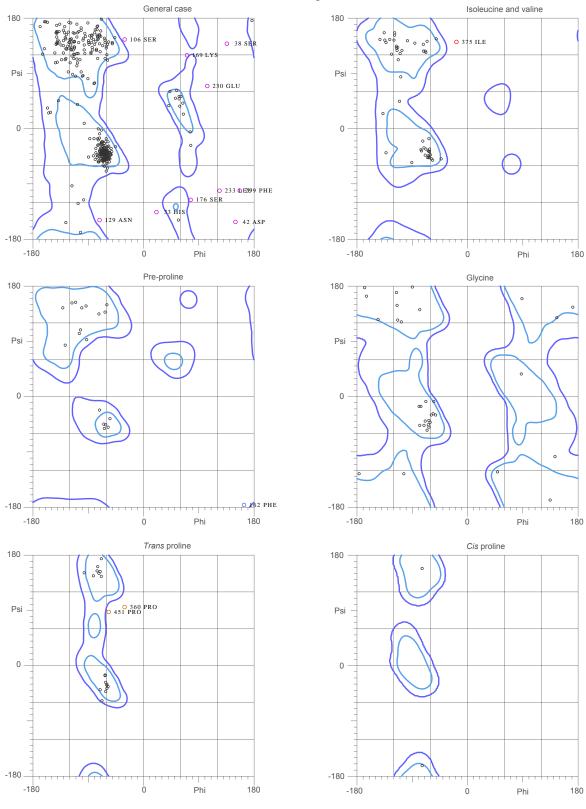
## MolProbity Ramachandran analysis

1ac5\_.B99990022.pdb, model 1



90,2% (434/481) of all residues were in favored (98%) regions.

33 HIS (21,2, -136,5) 38 SER (136,5, 139,5) 42 ASP (150,5, -152,6) 106 SER (-31,7, 146,0) 129 ASN (-72,7, -149,3) 162 PHE (164,3, -177,6)

106 SER (-31,7, 146,0) 129 ASN (-72,7, -149,3) 162 PHE (164,3, -177,6) 169 LYS (71,6, 120,7) 176 SER (77,7, -116,6) 230 GLU (104,9, 70,2) 233 LEU (124,8, -101,7) 199 PHE (157,2, -101,1) 160 PRO (-31,7, 96,2)