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## **Supporting Information**

## Preferential S/Se occupation in anisotropic ReS<sub>2(1-x)</sub>Se<sub>2x</sub> monolayer alloy

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**Fig. S1.** (a) STEM image of  $\text{ReS}_{1.4}\text{Se}_{1.6}$ , showing a Re4 atomic chains and local atomic coordination structure. (b) STEM image and local coordination structure.

## Supporting Note S1. Details on data analysis and image processing.

To extract the S/Se concentration at different atomic coordination sites, we carried out data analysis and image processing on different STEM images. The preliminary images were firstly filtered using Image J software for displaying clear atomic position. The data analysis was performed using a custom-written Matlab (MathWorks) code. The flow diagram of the analysis process is presented in Fig. S(2). After the loading of STEM image, which was firstly filtered by Image J. We can set the x, y range of the region we want to do the analysis. Then, we input the lattice parameter of a, b, the Theta between the a and b axis and the tilt angle of image. We can compare the lattice identification results on a Figure window in Matlab to check the precise of lattice generation. The main function for determining the lattice is shown as following:

```
function [ x_position, y_position, number_index, number_estimated
,a_vector,b_vector, x_position_ideal,y_position_ideal] =
STEM analysis lattice generate direct( a,b,theta,tilt angle,x0,y0,xmax,ymax )
widthmax=max((ymax-y0), (xmax-x0));
number estimated=(ymax-y0)*(xmax-x0)/a/a/sin(theta/180*pi);
number index=0;
column_number=round(1.1*(round(sqrt((widthmax)^2+(widthmax)^2)/min(a,b))+1));
for p=0:column number %a, hirozotal
for q=0:column number %b, vertical
xy distance=sqrt((p*a)^2+(q*b)^2-2*cos(pi-theta/180*pi)*a*p*b*q)
if p~=0
      theta to x axes=tilt angle/180*pi+acos(((p*a)^2+xy distance^2-
(q*b)^2)/2/xy_distance/p/a)
elsetheta to x axes=(theta+tilt angle)/180*pi
end
x temp=round(x0+xy distance*cos(theta to x axes));
y temp=round(y0+xy distance*sin(theta to x axes));
if p==0 & q==0
number index=1;
x position(number index, 1) = x0;
y position(number index,1)=y0;
a vector(number index,1)=0;
b_vector(number_index,1)=0;
x position ideal(number index,1) = x temp;
y position ideal(number index,1)=y temp;
else
```

```
р
q
x_temp
y temp
if (x_temp<=xmax&x_temp>=x0-1 &y_temp<=ymax&y_temp>=y0-1);
number index=number index+1;
x position(number index,1)=x temp;
y position(number index,1)=y temp;
a vector(number index,1)=p;
b vector(number index,1)=q;
x position ideal(number index,1) = x temp;
y position ideal(number index,1)=y temp;
end
end
end
end
```

In the analysis, we did not use the lattice generated by the first step directly, but searched a maximum intensity in the region near the generated position as the practical Re position and the detailed function is:

```
function [ x position 1D, y position 1D] =
STEM_analysis_lattice_generate_search(
image_data,x_position_1D,y_position_1D,ROI,int_range_min,int_range_max,min_p
ixel );
number index=size(x position 1D);
for i=1:number index
ROI array=image data((y position 1D(i)-
ROI): (y position 1D(i)+ROI), (x position 1D(i)-ROI): (x position 1D(i)+ROI));
             [ROI_1D_max ROI_y_position_line]=max(ROI_array);
             [ROI 2D max ROI x position]=max(ROI 1D max);
ROI y position=ROI y position line(ROI x position);
if ROI 2D max<=int range max& ROI 2D max>=int range min
                 x position 1D(i) = x position 1D(i) + ROI x position - ROI;
                 y position 1D(i)=y position 1D(i)+ROI y position-ROI;
end
end
```

As shown by the codes highlighted in blue, we carried out a searching process near the position of lattice and determined a point with maximum intensity as the position of Re atom, which can eliminate the deviation between the standard  $\text{ReS}_{2(1-x)}\text{Se}_{2x}$  lattice and practical STEM image. After the determination of the lattice, we set the offset values for S/Se sites compared with the pristine Re position in each unit cells. Next, we searched a maximum value near the firstly determined S/Se coordination sites as the intensity of the sites and the main function is shown as follow:

```
function [ atomassign_result , max_int_array] =
STEM analysis atom assignment(atomassign result,image data,atom site x,atom
site y, atomassign site selected, atomassign ROI, atomassign min, atomassign max, a
tomassign atom name, atom number );
%UNTITLED1 Summary of this function goes here
% Detailed explanation goes here
[n, m]=size(atom site x)
x_position=atom_site_x(1:atom_number,atomassign_site_selected);
y position=atom site y(1:atom number, atomassign site selected);
max(x position)
min(x position)
max(y position)
find(y position==1500)
for i=1:atom number
ROI_array=image_data((y_position(i)-
atomassign_ROI):(y_position(i)+atomassign_ROI),(x_position(i)-
atomassign_ROI):(x_position(i)+atomassign_ROI),1);
   ave int=sum(sum(ROI array))/(2*atomassign ROI+1)/(2*atomassign ROI+1);
max int=max(max(ROI array));
max int array(i) = max int;
ifmax int<=atomassign max&max int>=atomassign min
      atomassign_result(i,atomassign_site_selected)={atomassign_atom_name};
end
end
figure(2);
int array=0:65535/255:65535;
```

```
hist(max_int_array,int_array);
```

As shown in the highlight part, we extracted the maximum value in the region of each coordination sites as the intensity, which ensures the accuracy of our analysis results. The numbers marked in the Fig. S4 and S5 is added by the set position according to the ideal lattice, since extract the position of maximum intensity requires additional codes, which is detrimental to the running speed.



