MolProbity Ramachandran analysis
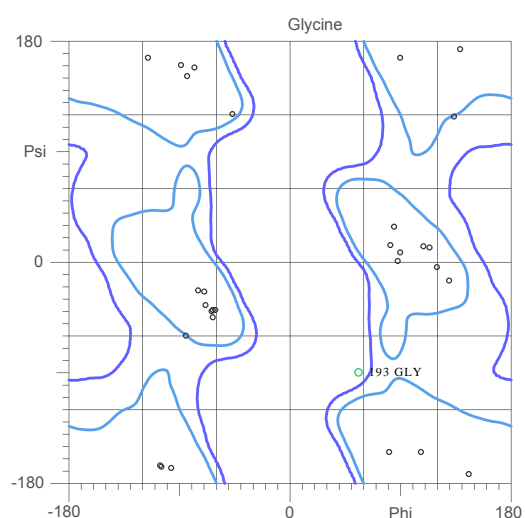
1ak5_.B99990002.pdb, model 1

General case

194 ARG
170 ASN
99 LYS
276 SER
101 HIS
100 ALA

Isoleucine and valine

192 ILE

Pre-proline

Glycine

193 GLY

Trans proline

Cis proline
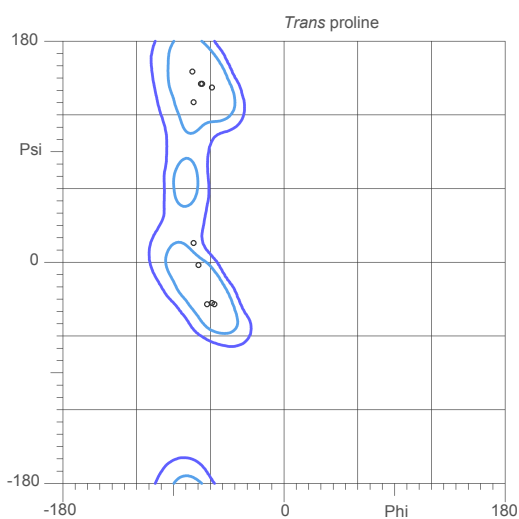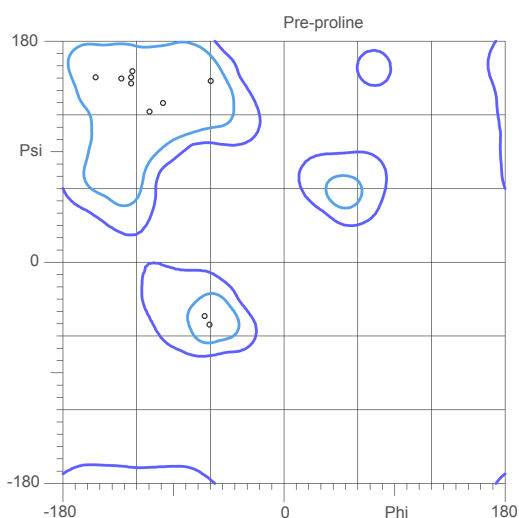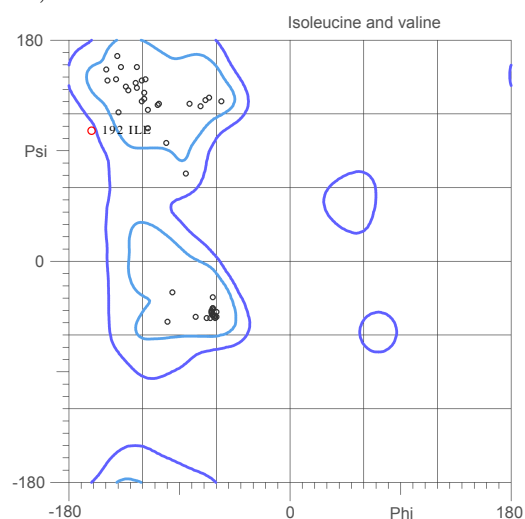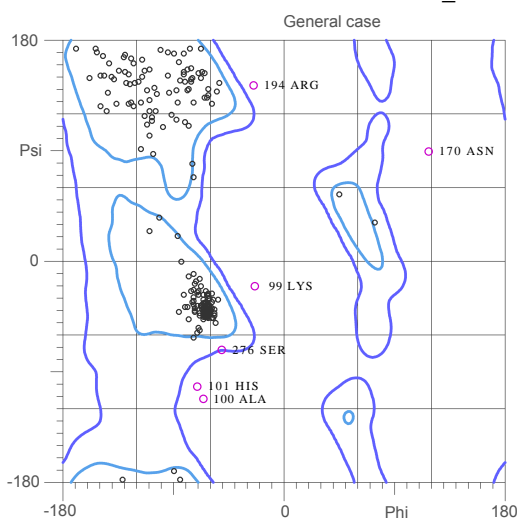
95,4% (312/327) of all residues were in favored (98%) regions.
97,6% (319/327) of all residues were in allowed (>99.8%) regions.

There were 8 outliers (phi, psi):
    99 LYS (-24,9, -20,2)
    100 ALA (-66,1, -112,0)
    101 HIS (-71,8, -102,4)
    170 ASN (118,2, 90,2)
    192 ILE (-162,6, 107,8)
    193 GLY (56,2, -90,2)
    194 ARG (-25,8, 144,6)
    276 SER (-51,3, -73,0)