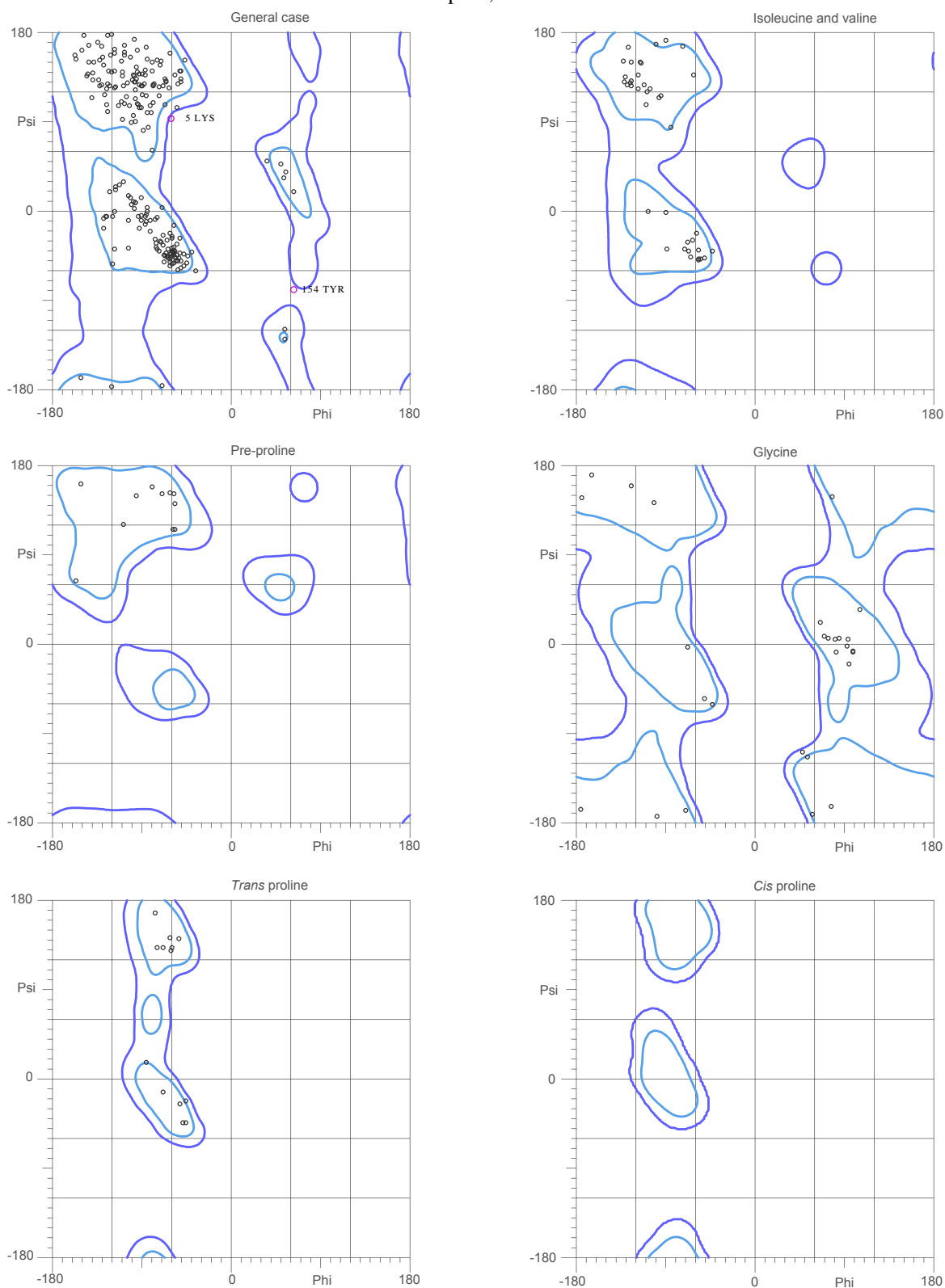


# MolProbity Ramachandran analysis

5re5.H.pdb, model 1



95.2% (296/311) of all residues were in favored (98%) regions.

99.4% (309/311) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

5 LYS (-61.7, 94.3)

154 TYR (63.3, -79.4)