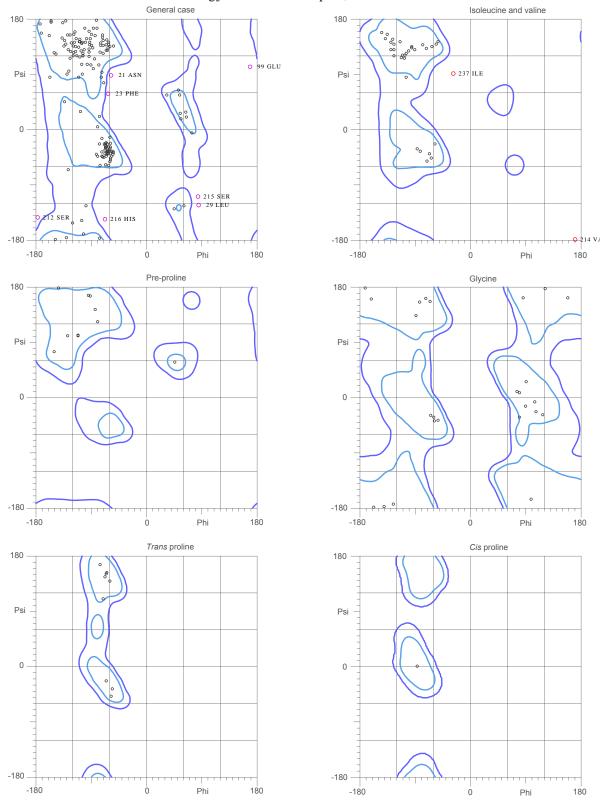
MolProbity Ramachandran analysis

1agjA.B99990033.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): 21 ASN (-58.5, 89.0) 23 PHE (-63.7, 59.2) 29 LEU (85.6, -123.6) 99 GLU (169.5, 103.6) 212 SER (-179.0, -143.6) 214 VAL (171.1, -179.7) 215 SER (84.7, -109.1) 216 HIS (-68,6, -146,4) 237 ILE (-28,3, 92,3)