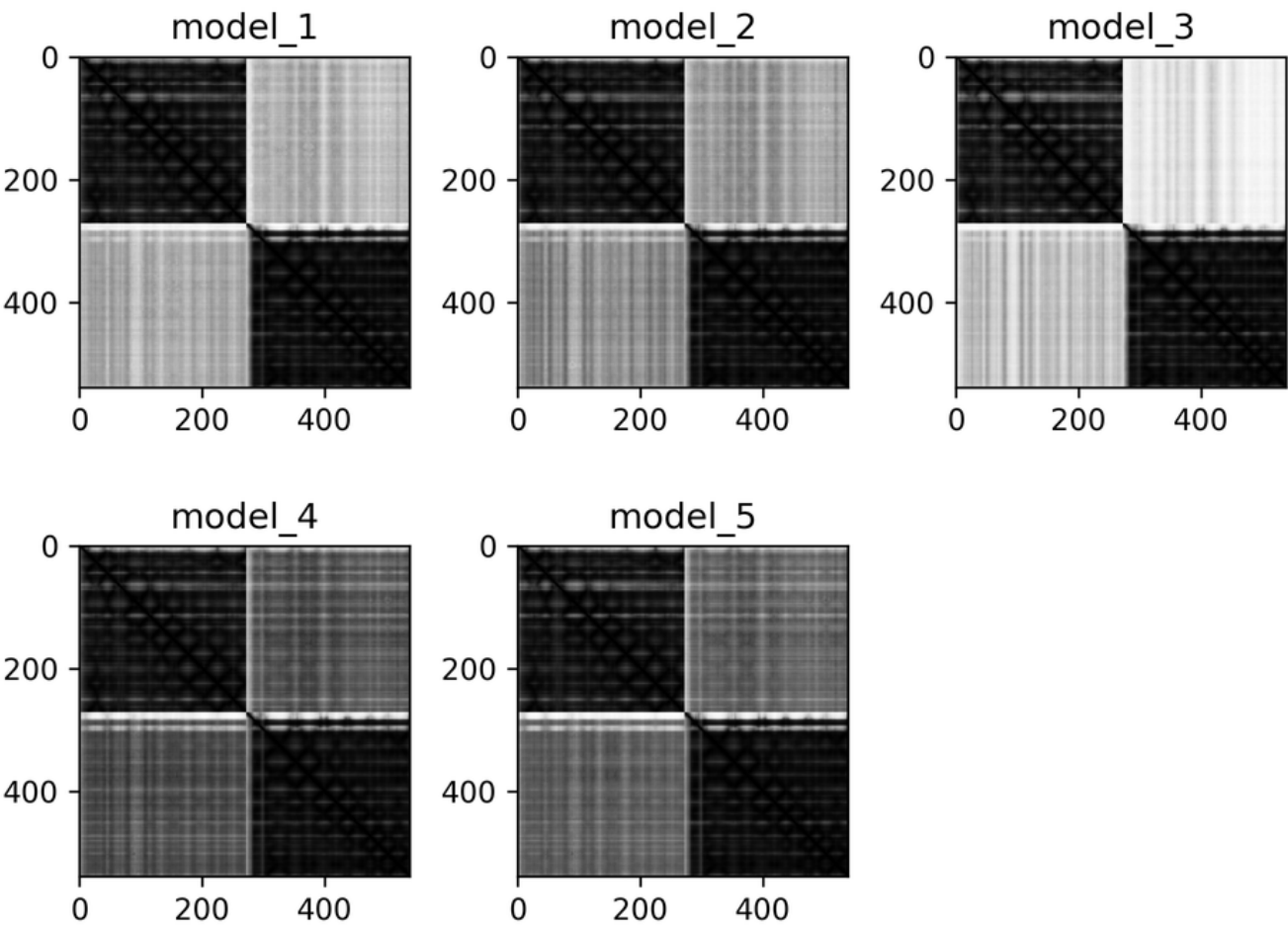


P0A927_P0A921 Tsx_OmPLA
residue length: 272 267

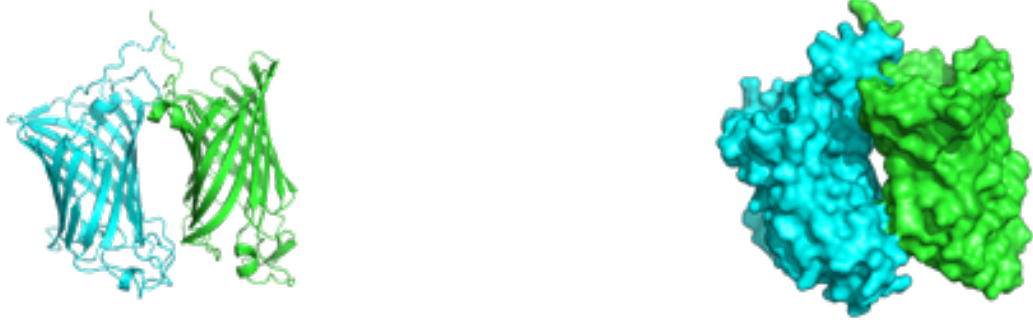


parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	92.36	92.43	91.49	93.13	92.81	92.444
inter-PAE	22.33	19.27	26.5	11.8	13.63	18.706
intra-PAE	4.49	4.43	4.88	4.49	4.56	4.57

