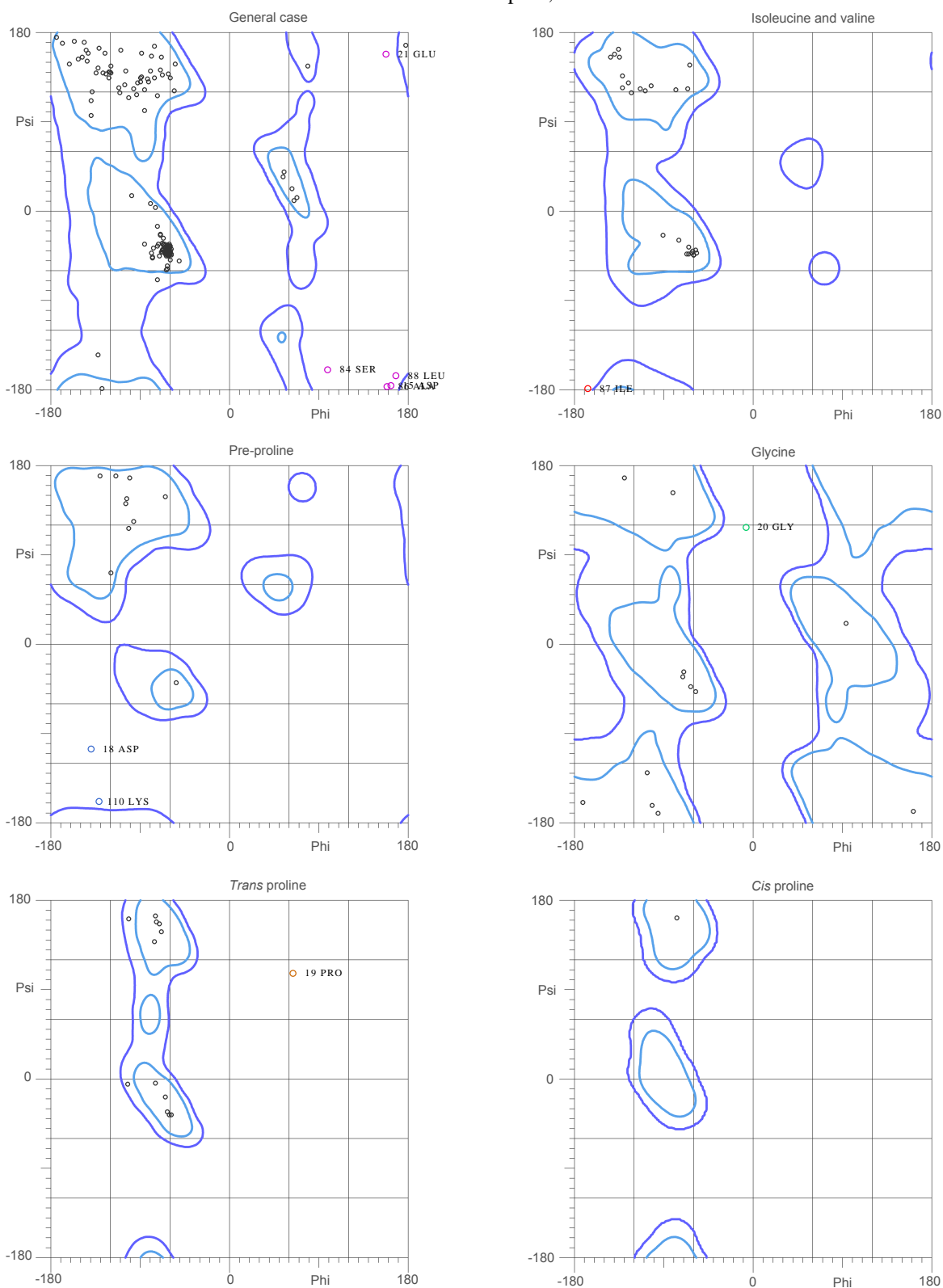# MolProbity Ramachandran analysis

1at3A.B99990013.pdb, model 1

92,1% (198/215) of all residues were in favored (98%) regions.
95,3% (205/215) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):
    15 ASP (163,2, -176,7)
    18 ASP (-140,3, -106,8)
    19 PRO (64,5, 107,7)
    20 GLY (-7,6, 118,7)
    21 GLU (158,6, 159,8)
    84 SER (99,8, -161,0)
    86 ALA (159,8, -177,7)
    87 ILE (-167,9, -179,8)
    88 LEU (168,8, -166,5)
    110 LYS (-132,4, -159,0)