

Figure 3. Cryo EM structure of the *Klebsiella aerogenes* alpha glucosidase trimer obtained to 3.5 Å resolution. **A** and **B**) 3D reconstruction of the trimer population. **C**) Dimer of trimers fit to the cryo-EM map. **D**) Side view of the dimer of trimers.

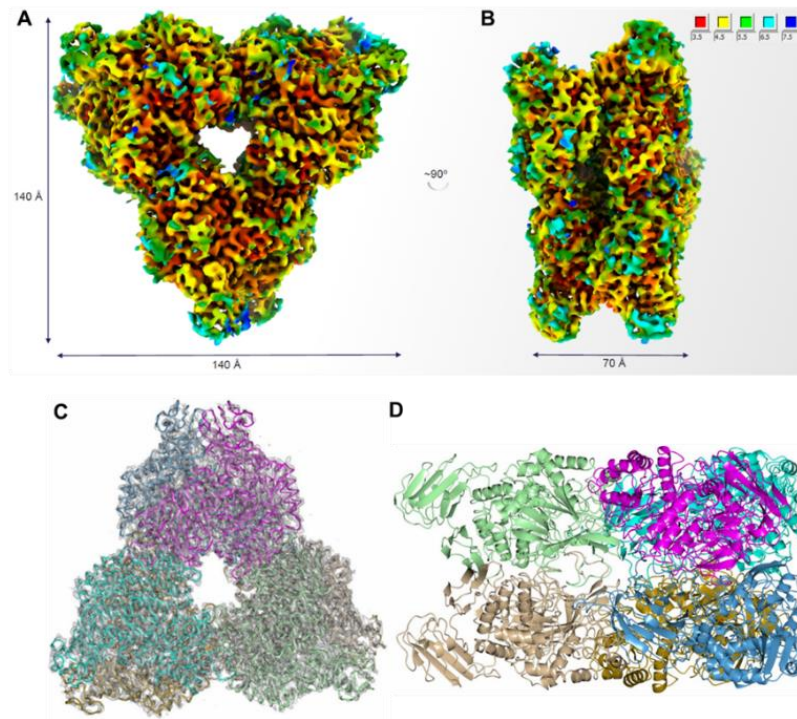


Figure 4. Cryo-EM structure of *M. tuberculosis* Valine--tRNA ligase. **A**) 3D reconstruction and **B**) model fit to the sharpened cryo-EM map (3.5 Å resolution).