model 1



model 2



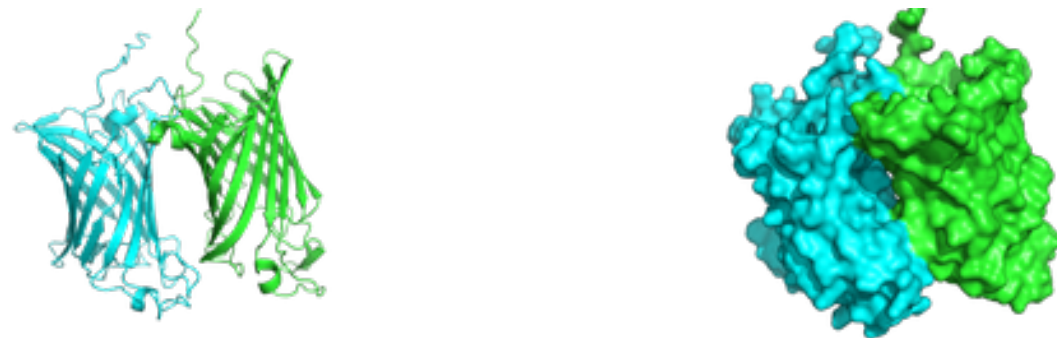
model 3



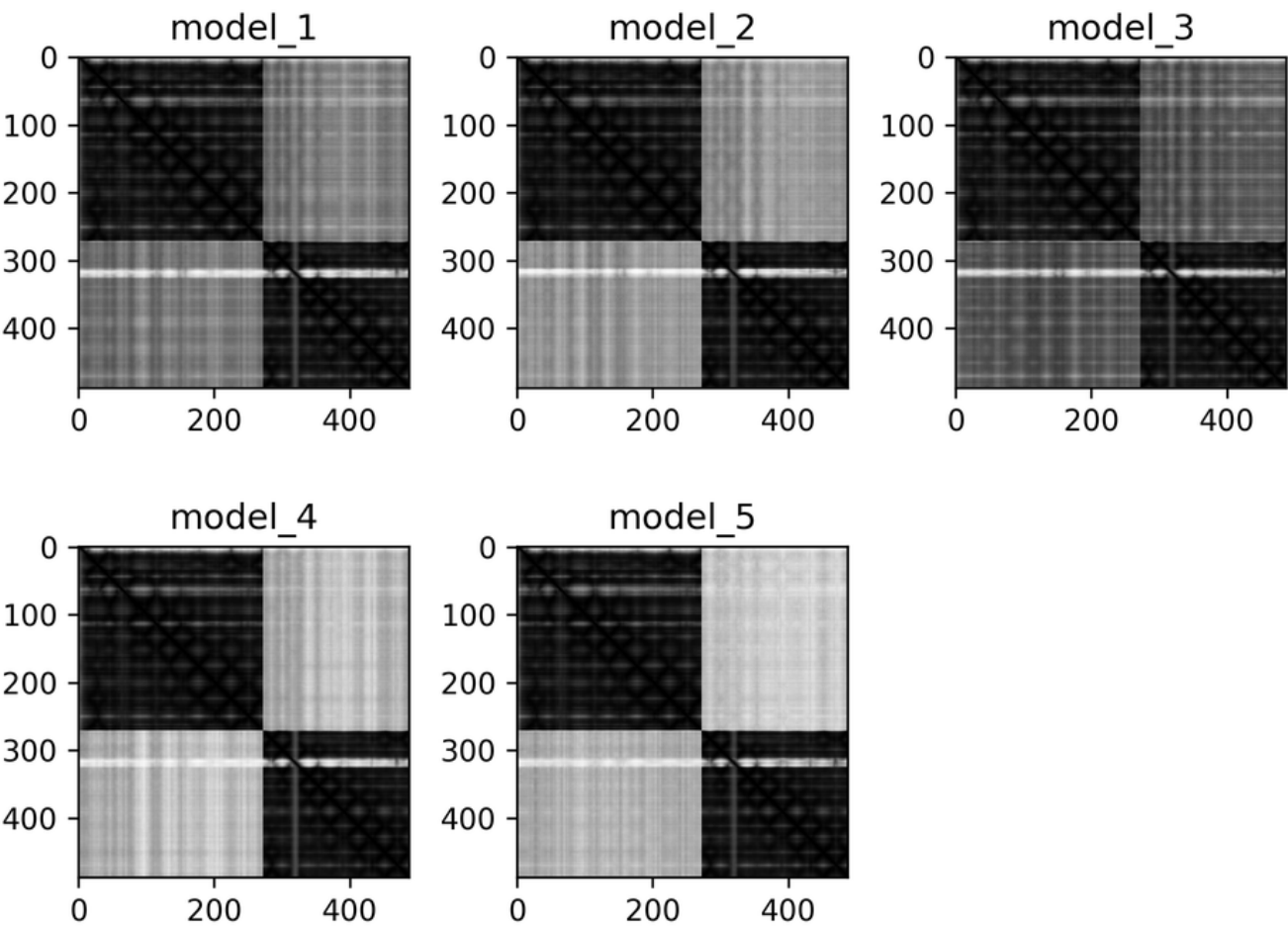
model 4



model 5



P0A927_P69856 Tsx_NanC
residue length: 272 216



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	91.19	91.37	91.19	91.98	92.02	91.55
inter-PAE	15.7	18.91	11.96	22.74	23.45	18.552
intra-PAE	4.44	4.27	4.48	4.38	4.46	4.406

