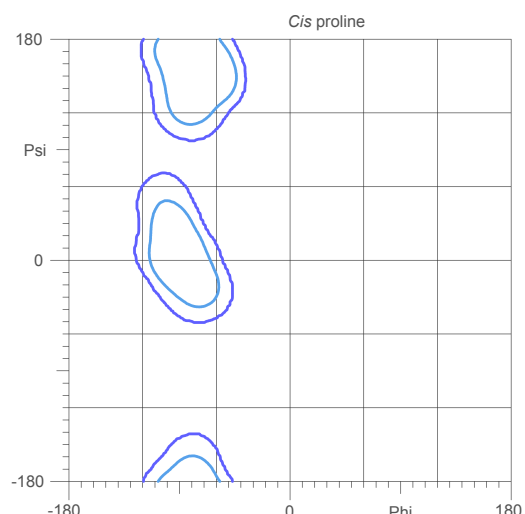
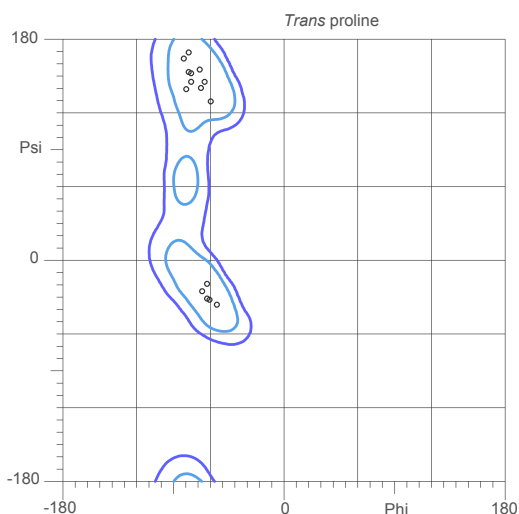
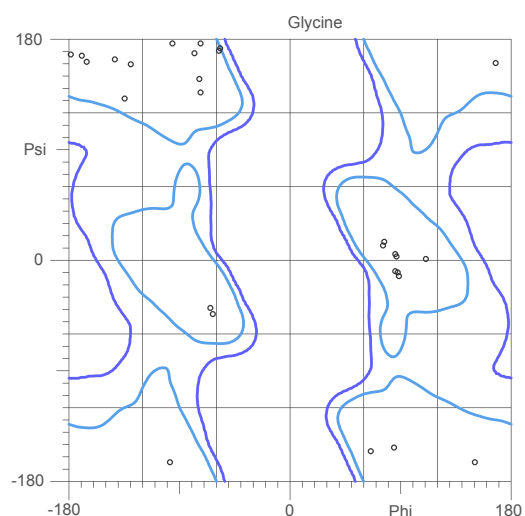
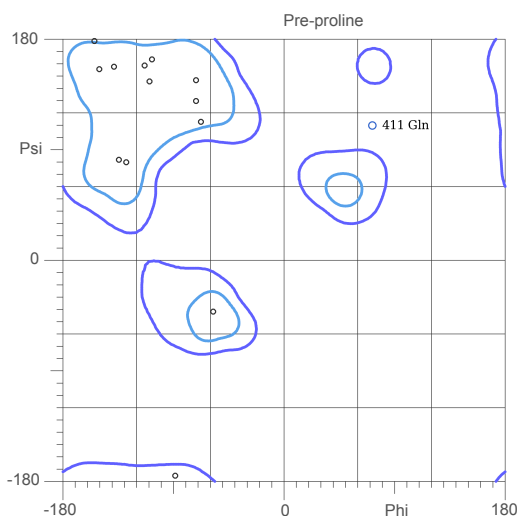
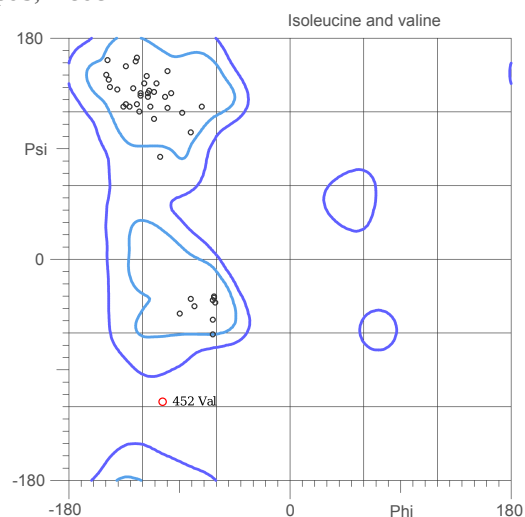
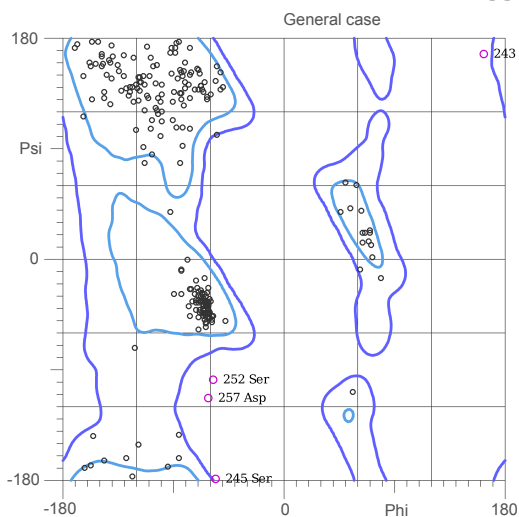


MolProbity Ramachandran analysis

TMPRSS2_HHPred.pdb, model 1



92.5% (307/332) of all residues were in favored (98%) regions.
98.2% (326/332) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):

243 Ser (163.2, 168.0)
245 Ser (-56.3, -179.4)
252 Ser (-58.6, -98.6)
257 Asp (-62.7, -113.5)
411 Gln (72.8, 110.1)
452 Val (-104.4, -117.0)