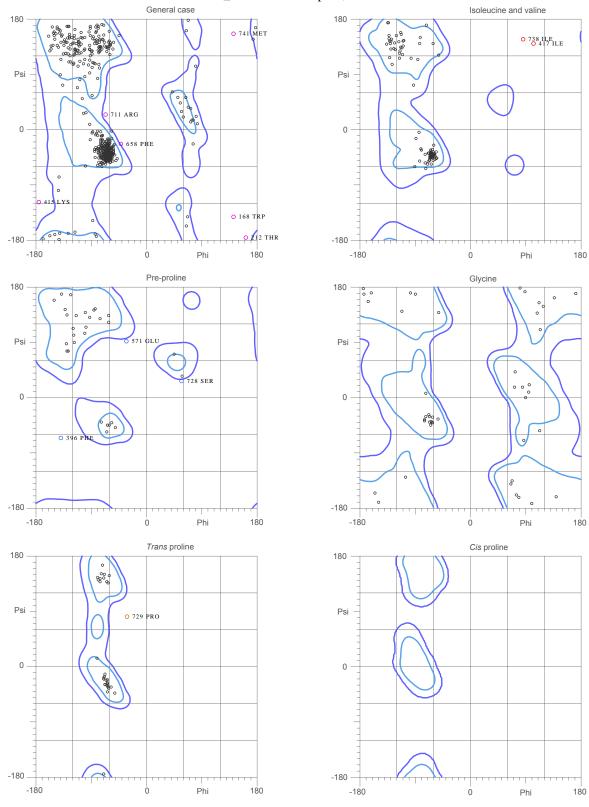
MolProbity Ramachandran analysis

1a8i_.B99990011.pdb, model 1



94,7% (767/810) 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, 168 TRP (142,9, -142,3) 212 THR (162,1, -176,4) 396 PHE (140,3, -66,6) 415 LYS (-176,2, -118,5) 417 ILE (103,8, 141,8) 571 GLU (-33,6, 92,1) 658 PHE (-42,6, -23,8) 711 ARG (-67,2, 25,4) 728 SER (57,5, 27,4) 729 PRO (-32,8, 82,8) 738 ILE (86,8, 148,6)