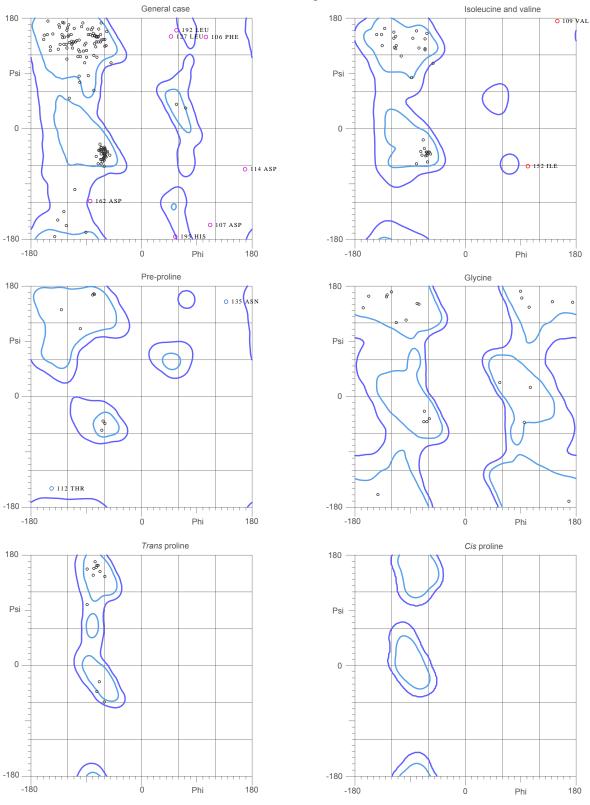
MolProbity Ramachandran analysis

1ad2_.B99990016.pdb, model 1



89.6%~(199/222) of all residues were in favored (98%) regions. 95.0%~(211/222) of all residues were in allowed (>99.8%) regions

There were 11 outliers (phi, psi): 106 PHE (105,9, 150,1) 107 ASP (112,8, -157,6) 109 VAL (150,3, 176,8) 112 THR (-148,0, -150,8) 114 THR (-148,0, -150,8) 114 ASP (169,8, -66,7) 127 LEU (48,5, 151,4) 135 ASN (138,8, 155,1) 152 ILE (102,3, -61,2) 162 ASP (-84,5, -118,6) 192 LEU (37,9, 161,6) 195 HIS (55,1, -177,0)