-3600i Plus). SCXRD data at room temperature were collected by using a Bruker D8 Venture diffractometer with Mo Ka radiation ( $\lambda = 0.77$  Å). The crystal structures of (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub> were solved by the direct methods and then refined with the full-matrix least-squares technique based on F2 using SHELXLTL software package.

### **Detector performance measurement**

To apply different bias voltages, we utilized the Keithley 6517B Source Meter. For evaluating X-ray detection performance, we employed the Amptek Mini-X2 tube with an Au target (Newton Scientific M237) as the X-ray source. Additionally, we utilised a horizontal displacement stage to move the samples in the X-Y direction. Dose rates were calibrated using accu-diode ionization chambers, including DDX6-W and DDX6-WL (Accu-Gold+, Radcal).

Table S1. Crystal data and structure refinement for (3AMPY)(MA)Pb<sub>2</sub>Br<sub>7</sub>.

Empirical formula	C <sub>7</sub> H <sub>16</sub> Br <sub>7</sub> N <sub>3</sub> Pb <sub>7</sub>
Formula weight	1115.98
Temperature/K	296(2)
Crystal system	Monoclinic
Space group	P2 <sub>1</sub> /c
a/Å	15.559(2)
b/Å	17.143(2)
c/Å	8.2450(11)
α/°	90
β/°	97.367(3)
γ/°	90
Volume/Å <sup>3</sup>	2181.1(5)
Z	4
ρ <sub>calc</sub> Mg/m <sup>3</sup>	3.398
μ/mm <sup>-1</sup>	28.233
F(000)	1952
Crystal size/mm <sup>3</sup>	0.220×0.200×0.180
θ Range for data collection/°	2.376 to 28.355
Index ranges	-20≤h≤18, -22≤k≤22, -11≤l≤10
Reflections collected	34760
Independent reflections	5417 [R(int) = 0.1281]
Data/restraints/parameters	5417 / 1 / 176
Goodness-of-fit on F <sup>2</sup>	1.006
Final R indexes [I≥2σ(I)]	R1 = 0.0576, wR2 = 0.1051
Final R indexes [all data]	R1 = 0.1489, wR2 = 0.1347
Largest diff. peak/hole/e Å <sup>-3</sup>	1.951 and -2.747

Figure S1. 3AMPYPbBr<sub>4</sub> crystal  $\mu\tau$  test results.