model 1



model 2



model 3



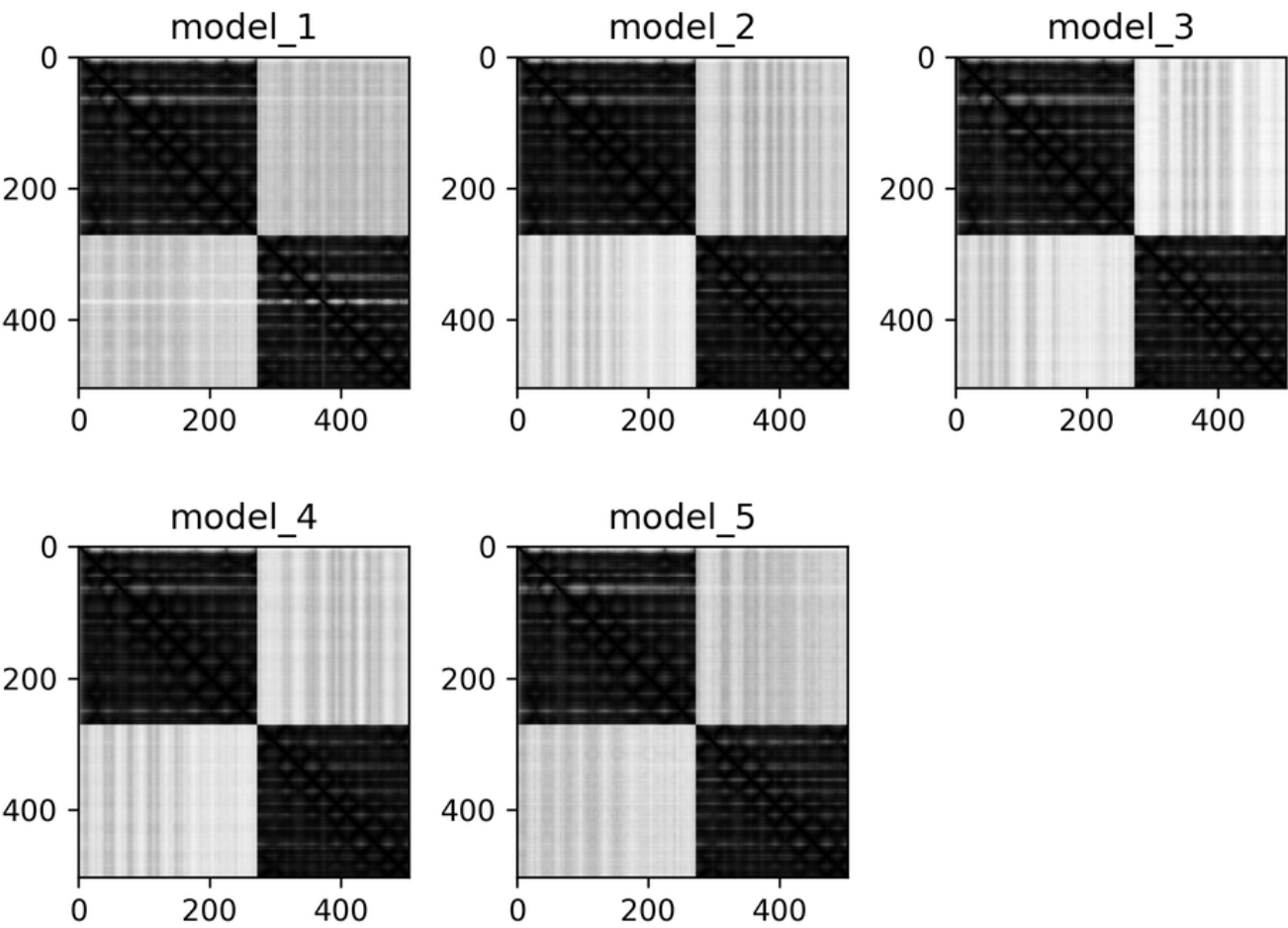
model 4



model 5



P0A927_P45564 Tsx_YfeN
residue length: 272 232



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	92.03	93.38	90.97	93.98	92.83	92.638
inter-PAE	24.59	25.48	27.36	26.15	25.26	25.768
intra-PAE	4.28	3.75	4.17	3.73	4.16	4.018

