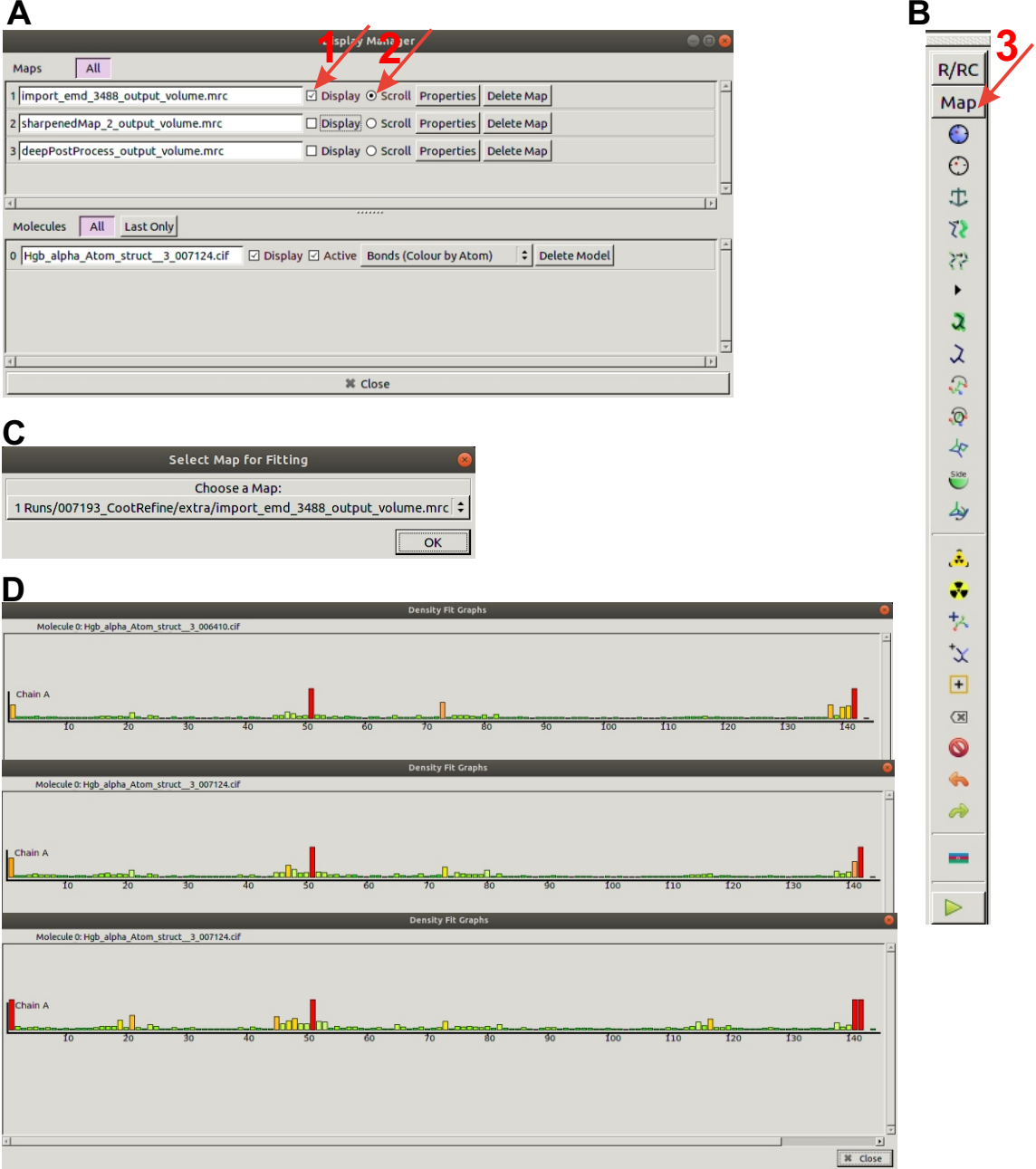
A
B
C
D
1
2
3
Display Manager
Maps
All
1 import_emd_3488_output_volume.mrc
Display
Scroll
Properties
Delete Map
2 sharpenedMap_2_output_volume.mrc
3 deepPostProcess_output_volume.mrc
Molecules
Last Only
0 Hgb_alpha_Atom_struct__3_007124.cif
Active
Bonds (Colour by Atom)
Delete Model
Close
R/RC
Map
Select Map for Fitting
Choose a Map:
1 Runs/007193_CootRefine/extra/import_emd_3488_output_volume.mrc
OK
Density Fit Graphs
Molecule 0: Hgb_alpha_Atom_struct__3_006410.cif
Molecule 0: Hgb_alpha_Atom_struct__3_007124.cif
Chain A
10
20
30
40
50
60
70
80
90
100
110
120
130
140