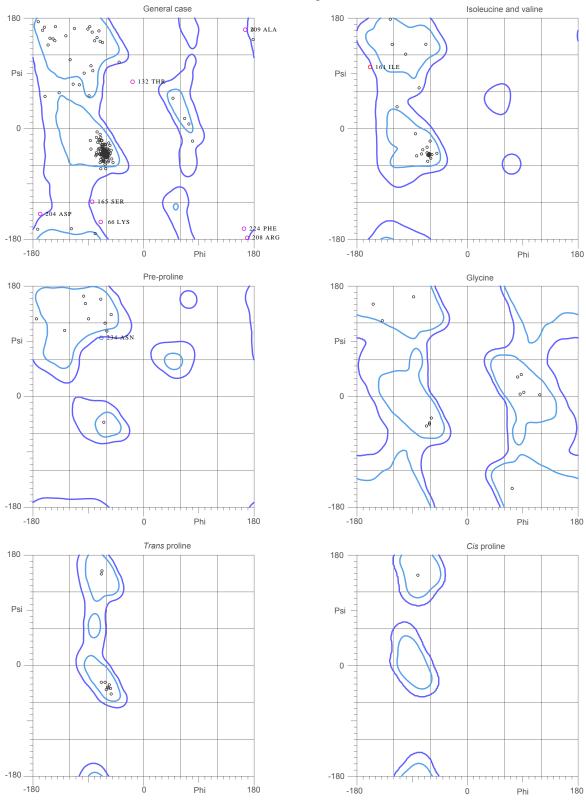
MolProbity Ramachandran analysis

1aoa_.B99990041.pdb, model 1



89,8%~(220/245) of all residues were in favored (98%) regions

96,3% (236/245) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 66 LYS (-70.9, -152.3) 132 THR (-18.4, 77.4) 161 ILE (-159.1, 101.0) 165 SER (-84.6, -119.1) 204 ASP (-169.7, -139.3) 208 ARG (169.3, -178.3) 209 ALA (166.1, 162.1) 274 PHE (164.8, -16.5 8) 224 PHE (164,8, -163,8) 234 ASN (-69,5, 96,4)