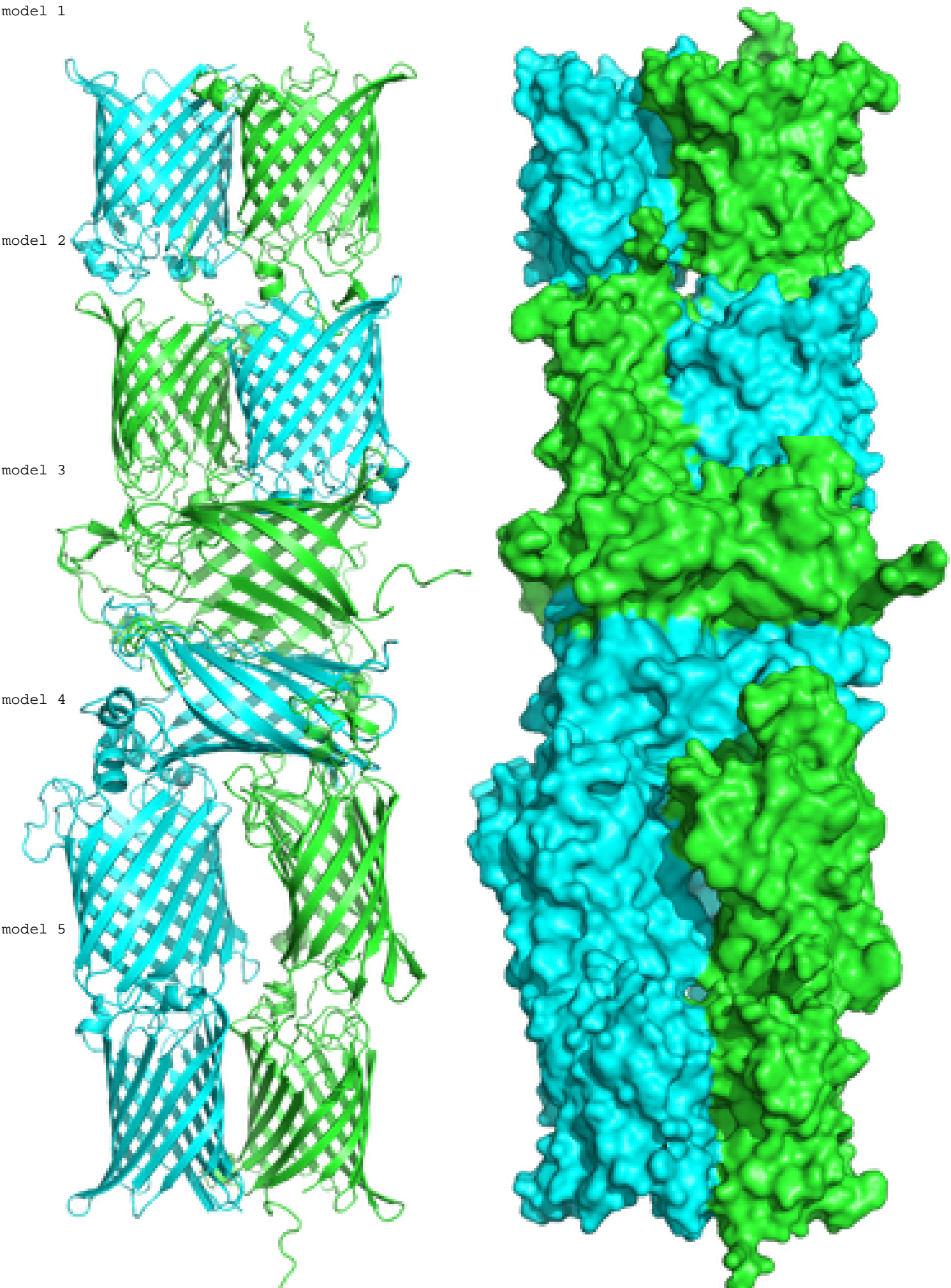
model 1

model 2

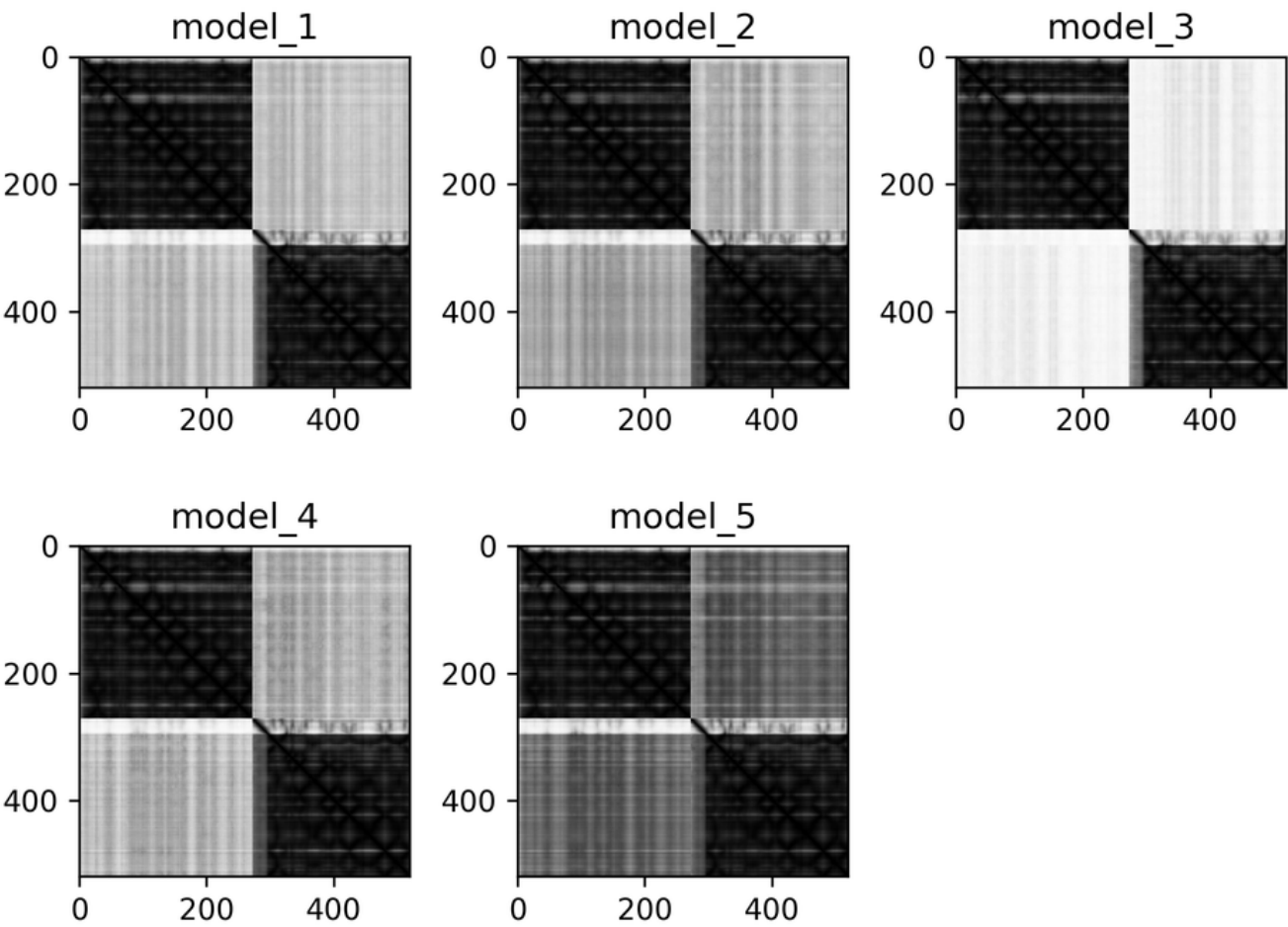
model 3

model 4

model 5



P0A927_P0A908 Tsx_MipA
residue length: 272 248



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	90.02	90.26	89.7	90.99	91.34	90.462
inter-PAE	24.19	21.94	29.49	22.42	13.8	22.368
intra-PAE	5.13	5.01	5.43	5.07	5.15	5.158

