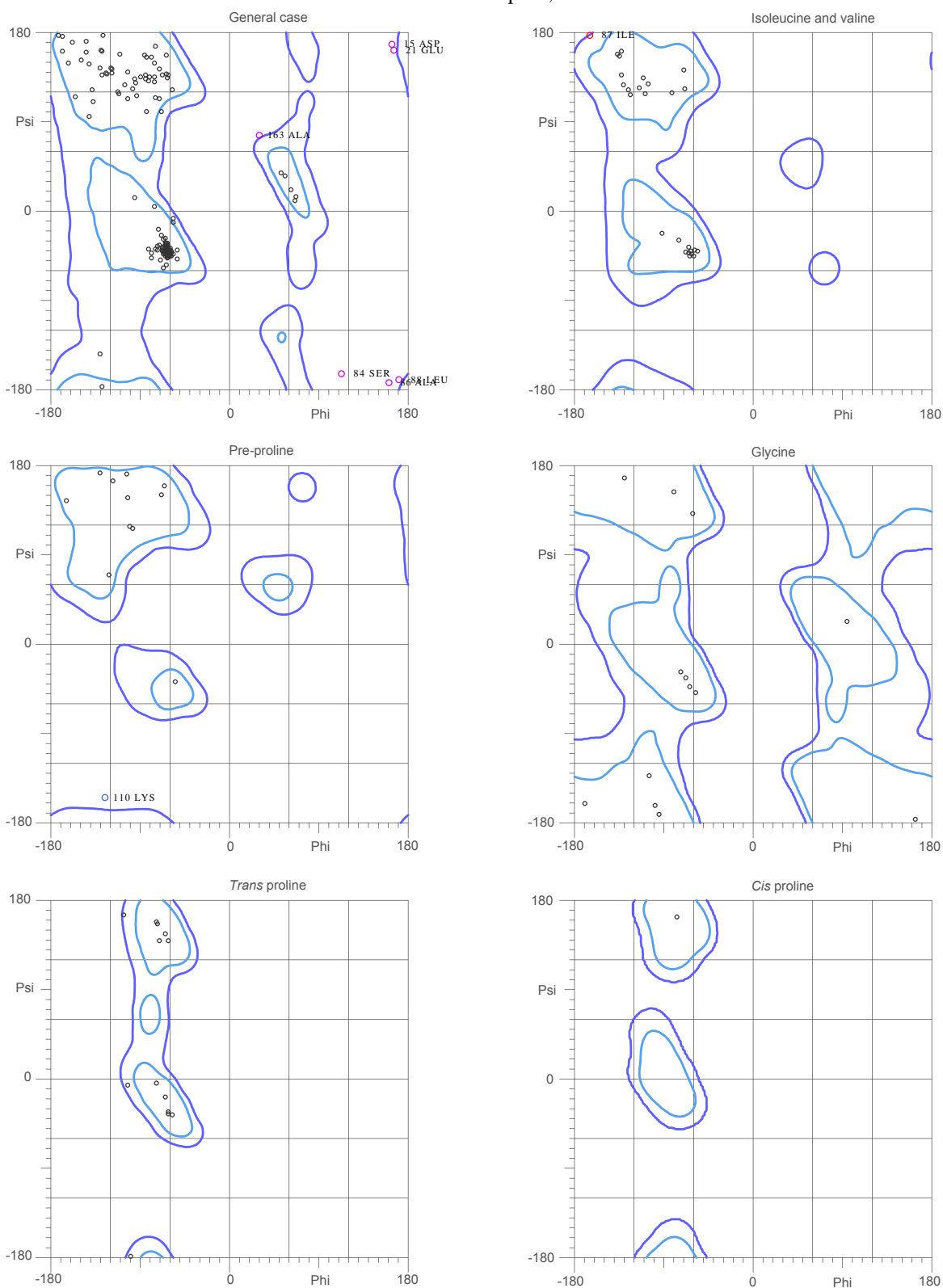


# MolProbity Ramachandran analysis

1at3A.B99990018.pdb, model 1



93,5% (201/215) of all residues were in favored (98%) regions.  
96,3% (207/215) of all residues were in allowed (>99,8%) regions.

There were 8 outliers (phi, psi):

15 ASP (164,4, 169,6)  
21 GLU (166,4, 163,5)  
84 SER (113,1, -164,2)  
86 ALA (161,4, -173,8)  
87 ILE (-165,5, 178,4)  
88 LEU (171,1, -170,1)  
110 LYS (-126,7, -155,3)  
163 ALA (30,7, 77,8)