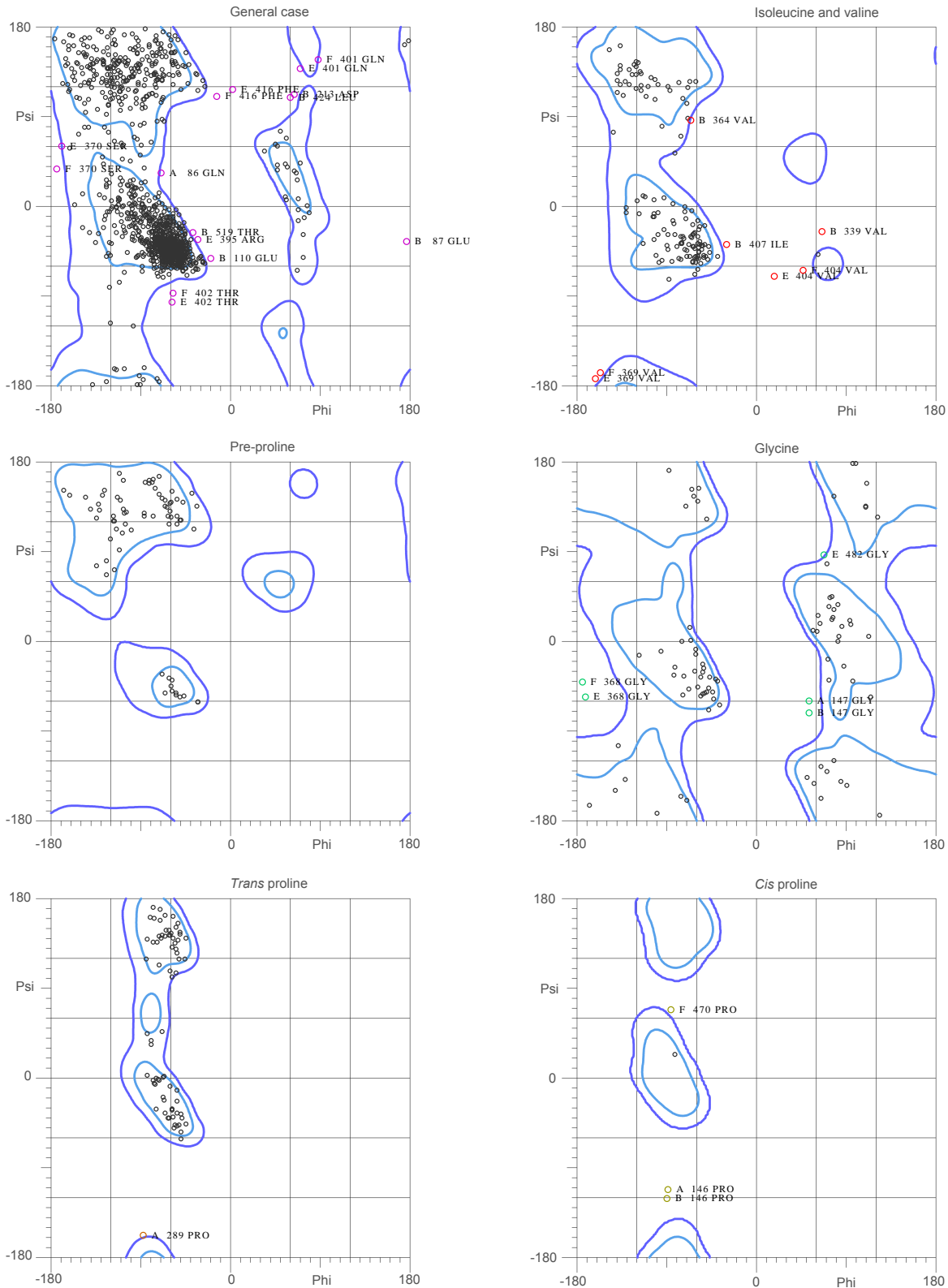


# MolProbity Ramachandran analysis

2ajf.H.pdb, model 1



87.0% (1249/1436) of all residues were in favored (98%) regions.  
10.0% (1436/1436) of all residues were in allowed (1.0%) regions.

There were 11 outliers (Phi, Psi):

- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)
- A 86 GLN (1.0, 1.0)