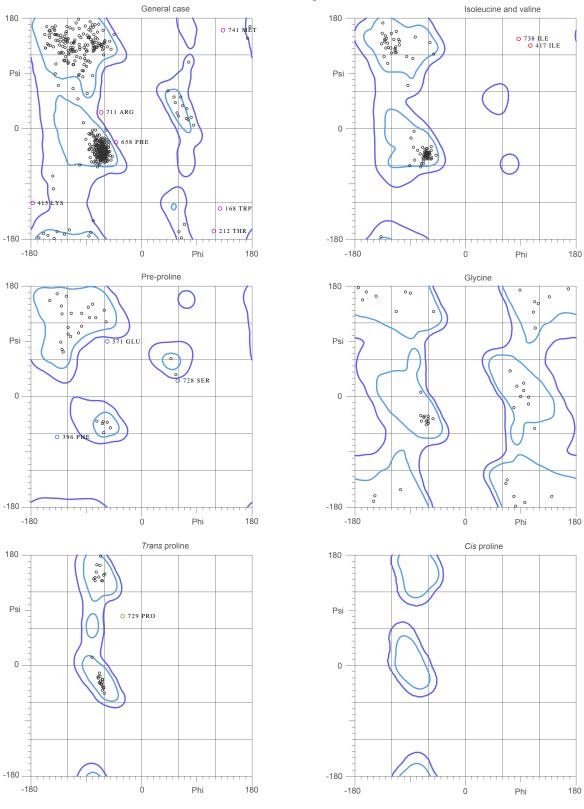
## MolProbity Ramachandran analysis

1a8i\_.B99990023.pdb, model 1



95,1% (770/810) of all residues were in favored (98%) regions. 98,5% (798/810) of all residues were in allowed (>99.8%) regions

.38 (198.10) of all residues we see were 12 outliers (phi, psi): 168 TRP (128.5, -130.3) 212 THR (118.0, -167.2) 336 PHE (128.3, -66.1) 415 LYS (-178.1, -121.1) 417 ILE (106.3, 136.5) 571 GLU (-56.8, 90.2) 658 PHE (-42.4, -32.0) 711 ARG (-66.7, 27.8) 728 ER (69.1, 26.7) 729 PRO (-31.2, 81.9) 738 ILE 67.6, 147.4) 741 MET (133.1, 161.3)