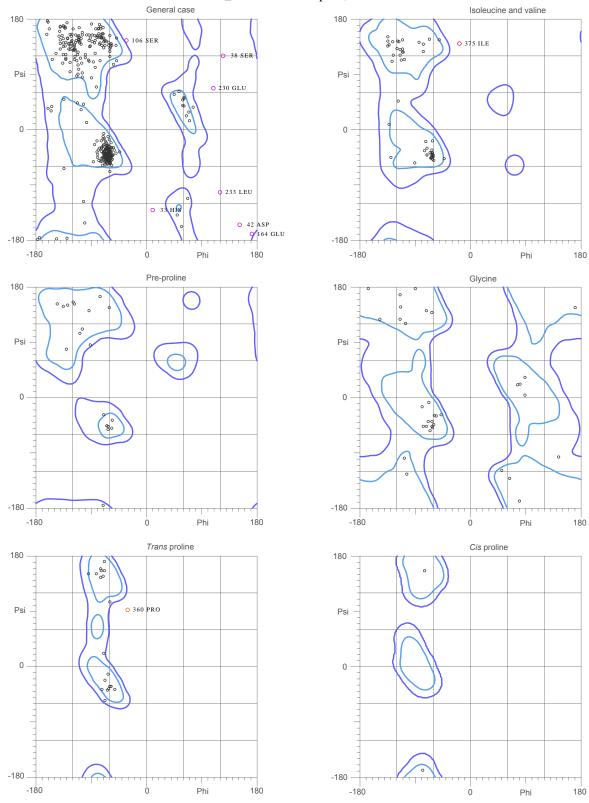
MolProbity Ramachandran analysis

1ac5_.B99990042.pdb, model 1



91,7%~(441/481) of all residues were in favored (98%) regions

98,1% (472/481) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 33 HIS (10,3, -131,3) 38 SER (125,9, 121,8) 38 SER (125,9, 121,8) 42 ASP (152,9, -155,5) 106 SER (-33,7, 146,5) 164 GLU (172,2, -170,9) 230 GLU (109,5, 68,8) 233 LEU (121,0, -102,0) 360 PRO (-31,1, 93,9) 375 ILE (-18,2, 141,4)