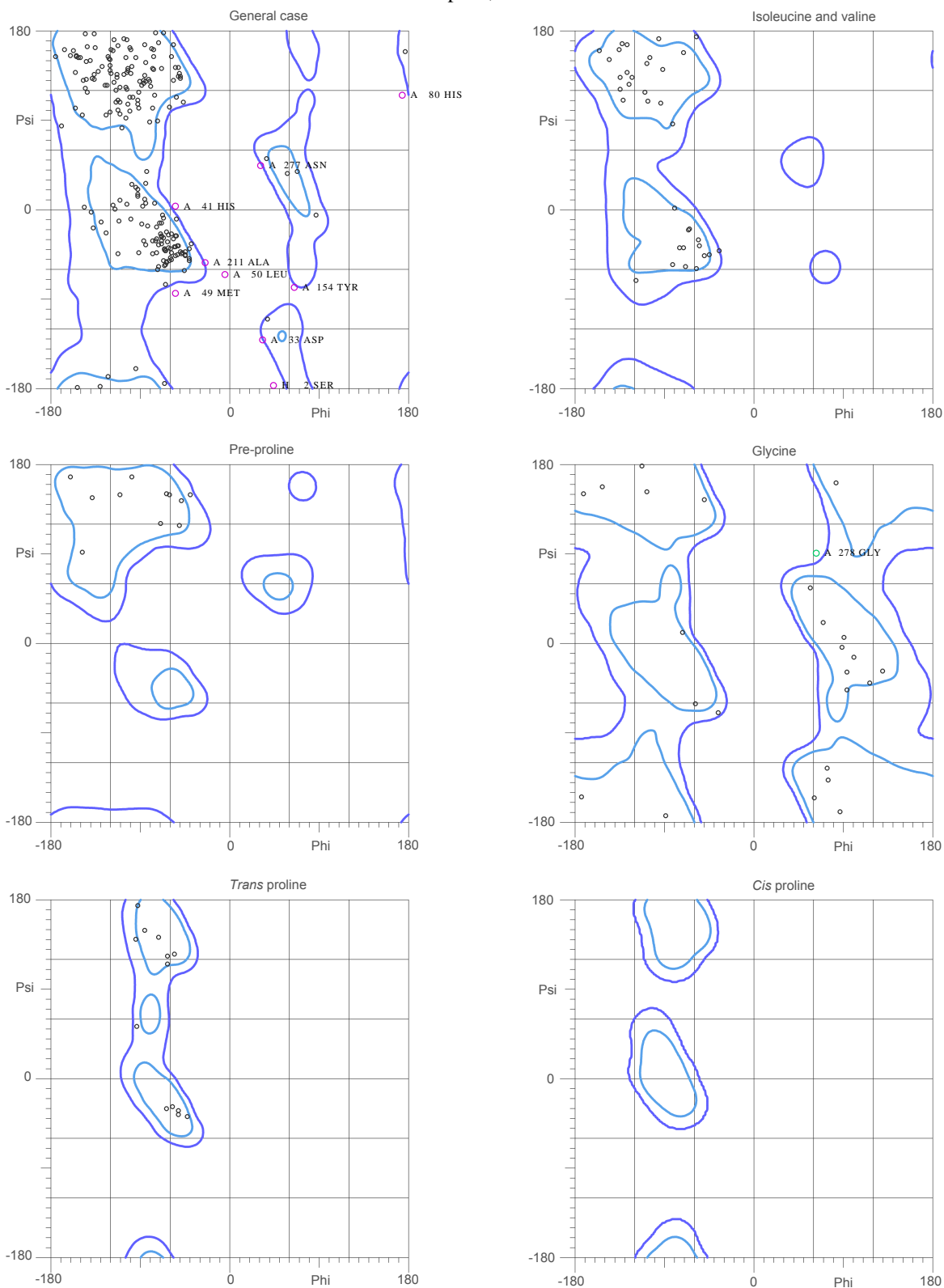


# MolProbity Ramachandran analysis

3sna.H.pdb, model 1



84.5% (256/303) of all residues were in favored (98%) regions.  
96.7% (293/303) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):

A 33 ASP (33.2, -131.2)  
A 41 HIS (-56.0, 4.7)  
A 49 MET (-55.2, -84.4)  
A 50 LEU (-5.7, -65.5)  
A 80 HIS (174.6, 117.0)  
A 154 TYR (66.0, -78.8)  
A 211 ALA (-25.6, -53.6)  
A 277 ASN (31.4, 45.1)  
A 278 GLY (63.0, 91.6)  
H 2 SER (44.0, -177.2)