model 1



model 2



model 3



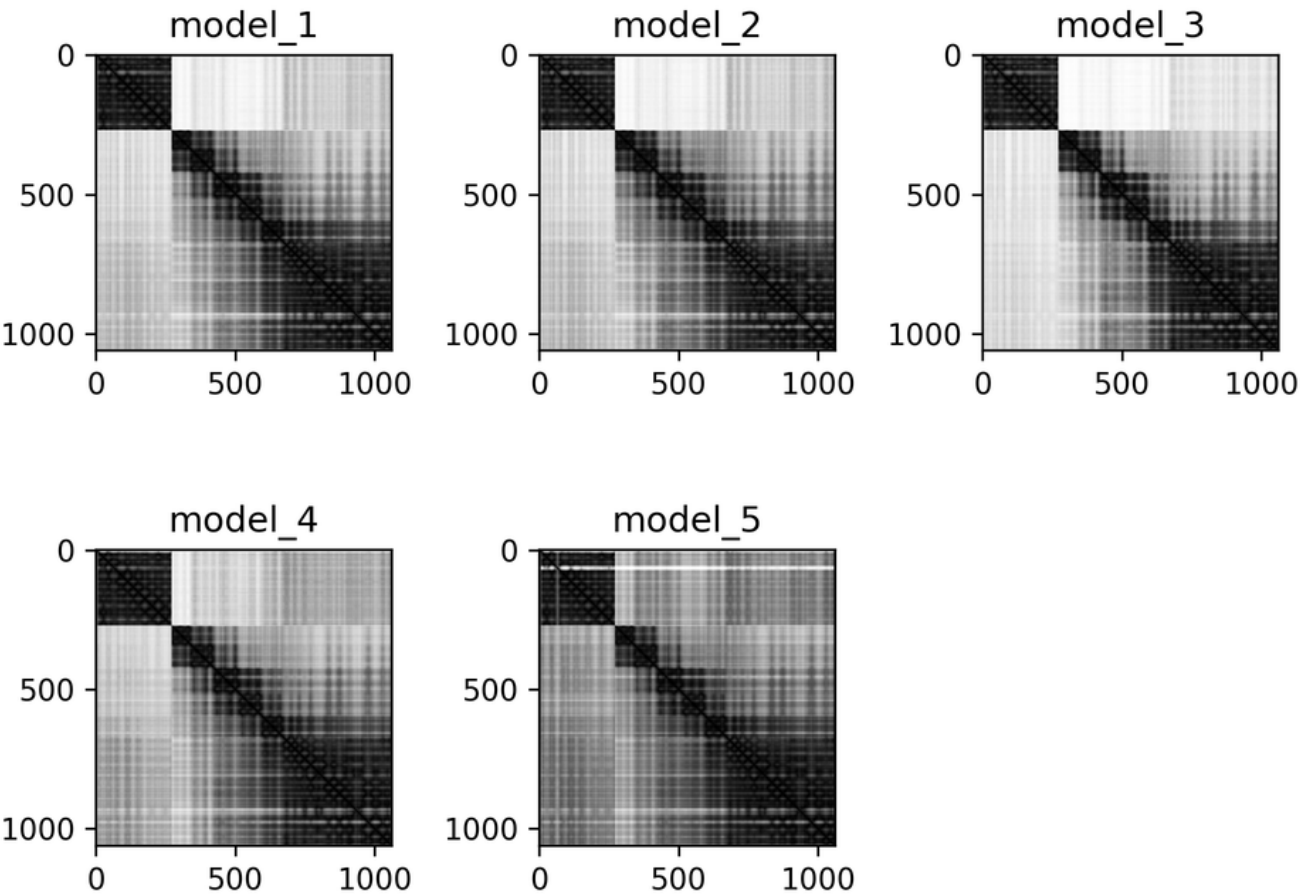
model 4



model 5



P0A927_P0A940 Tsx_BamA
residue length: 272 788



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	88.72	87.91	86.78	89.8	89.94	88.63
inter-PAE	26.5	26.66	28.76	23.34	18.12	24.676
intra-PAE	12.33	12.25	13.23	12.05	10.91	12.154

