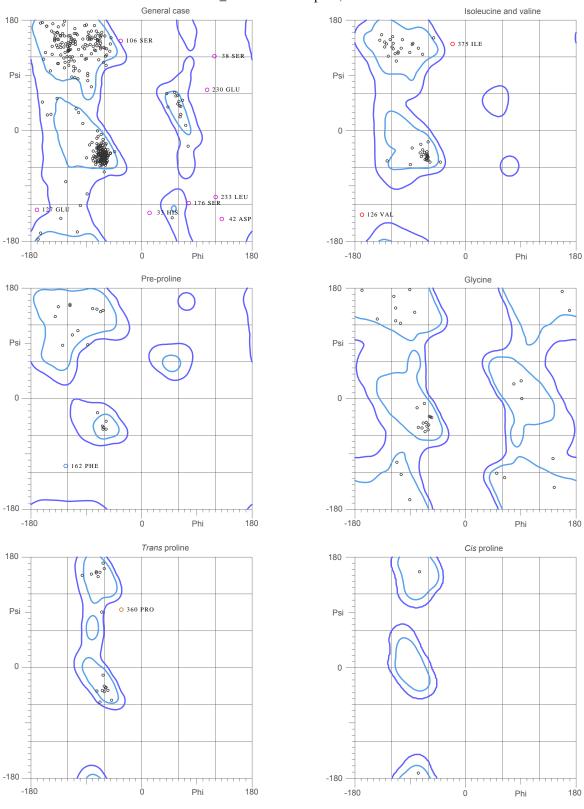
## MolProbity Ramachandran analysis

1ac5\_.B99990047.pdb, model 1



92,1% (443/481) of all residues were in favored (98%) regions. 97,5% (469/481) of all residues were in allowed (>99.8%) regions

7,5% (469/481) of all residues we here were 12 outliers (phi, psi): 33 HIS (134, -134.6) 38 EER (119.8, 122.5) 42 ASP (131.2, 144.5) 106 SER (34.9, 147.7) 126 VAL (169.4, -137.9) 127 GLU (+171,4, -129.8) 127 GLU (+171,4, -129.8) 129 EFR (77.4, -118.3) 230 GLU (107.9, 67.1) 233 LEU (121.7, -108.4) 360 PRO (33.0, 55.9) 375 ILE (-21.5, 143.0)