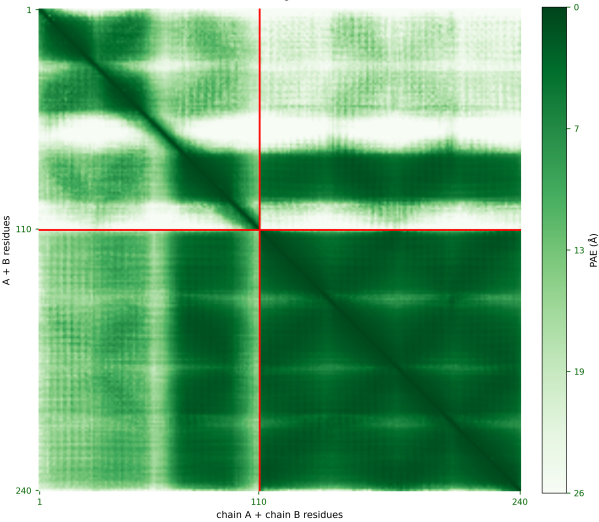
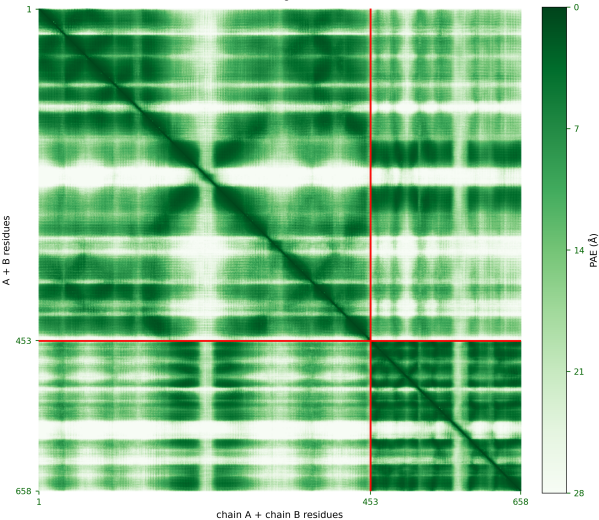


A2RU14 TMEM218 | Q9PON5 TMEM216
Best interchain (off-diagonal) PAE = 1.138 Å

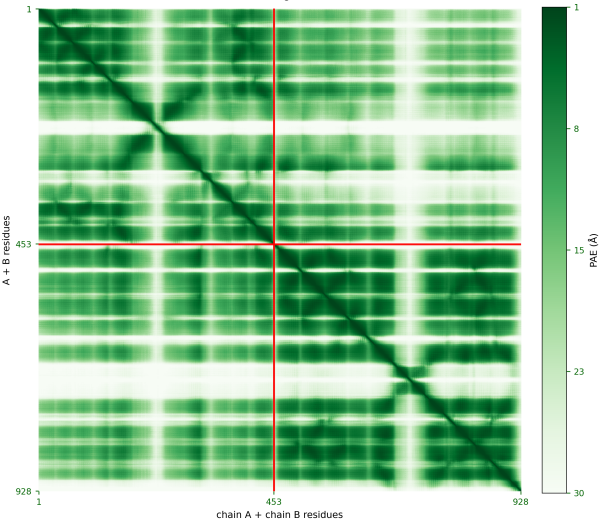


B7ZQAQ6 GPR89A | Q5HYI8 RABL3

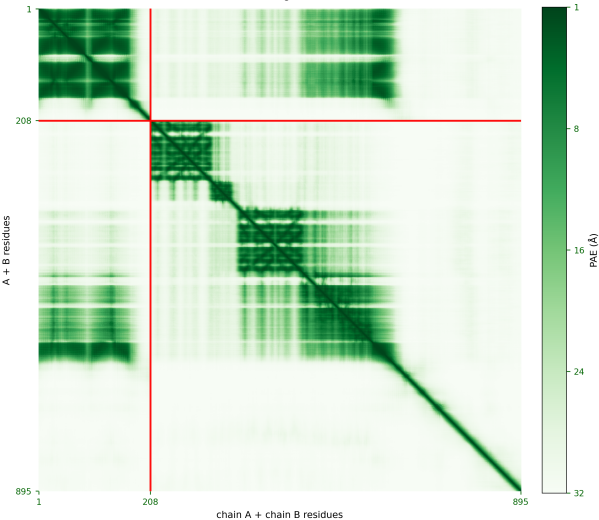
Best interchain (off-diagonal) PAE = 1.240 Å



