



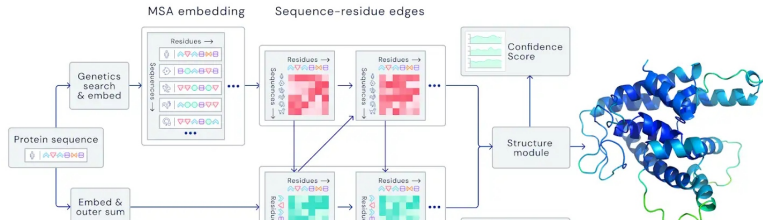
● Experimental result
● Computational prediction



T1037 / 6vvr4
90.7 GDT
(RNA polymerase domain)



T1049 / 6y4f
93.3 GDT
(adhesin tip)



Median Free-Modelling Accuracy

