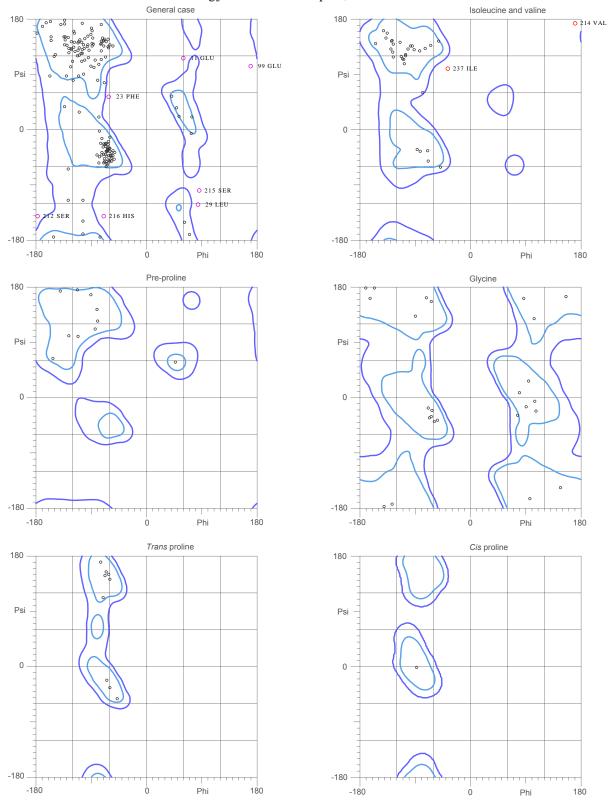
MolProbity Ramachandran analysis

1agjA.B99990039.pdb, model 1



89,2%~(214/240) 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 (60.4, 117.0)
23 PHE (-62.4, 54.9)
29 LEU (84.9, -122.1)
99 GLU (170.1, 104.3)
212 SER (-178.6, -141.9)
214 VAL (171.1, 174.4)
215 SER (66.8, 99.8)
216 HIS (-70.4, -141.4)
237 ILE (-37.4, 100.2)