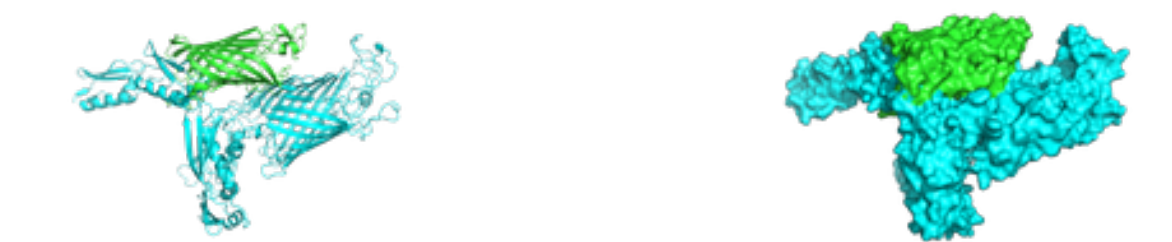
model 1



model 2



model 3



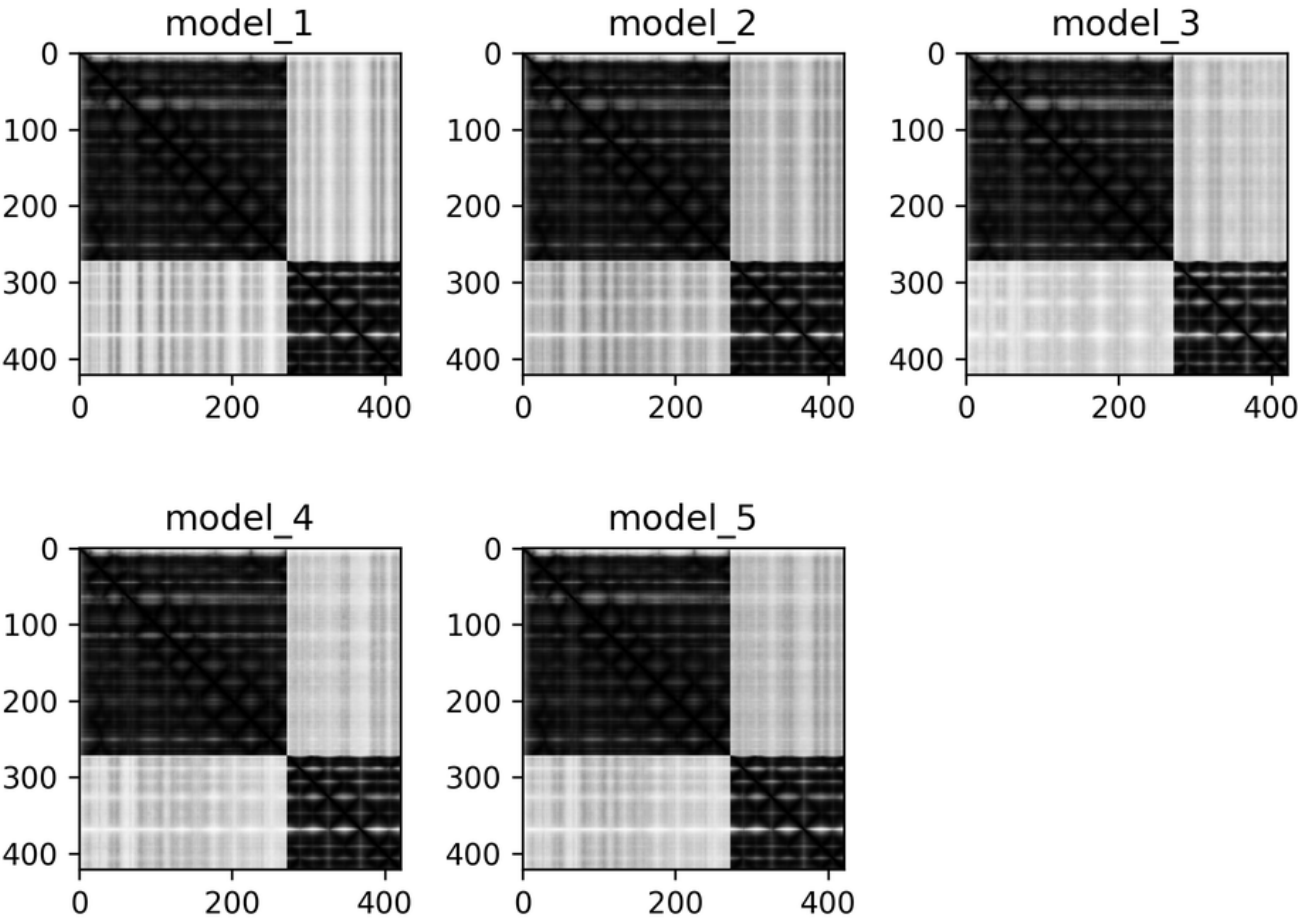
model 4



model 5



P0A927_P0A917 Tsx_OmpX
residue length: 272 149



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	91.63	91.16	88.61	91.81	91.75	90.992
inter-PAE	25.26	23.13	25.8	25.35	24.6	24.828
intra-PAE	4.69	4.77	5.13	4.76	4.77	4.824

model 1



model 2



model 3



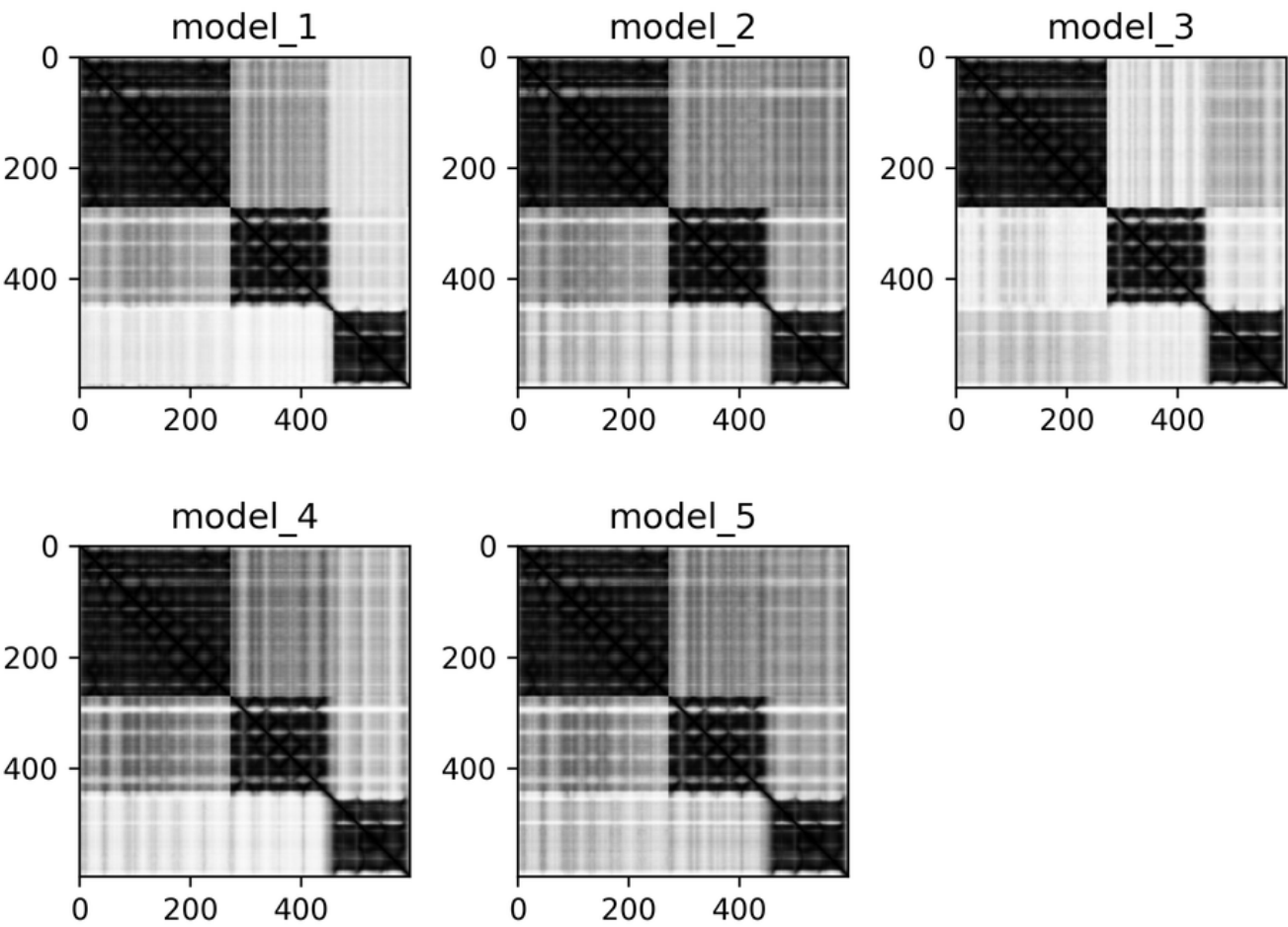
model 4



model 5



P0A927_P0A910 Tsx_OmpA
residue length: 272 324



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	85.83	87.01	84.06	87.65	87.09	86.328
inter-PAE	24.16	20.38	26.53	21.61	20.45	22.626
intra-PAE	12.28	10.9	12.41	11.77	10.95	11.662

