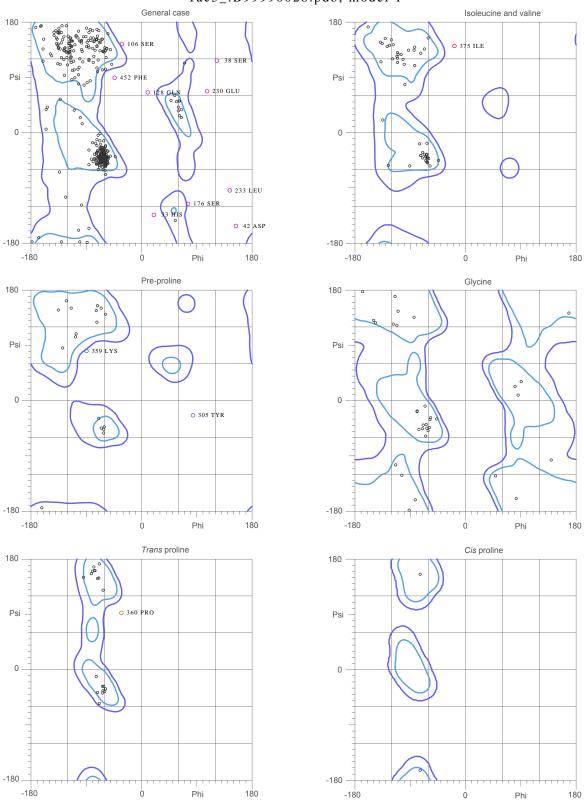
MolProbity Ramachandran analysis

1ac5_.B99990028.pdb, model 1



Se (468/48) of all residues w ere wer. 13 outliers (ph. ps.): 38 SIRs (123.6, 118.2) 42 ASP (154.2, 155.9) 106 SEP (22.1, 145.9) 107 SEP (23.1, 145.9) 108 SEP (23.1, 145.9) 108 SEP (23.1, 145.9) 109 SEP (23.1, 145.9) 109 SEP (24.1, 145.9) 109