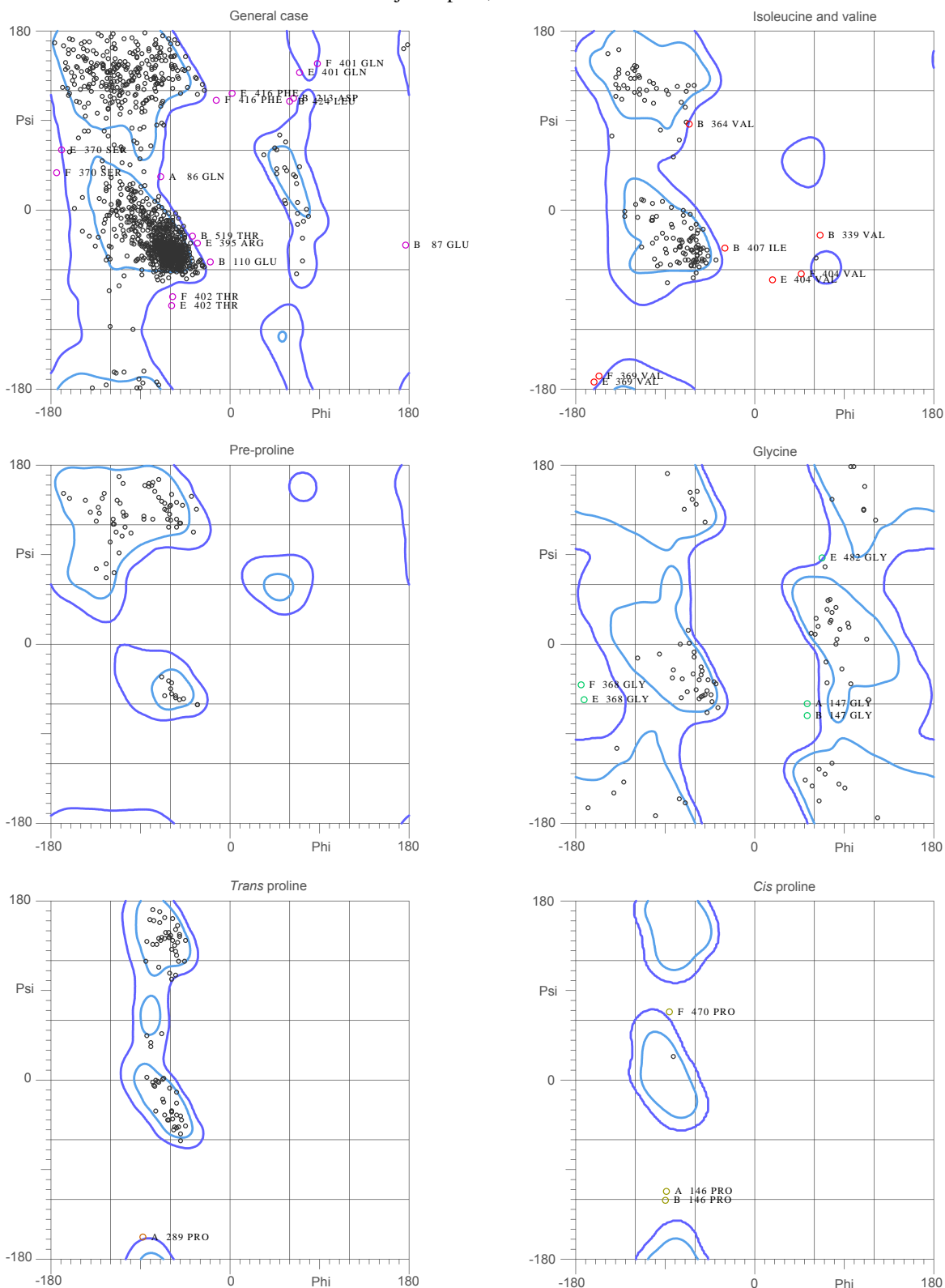


# MolProbity Ramachandran analysis

2ajf.H.pdb, model 1



87.0% (124/143) of all residues were in favored (98%) regions.  
10.0% (14/143) of all residues were in allowed (1.0%) regions.

There were 11 outliers (Phi, Psi):

A 86 GLN (Phi, Psi)  
B 519 THR (Phi, Psi)  
B 110 GLU (Phi, Psi)  
F 402 THR (Phi, Psi)  
E 402 THR (Phi, Psi)  
F 401 GLN (Phi, Psi)  
E 401 GLN (Phi, Psi)  
F 416 PHE (Phi, Psi)  
B 124 ASP (Phi, Psi)  
A 289 PRO (Phi, Psi)  
F 470 PRO (Phi, Psi)  
A 146 PRO (Phi, Psi)  
B 146 PRO (Phi, Psi)  
F 368 GLY (Phi, Psi)  
E 368 GLY (Phi, Psi)  
E 482 GLY (Phi, Psi)  
B 364 VAL (Phi, Psi)  
B 407 ILE (Phi, Psi)  
B 339 VAL (Phi, Psi)  
E 404 VAL (Phi, Psi)  
F 369 VAL (Phi, Psi)  
E 369 VAL (Phi, Psi)