model 1



model 2



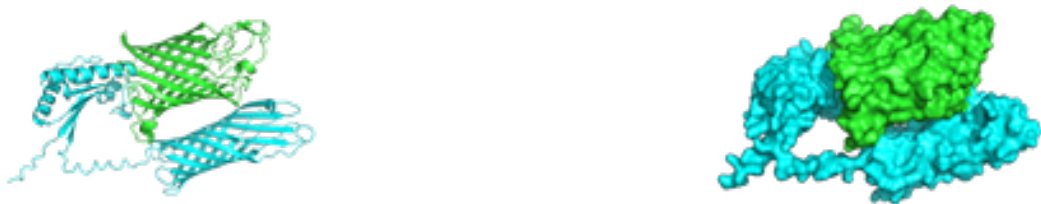
model 3



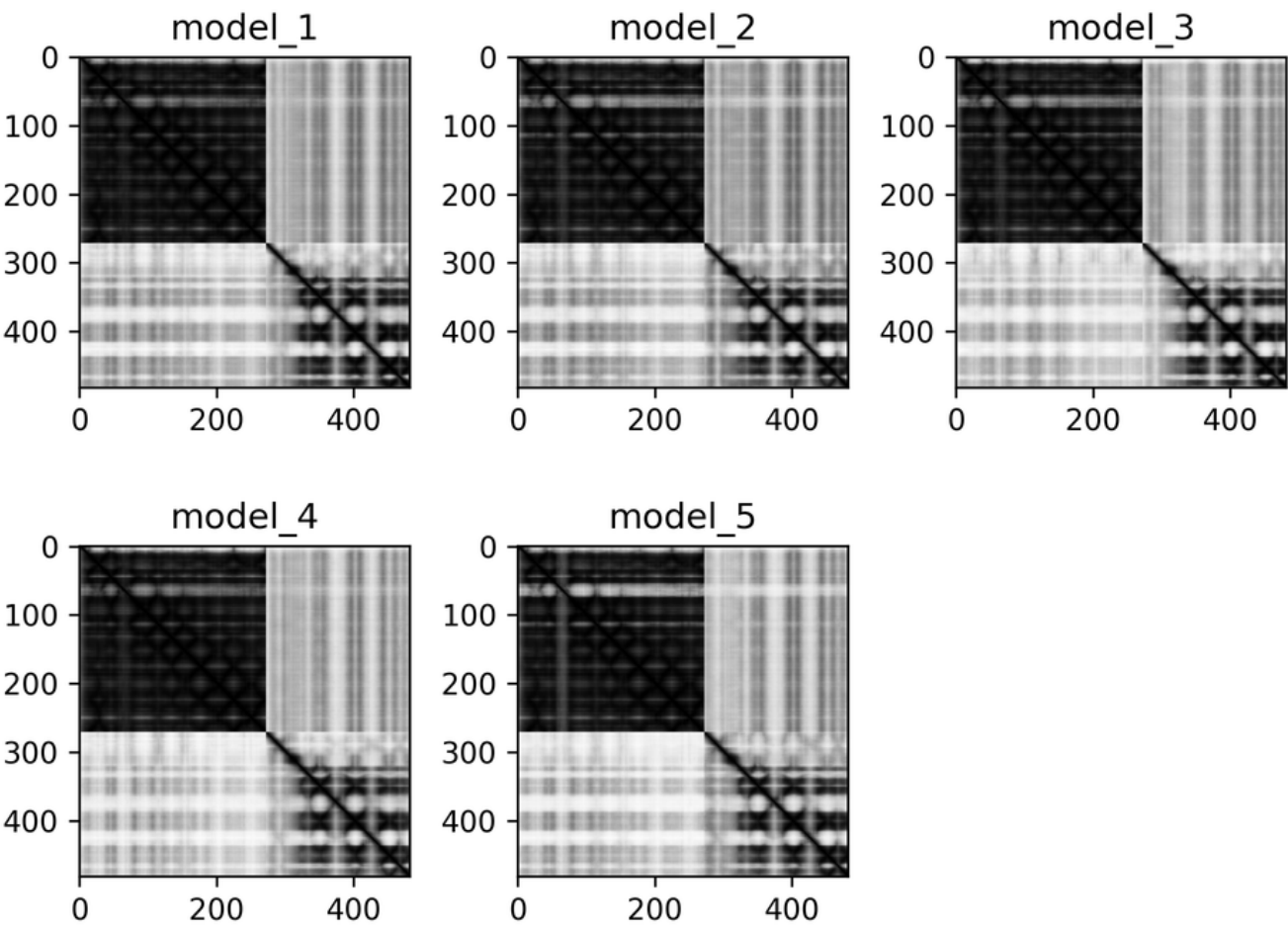
model 4



model 5



P0A927_P37663 Tsx_YhjY
residue length: 272 210



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	76.88	76.09	72.74	79.8	78.33	76.768
inter-PAE	22.65	22.31	23.75	24.23	24.27	23.442
intra-PAE	8.71	9.24	9.74	9.25	9.69	9.326

