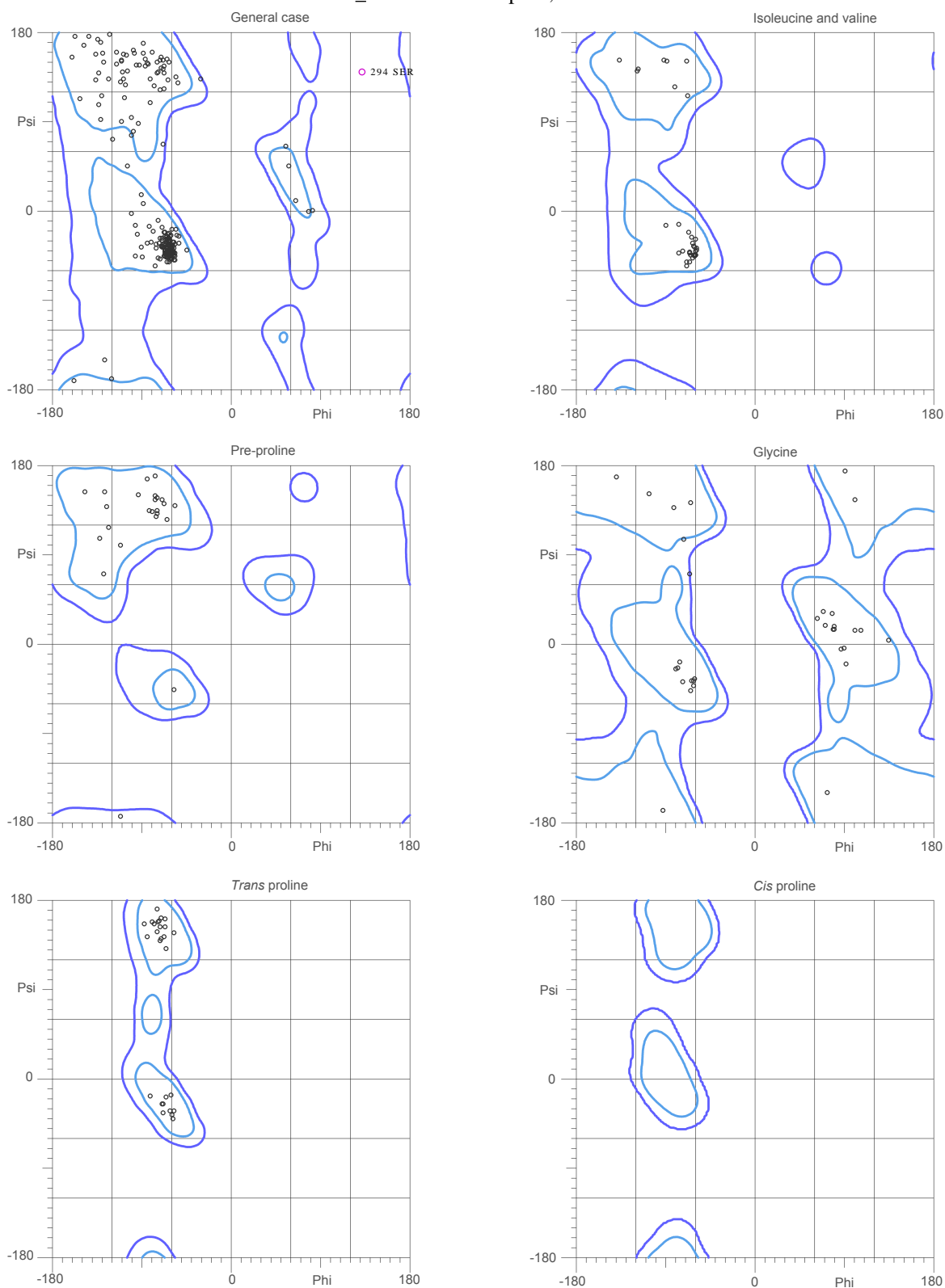


# MolProbity Ramachandran analysis

1aru\_.B99990031.pdb, model 1



96,4% (322/334) of all residues were in favored (98%) regions.

99,7% (333/334) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

294 SER (132,3, 141,8)