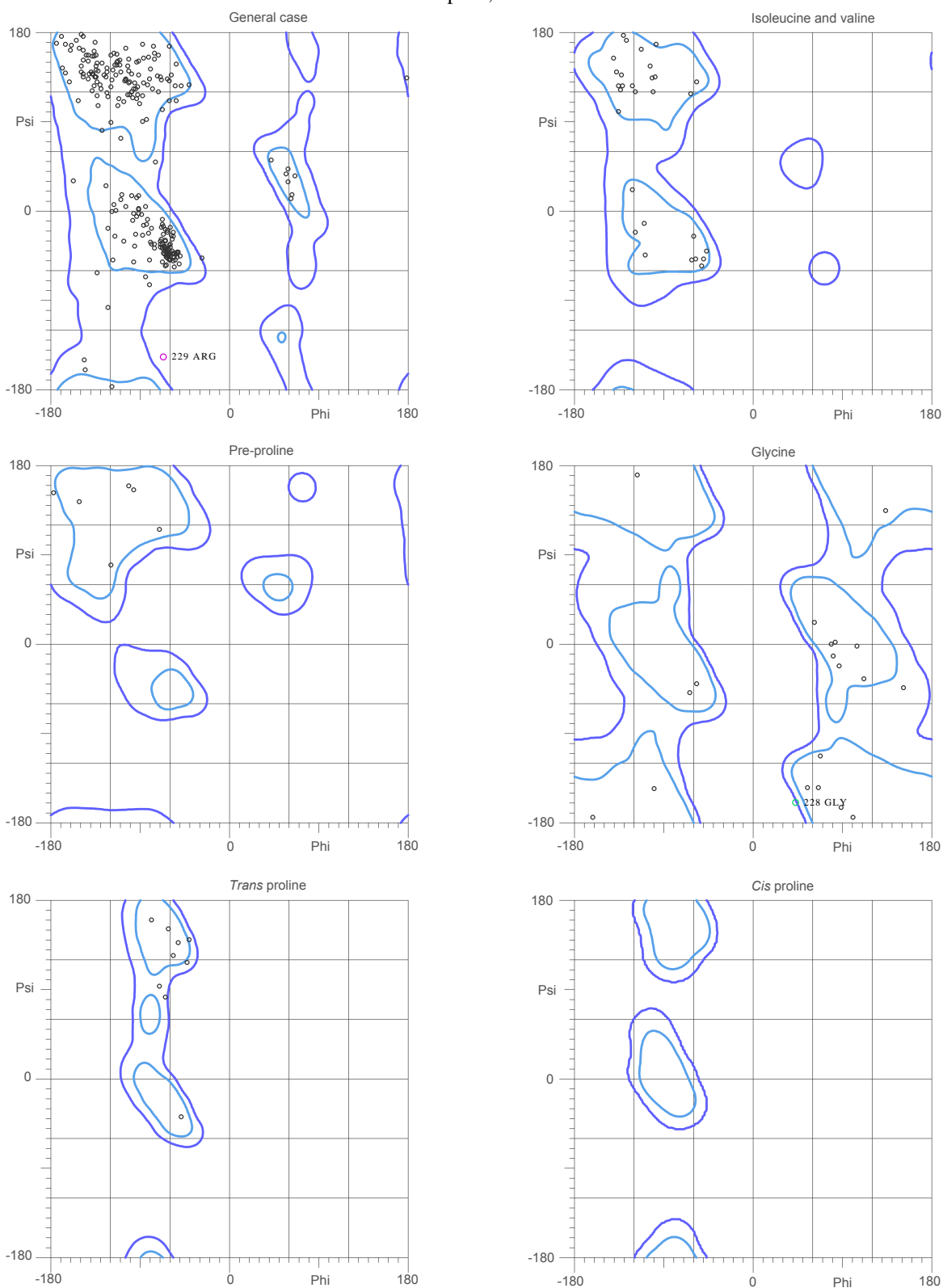


# MolProbity Ramachandran analysis

3e9s.H.pdb, model 1



93.0% (292/314) of all residues were in favored (98%) regions.

99.4% (312/314) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

228 GLY (43.0, -160.6)

229 ARG (-67.3, -147.2)