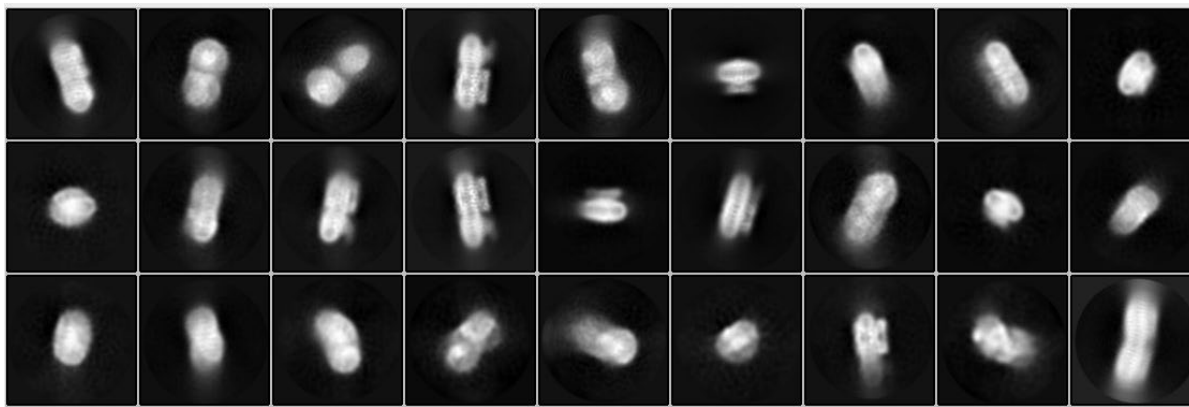
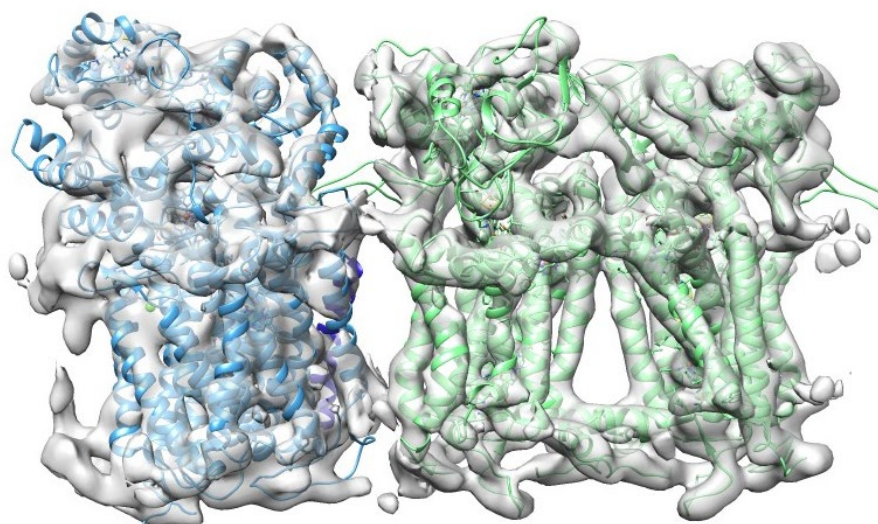


Selected 2D class averages of ~ 500.000 particles:



CryoEM map at 6.2 Å resolution with fitted models of CIII (green, x-ray structure, PDB: 1ZRT) and CIV (blue, homology model):



Local resolution map:

