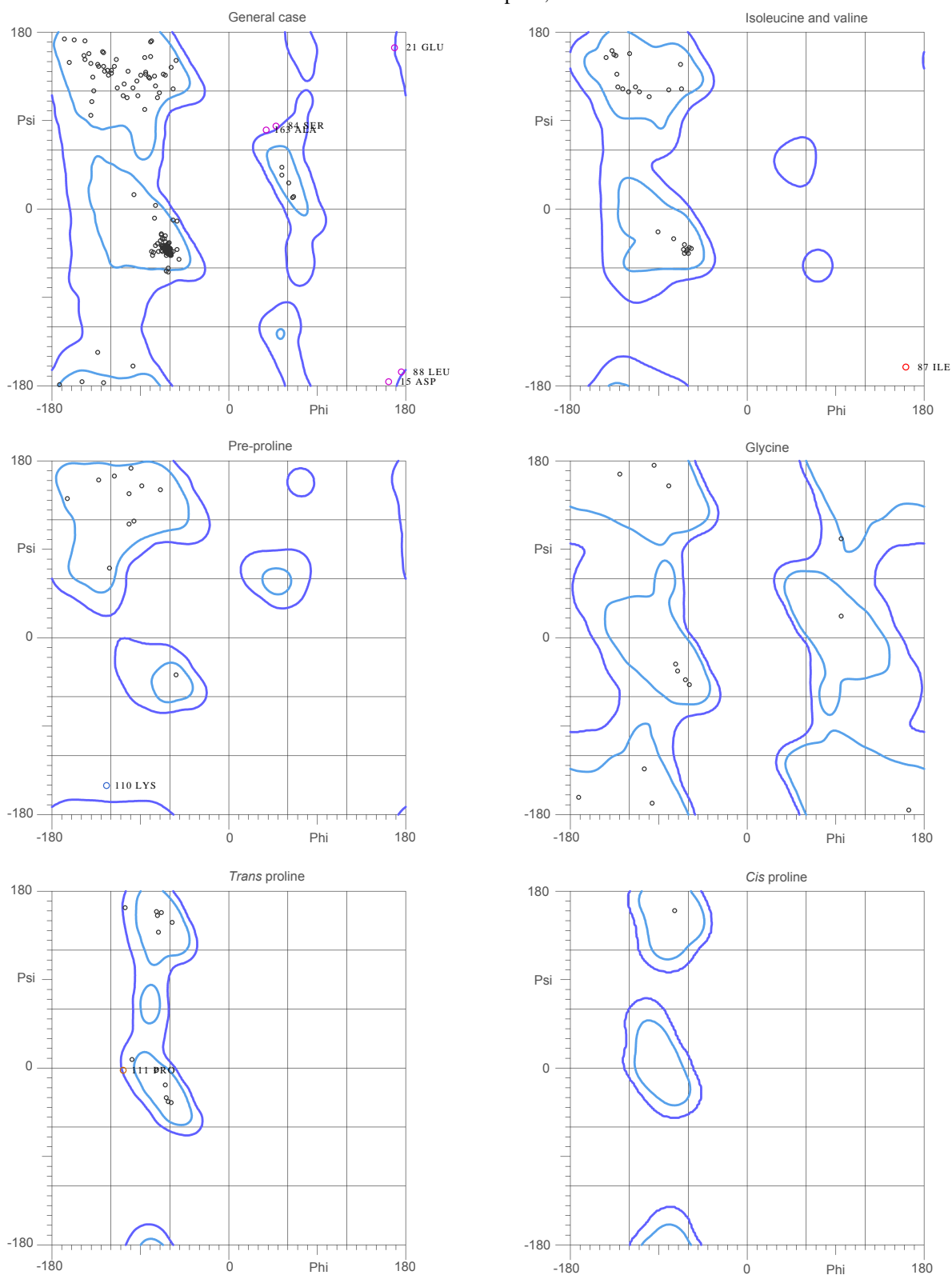


# MolProbity Ramachandran analysis

1at3A.B99990036.pdb, model 1



92,6% (199/215) 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 (163,4, -176,5)  
21 GLU (169,3, 165,3)  
84 SER (48,7, 85,0)  
87 ILE (162,9, -161,6)  
88 LEU (176,7, -166,9)  
110 LYS (-125,7, -151,0)  
111 PRO (-108,8, -2,1)  
163 ALA (38,4, 81,8)