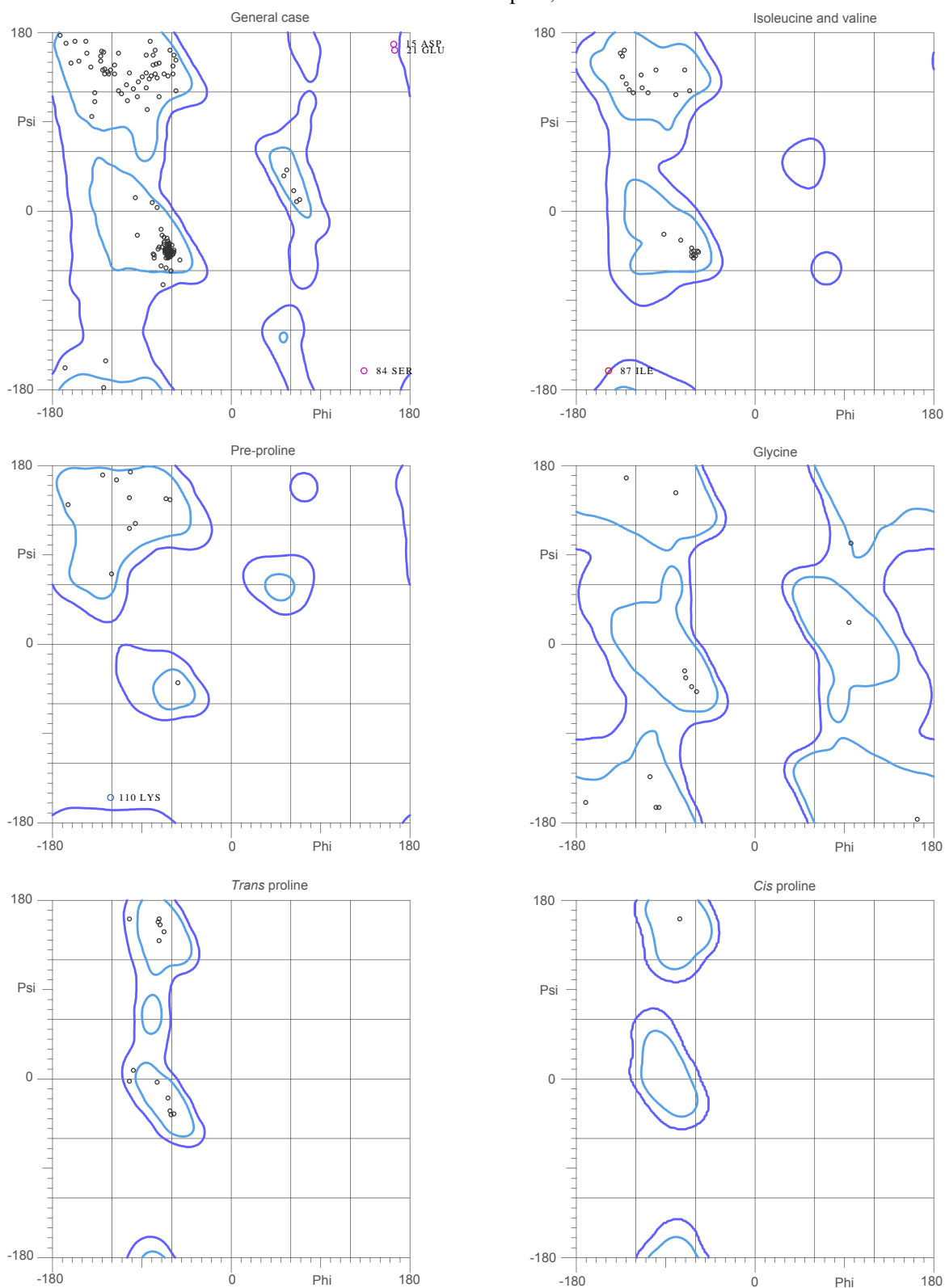# MolProbity Ramachandran analysis

1at3A.B99990016.pdb, model 1

94,9% (204/215) of all residues were in favored (98%) regions.

97,7% (210/215) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

    15 ASP (164,9, 169,7)
    21 GLU (165,3, 163,6)
    84 SER (134,7, -161,6)
    87 ILE (-148,2, -161,5)
    110 LYS (-122,7, -155,8)