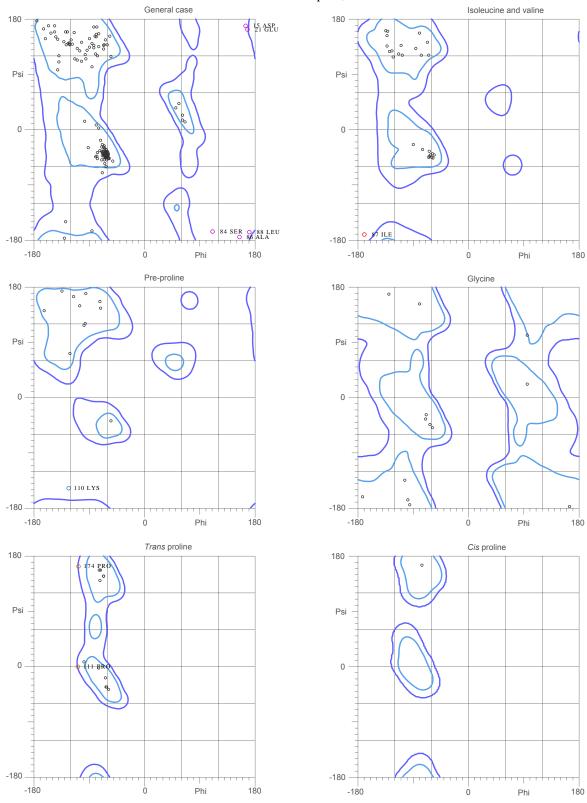
## MolProbity Ramachandran analysis

1at3A.B99990029.pdb, model 1



93,5%~(201/215) of all residues were in favored (98%) regions

95,8% (206/215) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi):
15 ASP (165.4, 170.2)
21 GLU (168.9, 164.8)
84 SER (111.8, -167.0)
86 ALA (155.1, -175.0)
87 ILE (-170.1, -171.1)
88 LEU (171.3, -167.6)
110 LYS (-124.2, -148.2) 111 PRO (-109,8, -0,7) 174 PRO (-108,5, 164,5)