

Average E-value, SeqID <= 25%							
Strands per Chain	8	10	12	14	16	18	22
8	6.40e-15	4.43e-05	1.69e-06	1.83e-06	3.11e-05		
10	NA	1.10e-14	1.41e-04	9.40e-04	NA	NA	NA
12			4.40e-27	1.39e-04	7.49e-06		6.16e-04
14	NA	NA	NA	1.61e-35	6.89e-11	NA	1.23e-04
16					6.53e-37	2.29e-05	
18	NA	NA	NA	NA	NA	6.12e-32	NA
22							3.13e-52