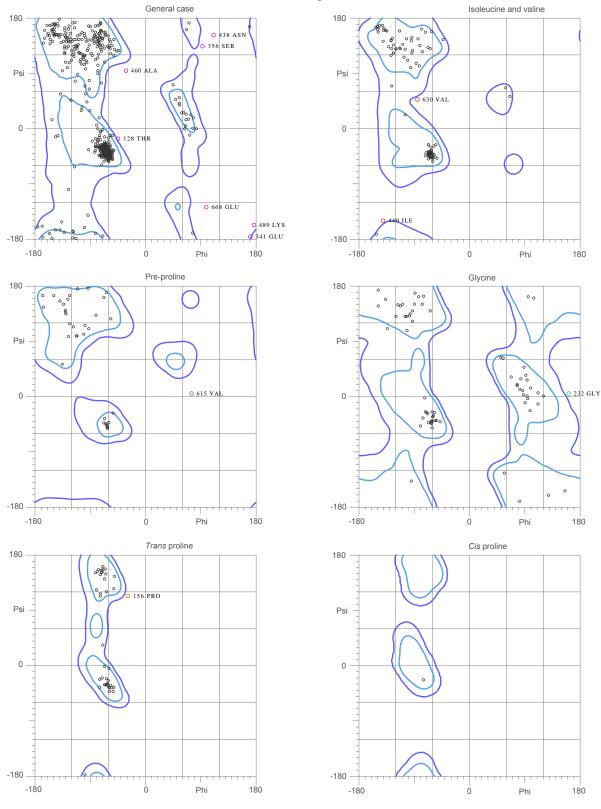
MolProbity Ramachandran analysis

1aco_.B99990005.pdb, model 1



92,5% (695/751) of all residues were in favored (98%) regions. 98,4% (739/751) of all residues were in allowed (>99.8%) regions

98,4% (739751) of all residues we There were 12 outliers (phi, psi): 128 THR (-46,0, -16,1) 156 TRO (-29,5, 114,2) 222 GLY (162,3, 5,3) 341 GLU (171,6, -176,3) 356 SER (31, 135,9) 438 ASN (111,6, 153,5) 440 ILE (-141,8, -150,6) 460 ILA (-32,7, 95,1) 489 LY\$ (177, -157,4) 615 VAL (75,7, 5,5) 630 VAL (-858,8,48) 668 GLU (199,8, -128,5)