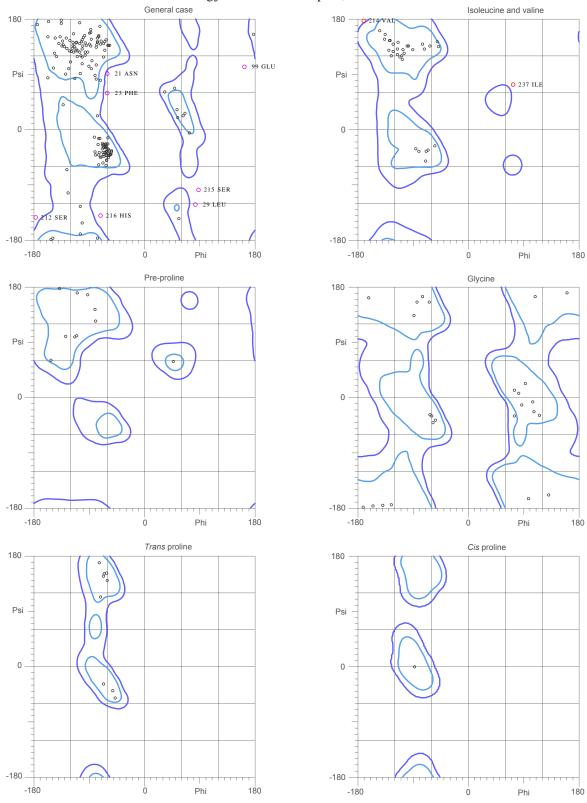
MolProbity Ramachandran analysis

1agjA.B99990009.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 (-61,2, 92,0) 23 PHE (-61,4, 60,7) 29 LEU (83,5, -122,7) 99 GLU (163,5, 103,8) 212 SER (-178,1, -143,1) 215 SER (88,8, 98,9) 216 HS (-72, 1 -140,9) 216 HIS (-72,1, -140,9) 237 ILE (73,5, 74,1)