Electronic Supplementary Information

A 9R-Like 2D Hybrid Metal Halide with Remarkably Low Melting Temperature

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Experimental section

Starting materials. 2-(methylamino)ethan-1-ol (98%), lead bromide (PbBr₂, 99.9%), and hydrobromic acid (HBr, 48 *wt*.% in H₂O) were commercially available and used as received without further purification.

Synthesis of (MHEA)4Pb3Br10. PbBr₂ powder (9 mmol, 3.303 g) was dissolved in a solution comprising 4 mL of aqueous HBr. Following this, 2-(methylamino)ethan-1-ol (12 mmol, 0.901 g) was introduced into the solution. Subsequently, the clear solution was cooled and kept at 0 ℃ without any interference. During the process, colorless rod-like crystals began to precipitate at the bottom of the beaker. The solution was left for several days to facilitate further crystal growth. Afterward, the crystals were collected via suction filtration and dried in ambient air.

General Characterizations: PXRD patterns were obtained using a Rigaku SmartLab X-ray diffraction system. TGA was conducted with a NETZSCH TG209 F3 instrument at a heating rate of 10° C min⁻¹ in a nitrogen atmosphere. DSC measurements were performed using a NETZSCH DSC 214 Polyma instrument, also at a standard scan rate of 10° C min⁻¹, unless noted otherwise. Ultraviolet-visible diffuse reflection spectra were recorded with a Shimadzu UV2600 spectrophotometer equipped with an ISR-2600Plus integrating sphere. Fourier-transform infrared spectra were collected using a Nicolet iS50 instrument through a transmission method. The viscosity was measured using a Thermo HAAKE MARS 60 with a 25 mm parallel plate, the cooling and shear rates are 10 K/min and 10 s^{-1} , respectively.

Single-Crystal X-ray Diffraction. Crystallographic data for the compound was gathered using a Rigaku Oxford Diffraction Supernova Dual Source, Cu at Zero, which is equipped with an AtlasS2 CCD and an XtaLAB Synergy R, DW system, HyPix diffractometer utilizing Mo Kα radiation. The data collection, cell refinement, and reduction were performed using the Rigaku CrysAlisPro software. Structure solution was achieved via direct methods using the SHELXL-2014 package, with all nonhydrogen atoms refined anisotropically. The crystallographic data can be found in the Cambridge Crystallographic Data Centre (CCDC number 2391828).

Theoretical calculations. Density Functional Theory (DFT) calculations were conducted using the

Vienna ab initio Simulation Package $(VASP)^1$. The input structures were derived from the crystallographic data and subsequently optimized with the cell fixed. Thresholds of convergence criteria are set as 10^{-7} eV and 0.02 eV/Å for energy differences during electronic steps and maximum forces in geometry optimization. The Perdew–Burke–Ernzerhof (PBE) functional, implemented within the generalized gradient approximation (GGA), was utilized to model the exchange-correlation interactions. 2,3 DFT-D3 method was used to evaluate the van der Waals interactions. ⁴ Ion-electron interactions were represented using projector augmented wave (PAW) potentials with an energy cutoff of 500 eV. Postprocessing analysis was performed using VASPKIT.⁵

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Fig. S1 Bulk single crystals of (MHEA)₄Pb₃Br₁₀

Fig. S2 PXRD pattern of (MHEA)₄Pb₃Br₁₀ powders.

Fig. S3 Conformation analysis of MHEA cation.

Fig. S4 Diagrams of supramolecular frameworks of crystal (MHEA)₄Pb₃Br₁₀ with hydrogen bonds.

Fig. S5 Images of melts, and recrystallized crystals (from melts) under unpolarized light (i, ii) and cross-polarized light (iii, iv), respectively. Scale bar: 200 μm.

Fig. S6 IR spectra of fresh crystal and melt-processed powders.

Fig. S7 Excitation and emission spectra of (MHEA)₄Pb₃Br₁₀.

Fig. S8 Density of states of (MHEA)₄Pb₃Br₁₀.

Table S1 Crystallographic data and refinement parameters of (MHEA)₄Pb₃Br₁₀.

 $[a]$ $R_1 = \sum ||F_o| - |F_c||/\sum |F_o|$.

