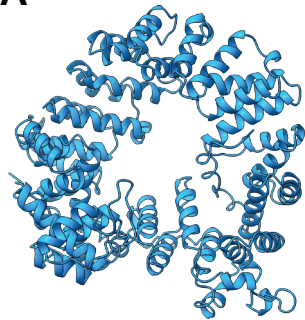
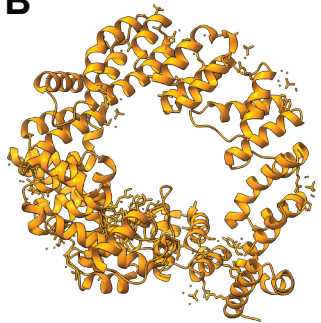
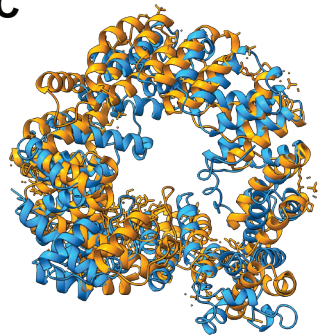


A

BB0259
Foldseek generated

B

E. coli Slt70
PDB 1QSA

C

BB0259-Slt70 alignment
RMSD 1.302
(Needleman-Wunsch algorithm)