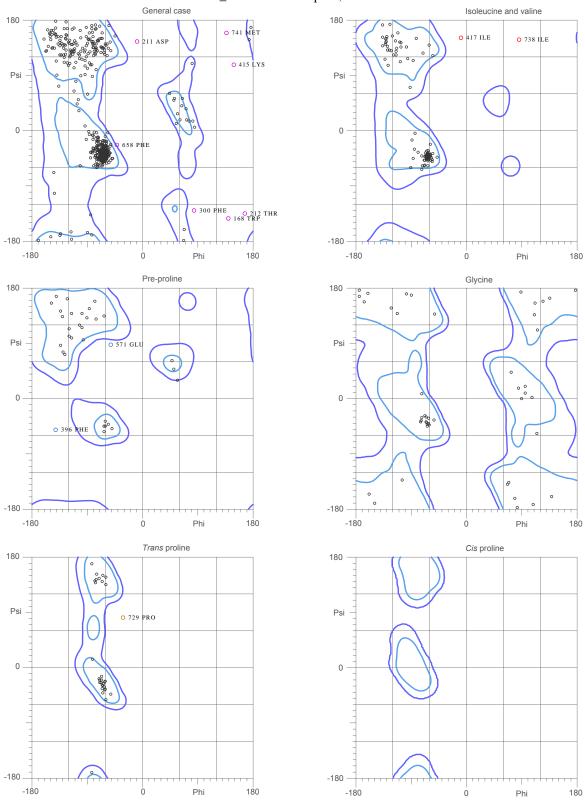
MolProbity Ramachandran analysis

1a8i_.B99990004.pdb, model 1



94,9% (769/810) 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 (140, . 143,4) 211 ASP (-9.7, 146,9) 212 TRR (167, -135,1) 300 PHE (84.9, -130,6) 300 PHE (84.8, -1.30,6) 350 PHE (142.8, -51.8) 415 LYS (149.2, 108.3) 417 LEC (9.1, 152.0) 571 GLU (52.8, 88.8) 68 PHE (42.9, -2.3.0) 729 PRO (32.4, 82.5) 738 ILE (86.0, 149.3) 741 MET (137.8, 160.4)