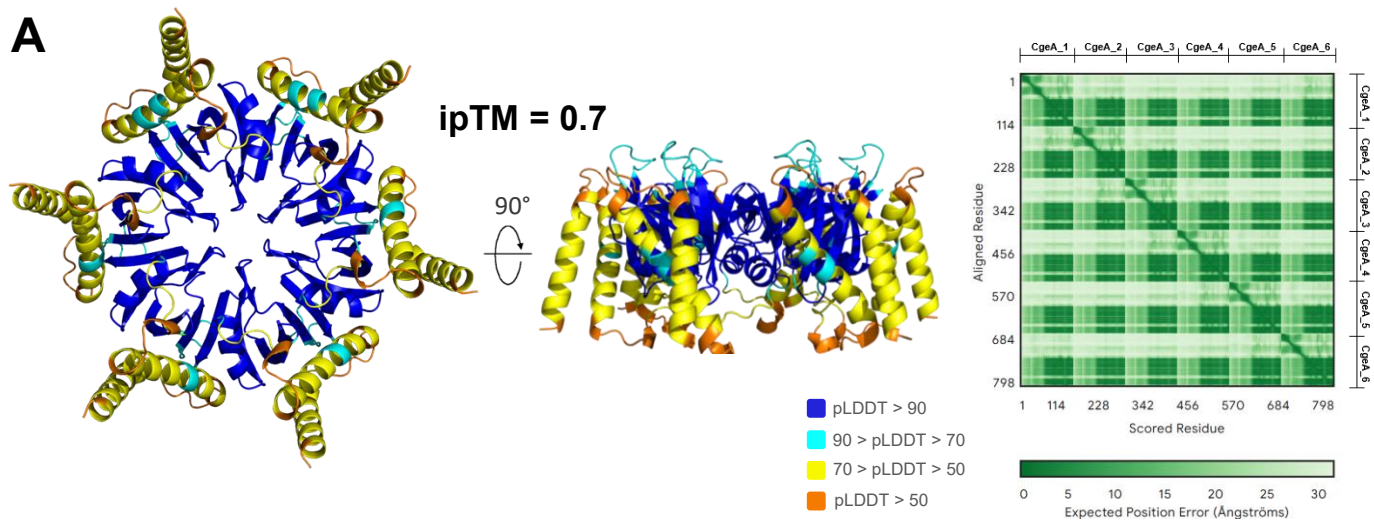
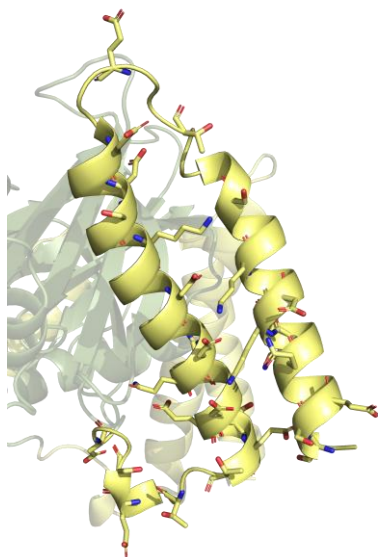


A



B



1 MSSENAQLKKDLIKAVLSPLFPTATEG
 28 GENMDSNLKALLDAIDQKVDESETVT
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 109 SSVTPAGQATTLIDVDKVESVTFFS