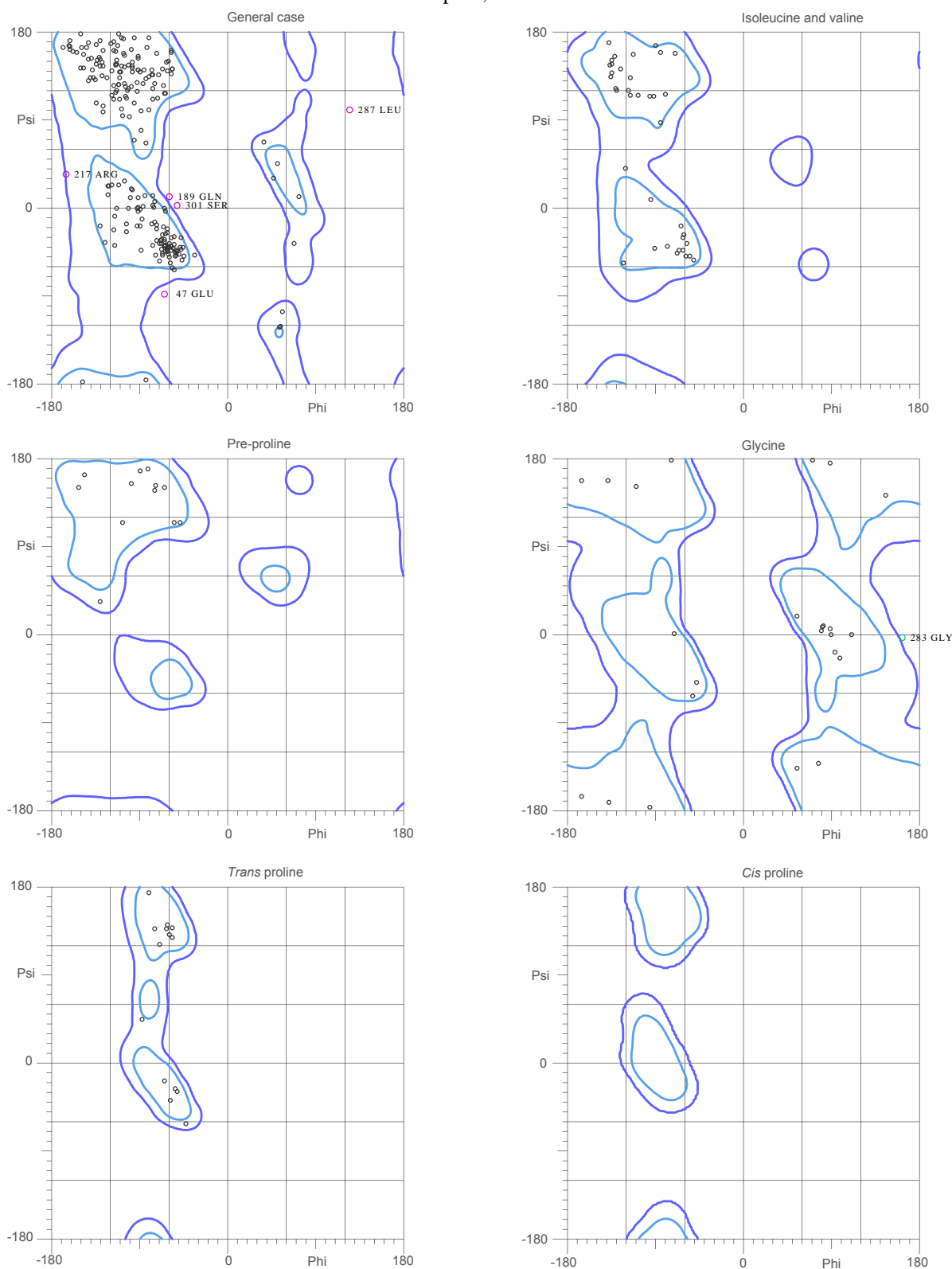


# MolProbity Ramachandran analysis

2alv.H.pdb, model 1



94.0% (300/319) of all residues were in favored (98%) regions.  
98.1% (313/319) of all residues were in allowed (>99.8%) regions.

There were 6 outliers (phi, psi):

47 GLU (-65.5, -88.8)  
189 GLN (-60.3, 12.0)  
217 ARG (-166.5, 35.2)  
283 GLY (163.0, -3.4)  
287 LEU (125.9, 101.9)  
301 SER (-53.0, 3.5)