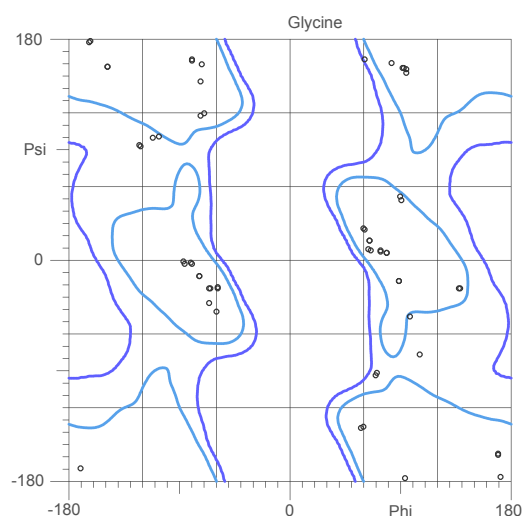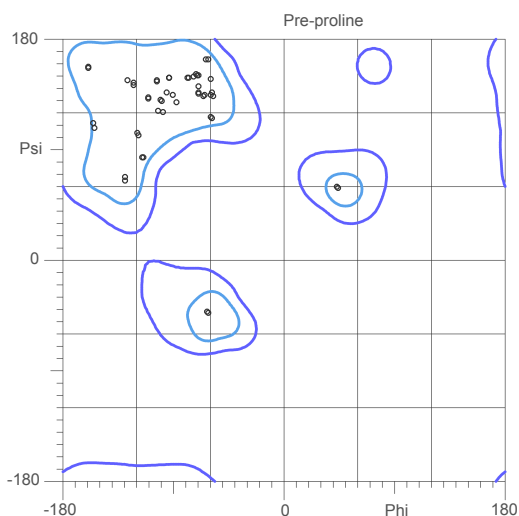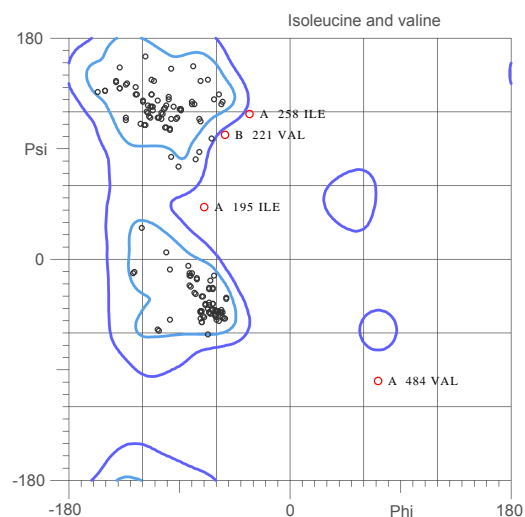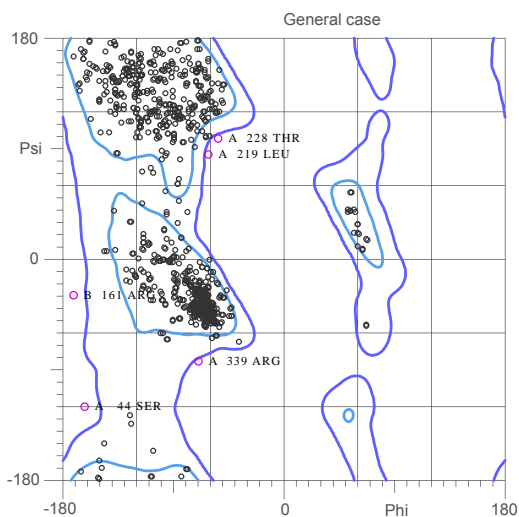# MolProbity Ramachandran analysis

5rlu.H.pdb, model 1
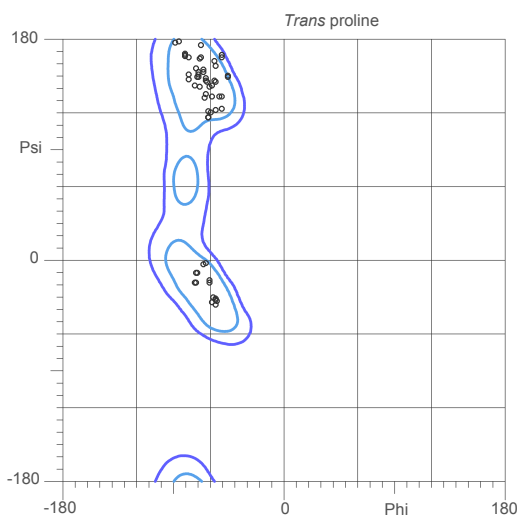
94.7% (1085/1146) of all residues were in favored (98%) regions.
99.2% (1137/1146) of all residues were in allowed (>99.8%) regions.

There were 9 outliers (phi, psi):
    A   44 SER (-163.1, -120.8)
    A 195 ILE (-70.2, 43.1)
    A 219 LEU (-62.0, 86.9)
    A 228 THR (-54.9, 99.1)
    A 258 ILE (-33.1, 119.6)
    A 339 ARG (-70.6, -83.6)
    A 484 VAL (72.5, -99.7)
    B 161 ARG (-172.1, -29.8)
    B 221 VAL (-53.8, 102.1)