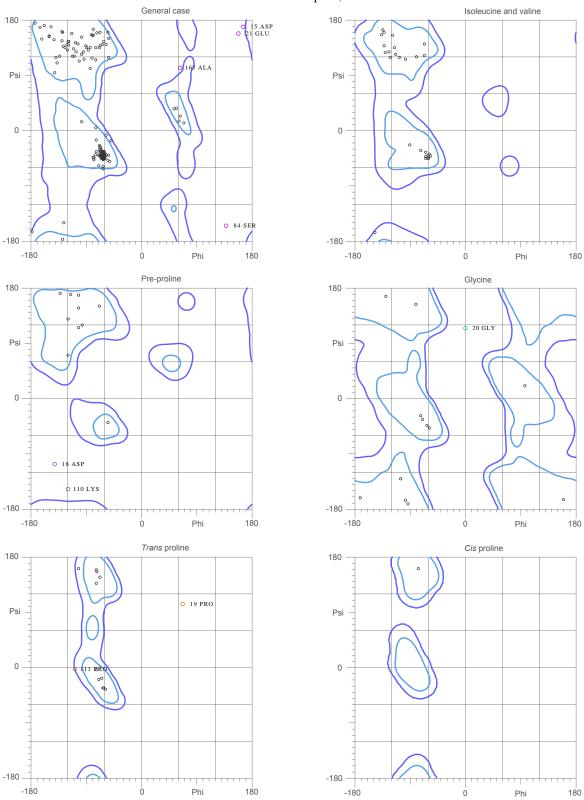
MolProbity Ramachandran analysis

1at3A.B99990025.pdb, model 1



92,6%~(199/215) of all residues were in favored (98%) regions

95,8% (206/215) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 15 ASP (166,5, 170,5) 18 ASP (-142,4, -107,0) 18 ASP (-142,4, -107,0) 19 PRO (67,5, 104,5) 20 GLY (0,2, 115,4) 21 GLU (158,6, 159,4) 84 SER (138,0, -155,7) 110 LYS (-120,4, -148,9) 111 PRO (-109,2, -2,1) 163 ALA (63,4, 103,2)