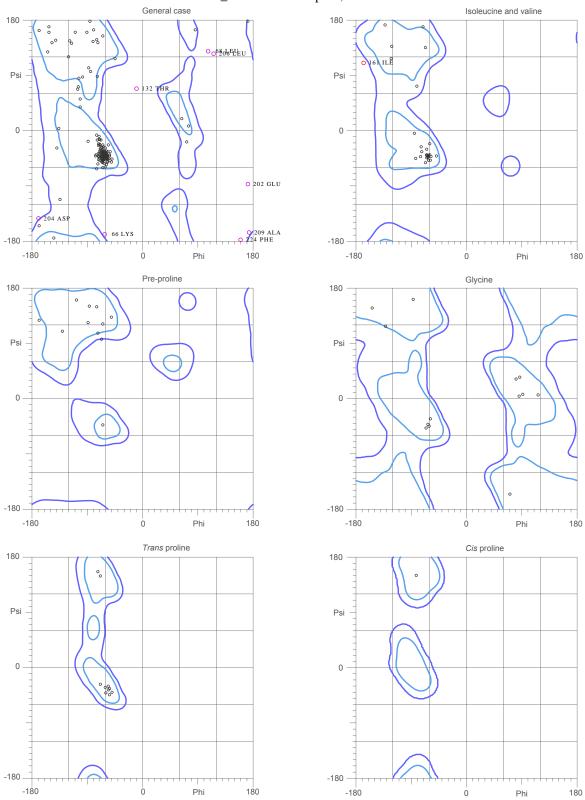
MolProbity Ramachandran analysis

1aoa_.B99990011.pdb, model 1



 $89,\!4\%$ (219/245) 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 (-62,2, -169,6) 68 LEU (107,6, 130,6) 132 THR (-10,1,69,9) 161 ILE (-168,1, 111,8) 202 GLU (172,6, -87,4) 204 ASP (-170,2, -143,1) 206 LEU (116,5, 126,3) 209 ALA (174,0, -166,1) 209 ALA (174,0, -166,1) 224 PHE (160,1, -178,5)