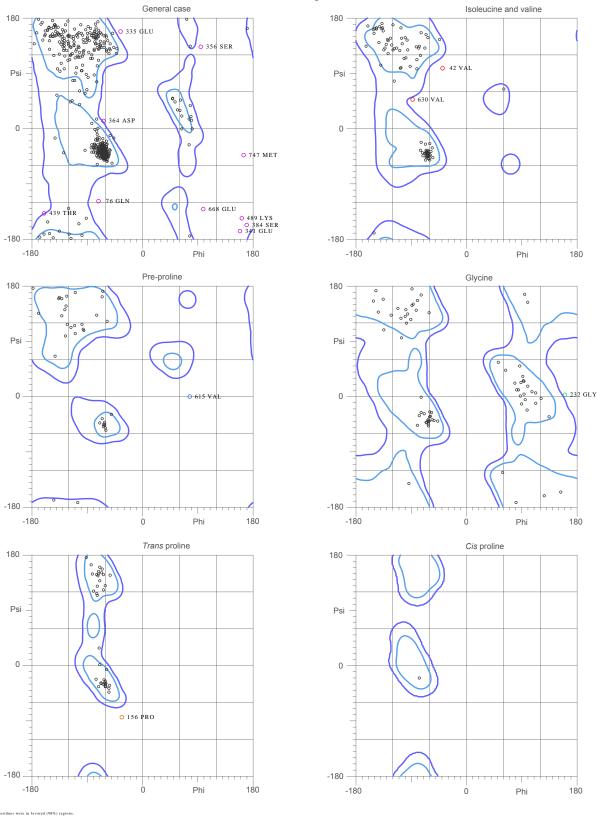
MolProbity Ramachandran analysis

1aco_.B99990031.pdb, model 1



92,3% (693/751) of all residues were in favored (98%) regions. 98,0% (736/751) of all residues were in allowed (>99.8%) region

there were 15 outliers (phi, p. 42 VAL (-39.7, 99.4) 76 GLN (-7.9, -118.7) 156 PRO (-34.3, -34.0) 156 PRO (-34.3, -34.0) 222 GLY (161.8, -5.5) 335 GLU (-36.4, 159.6) 345 GER (95.9, 134.1) 364 ASP (-64.4, 15.1) 348 ASER (170.4, -157.2) 439 THR (-161.0, -138.9) 439 LYS (162.4, -146.2) 615 VAL ((77.5, 0.3) 630 VAL (63.4, 48.5)