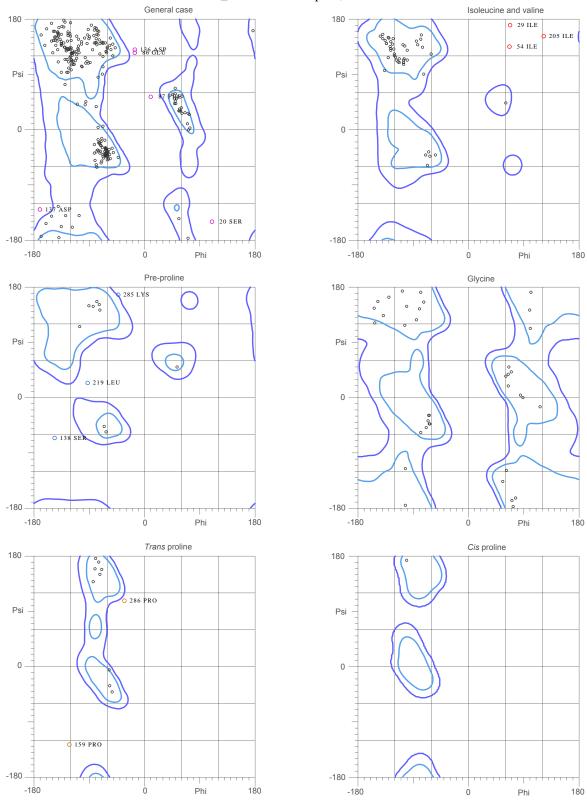
MolProbity Ramachandran analysis

1air_.B99990050.pdb, model 1



89,4% (313/350) of all residues were in favored (98%) regions.

there were 13 outliers (phi, p 20 SER (110,4, -150,1) 29 ILE (68,7, 171,3) 54 ILE (67,5, 136,1) 86 GLU (-16,2, 126,1) 87 PHE (10,5, 54,5) 136 ASP (-16,5, 131,3) 137 ASP (-171,2, -130,6) 138 SER (-147,6, -66,7) 159 PRO (-123,6, -127,4) 205 ILE (123,7, 153,2) 219 LEU (-94,0, 24,5) 285 LYS (-431, 168,4)