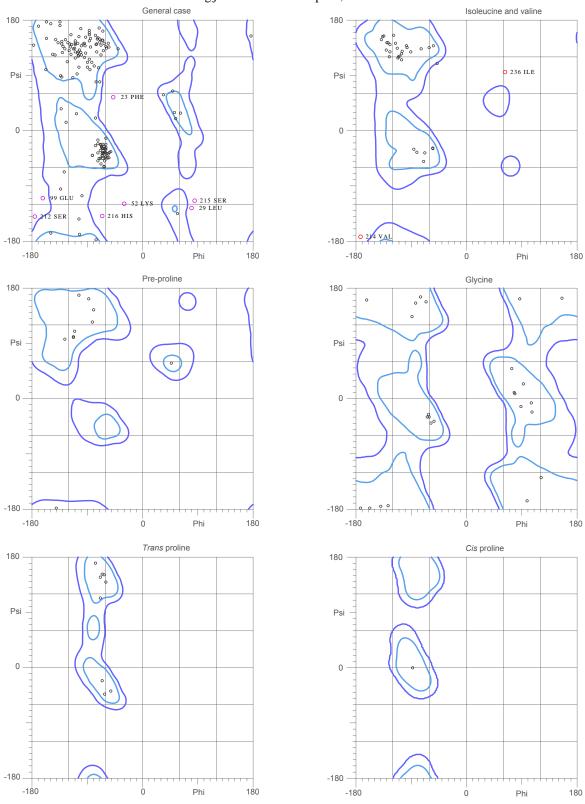
MolProbity Ramachandran analysis

1agjA.B99990011.pdb, model 1



89,6%~(215/240) 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): 23 PHE (-48,0,55,8) 29 LEU (80,6,-126,4) 52 LYS (-30,4,-119,9) 99 GLU (-163,5,-110,2) 212 SER (-176,1,-141,0) 214 VAL (-173,7,-173,4) 215 SER (85,3,-114,8) 216 HS (-66,8-139,1) 216 HIS (-66,8, -139,1) 236 ILE (63,1, 97,0)