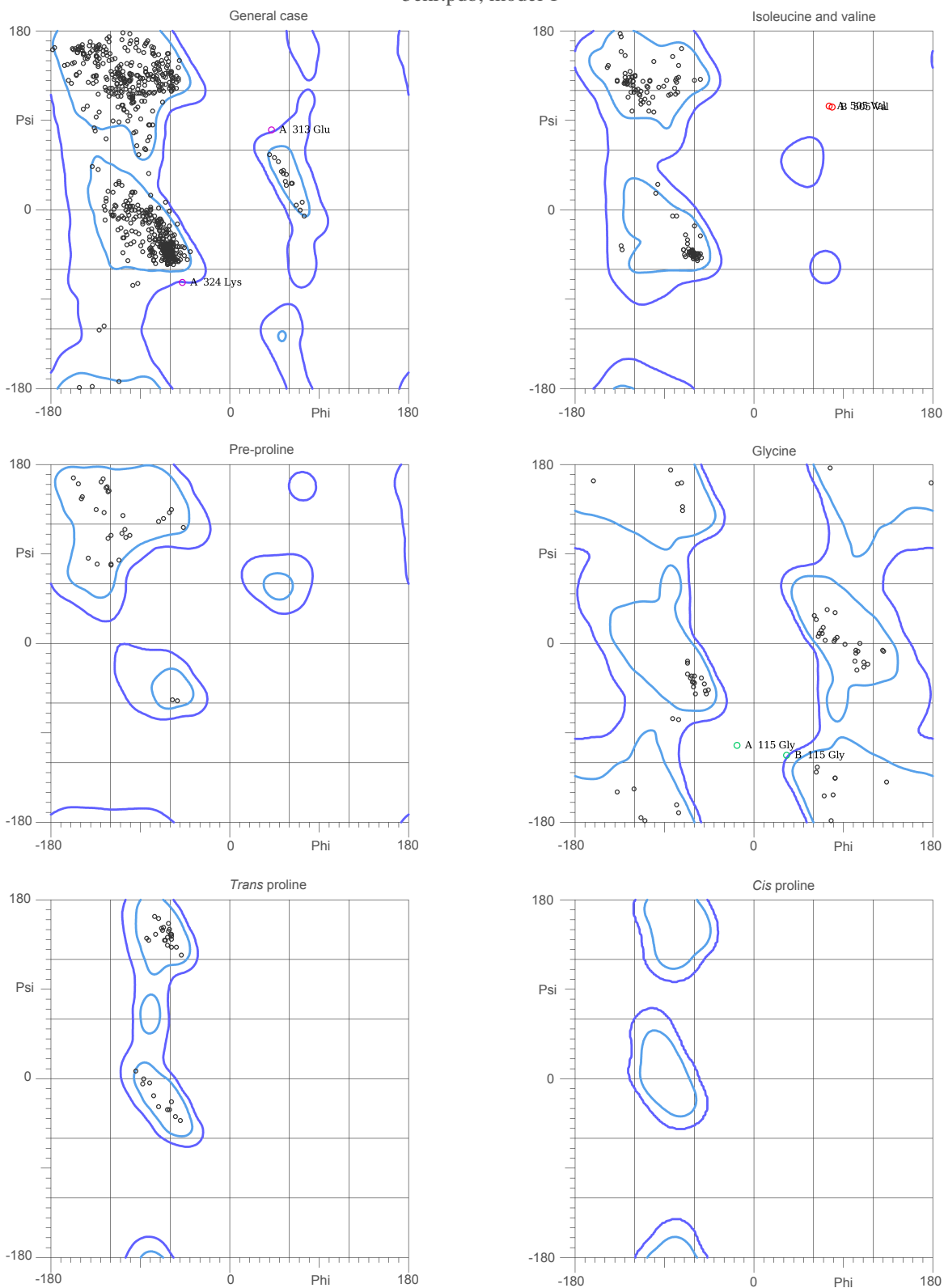


# MolProbity Ramachandran analysis

5ehr.pdb, model 1



97.1% (907/934) of all residues were in favored (98%) regions.  
99.4% (928/934) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):

- A 115 Gly (-17.6, -103.8)
- A 313 Glu (42.4, 81.1)
- A 324 Lys (-48.5, -73.6)
- A 505 Val (76.8, 105.4)
- B 115 Gly (33.9, -113.3)
- B 505 Val (79.3, 104.7)