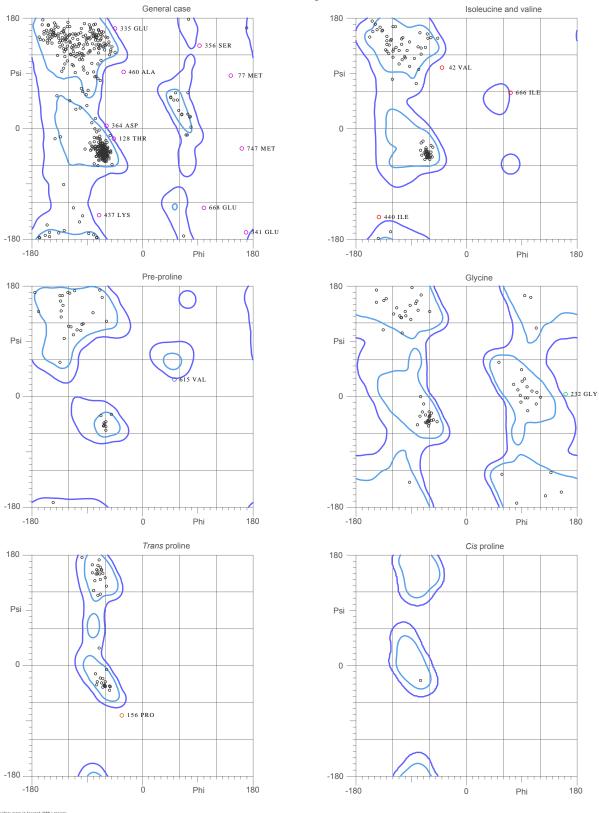
MolProbity Ramachandran analysis

1aco_.B99990034.pdb, model 1



92,7% (696/751) of all residues were in favored (98%) regions. 97,9% (735/751) of all residues were in allowed (>99.8%) regio

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here were 16 outliers (phi, 42 VAL (-40,4, 100,3) 77 MET (144,0, 87,0) 128 THR (-47,2, -16,7) 156 PRO (-34,3, -81,9) 232 GLY (162,2, 5,0) 335 GLU (-45,0, 164,2) 341 GLU (169,4, -169,4) 356 SER (93,2, 136,1) 364 ASP (-59,7, 5,6) 437 LYS (-71,4, -141,6) 440 ILE (-143,3, -145,0)