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# Model building, validation, and visualization

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Protocol status: Working

We use this protocol and it's working

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### **Abstract**

Model building, validation, and visualization

## Troubleshooting



### Model building

- 1 Generate a model by
  - [1] existing model in PDB
  - [2] creating a homology model by SWISS-MODEL
  - [3] creating a ab-initial model by AlphaFold
- 2 Do the flexible model fitting by [ISOLDE] Roughly fit the secodary structures into the map
- 3 Do model fitting manually in [Coot] or automatically in [Phenix] Repeat the model fitting until the coordinates fit the map resonably

#### Validation

4 Validate the quality of the models by using the validation tools in Phenix and the online validation service provided by wwPDB (https://www.wwpdb.org/validation/validationreports)

Make sure the statistics, particularly the Ramachandran plot, Rotamers, R.m.s. deviations and B-factors are resonable.

#### Visualization

5 Visualize the models and maps by [ChimeraX] Create figures by [Adobe Illustrator] Create movies by [iMovie]