B7ZAQ6 GPR89A | Q6UX01 LMBR1L
Best interchain (off-diagonal) PAE = 2.016 Å

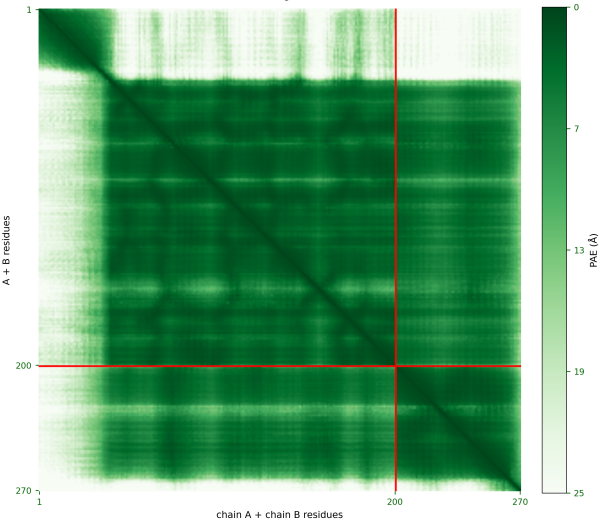


O75204 TMEM127 | Q7Z4F1 LRP10
Best interchain (off-diagonal) PAE = 1.489 Å



O75880 SCO1 | Q5JTJ3 COA6

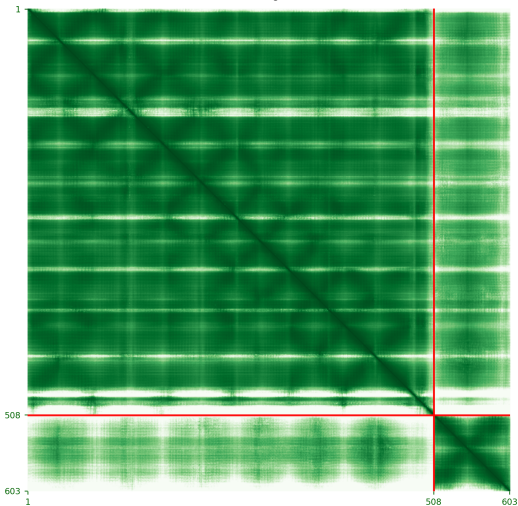
Best interchain (off-diagonal) PAE = 1.088 Å



P00395 MT-CO1 | Q7Z7K0 CMC1

Best interchain (off-diagonal) PAE = 1.665 Å

A + B residues



PAE (Å)

