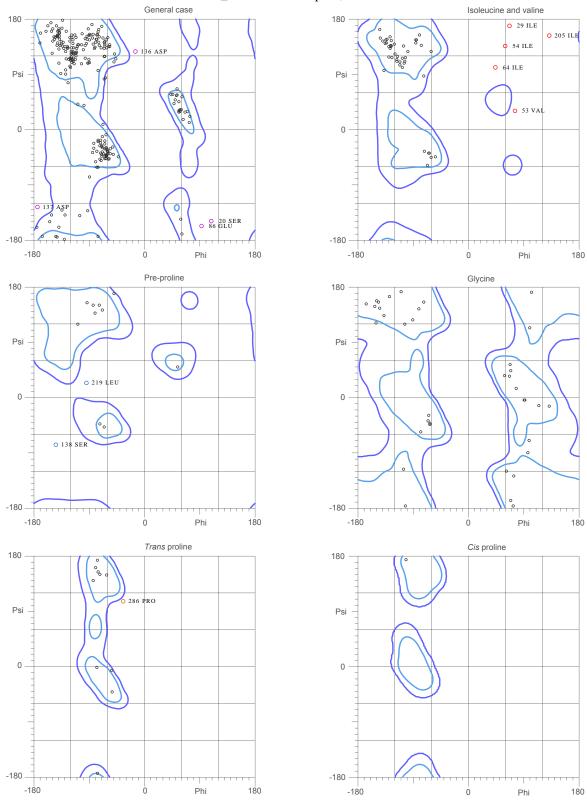
MolProbity Ramachandran analysis

1air_.B99990004.pdb, model 1



88,3% (309/350) of all residues were in favored (98%) regions. 96,6% (338/350) of all residues were in allowed (>99.8%) regions

96,6% (338/350) of all residues were in allowed (>99.8%) region

0.6% (53.8/30) of all residues we here were 12 outliers (phi, pai): 20 SER (109.0, -150.0) 29 ILE (67.7, 170.5) 53 VAL (76.8, 31.9) 54 ILE (60.8, 137.2) 64 ILE (60.8, 137.2) 66 ILU (93.6, -157.7) 136 ASP (15.0, 128.0) 137 ASP (175.3, -126.3) 138 SER (145.7, 77.2) 205 ILE (132.9, 154.3) 219 ILE U (95.2, 24.8) 286 PRO (-36.0, 107.1)