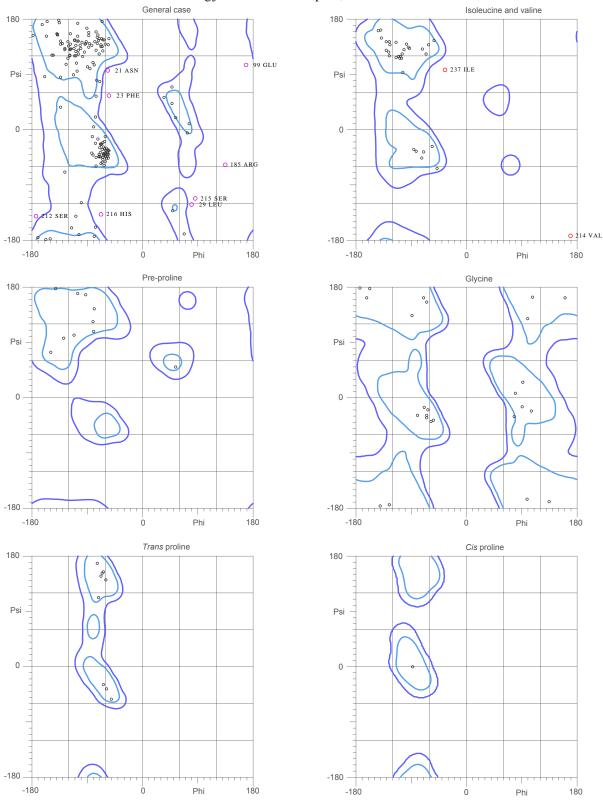
## MolProbity Ramachandran analysis

1agjA.B99990002.pdb, model 1



87,1%~(209/240) of all residues were in favored (98%) regions. 95,8%~(230/240) of all residues were in allowed (>99.8%) regions

There were 10 outliers (phi, psi):
21 ASN (-58,0, 97,3)
23 PHE (-55,9, 56,9)
29 LEU (80,2 - 122,4)
99 GLU (170,0, 106,4)
185 ARG (135,1, -57,4)
212 SER (-174,4, -141,4)
214 VAL (171,0, -173,8)
215 SER (86,8, -112,1)
216 HIS (-68,6, -138,5)
237 ILE (-35,8, 98,8)