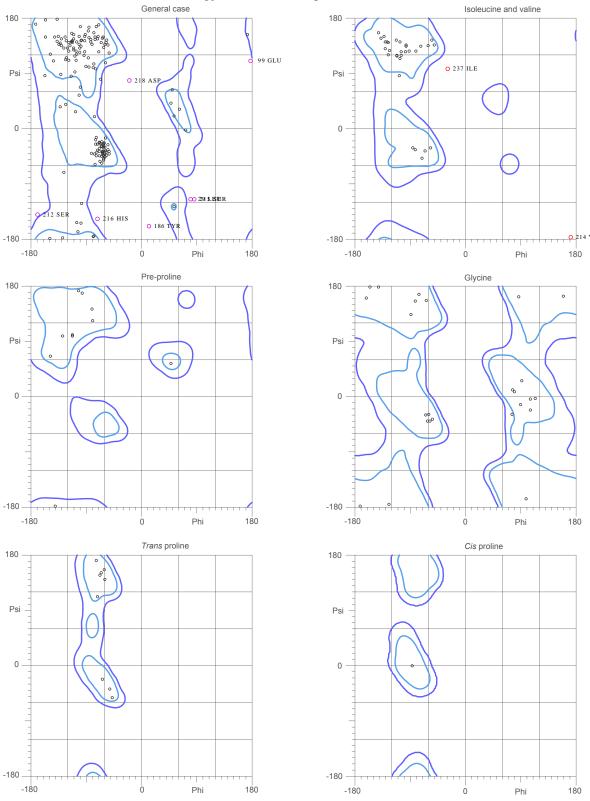
MolProbity Ramachandran analysis

1agjA.B99990018.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): 29 LEU (80,2, -116,0) 99 GLU (178,7, 111,1) 99 GLU (178,7, 111,1) 186 TYR (13,0, -159,8) 212 SER (-170,9, -140,8) 214 VAL (172,5, -177,1) 215 SER (86,6, -115,3) 216 HIS (-72,3, -147,7) 218 ASP (-20,7, 79,1) 237 ILE (-29,2, 98,9)