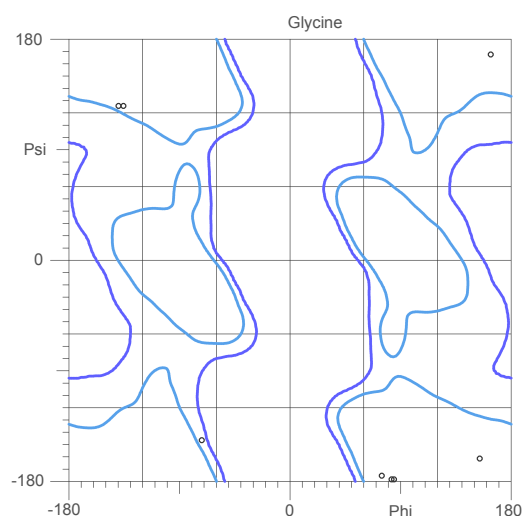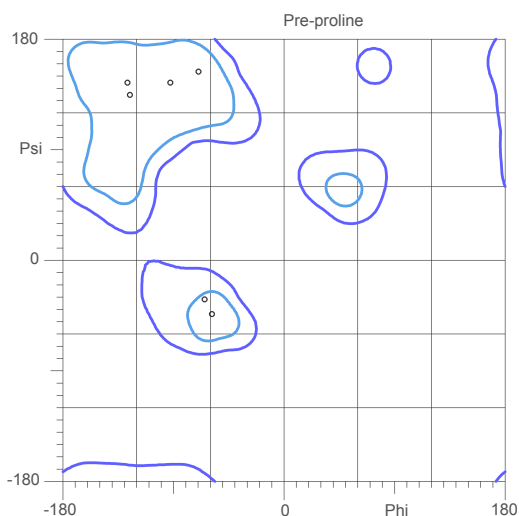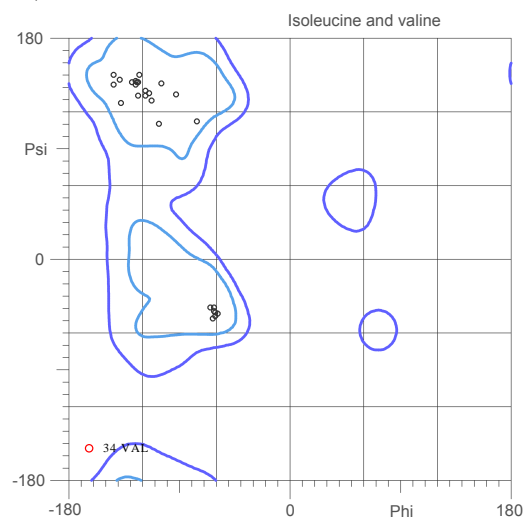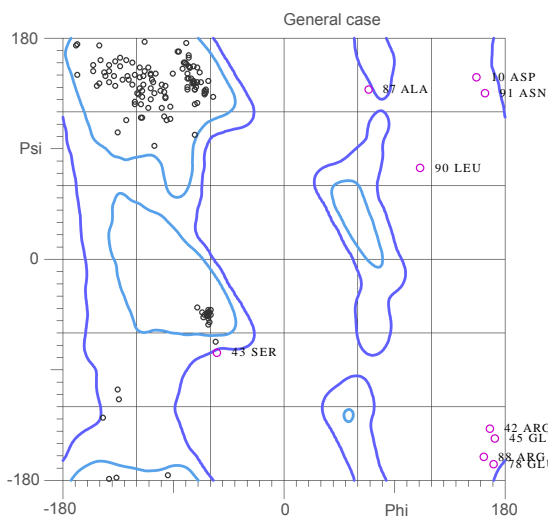# MolProbity Ramachandran analysis

1aisB.B99990031.pdb, model 1
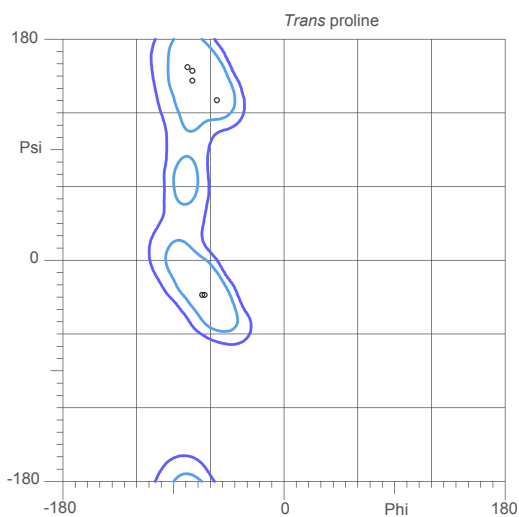
92,1% (176/191) of all residues were in favored (98%) regions.
94,8% (181/191) of all residues were in allowed (>99.8%) regions.

There were 10 outliers (phi, psi):
  10 ASP (157,2, 149,8)
  34 VAL (-164,9, -154,2)
  42 ARG (168,4, -138,6)
  43 SER (-55,3, -76,4)
  45 GLU (172,9, -146,9)
  78 GLU (171,5, -167,4)
  87 ALA (69,2, 139,5)
  88 ARG (163,9, -161,8)
  90 LEU (111,1, 75,7)
  91 ASN (164,3, 136,4)