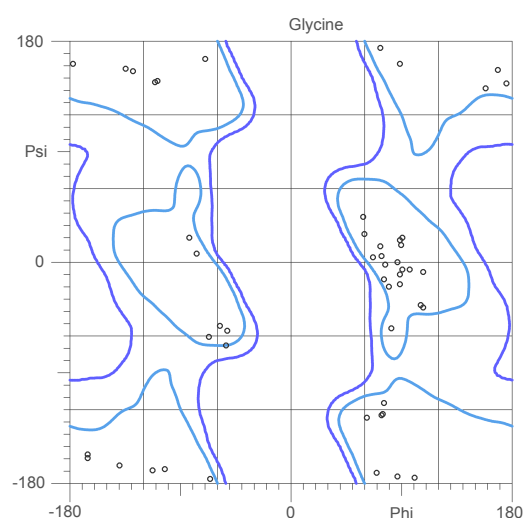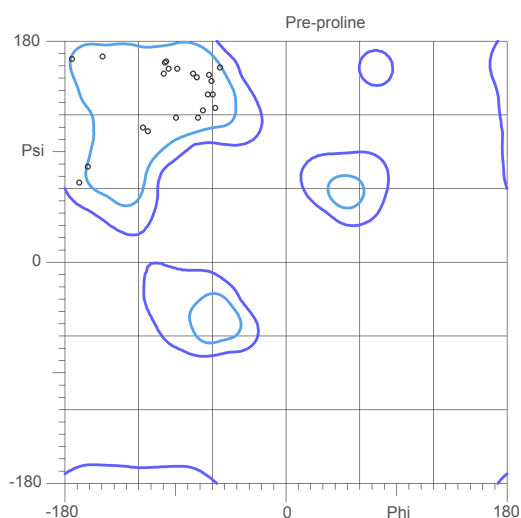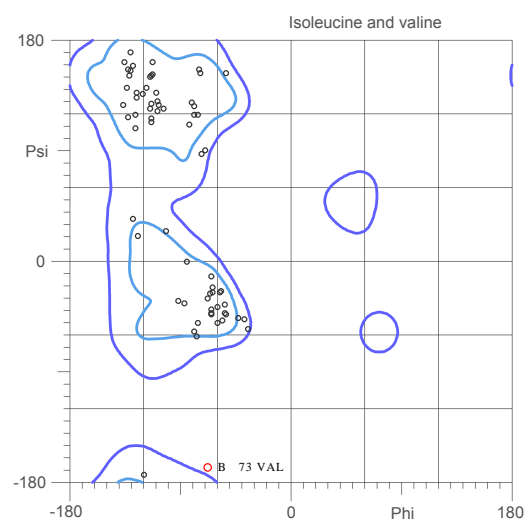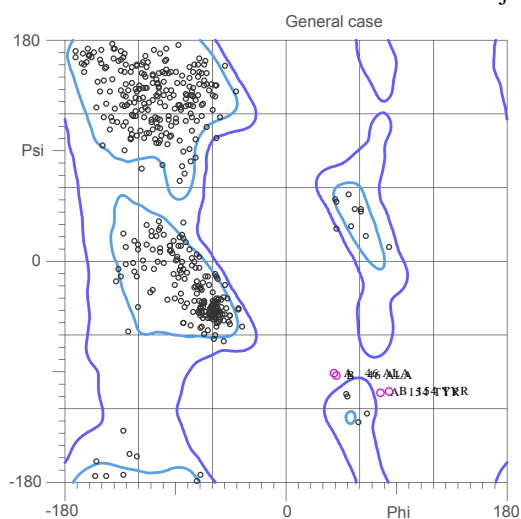# MolProbity Ramachandran analysis

3eaj.H.pdb, model 1
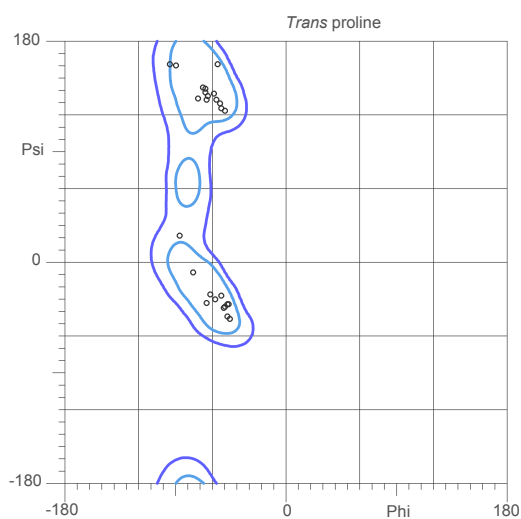
92.0% (552/600) of all residues were in favored (98%) regions.
99.2% (595/600) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    A   46 ALA (39.6, -91.8)
    A 154 TYR (77.3, -107.7)
    B   46 ALA (41.0, -93.9)
    B   73 VAL (-68.8, -168.5)
    B 154 TYR (84.4, -106.3)