MolProbity Ramachandran analysis

1air_.B99990022.pdb, model 1



89,1% (312/350) of all residues were in favored (98%) regions.

there were 13 outliers (phi., p 20 SER (111.4, -146.9) 29 ILE (67.9, 173.1) 36 ALA (34.7, 80.5) 53 VAL (74.5, 34.6) 54 ILE (59.6, 136.8) 109 LYS (78.8, -169.3) 136 ASP (110.9, -103.0) 205 ILE (128.3, 153.0) 219 LEU (-95.7, 23.7) 267 ARG (-179.9, -137.5) 268 TYR (34.7, 88.2) 285 LYS (-70.6, -159.9)