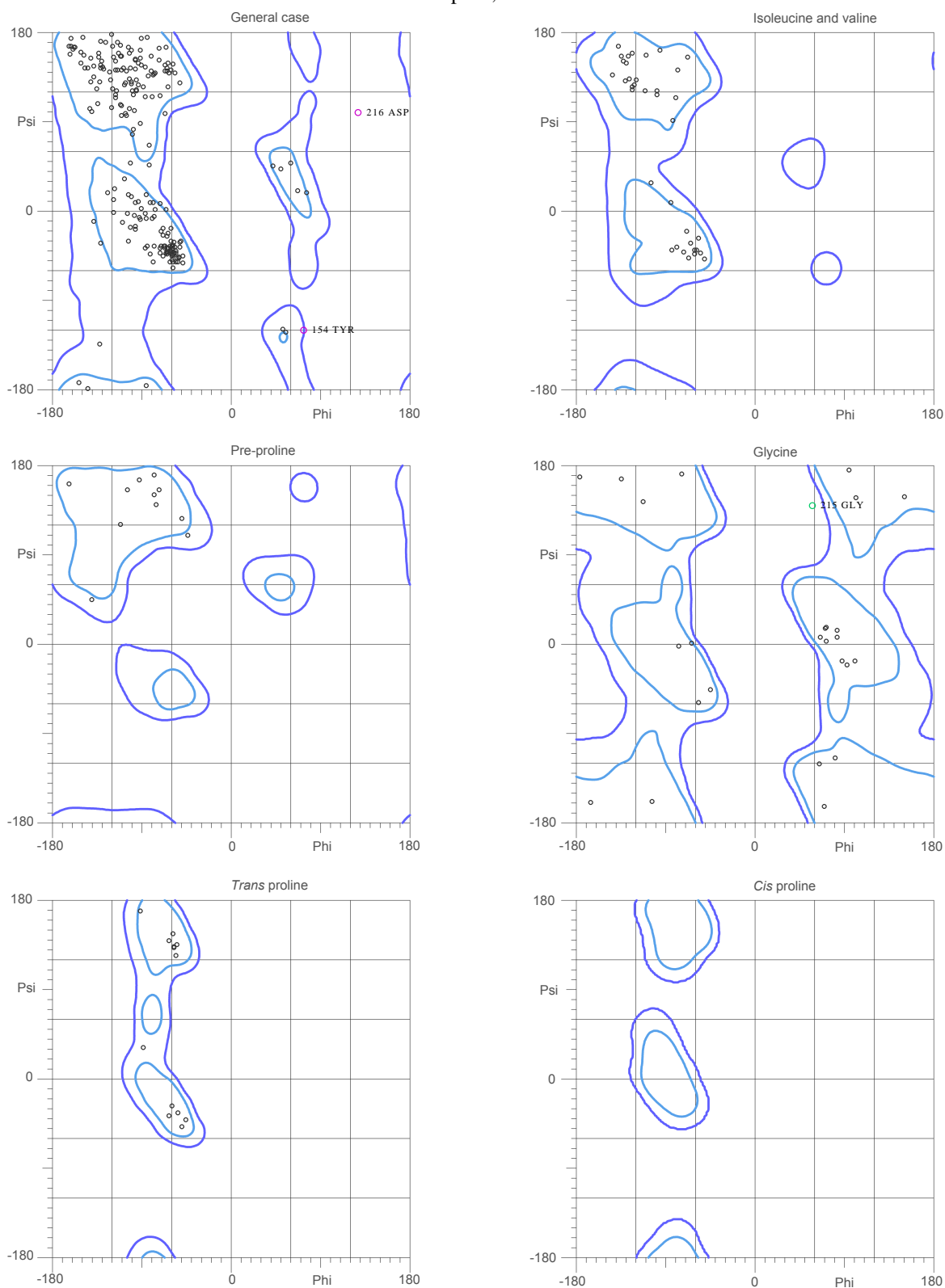


# MolProbity Ramachandran analysis

2zu4.H.pdb, model 1



93.8% (285/304) of all residues were in favored (98%) regions.  
99.0% (301/304) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

154 TYR (73.8, -120.7)  
215 GLY (58.7, 140.4)  
216 ASP (128.2, 100.3)