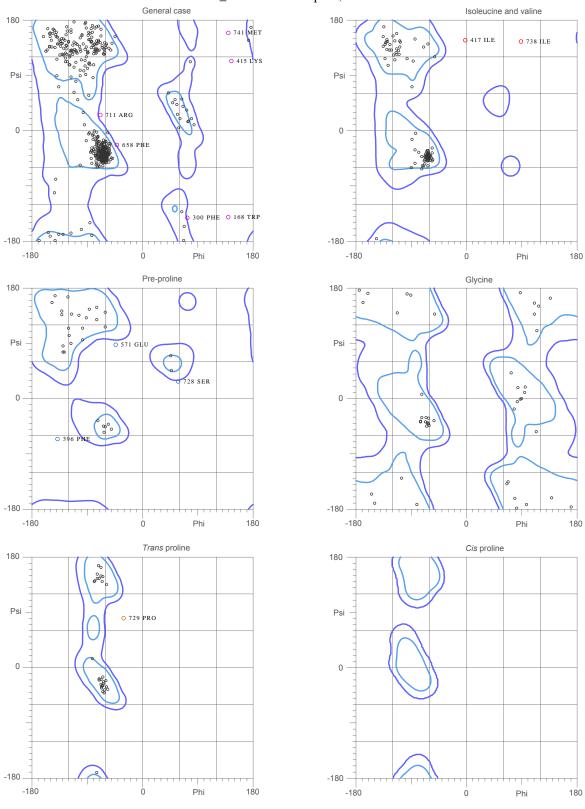
## MolProbity Ramachandran analysis

1a8i\_.B99990040.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

There were 12 outliers (phi, p 168 TRP (140,7, -141,9) 300 PHE (73,6, -142,3) 396 PHE (-139,3, -66,0) 415 LYS (145,2, 114,6) 417 ILE (-2,5, 148,7) 571 GLU (-44,4, 88,4)

115 LYS (145.2, 114.6) 117 ILE (-2.5, 148.7) 571 GLU (-44.4, 88.4) 558 PHE (-42.8, -23.1) 711 ARG (-69.4, 26.6) 728 SER (58.0, 28.3) 729 PRO (-31.7, 81.9) 738 ILE (89.8, 146.5)