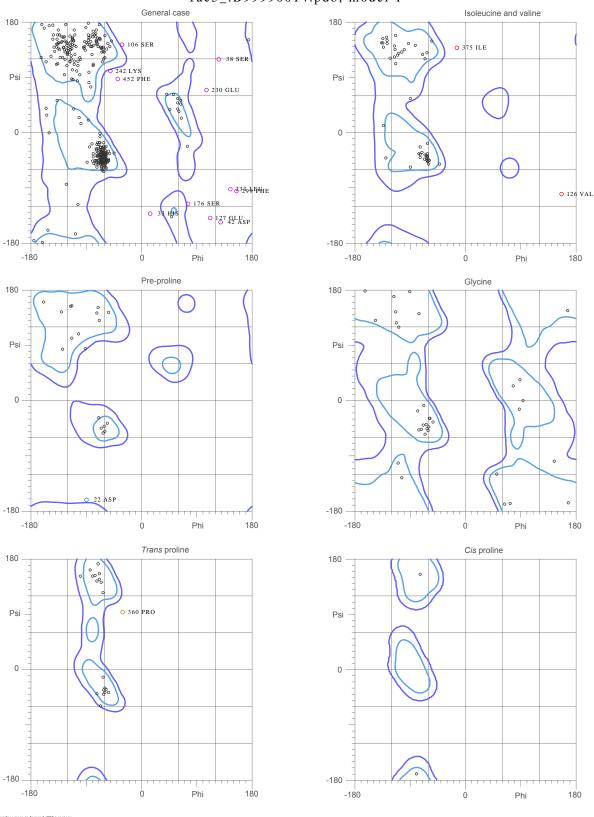
MolProbity Ramachandran analysis

1ac5_.B99990014.pdb, model 1



91,5% (440/481) of all residues were in favored (98%) regions. 96,9% (466/481) of all residues were in allowed (>99.8%) regio

There were 15 outliers (ph. 22 ASP (9.03, .162.4) 33 HIS (14.6, .132.9) 38 SER (12.6, 2.120.2) 38 SER (12.6, 2.120.2) 106 SER (.33.0, 144.9) 102 VAL (157.5, .106.8) 127 GLU (112.1, .193.3) 127 GLU (112.1, .193.3) 123 LEU (14.7, .195.3) 125 LEU (14.7, .195.3) 126 DEP (14.7, .195.3) 146 DEP (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .195.3) 147 (14.7, .196.3) 147 (14.7, .