0

5

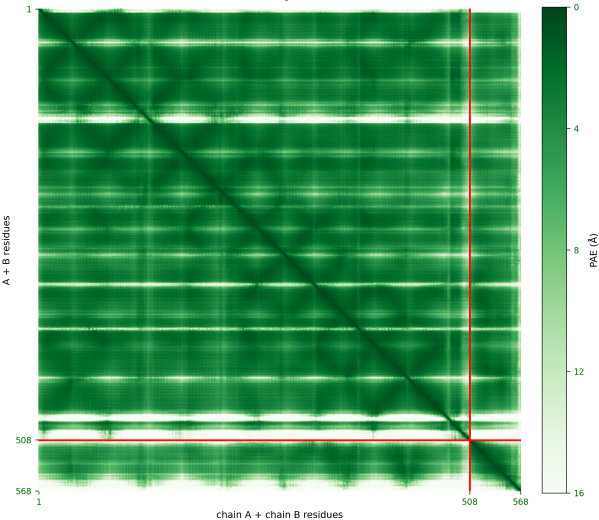
10

14

19

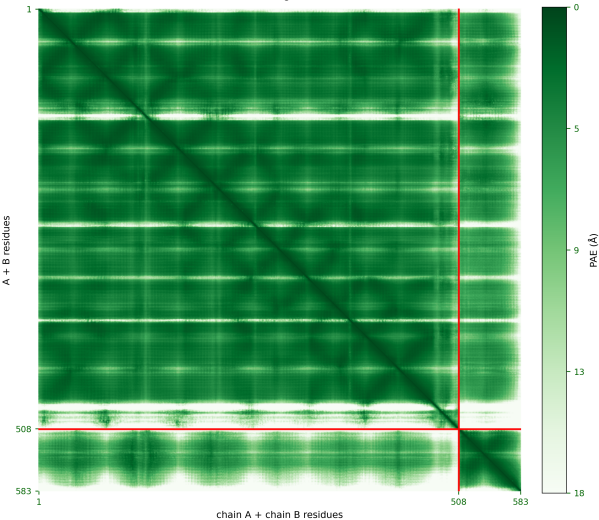
chain A + chain B residues

P00395 MT-CO1 | Q8N5G0 SMIM20
Best interchain (off-diagonal) PAE = 1.063 Å

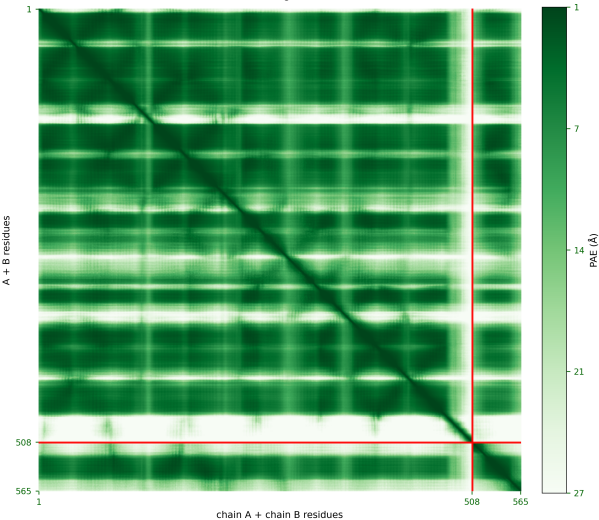


P00395 MT-CO1 | Q9NRP2 CMC2

Best interchain (off-diagonal) PAE = 1.305 Å

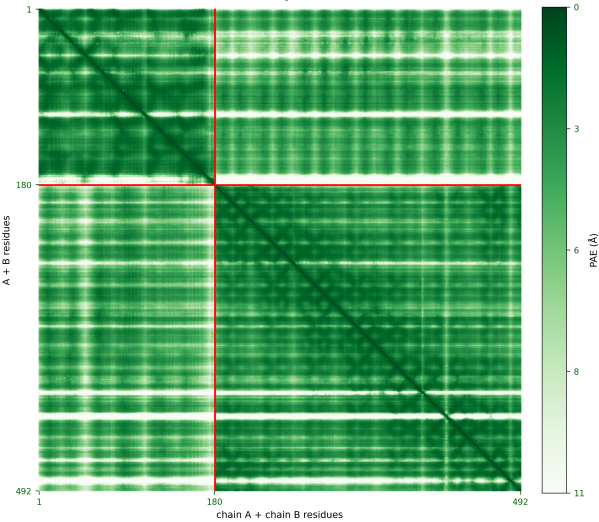


P00395 MT-CO1 | Q9Y2R0 COA3
