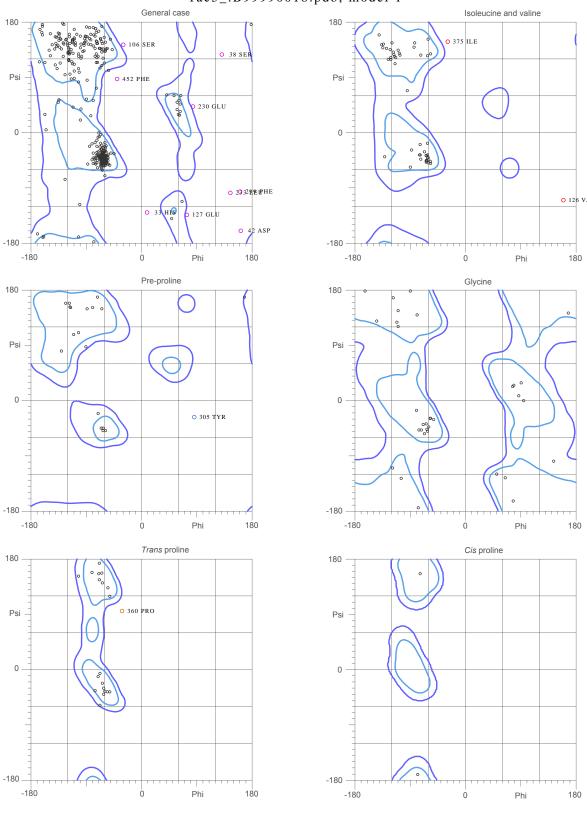
MolProbity Ramachandran analysis

1ac5_.B99990018.pdb, model 1



91,3% (439/481) of all residues were in favored (98%) regions.

There were 13 outliers (phi, pai):
33 HIS (9.1, -130.9)
38 SER (131.4 (28.9)
42 ASP (162.2, -160.1)
105 SER (-30.8, 144.7)
120 VAL (160.5, -110.3)
127 GLU (74.2, -134.3)
230 GLU (84.3, -34.6)
231 LEU (140.0, -98.3)
299 PHE (161.0, -96.6)
305 TYR (86.3, -27.7)
360 PRO (-32.1, 97.0)
375 ILE (-29.8, -149.6)