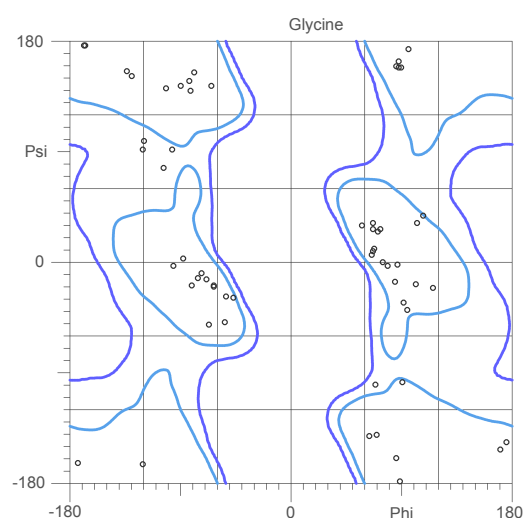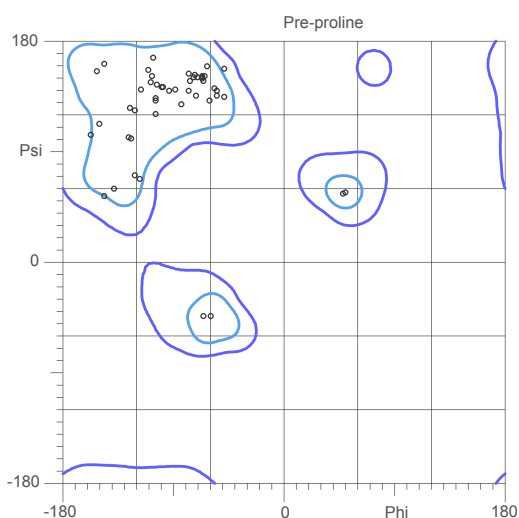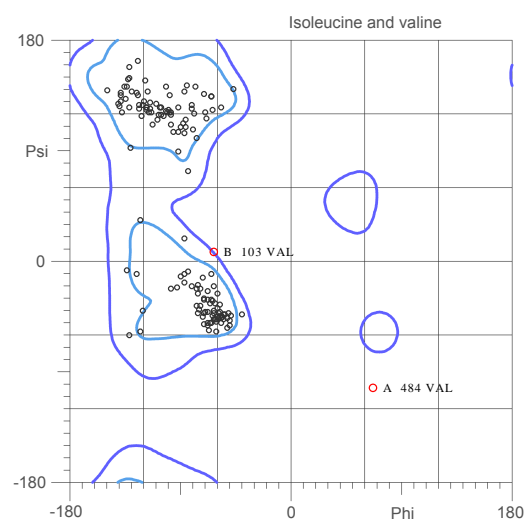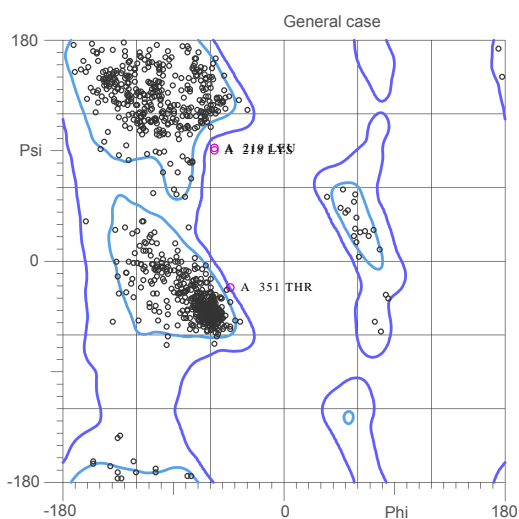# MolProbity Ramachandran analysis

5rm2.H.pdb, model 1
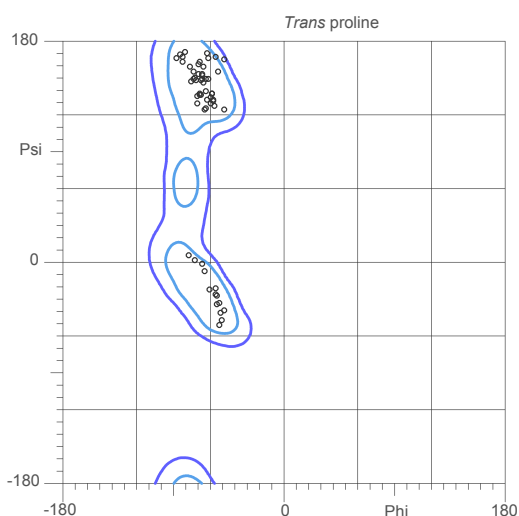
94.9% (1088/1146) of all residues were in favored (98%) regions.
99.6% (1141/1146) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    A 218 LYS (-57.3, 91.0)
    A 219 LEU (-57.6, 93.8)
    A 351 THR (-44.2, -21.7)
    A 484 VAL (67.3, -103.7)
    B 103 VAL (-63.9, 8.7)