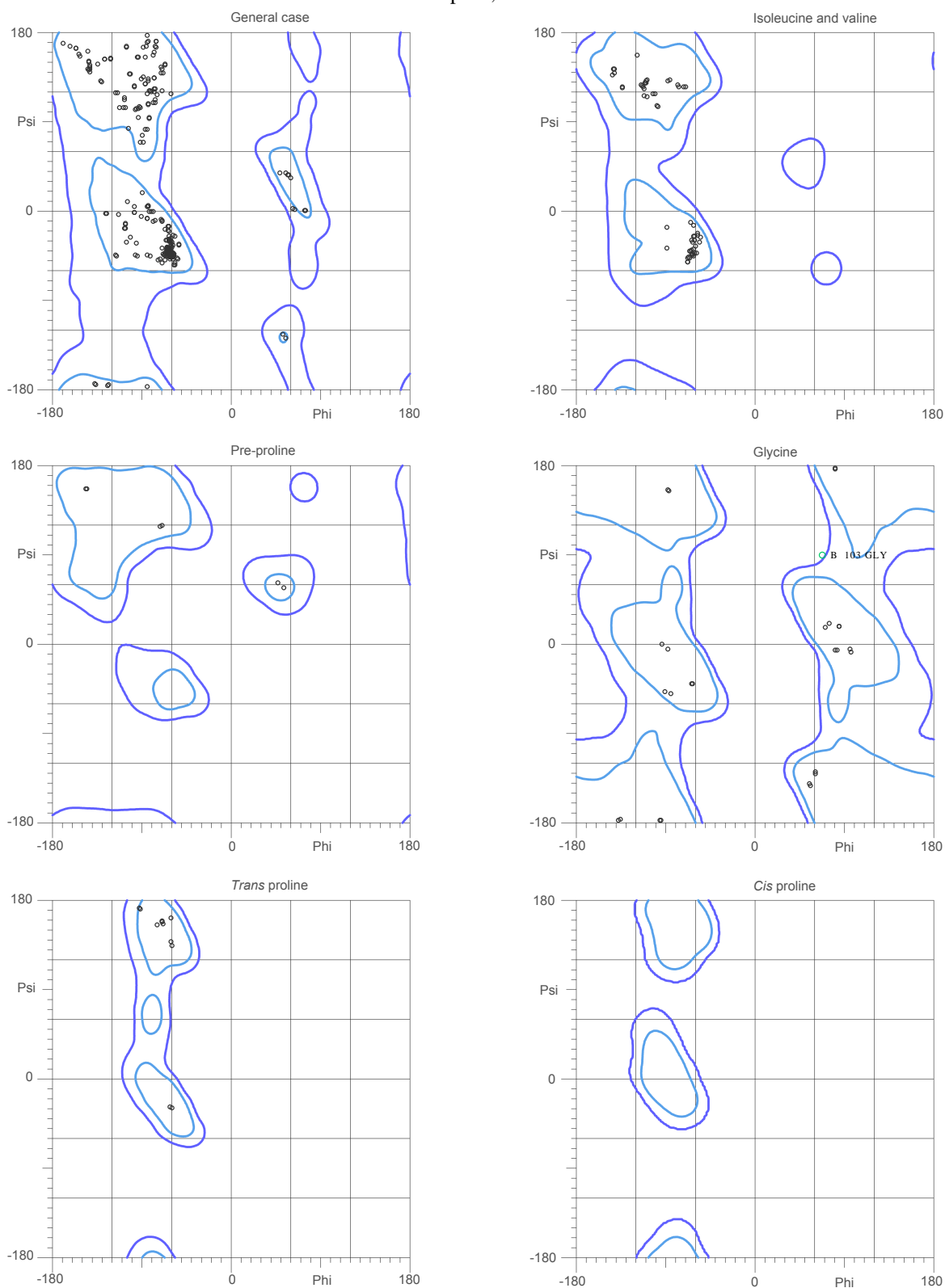


MolProbity Ramachandran analysis

5s3d.H.pdb, model 1



98.8% (338/342) of all residues were in favored (98%) regions.

99.7% (341/342) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

B 103 GLY (68.4, 90.1)