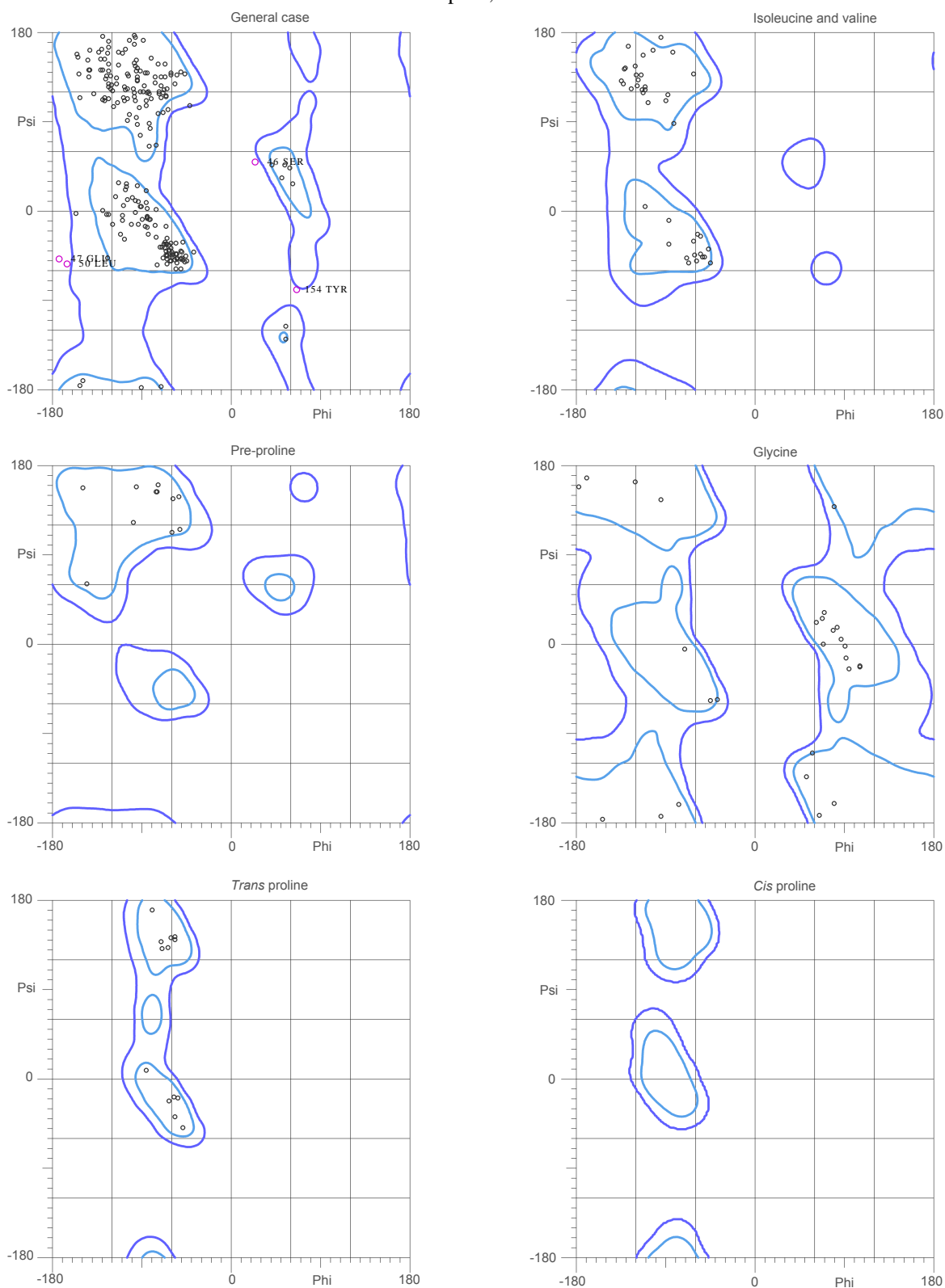


# MolProbity Ramachandran analysis

5ren.H.pdb, model 1



95.8% (298/311) of all residues were in favored (98%) regions.  
98.7% (307/311) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):

46 SER (24.9, 50.9)  
47 GLU (-174.7, -48.4)  
50 LEU (-166.8, -53.7)  
154 TYR (66.3, -79.6)