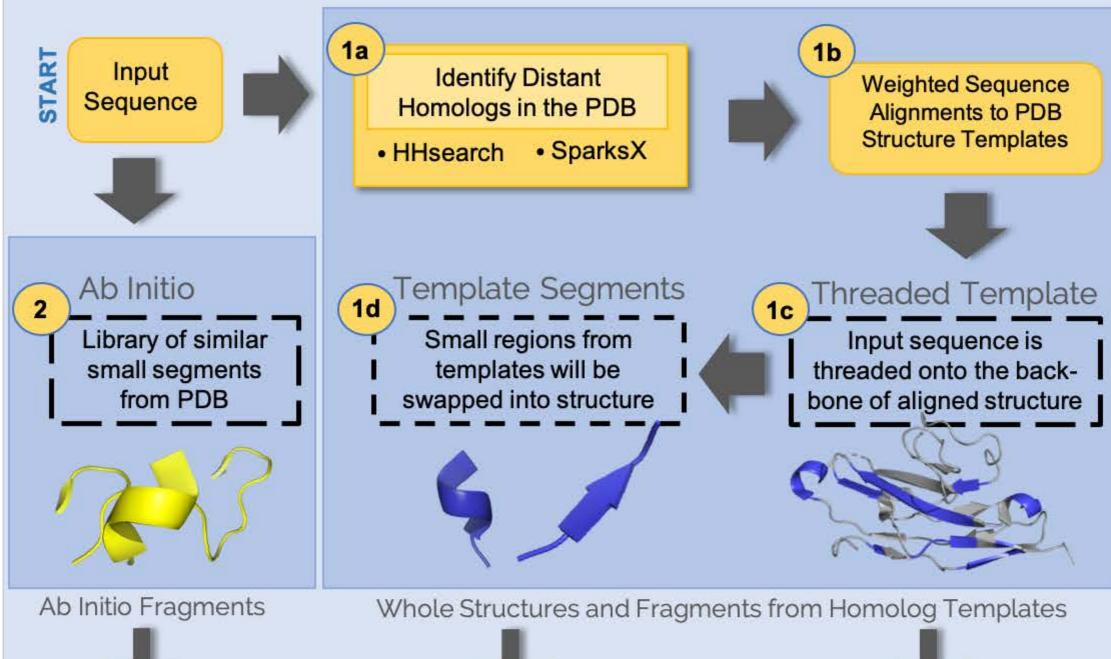


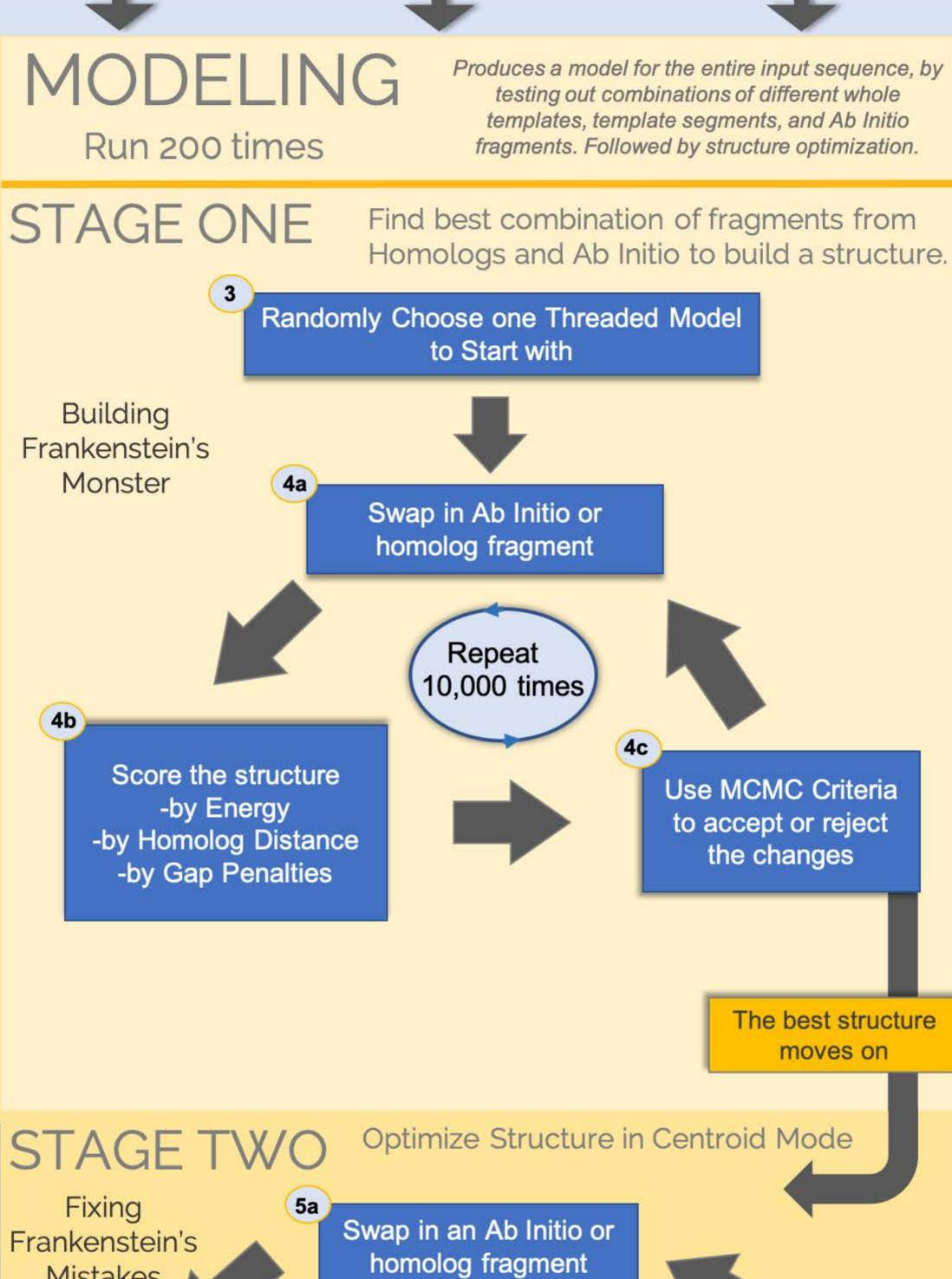
HOMOLOGY MODELING

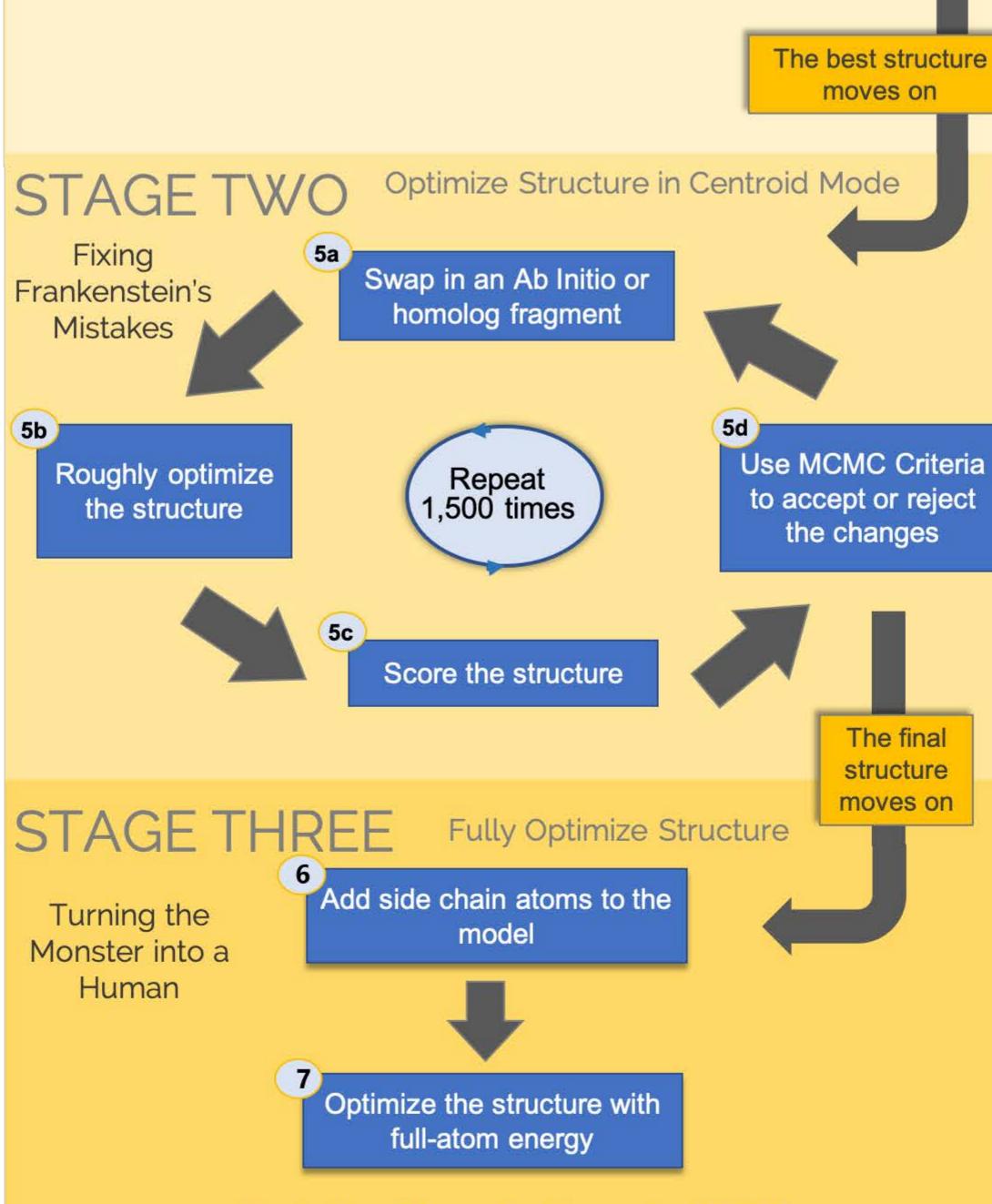
From Sequence to Predicted Structure

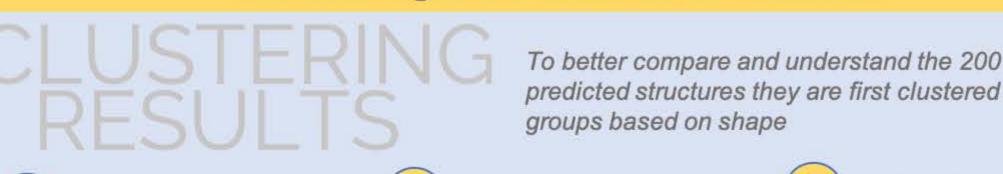


Input sequence is used to find similar structures to use as conformational sampling guides



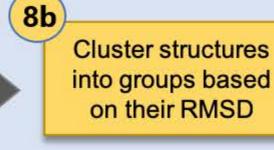






predicted structures they are first clustered into

8a Calculate the backbone RMSD between each of the 200 structures



Modeling Phase is Repeated 200X,

Generating 200 Predicted Structures-

8c The best structure from each of the top five clusters is presented in HM results