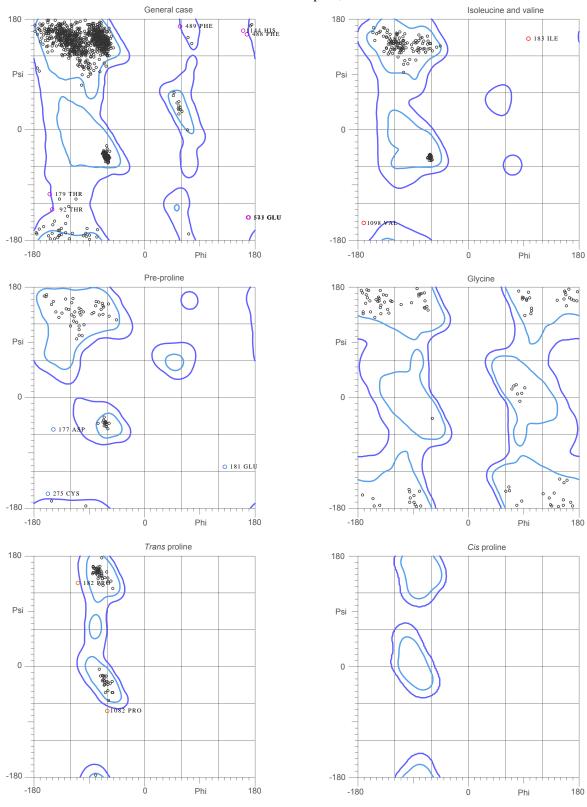
## MolProbity Ramachandran analysis

6abu.B99990043.pdb, model 1



96,1% (1251/1302) of all residues were in favored (98%) regions 98.9% (1288/1302) of all residues were in allowed (>99.8%) regi

There were 14 outliers (phi, p 92 THR (-151,8, -130,4) 171 GLU (169,9, -143,4) 177 ASP (-149,0, -52,1) 179 THR (-155,3, -105,6) 181 GLU (131,6, -113,1) 182 PRO (-109,2, 137,2) 183 ILF (98,1, 142,2)

181 GLU (131,6, -113,1) 182 PRO (-109,2, 137,2) 183 ILE (98,1, 149,2) 175 CYS (-158,6, -157,2) 188 PHE (167,7, 156,6) 189 PHE (58,1, 169,0) 143 GLU (170,0, -143,6) 082 PRO (-61,8, -72,1)