MolProbity Ramachandran analysis

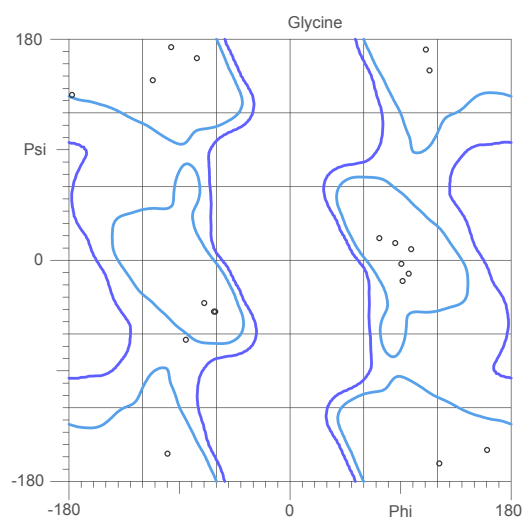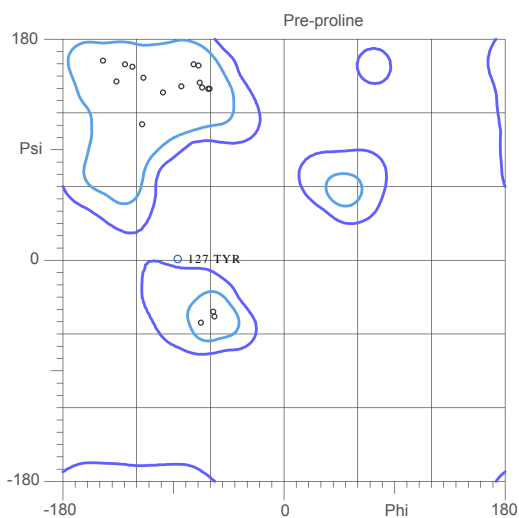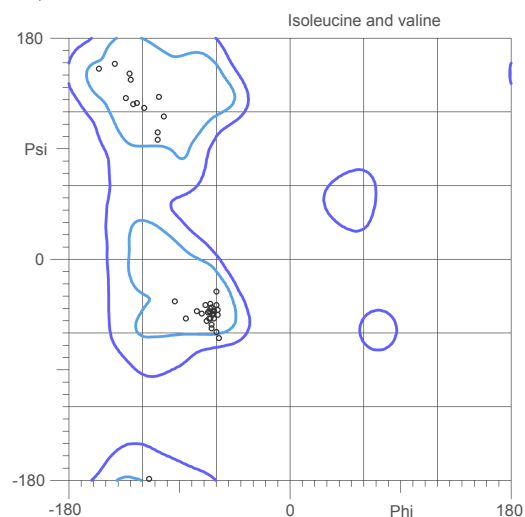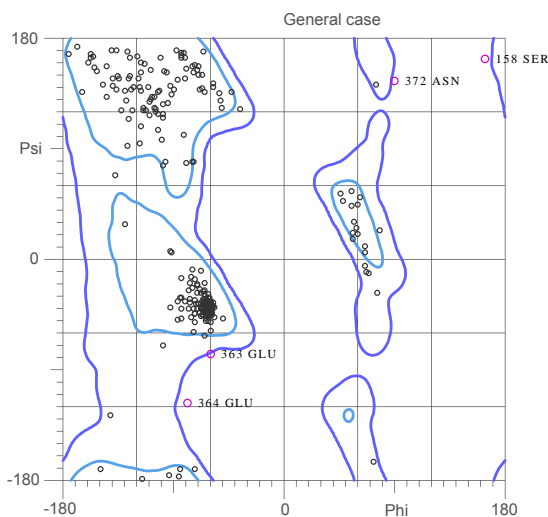1auiA.B99990003.pdb, model 1
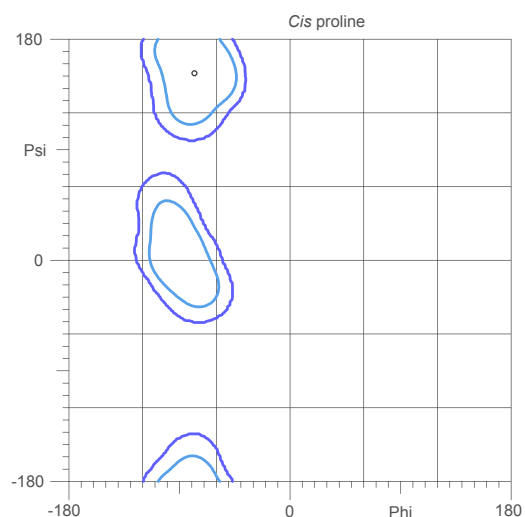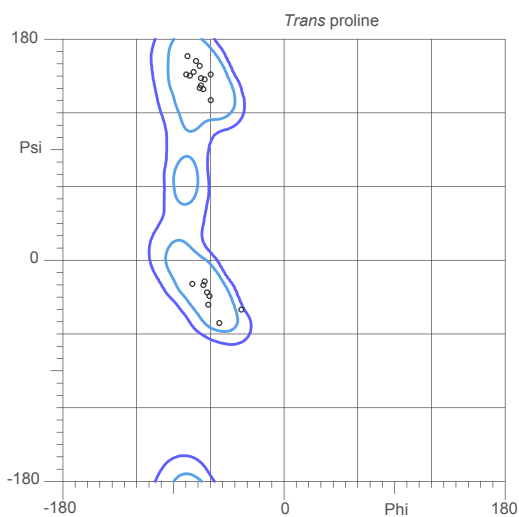
94,1% (354/376) of all residues were in favored (98%) regions.
98,7% (371/376) of all residues were in allowed (>99.8%) regions.

There were 5 outliers (phi, psi):
  127 TYR (-87,0, 2,0)
  158 SER (164,9, 164,6)
  363 GLU (-61,0, -77,7)
  364 GLU (-79,0, -117,3)
  372 ASN (90,6, 146,3)