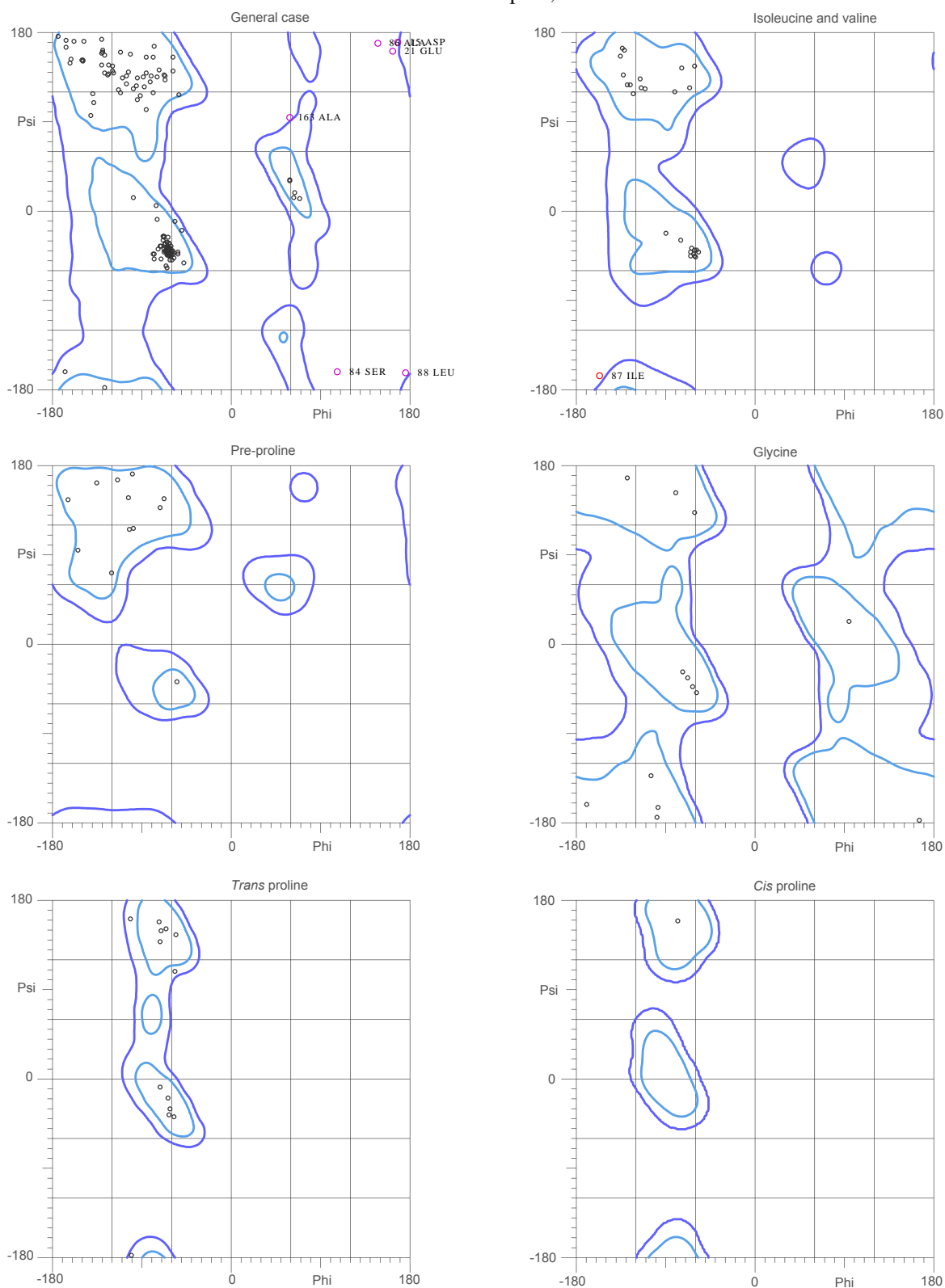# MolProbity Ramachandran analysis

1at3A.B99990010.pdb, model 1

93,5% (201/215) of all residues were in favored (98%) regions.
96,7% (208/215) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):
        15 ASP (168,2, 171,2)
        21 GLU (163,3, 162,8)
        84 SER (107,9, -162,6)
        86 ALA (148,3, 170,3)
        87 ILE (-157,5, -166,8)
        88 LEU (176,1, -163,6)
        163 ALA (59,0, 95,3)