MolProbity Ramachandran analysis

16pk_.B99990039.pdb, model 1
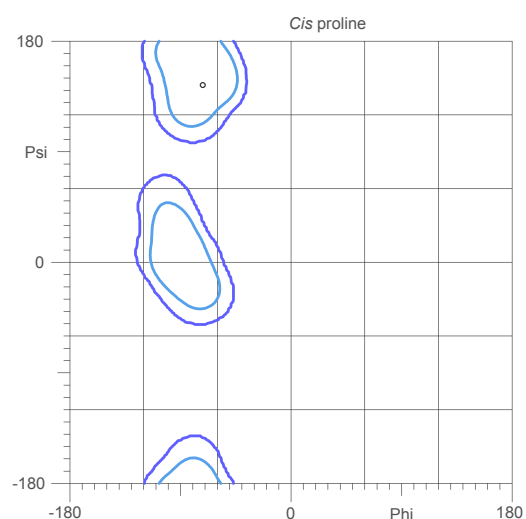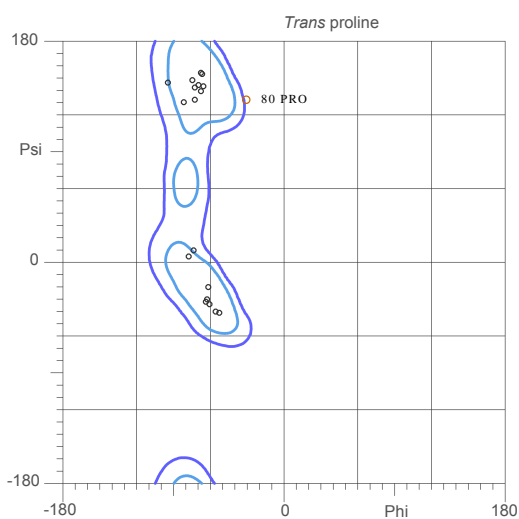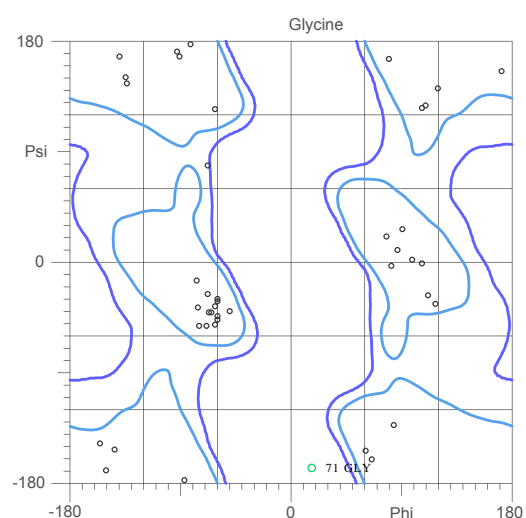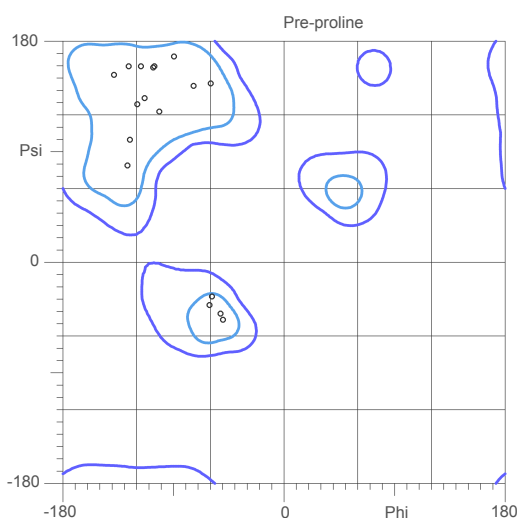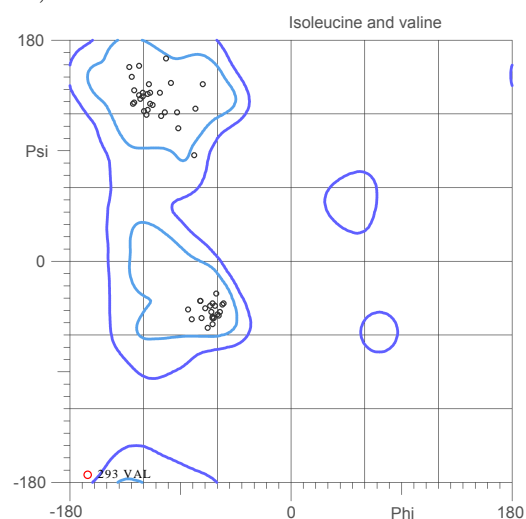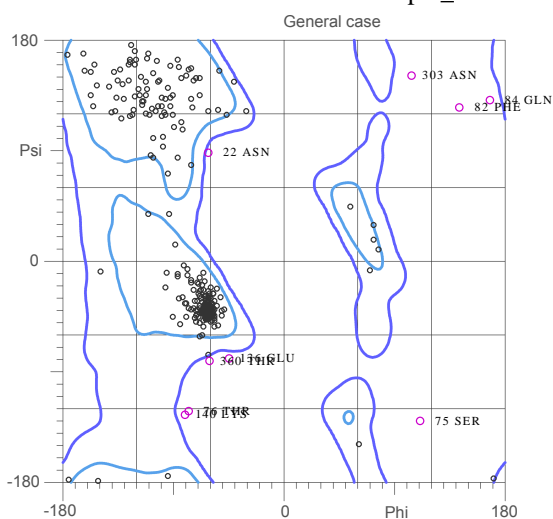

93,5% (386/413) of all residues were in favored (98%) regions.
97,1% (401/413) of all residues were in allowed (>99.8%) regions.

There were 12 outliers (phi, psi):
    22 ASN (-62,5, 89,9)
    71 GLY (17,9, -168,6)
    75 SER (111,0, -130,5)
    76 THR (-78,1, -122,4)
    80 PRO (-31,1, 133,6)
    82 PHE (143,2, 126,6)
    84 GLN (168,2, 132,2)
    136 GLU (-45,9, -80,0)
    140 LYS (-81,6, -125,7)
    293 VAL (-166,6, -174,3)
    303 ASN (104,8, 152,9)
    360 THR (-61,5, -81,7)