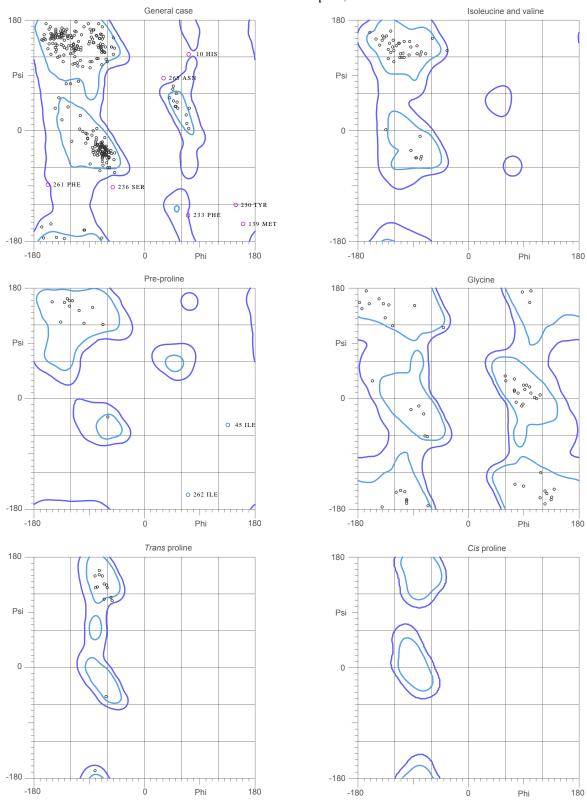
## MolProbity Ramachandran analysis

1a12A.B99990047.pdb, model 1



92,2%~(368/399) of all residues were in favored (98%) regions

97,7% (390/399) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 10 HIS (72,6, 125,5) 45 ILE (136,4, -43,7) 45 ILE (136,4, -43,7) 139 MET (161,4, -152,4) 230 TWR (149,1, -121,3) 233 PHE (71,8, -138,9) 236 SER (-52,2, -92,6) 261 PHE (-158,6, -89,0) 262 ILE (71,1, -157,8) 265 ASN (31,3, 86,6)