model 1



model 2



model 3



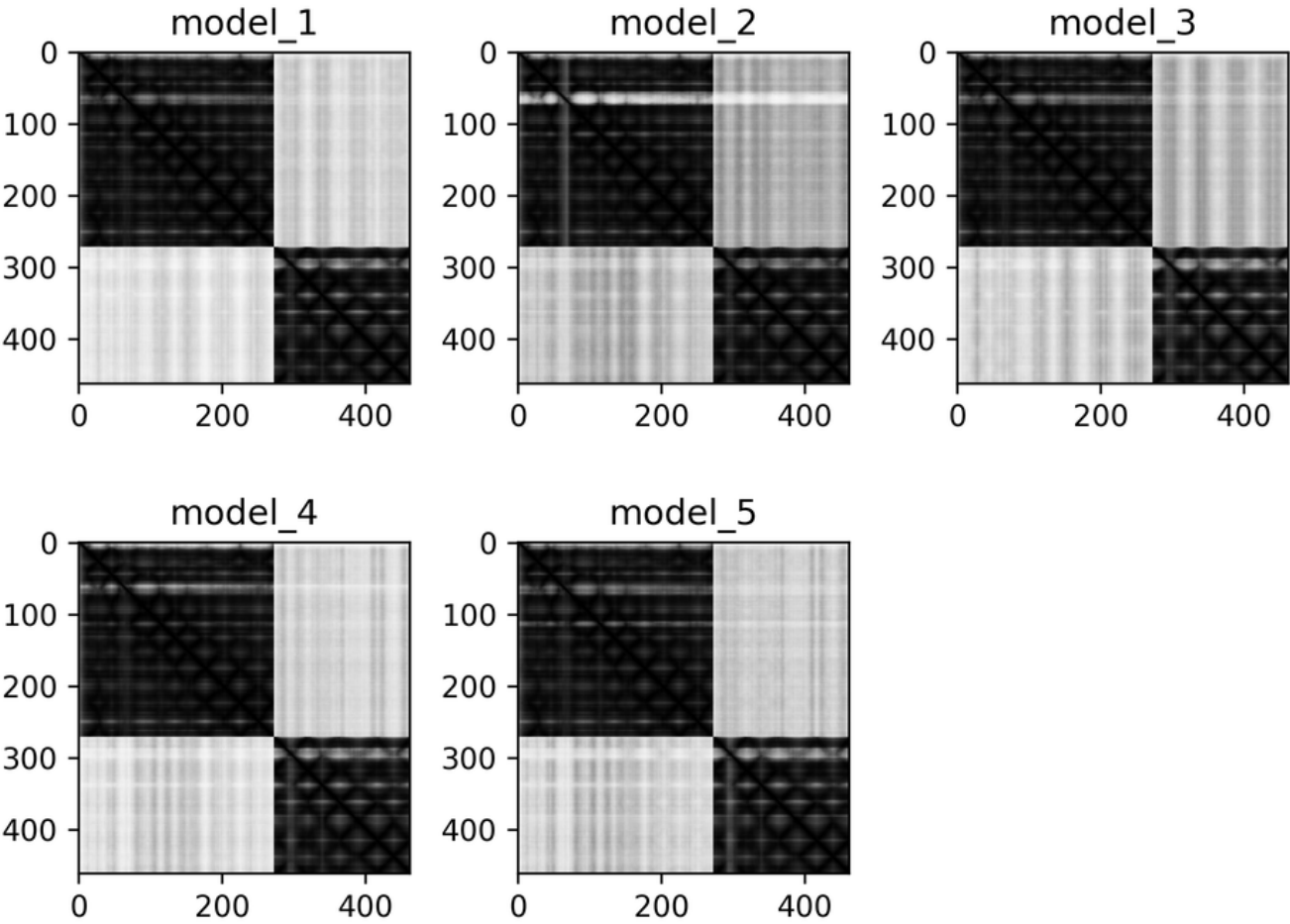
model 4



model 5



P0A927_P0A915 Tsx_OmpW
residue length: 272 190



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	92.14	91.17	89.58	92.32	92.13	91.468
inter-PAE	26.97	22.81	24.55	26.37	25.41	25.222
intra-PAE	4.47	4.95	4.79	4.76	4.81	4.756

model 1



model 2



model 3



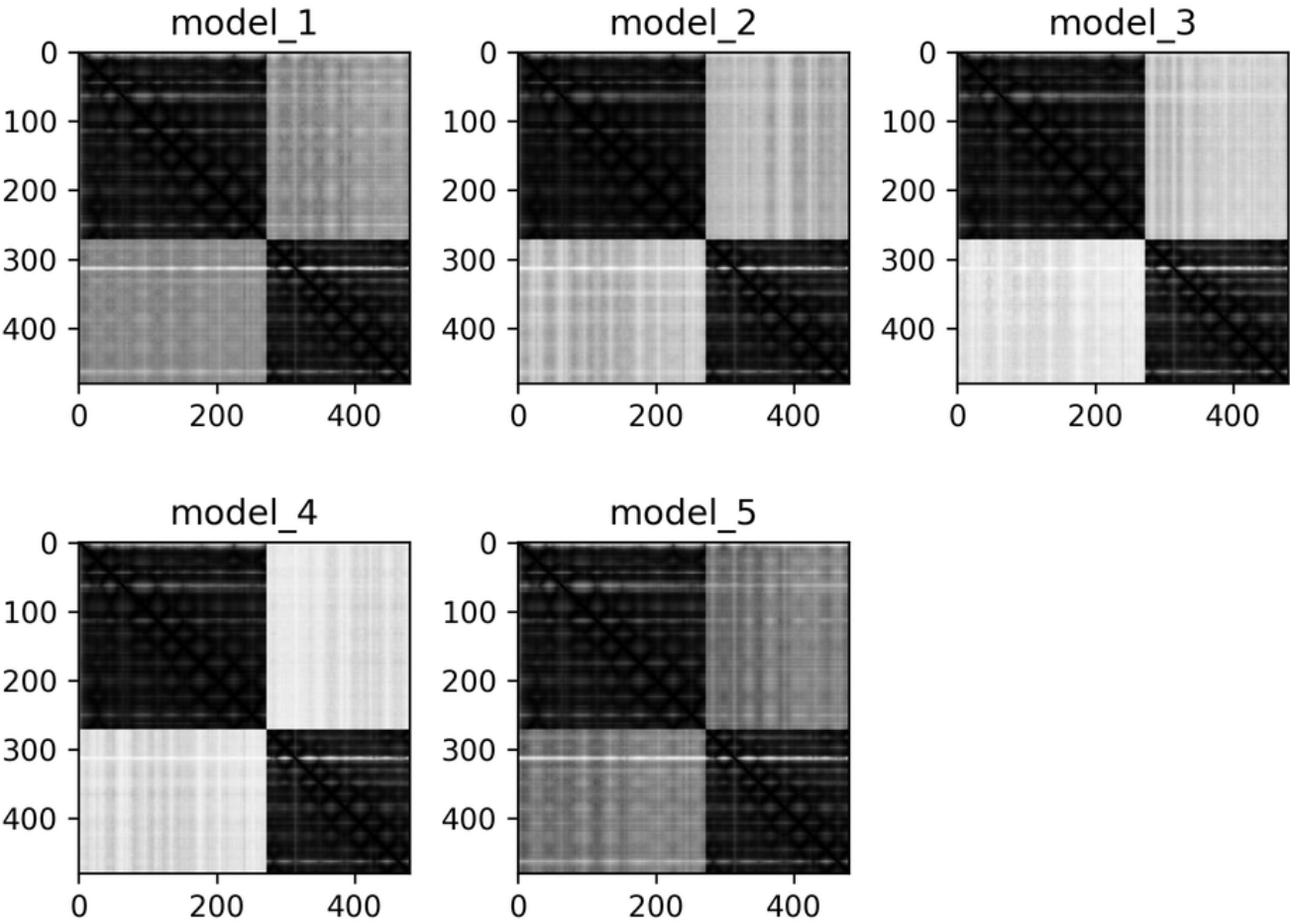
model 4



model 5



P0A927_P76773 Tsx_OmpL
residue length: 272 208



parameter	model_1	model_2	model_3	model_4	model_5	AVERAGE
pLLDT	92.15	92.08	91.11	93.14	92.75	92.246
inter-PAE	18.13	22.43	26.29	27.27	15.85	21.994
intra-PAE	4.08	3.78	4.24	4.03	4.25	4.076

model 1



model 2



model 3



model 4



model 5