Best interchain (off-diagonal) PAE = 1.085 Å



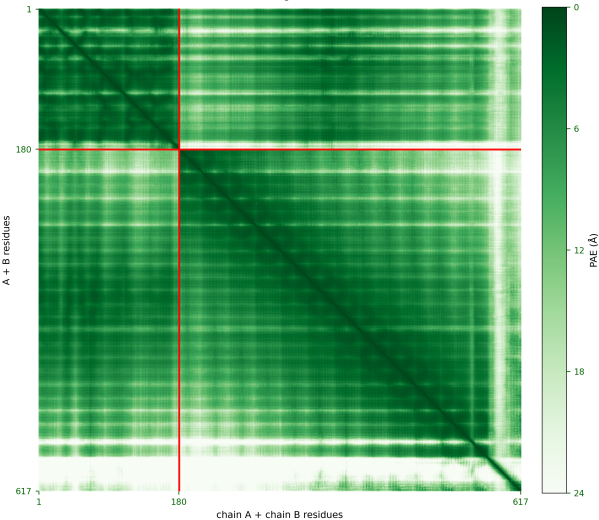
P51157 RAB28 | Q8NA23 WDR31

Best interchain (off-diagonal) PAE = 1.222 Å



P51157 RAB28 | Q96RK4 BBS4

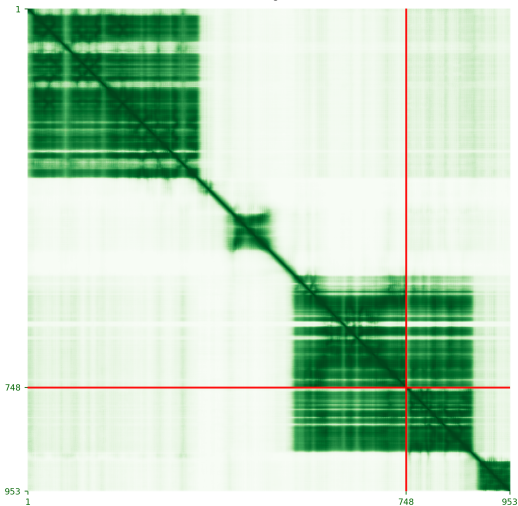
Best interchain (off-diagonal) PAE = 1.200 Å



P54277 PMS1 | Q9BTP7 FAAP24

Best interchain (off-diagonal) PAE = 1.394 Å

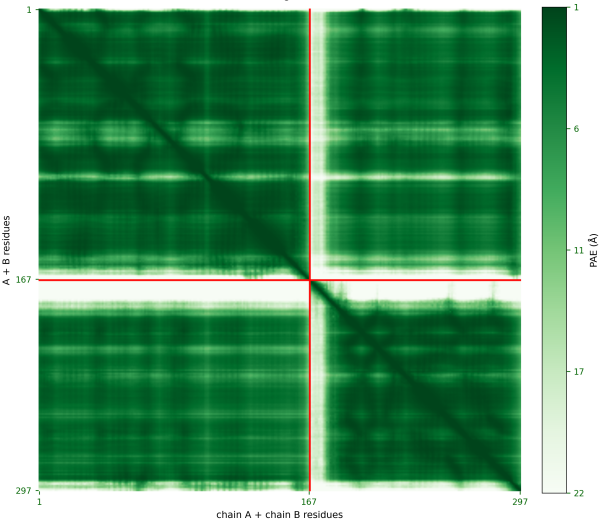
A + B residues



PAE (Å)

