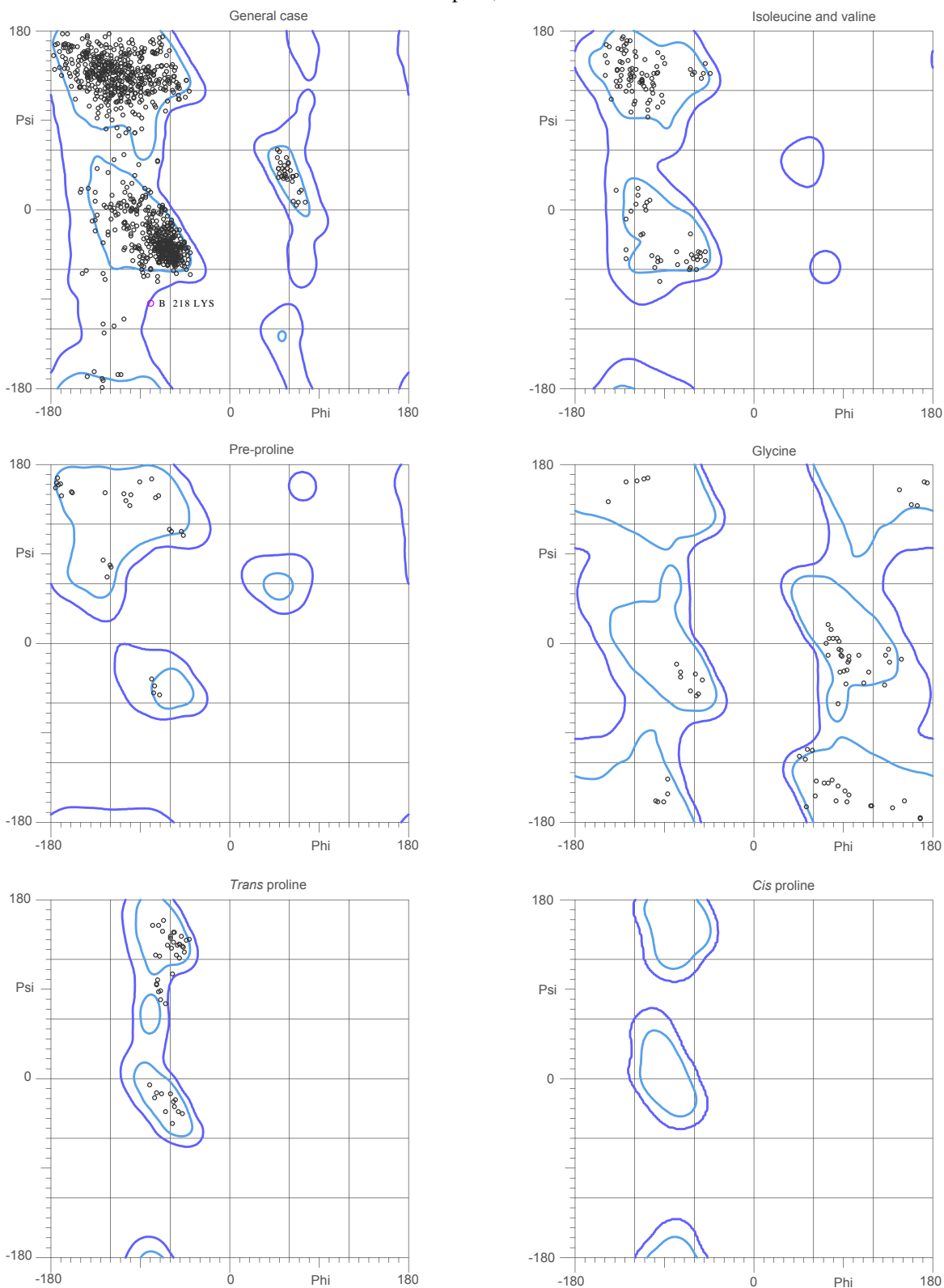


# MolProbity Ramachandran analysis

7cmd.H.pdb, model 1



93.5% (1157/1238) of all residues were in favored (98%) regions.

99.9% (1237/1238) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

B 218 LYS (-80.5, -94.8)