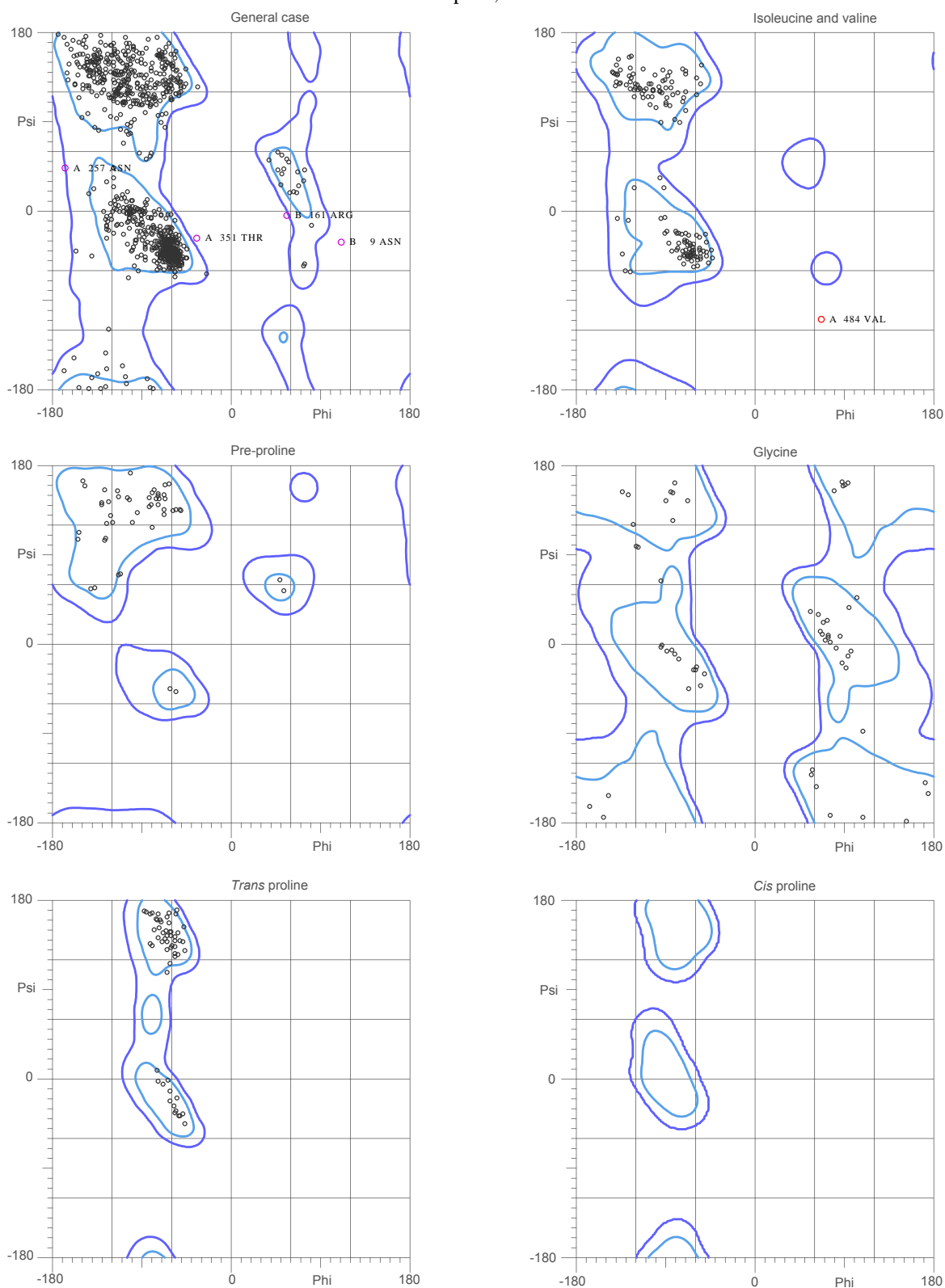


# MolProbity Ramachandran analysis

5rmd.H.pdb, model 1



94.7% (1085/1146) of all residues were in favored (98%) regions.  
99.6% (1141/1146) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

- A 257 ASN (-168.9, 44.4)
- A 351 THR (-35.7, -27.9)
- A 484 VAL (68.0, -109.2)
- B 9 ASN (111.2, -31.0)
- B 161 ARG (57.0, -4.7)