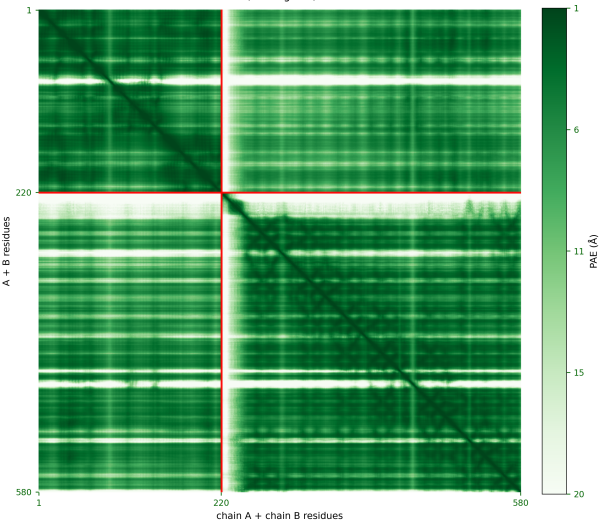
P58557 YBEY | P82912 MRPS11

Best interchain (off-diagonal) PAE = 0.880 Å

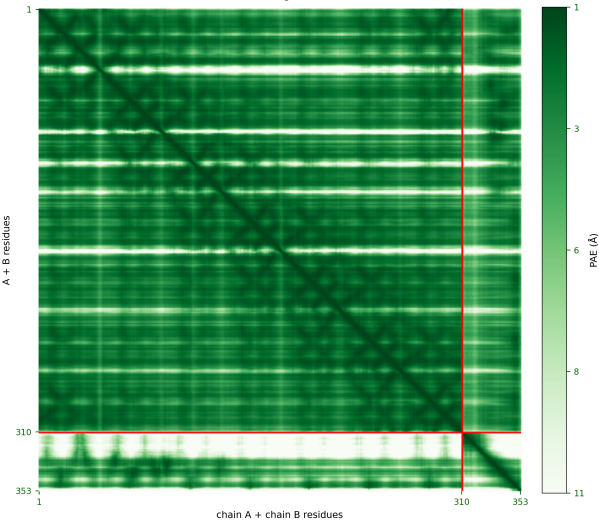


Q13216 ERCC8 | P49842 STK19

Best interchain (off-diagonal) PAE = 0.943 Å



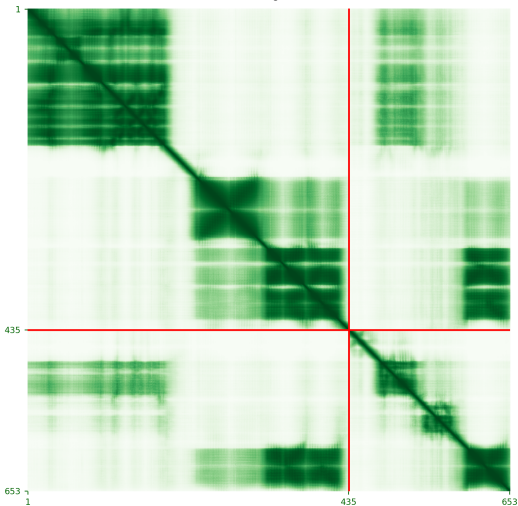
Q5I0X4 C6orf226 | O00628 PEX7
Best interchain (off-diagonal) PAE = 0.919 Å



Q5TKA1 LIN9 | Q9Y4I5 TESMIN

Best interchain (off-diagonal) PAE = 1.397 Å

A + B residues

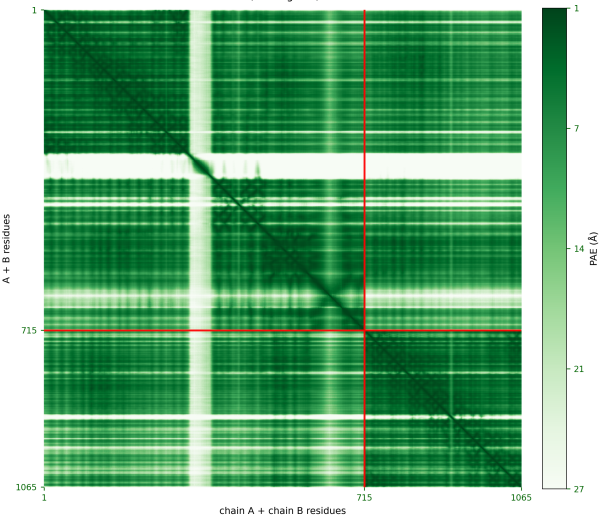


PAE (Å)

