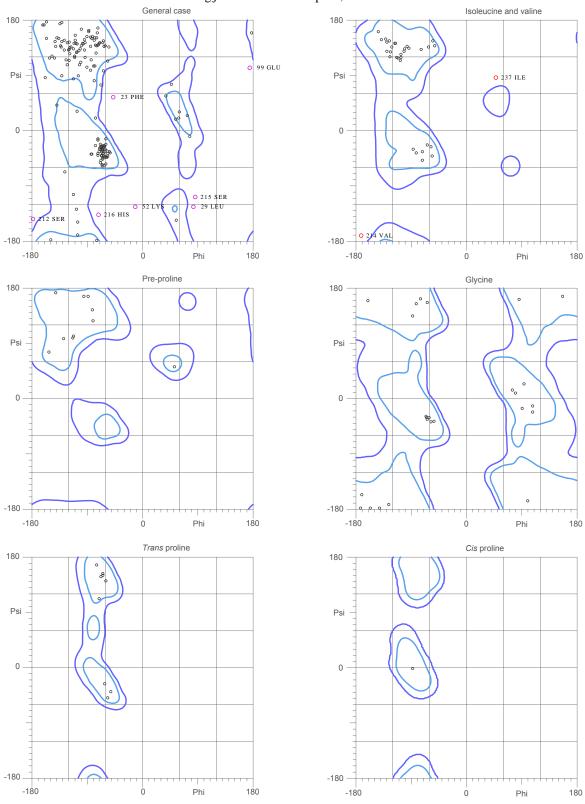
## MolProbity Ramachandran analysis

1agjA.B99990045.pdb, model 1



90,0%~(216/240) of all residues were in favored (98%) regions

96,2% (231/240) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 23 PHE (-48,5, 55,2) 29 LEU (83,2, -124,1) 52 LYS (-123, -124,6) 99 GLU (175,8, 103,9) 212 SER (-179,3, -144,2) 214 VAL (-172,3, -171,8) 215 SER (86,5, -108,5) 216 HS (-72,6, -137,4) 216 HIS (-72,6, -137,4) 237 ILE (48,2, 87,8)