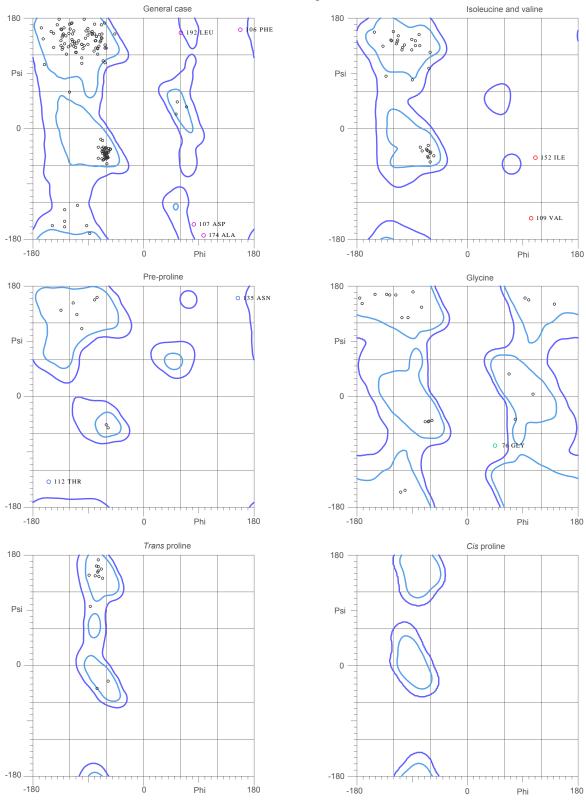
MolProbity Ramachandran analysis

1ad2_.B99990005.pdb, model 1



 $88,3\%\ (196/222)$ of all residues were in favored (98%) regions

95,9% (213/222) of all residues were in allowed (>99.8%) regions

There were 9 outliers (phi, psi): 76 GLY (45.9, -80.7) 106 PHE (158.6, 162.5) 107 ASP (82.2, -156.1) 109 VAL (104.1, -146.1) 112 THR (-155.6, -139.0) 135 ASN (154.1, 161.7) 152 ILE (111.4, -47.5) 174 JAI (4) 83 - 174 4) 174 ALA (98,3, -174,4) 192 LEU (61,6, 157,5)