 $[b]$ $wR_2 = [\Sigma w (F_0^2 - F_c^2)^2 / \Sigma w (F_0^2)^2]^{1/2}.$

[c] Maximum and minimum residual electron density.

Table S2 Bond angles and bond lengths of $(MHEA)_4Pb_3Br_{10}$.

Bond angles	Angle $/$ \circ	Bond lengths	Length $/\AA$
$Br(2)-Pb(1)-Br(3)$	84.12(3)	$Pb(1)-Br(2)$	2.9331(10)
$Br(2)-Pb(1)-Br(5)$	91.25(3)	$Pb(1)-Br(3)$	3.0920(11)
$Br(2)-Pb(1)-Br(5)^{i}$	177.21(3)	$Pb(1)-Br(4)$	2.8419(11)
$Br(3)-Pb(1)-Br(5)^1$	93.17(3)	$Pb(1)-Br(5)$	2.9737(10)
$Br(4)-Pb(1)-Br(2)$	93.18(3)	$Pb(1)-Br(5)^{i}$	3.1265(10)
$Br(4)-Pb(1)-Br(3)$	93.14(3)	$Pb(2)-Br(1)$	3.0307(11)
$Br(4)-Pb(1)-Br(5)$	92.82(3)	$Pb(2)-Br(2)^{ii}$	3.0595(12)
$Br(4)-Pb(1)-Br(5)^{i}$	87.63(3)	$Pb(2)-Br(3)$	3.0044(12)
$Br(5)-Pb(1)-Br(3)$	172.64(3)		
$Br(5)-Pb(1)-Br(5)^{i}$	91.375 (10)		
$Br(1)-Pb(2)-Br(1)ii$	170.94(6)		
$Br(1)-Pb(2)-Br(2)^{ii}$	102.35(3)		
$Br(1)-Pb(2)-Br(2)$	84.07(3)		
$Br(2)-Pb(2)-Br(2)^{ii}$	91.28(4)		
$Br(3)-Pb(2)-Br(1)$	83.87(3)		
$Br(3)-Pb(2)-Br(1)ii$	90.46(3)		
$Br(3)-Pb(2)-Br(2)$	83.50(3)		
$Br(3)^{ii} - Pb(2) - Br(2)$	171.48(3)		
$Br(3)-Pb(2)-Br(3)ii$	102.45(5)		
$Pb(1)-Br(2)-Pb(2)$	83.43(3)		
$Pb(2)-Br(3)-Pb(1)$	81.71 (3)		
$Pb(1)-Br(5)-Pb(1)$ ⁱⁱⁱ	168.39(4)		

Symmetry codes: (i) −*x*−1/2, *y*−1/2, −*z*+1/2; (ii) −*x*, *y*, −*z*+1/2; (iii) −*x*−1/2, *y*+1/2, −*z*+1/2.

Hydrogen bonds for $(MHEA)_4Pb_3Br_{10}$						
$D-H\cdots A$	$D-H/A$	$H \cdots A / A$	$D \cdots A / \AA$	\angle DHA / \circ		
$O(1) - H(1) \cdots N(1)$	0.82	2.58	2.933(15)	107		
$N(1) - H(1A) \cdots Br(5)$	0.89	2.55	3.378(8)	156		
$N(1) - H(1A) \cdots O(1)$	0.89	2.58	2.933(15)	104'		

Table S3. Summary of hydrogen bonds.

$N(1) - H(1B) \cdots Br(4)^{1}$	0.89	2.61	3.365(9)	143	
$O(2)$ -H(2)···Br(3) ¹¹	0.82	2.70	3.468(11)	157	
$N(2) - H(2C) \cdots O(1)$	0.89	2.15	2.942(16)	148	
$N(2)$ -H(2C) \cdots O(2) ^m	0.89	2.39	2.863(14)	114'	
$N(2) - H(2D) \cdots O(2)$	0.89	2.38	2.779(16)	107	
$N(2)$ -H(2D) \cdots O(2) ⁱⁱⁱ	0.89	2.50	2.863(14)	105'	
Symmetry codes: (i) x,2-y,1/2+z; (ii) -x,1+y,1/2-z; (iii) -x,3-y,1-z					

Table S4 Summary of the thermal properties of some reported 2D HMHs.

Reference

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