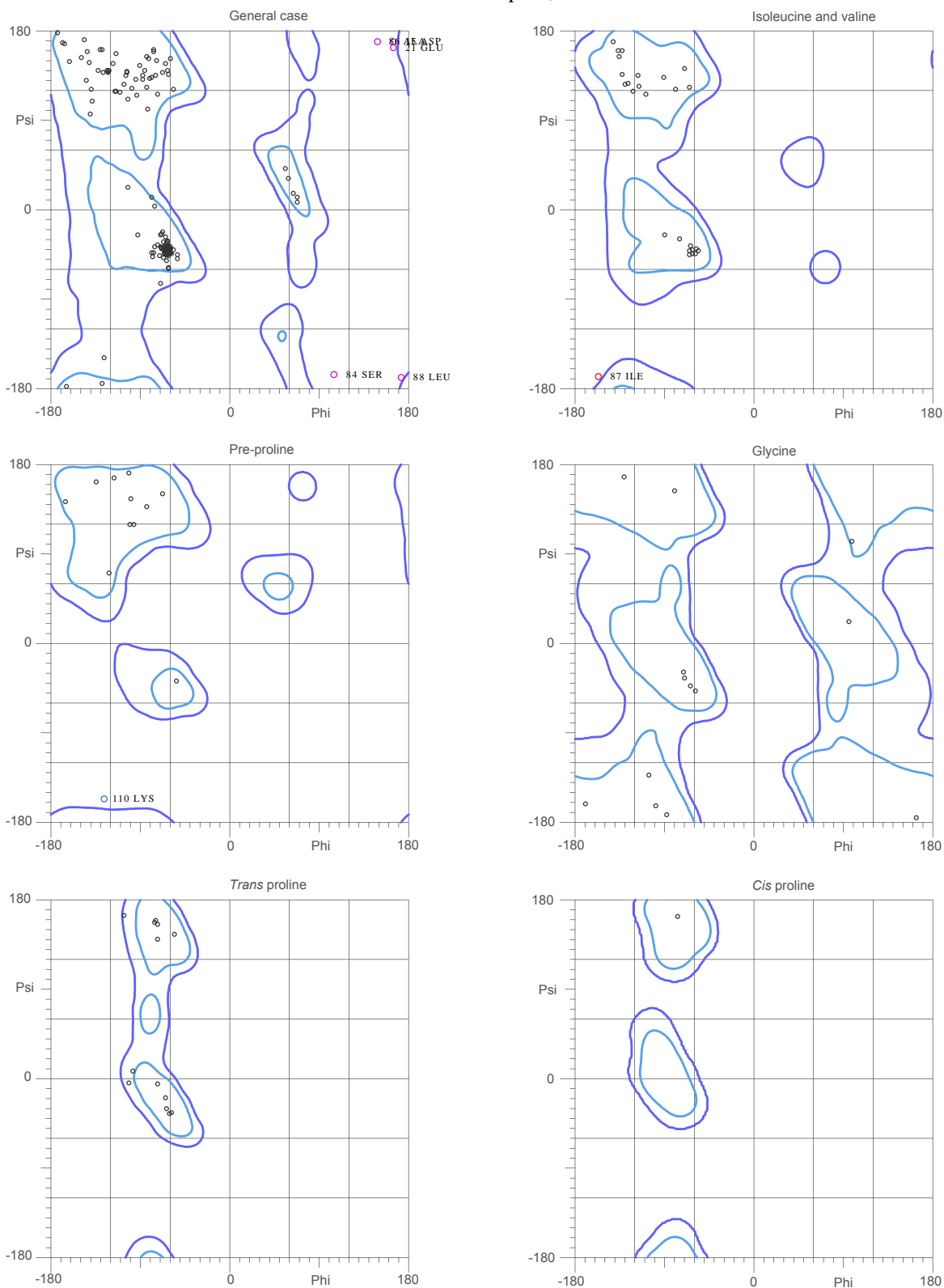


# MolProbity Ramachandran analysis

1at3A.B99990049.pdb, model 1



93,5% (201/215) of all residues were in favored (98%) regions.  
96,7% (208/215) of all residues were in allowed (>99.8%) regions.

There were 7 outliers (phi, psi):

15 ASP (165,6, 170,3)  
21 GLU (165,8, 164,8)  
84 SER (105,3, -166,2)  
86 ALA (149,3, 170,1)  
87 ILE (-157,4, -168,1)  
88 LEU (173,6, -170,0)  
110 LYS (-127,1, -157,5)