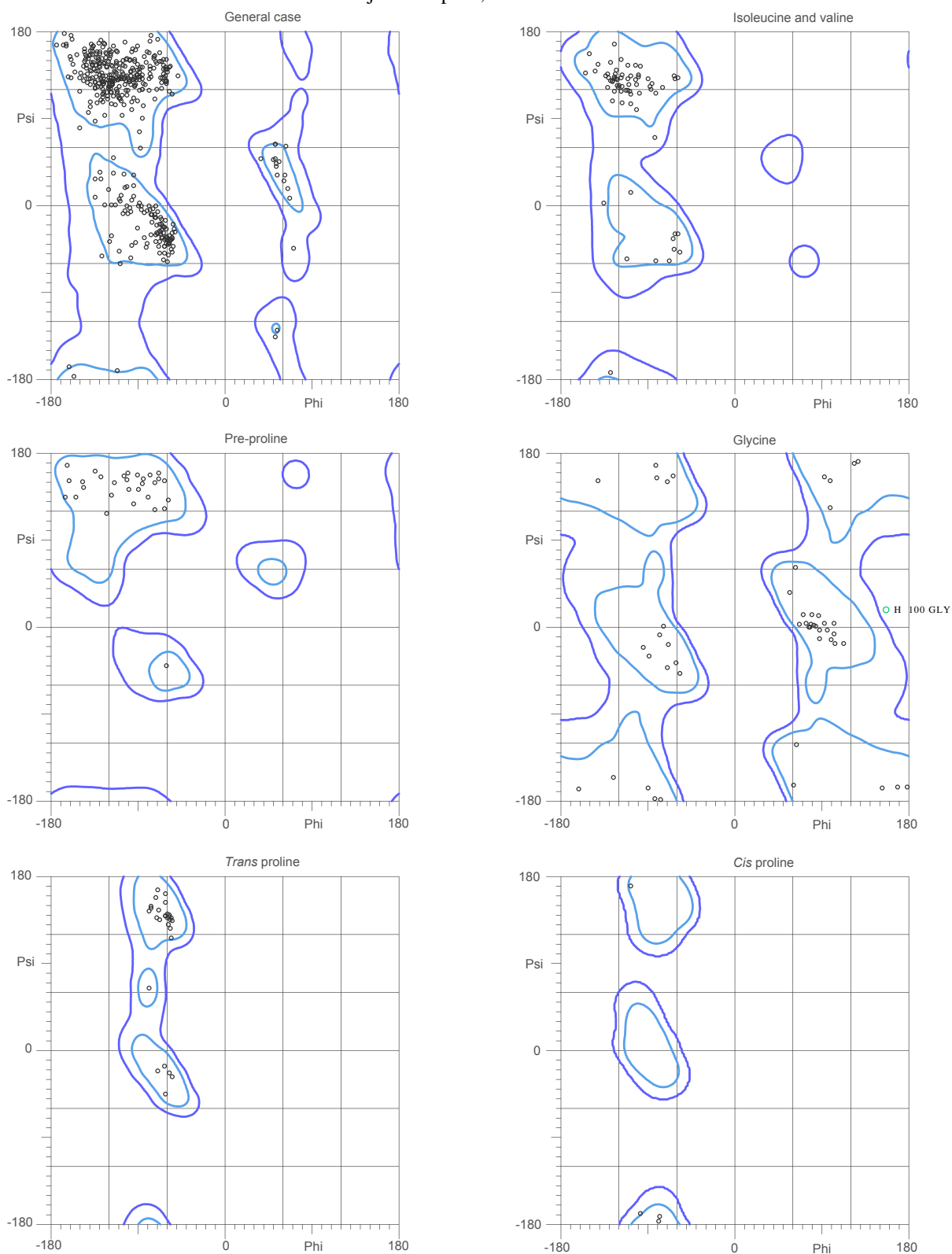


# MolProbity Ramachandran analysis

7jmo.H.pdb, model 1



97.1% (596/614) of all residues were in favored (98%) regions.

99.8% (613/614) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

H 100 GLY (157.1, 18.8)