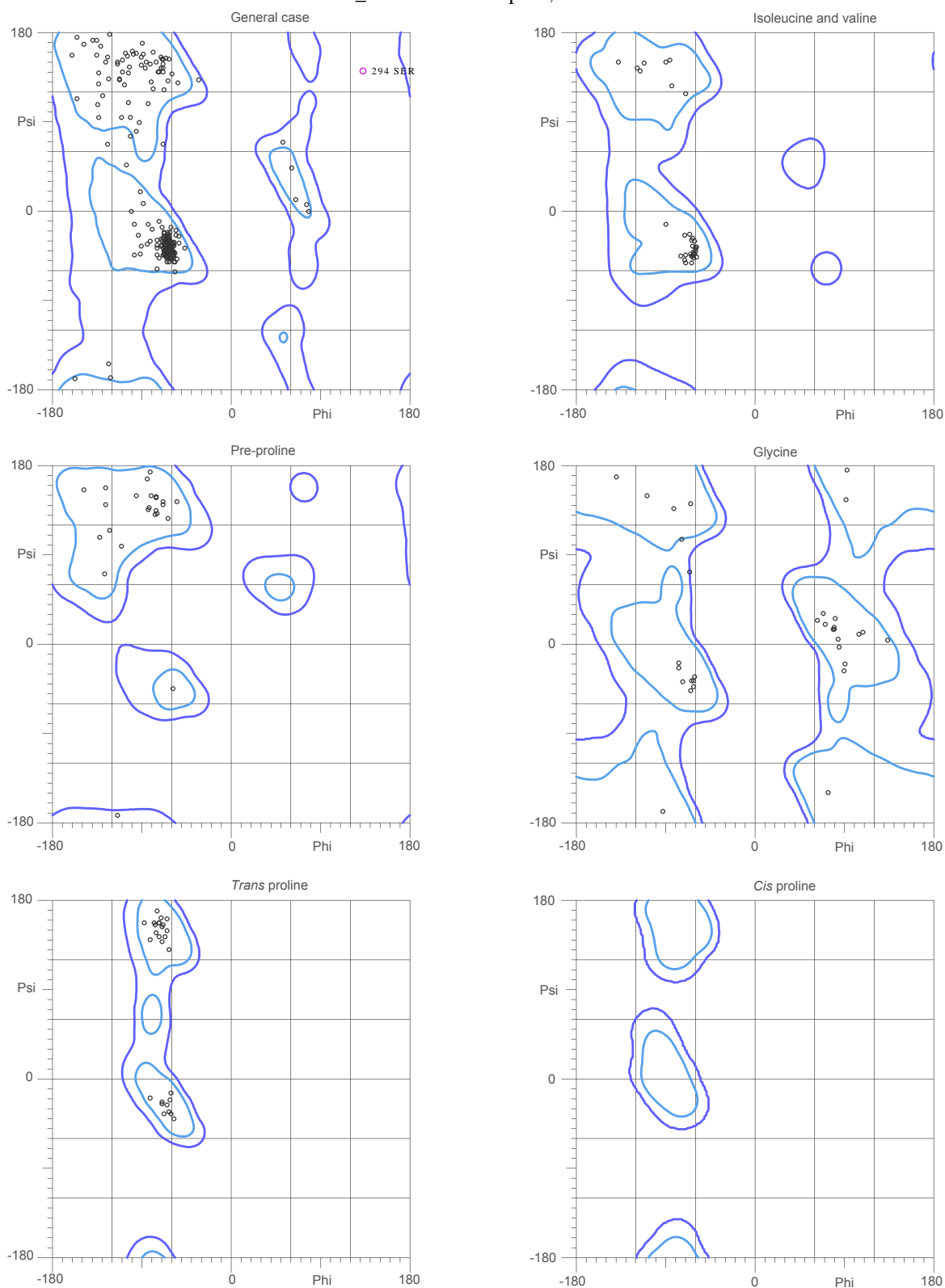


MolProbity Ramachandran analysis

1aru_.B99990023.pdb, model 1



96,7% (323/334) 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 (133,2, 142,9)