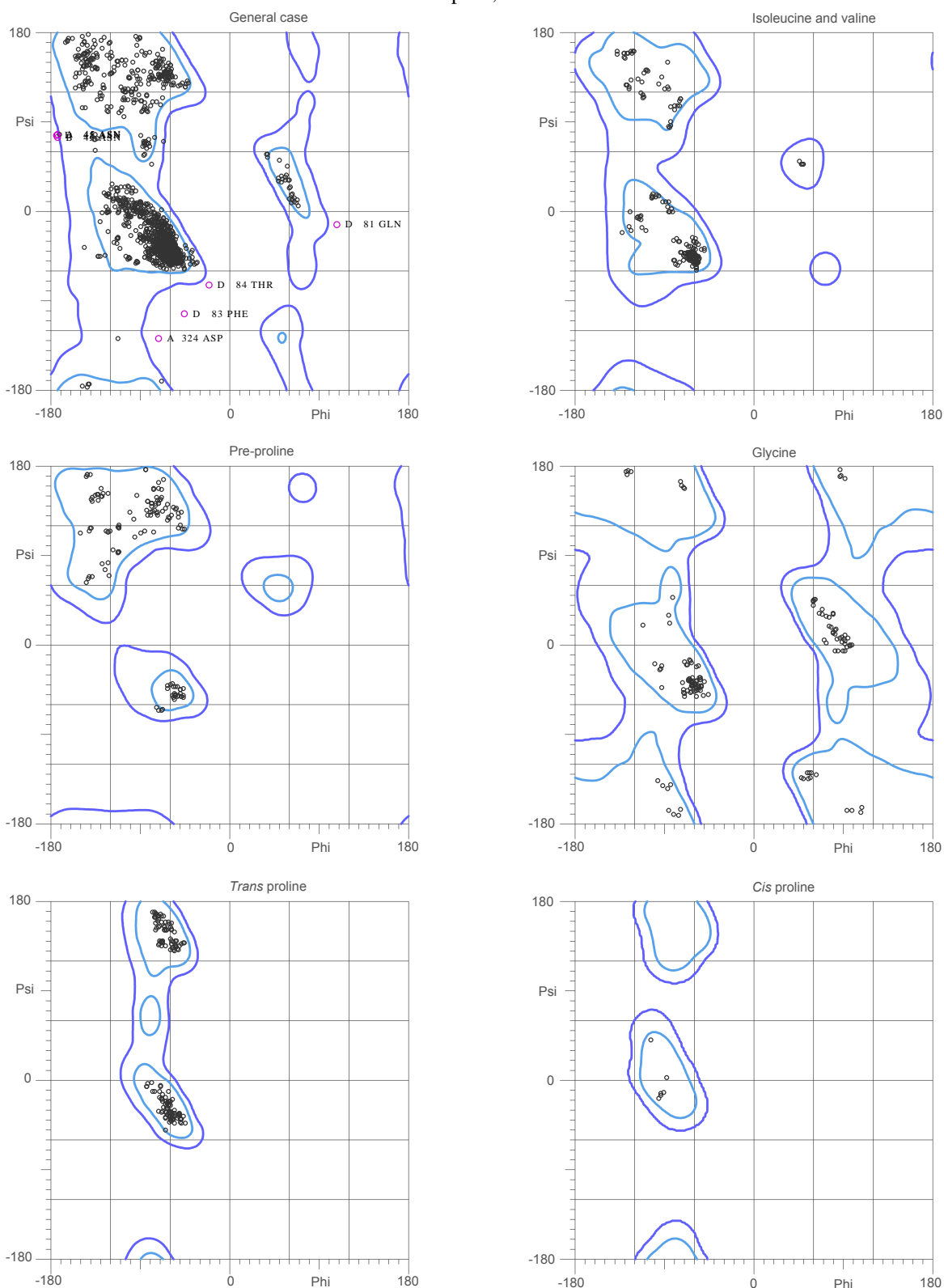


# MolProbity Ramachandran analysis

5ama.H.pdb, model 1



98.2% (2366/2410) of all residues were in favored (98%) regions.  
99.7% (2403/2410) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):

- A 45 ASN (-174.8, 78.2)
- A 324 ASP (-72.9, -128.6)
- B 45 ASN (-174.8, 75.1)
- D 45 ASN (-175.7, 77.6)
- D 81 GLN (108.7, -13.2)
- D 83 PHE (-46.1, -103.2)
- D 84 THR (-21.2, -74.7)