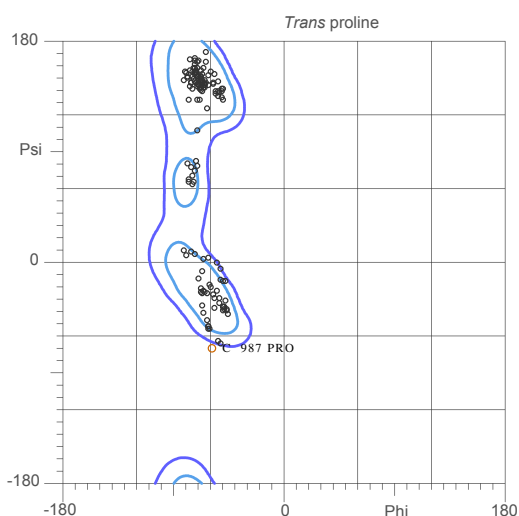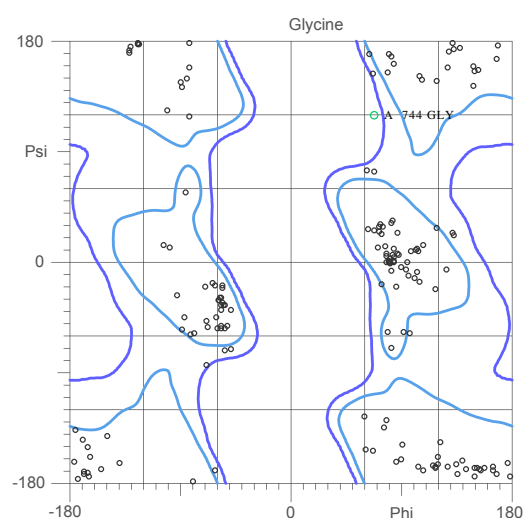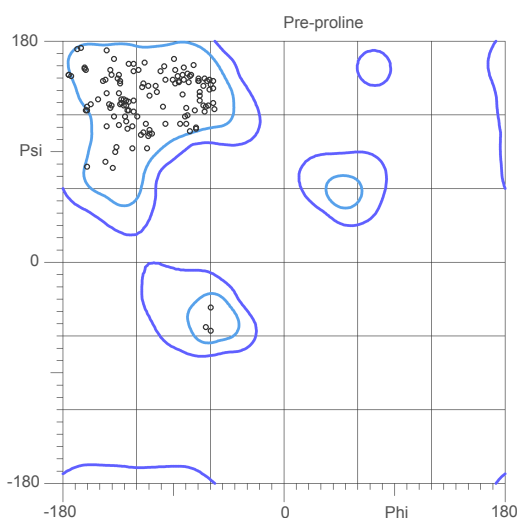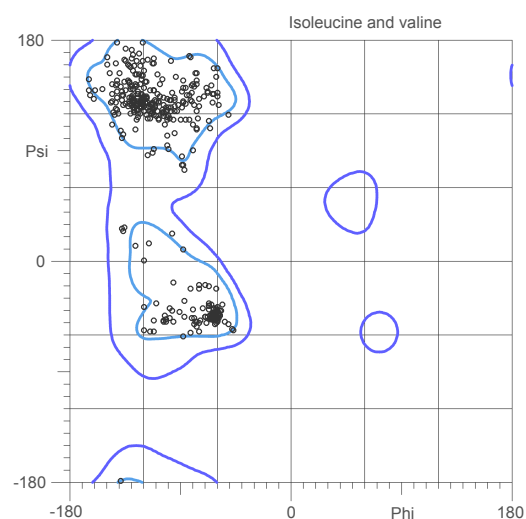# MolProbity Ramachandran analysis

6vsb.H.pdb, model 1

94.6% (2670/2821) of all residues were in favored (98%) regions.
99.8% (2816/2821) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
    A 744 GLY (68.1, 120.1)
    A 888 PHE (-50.5, -7.2)
    B 543 PHE (-59.4, 81.9)
    C 617 CYS (-64.2, 84.1)
    C 987 PRO (-59.9, -70.3)