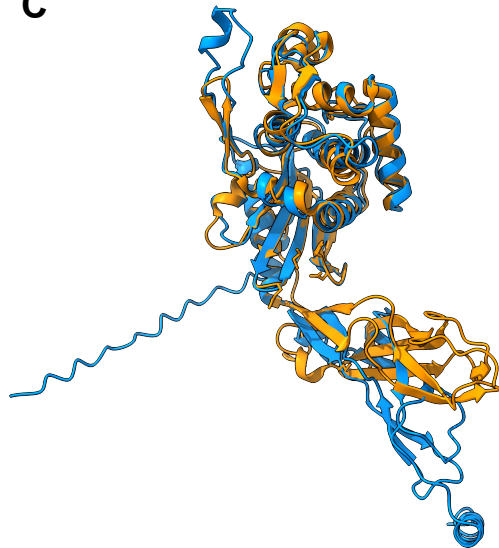


A

BB0605
Foldseek generated

B

E. coli DacA
PDB 1NZO

C

BB0605-DacA alignment
RMSD 0.908
(Needleman-Wunsch algorithm)