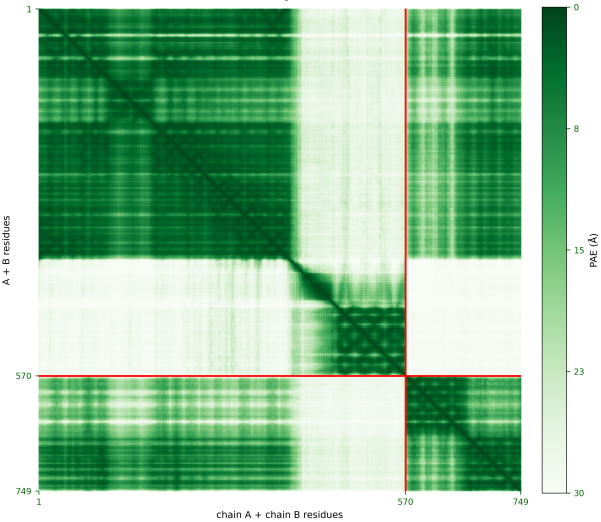
1
8
16
23
31

Q8IWZ6 BBS7 | Q8NEZ3 WDR19

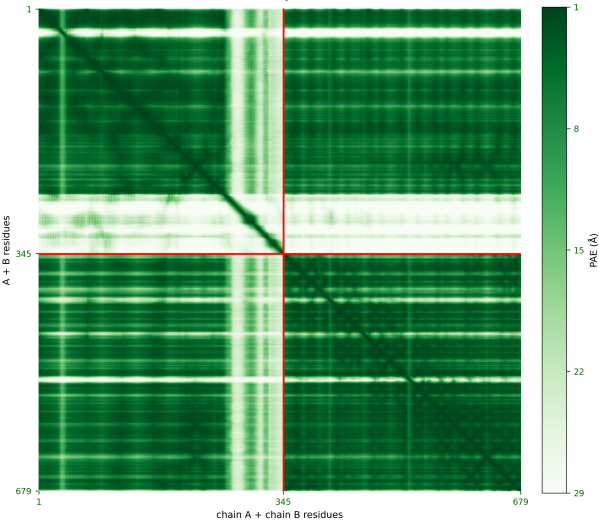
Best interchain (off-diagonal) PAE = 1.281 Å



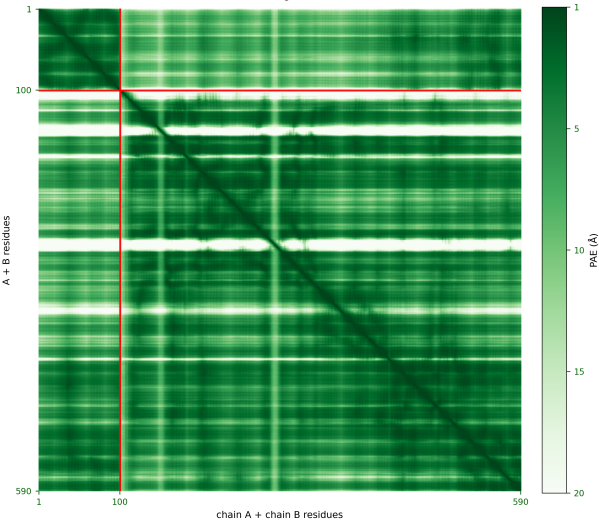
Q8NFJ9 BBS1 | Q9NVQ4 FAIM
Best interchain (off-diagonal) PAE = 1.015 Å



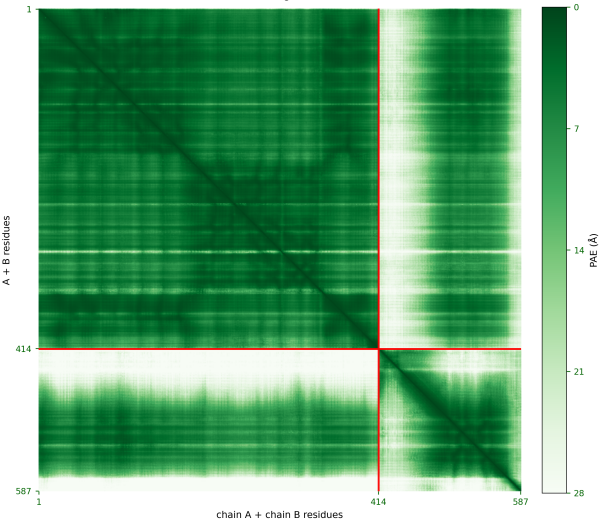
Q8WVF1 OSCP1 | Q9H977 WDR54
Best interchain (off-diagonal) PAE = 0.802 Å



Q99622 C12orf57 | Q9ULJ7 ANKRD50
Best interchain (off-diagonal) PAE = 1.086 Å



Q9BV19 C1orf50 | P23526 AHCY
Best interchain (off-diagonal) PAE = 1.276 Å



Q9BX26 SYCP2 | Q86X24 HORMAD1
Best interchain (off-diagonal) PAE = 1.132 Å

A + B residues

