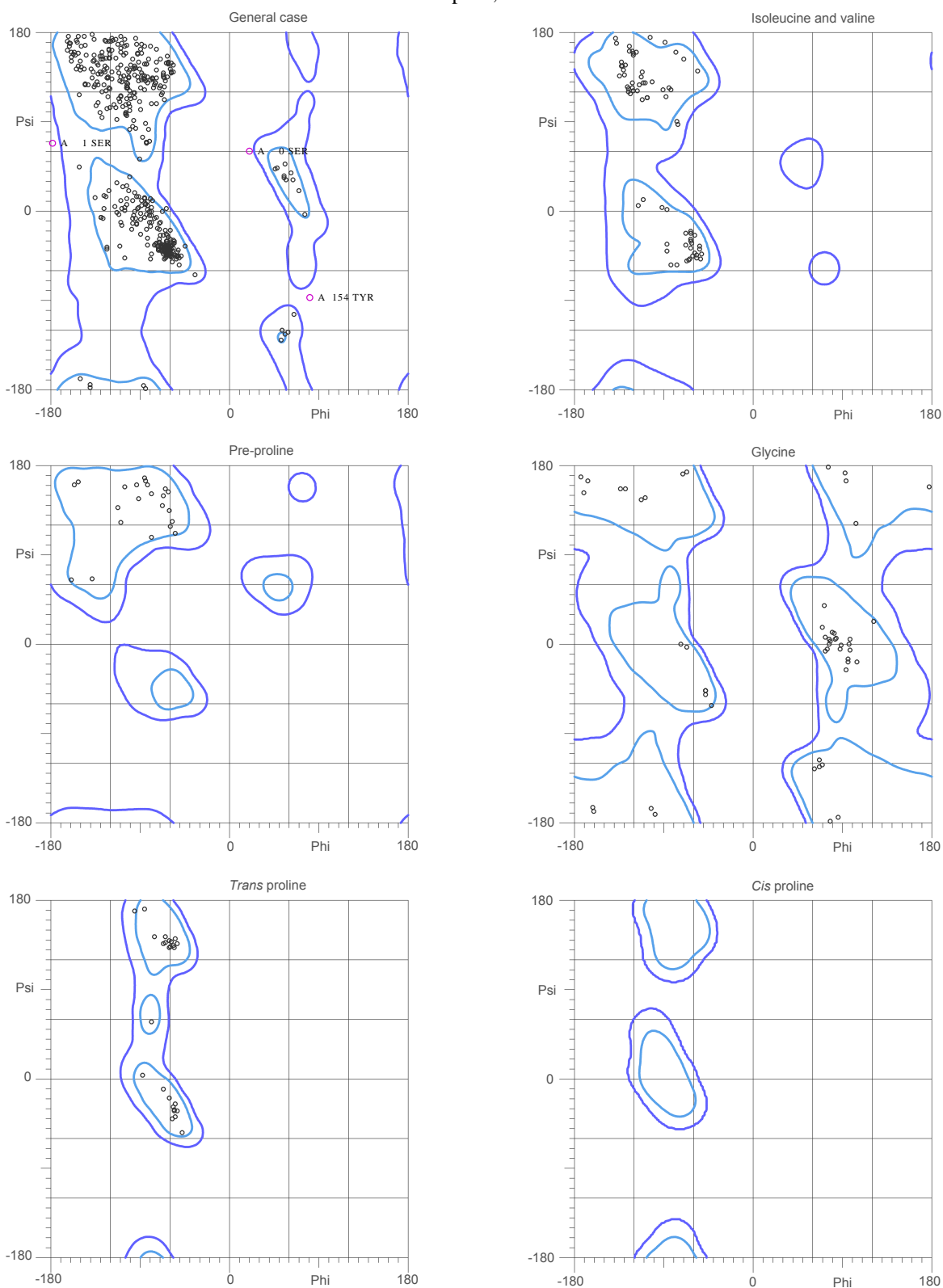


# MolProbity Ramachandran analysis

2amd.H.pdb, model 1



97.5% (594/609) of all residues were in favored (98%) regions.  
99.5% (606/609) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):

- A 0 SER (20.3, 61.0)
- A 1 SER (-179.8, 69.0)
- A 154 TYR (81.0, -87.1)