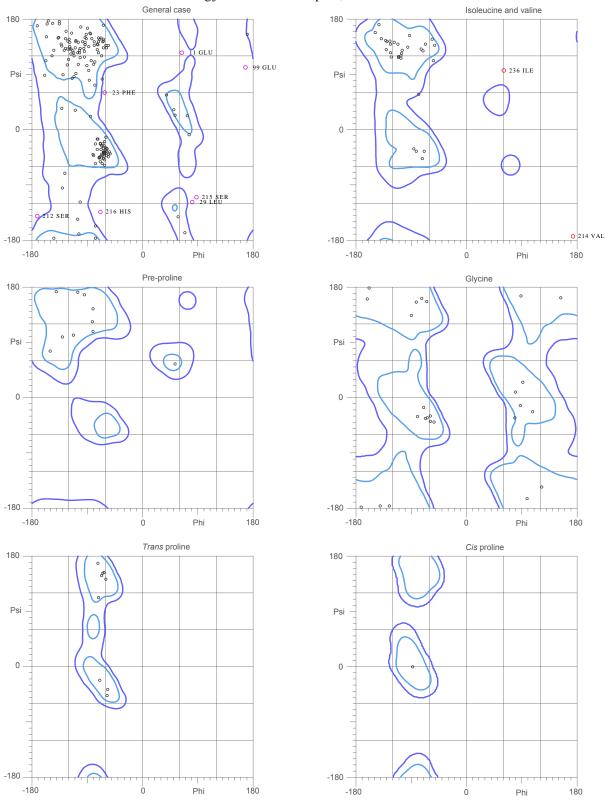
MolProbity Ramachandran analysis

1agjA.B99990024.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):
11 GLU (64.3, 126.1)
23 PHE (-62.9, 61.1)
29 LEU (81.8, -118.2)
99 GLU (168.3, 102.6)
212 SER (-172.6, -141.6)
214 VAL (174.2, -174.4)
215 SER (88.1, -110.9)
216 IMS (6-03 - 135.0) 216 HIS (-69,3, -135,0) 236 ILE (61,7, 97,9)