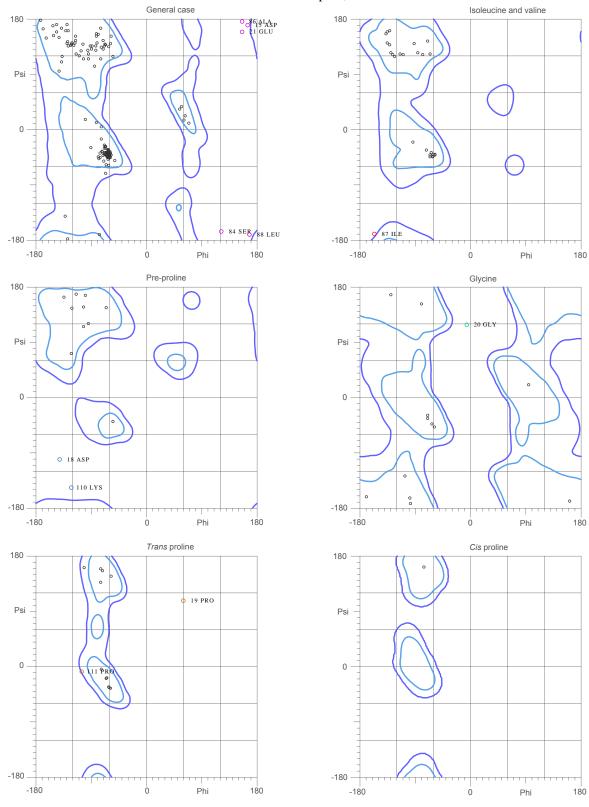
MolProbity Ramachandran analysis

1at3A.B99990034.pdb, model 1



93,5% (201/215) of all residues were in favored (98%) regions. 94,9% (204/215) of all residues were in allowed (>99.8%) regions

There were 11 outliers (phi., psi):
15 ASP (165,2, 171,0)
18 ASP (-142,4, -101,4)
19 PRO (604, 108.7)
20 GLY (-61, 119,3)
21 GLU (156,5, 166,8)
84 SER (122,3, -166,1)
86 ALA (156,1, 177,5)
87 ILE (-157,9, -170,8)
88 LEU (168,7, -171,0)
110 LYS (-123,9, -147,8)
111 PRO (-106,9, -8,4)