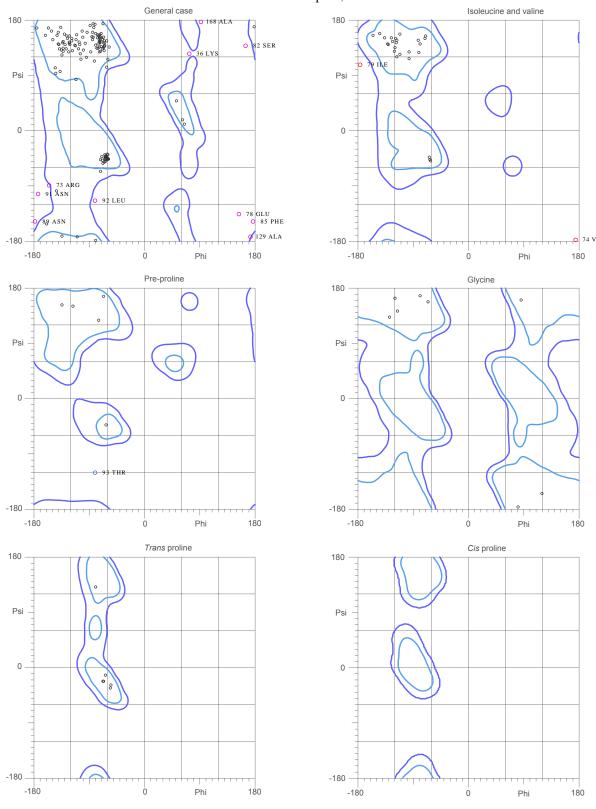
MolProbity Ramachandran analysis

1aisB.B99990013.pdb, model 1



90.1% (172/191) of all residues were in favored (98%) regions.

nere were 13 outliers (phi, 36 LYS (73.3, 126.6) 36 LYS (73.3, 126.6) 73 ARG (1-16.3, 389.7) 74 VAL (175.9, -178.4) 79 LE (-177.7, 108.4) 82 SER (165.4, 139.6) 85 PHE (177.4, -148.2) 89 ASN (-179.2, -148.9) 91 ASN (-179.2, -148.9) 92 LEU (-81.3, -114.4) 93 THR (-81.8, -121.7) 29 ALA (173.3, -173.2)