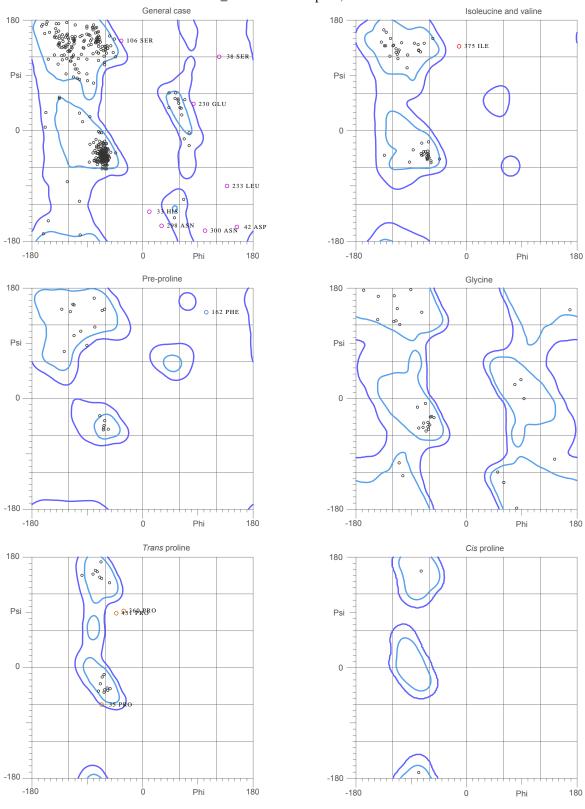
MolProbity Ramachandran analysis

1ac5_.B99990043.pdb, model 1



92,1% (443/481) of all residues were in favored (98%) regions.

79.73% (468/481) of all residues w There were 13 outliers (ph., ps): 33 HIS (11,5-122.3) 33 FRO (4-78, -00.6) 33 SER (125-3, -10.6) 42 ASP (154.8, -157.2) 106 SER (5-35.4, 121.9) 162 PHE (104.6, 141.4) 220 GLU (33, 74.48) 223 LEU (138.1, -90.9) 298 ASN (31,9-155.6) 300 ASN (102.2, -163.5) 300 FRO (-31.5, 93.8) 375 ILE (-12.3, 138.4) 451 PRO (-43.8, 89.8)