MolProbity Ramachandran analysis

1at3A.B99990009.pdb, model 1
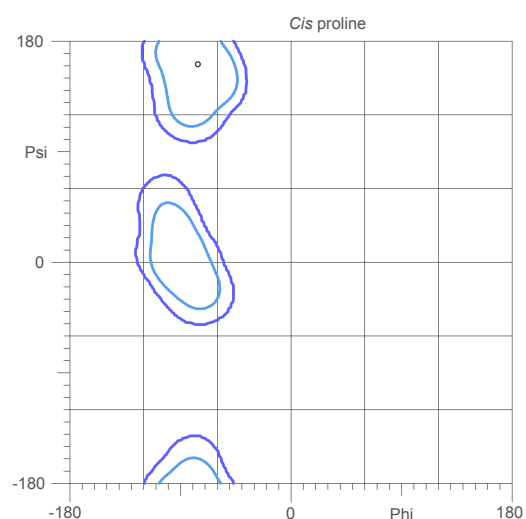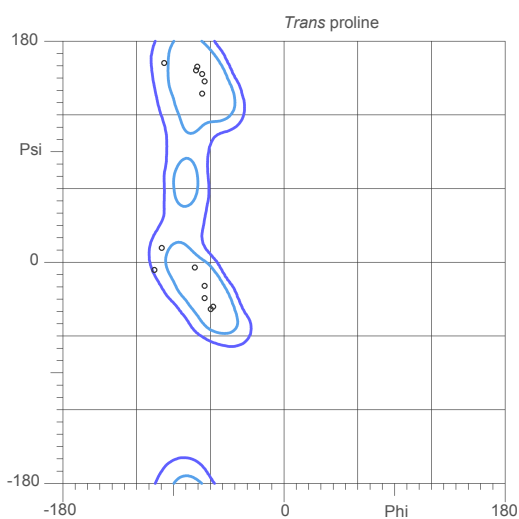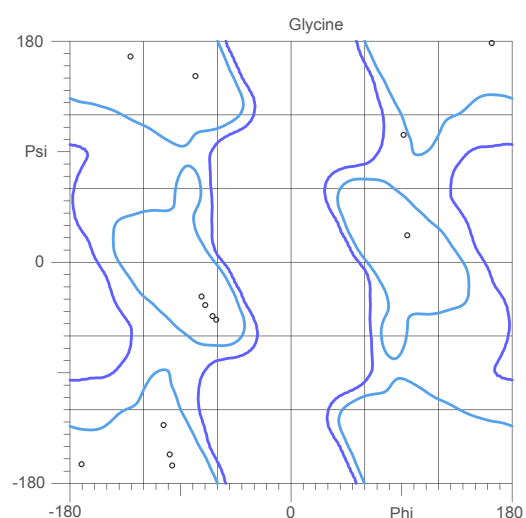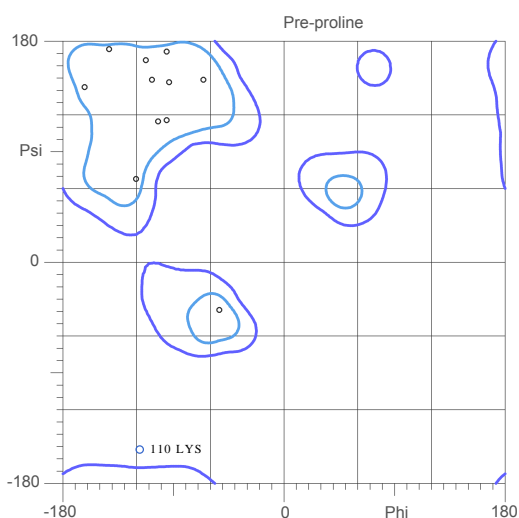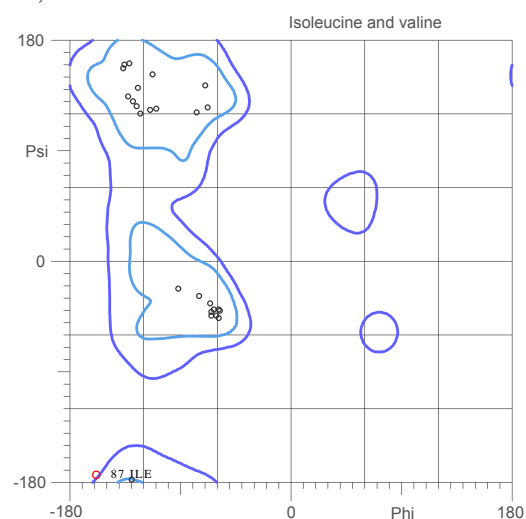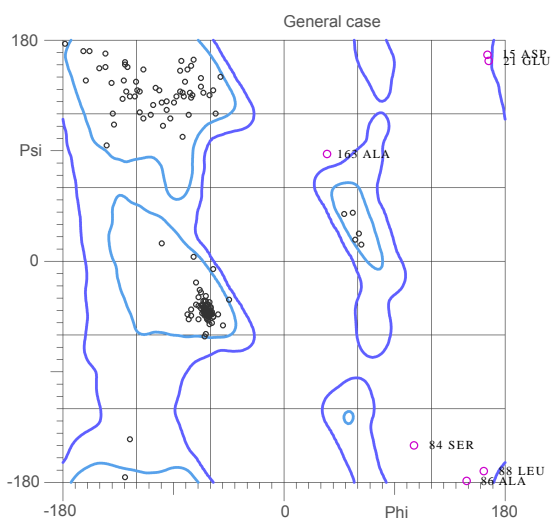

93,0% (200/215) of all residues were in favored (98%) regions.
96,3% (207/215) of all residues were in allowed (>99.8%) regions.

There were 8 outliers (phi, psi):
    15 ASP (166,5, 169,7)
    21 GLU (167,2, 164,1)
    84 SER (106,4, -150,1)
    86 ALA (149,1, -179,8)
    87 ILE (-159,4, -174,6)
    88 LEU (163,9, -171,9)
    110 LYS (-118,1, -153,8)
    163 ALA (35,7, 88,2)