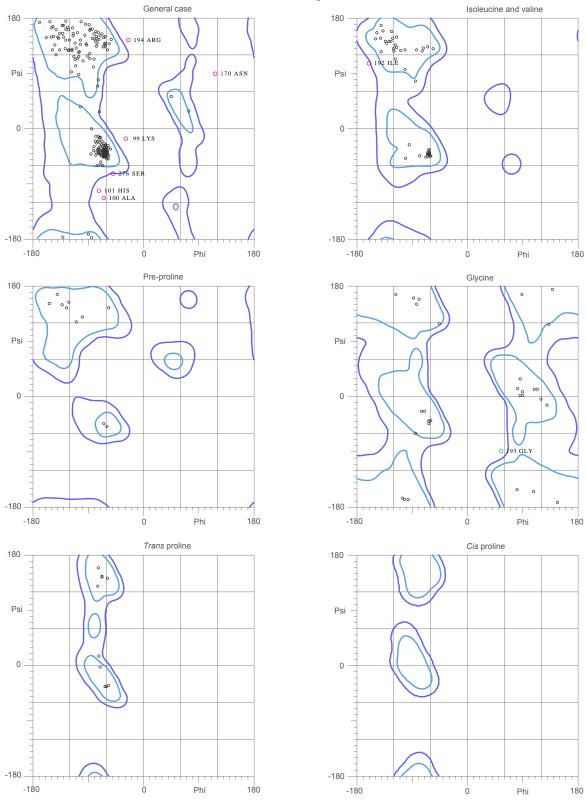
MolProbity Ramachandran analysis

1ak5_.B99990025.pdb, model 1



95,7% (313/327) of all residues were in favored (98%) regions. 97,6% (319/327) of all residues were in allowed (>99.8%) regions.

97,6% (319/327) of all residues were in allowed (>99.8%) regions

There were 8 outliers (phi, psi):
99 LYS (-29,1, -16,8)
100 ALA (-65,1, -113,4)
101 HIS (-73,1, -101,8)
170 ASN (117,4, 90,0)
192 ILE (-161,9, 107,4)
193 GLY (55,1, -89,9)
194 ARG (-25,1, 145,1)
276 SER (-50,9, -73,8)