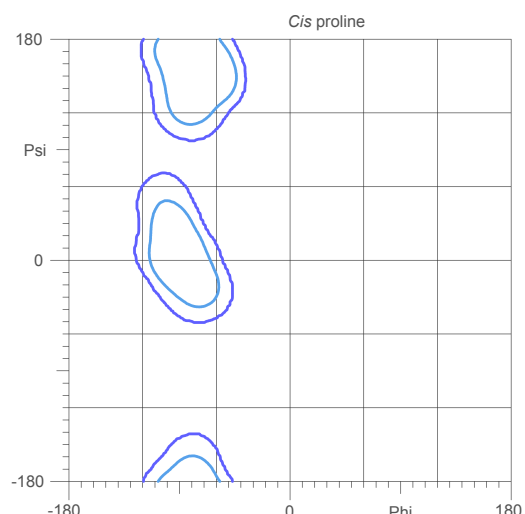
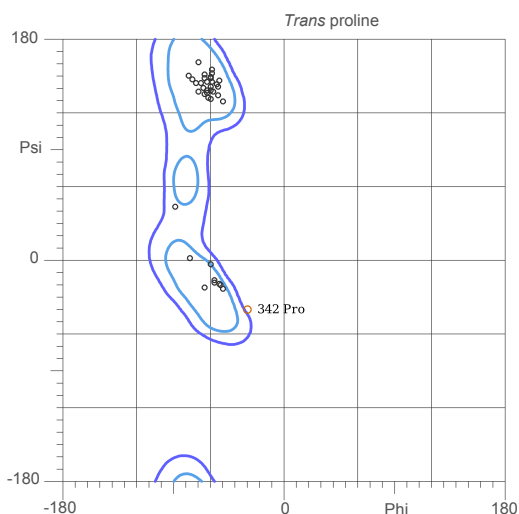
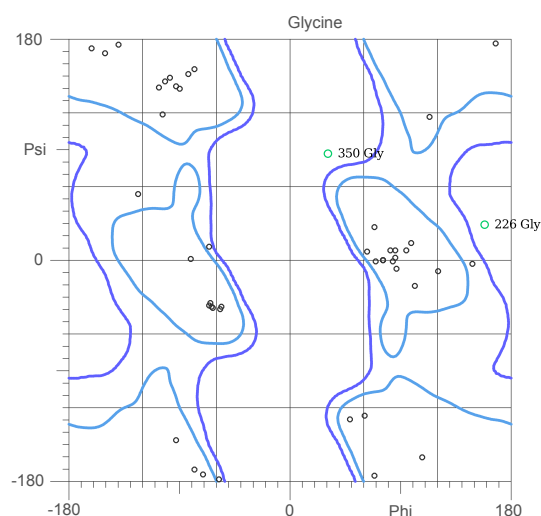
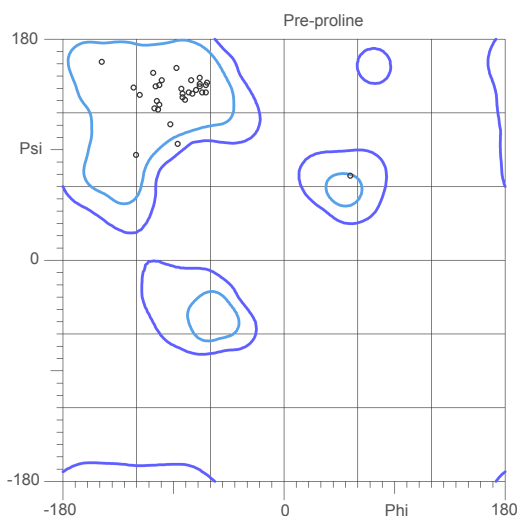
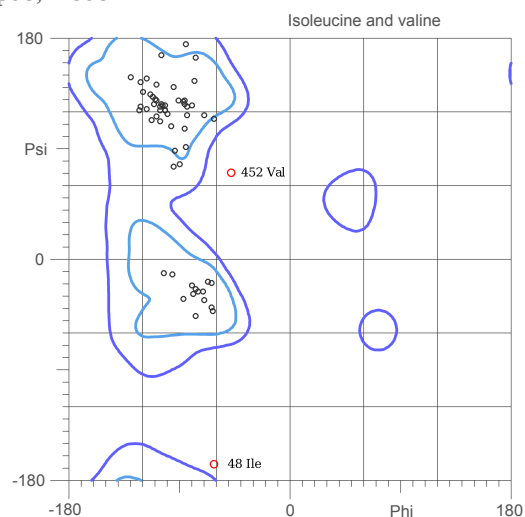
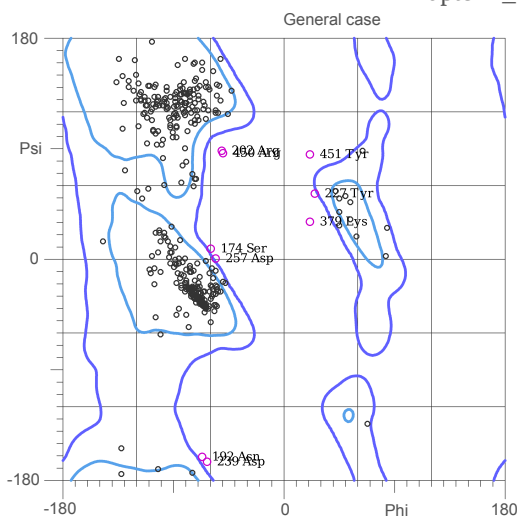


MolProbity Ramachandran analysis

RaptorX_TMPRS2.pdb, model 1



91.5% (482/527) of all residues were in favored (98%) regions.
97.3% (513/527) of all residues were in allowed (>99.8%) regions.

There were 14 outliers (phi, psi):

48 Ile (-63.0, -167.4)	342 Pro (-30.5, -40.5)
174 Ser (-60.3, 9.1)	350 Gly (31.5, 87.4)
192 Asn (-67.8, -161.7)	379 Lys (21.3, 31.1)
202 Arg (-51.9, 89.6)	450 Arg (-50.7, 87.5)
226 Gly (159.5, 29.2)	451 Tyr (21.4, 86.3)
227 Tyr (25.0, 54.8)	452 Val (-48.2, 72.0)
239 Asp (-63.6, -165.2)	
257 Asp (-56.8, 1.2)	