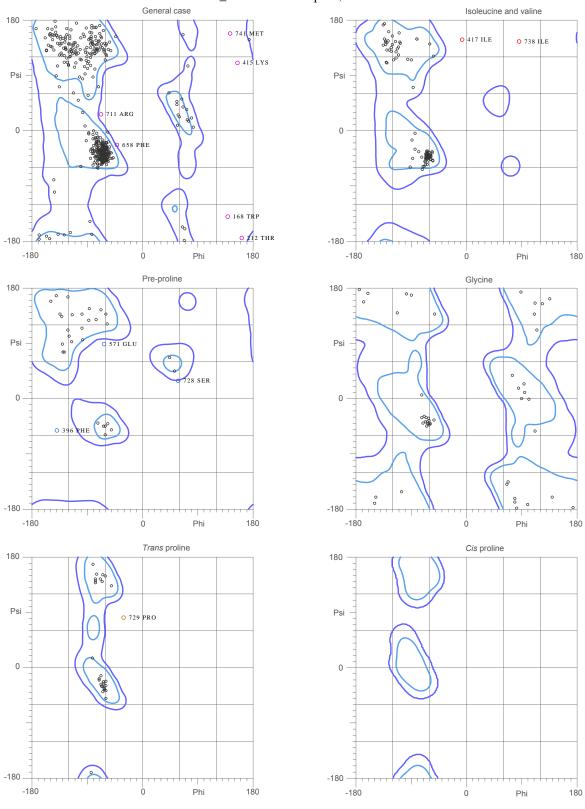
## MolProbity Ramachandran analysis

1a8i\_.B99990001.pdb, model 1



94.8% (768/810) of all residues were in favored (98%) regions. 98,5% (798/810) of all residues were in allowed (>99.8%) regions

.38 (1984.19) of all residues we see were 12 outliers (phi, psi): 168 TRP (140,0, -140,2) 212 THR (163,0, -175,6) 336 PHE (140,0, -52,5) 415 LYS (155,1, 111,6) 417 LE (-7,5, 149,7) 571 GLU (-63,1, 89,2) 658 PHE (-42, -32,5) 711 ARG (-68,9, 28,0) 728 SER ((58,7, 29,4) 729 PRO (-31,8, 82,6) 738 ILE (68,6, 146,7) 741 MET (143,6, 159,1)