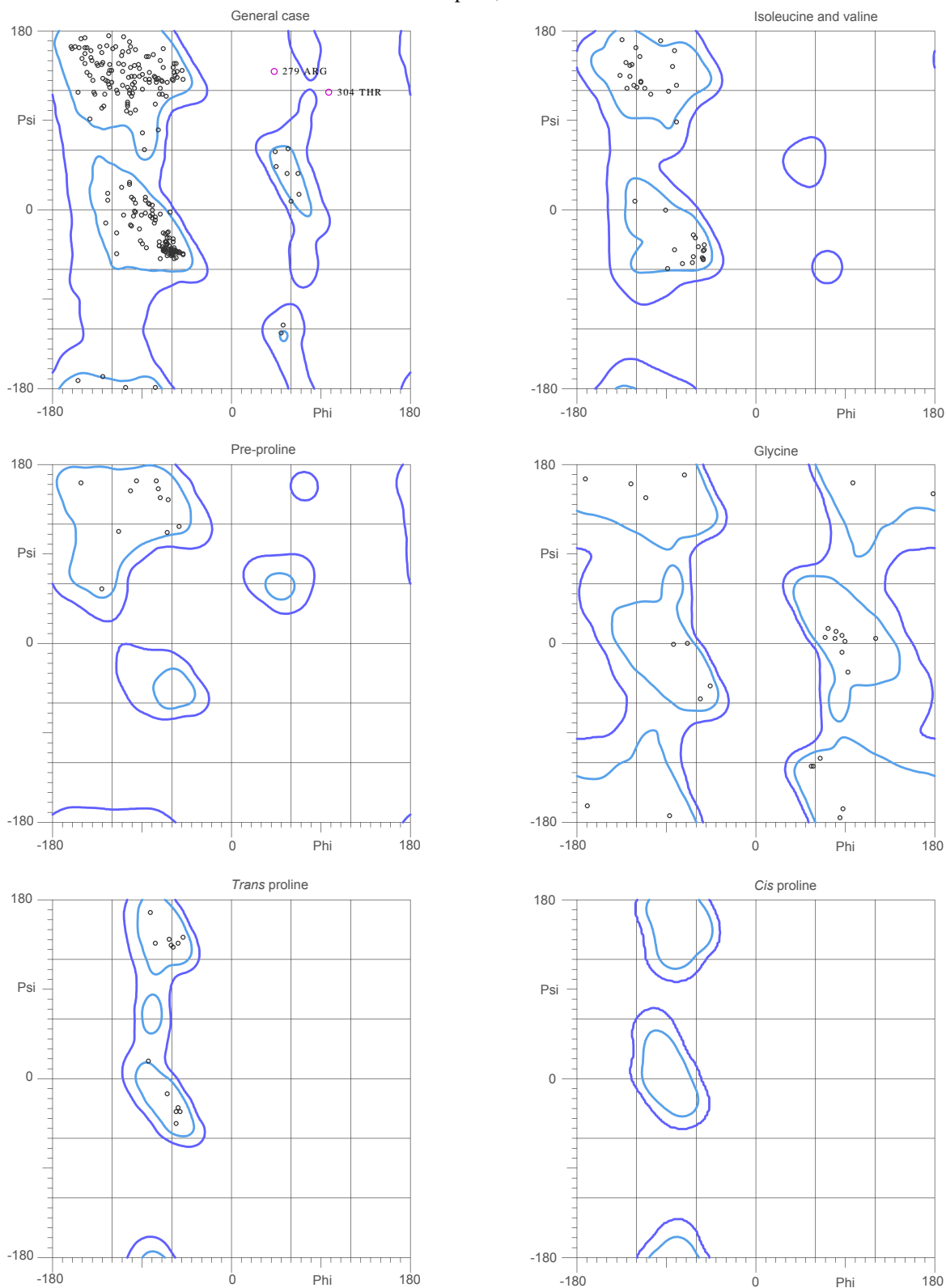


# MolProbity Ramachandran analysis

3v3m.H.pdb, model 1



97.1% (305/314) 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):

279 ARG (43.1, 140.6)

304 THR (98.8, 119.3)