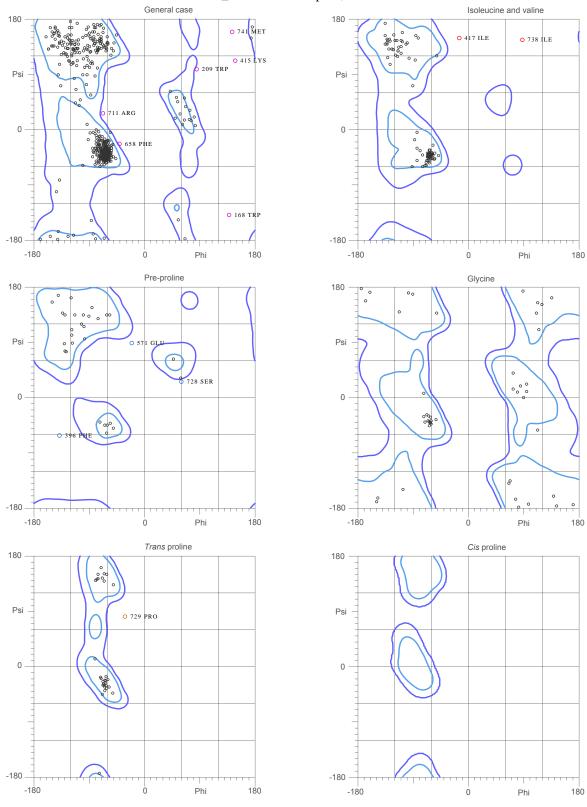
MolProbity Ramachandran analysis

1a8i_.B99990012.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

98,5% (798/810) of all residues were in allowed (>99.8%) region

5.5% (7.9% tu) of all residues we here were 12 outliers (phi, pai): 168 TRP (138.3, -139.9) 209 TRP (83.3, 99.1) 309 FIRE (139.4, -62.4) 415 LYS (148.7, 113.9) 415 LYS (148.7, 113.9) 571 GLU (-22.0, 89.9) 658 FIRE (-419, -23.8) 711 ARG (-68.0, 27.8) 728 SER (60.2, 26.7) 729 FRO (32.2, 82.7) 738 ILE (856, 148.0) 741 MET (143.6, 160.2)