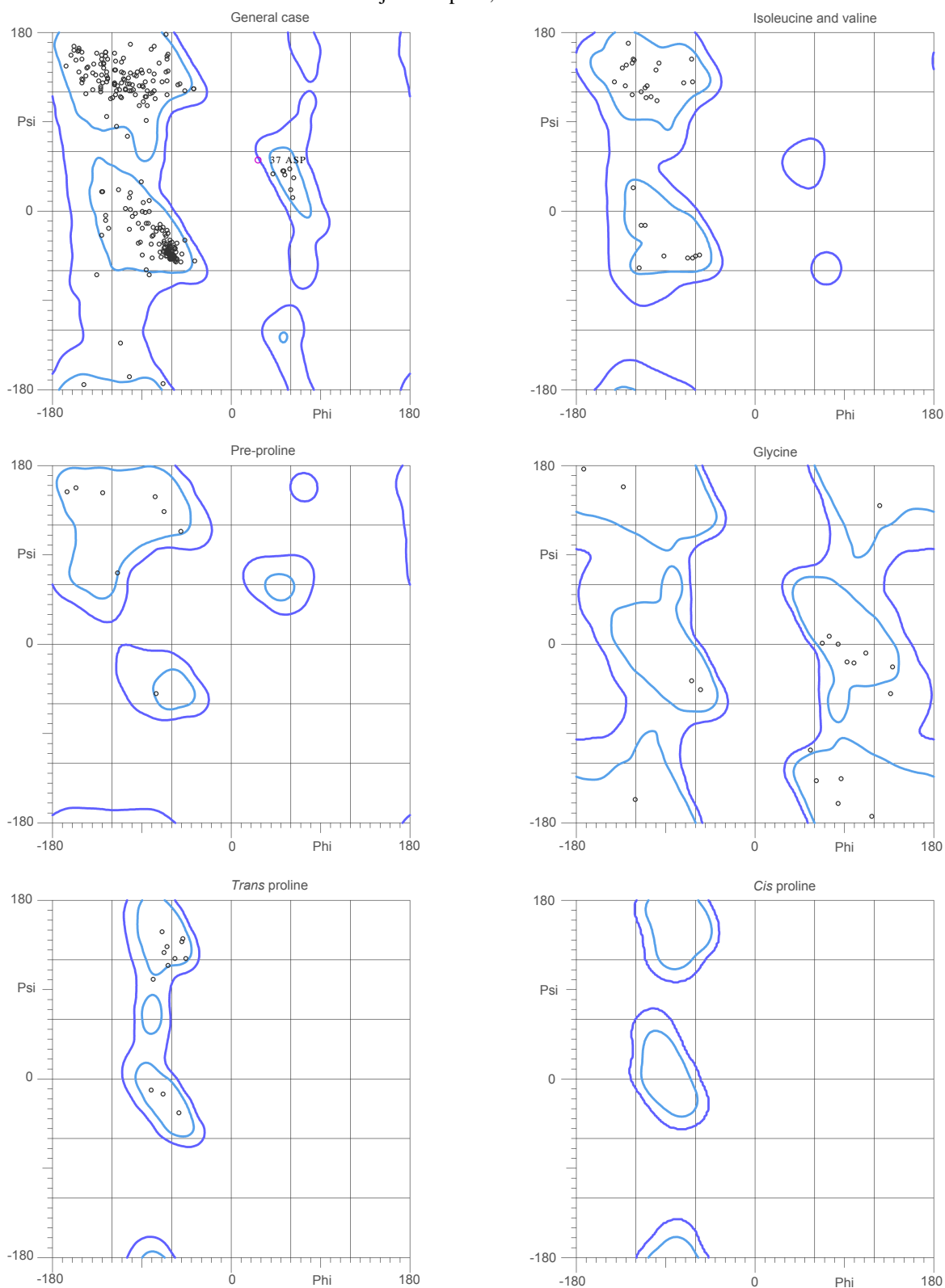


# MolProbity Ramachandran analysis

7jn2.H.pdb, model 1



94.9% (299/315) of all residues were in favored (98%) regions.

99.7% (314/315) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

37 ASP (27.9, 52.0)