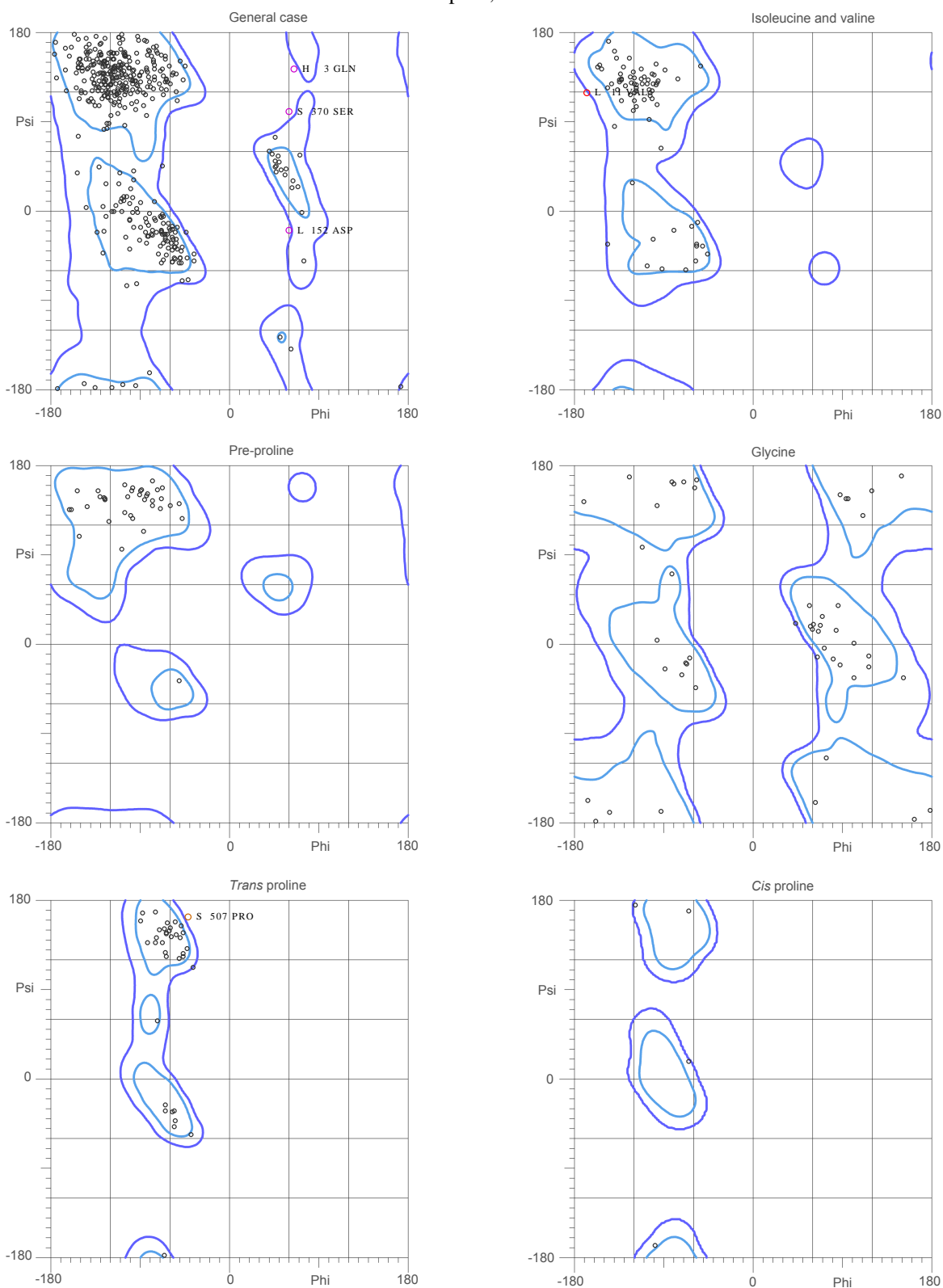


# MolProbity Ramachandran analysis

2dd8.H.pdb, model 1



92.1% (569/618) of all residues were in favored (98%) regions.  
99.2% (613/618) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):

H 3 GLN (65.1, 144.6)  
L 11 VAL (-168.1, 120.4)  
L 152 ASP (60.1, -19.3)  
S 370 SER (60.1, 101.2)  
S 507 PRO (-42.9, 164.6)