Figure S2. A flow diagram for analysis of S/Se coordination in STEM image.

•		STEM_analysis		- 🗆 🗙
Ele Edit View Insert Iools Ele Edit View Insert Iools DARX2 STEMselected Lattice generale x range 50 500 y 40 500 g 40 500 b 38.0 + f 0 Theta 114, tit 25.2 Direct m ▼ RO(+/- 6 int. range 400 655 tit. range 400 65	Desktop Window Help     Window Help     Pot outside      Add atom site     # of sets     9     Offset value 1     0     Offset value 2     -21     15     Offset value 3     -8     15     Offset value 4     -6     7     Offset value 5     7     7     Offset value 7     5     30     Offset value 8     12     16     Offset value 9     -5     16	r       Set marker color         Atom assignment         Select site         RO(+/-         2         Ave. int.         0         Solo         Atom name         Re1         Assign         Clear site         Show site!         Show site!	2D Guassian fit         Max delta x       Edit Text         Max delta y       Edit Text         Peak int. range       255         Peak int. range       20         Max search times       Edit Text         GOF criterion       Edit Text         Generate       Show         Coordination analysis       Center site         1       1	Indivudal atom correction Atom SN# 155 a vector 0 b vector 0 Site # 1 Select Show selected X 0 349 y 0 741 Corr. Atom name Z Z Corr.
Show po Show Show a Show b	Offset value 0 0 Generate Show Show Show Show Show Show Show Show Show Show sit		Center element Re1 Coord. site 1 Coord. element Re1 Distance 0 Dist. +/- (%) 5 Calculate Accept ✓ Show all Show spec.	Edit Text Save original data Edit Text run soript Copyright reserved @ Xielab at

Figure S3.Matlab GUI of the STEM coordination analysis.



**Fig. S4.** The extraction of local coordination sites from STEM images.(a)STEM image with determined position of unit cells. The position of unit cell is marked with a red 'x' on a Re atom. (b) STEM image with atomic sites marked with different numbers that represents different coordination sites shown in Figure 3(a) in main text. (c) Zoom-in STEM image presents clear coordination sites. (d) The distribution of grey scale value at position of different sites, which illustrates an obvious contrast between S and Se atoms.



**Fig. S5.** Extraction of local atomic coordination at another STEM region. (a) STEM image with determined position and lattice parameter of unit cells. (b) STEM image with determined atomic sites. The different atomic coordination sites are marked with numbers of 1-8. (c) Zoom-in STEM images, presenting clear position of coordination sites. The distribution of grey values, that represents the intensity of sites, is presented in Figure 3(b). The concentration of S and Se atoms at different coordination sites can be extracted.



Fig. S6. Raman spectrum of monolayer ReS<sub>1.4</sub>Se<sub>0.6.</sub>

## Supporting Note S2. Details on calculation on the relative occupation probability and Se percentage at growth temperature.

Since Se has different energy at different site, the relative occupation probability of Se at a specific site *i*,  $P_{(Se,i)}$  can be calculated as

$$P_{(Se,i)} = \exp(\frac{-E_i}{k_B T})$$

where  $E_i$  is the Se substitution energy at site *i*,  $k_B$  is the Boltzmann constant and *T* is the growth temperature (*T*=1303 K). From the calculated energy difference, the  $P_{(Se,i)}$  can be calculated. Then Se percentage at different site, *Content*<sub>(Se,i)</sub>, can be calculated by

$$Content_{(Se,i)} = \frac{P_{(Se,i)}}{\left(\sum_{j=1}^{8} P_{(Se,i)}\right)/8} \bullet Content_{(Se,average)}$$

where  $Content_{(Se, average)}$  is the average Se content (0.30). According to the calculated results shown in Table S1, the probability of Se substitution of S at the growth temperature is well consistent with experimental results.

**Table S1.**The calculated relative energy, relative occupation probability, Sepercentage and experimental Se percentage.

Sites	Relative energy	Calculated	Calculated Se	Experimental
	for Se	relative	percentage at	Se percentage
	substitution (eV)	occupation	different sites,	from STEM
	at room	probability,	$Content_{(Se,i)}$ at	imaging at
	temperature	$P_{(Se,i)}$ at 1303 K	1303 K	different sites
				(data derived
				from Fig. 3c)
1,2	0.057	0.602	0.217	0.212
3,4	0.026	0.794	0.287	0.279
5,6	0	1	0.362	0.363
7,8	0.009	0.923	0.334	0.319