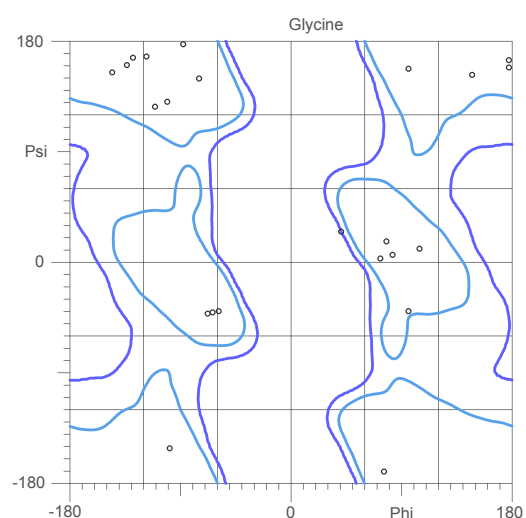# MolProbity Ramachandran analysis

1ad2_.B99990036.pdb, model 1

General case

Isoleucine and valine

○ 192 LEU
○ 127 LEU
○ 106 PHE
○ 109 VAL

○ 107 ASP
○ 174 ALA

Pre-proline

Glycine

○ 135 ASN

○ 112 THR

Trans proline

Cis proline
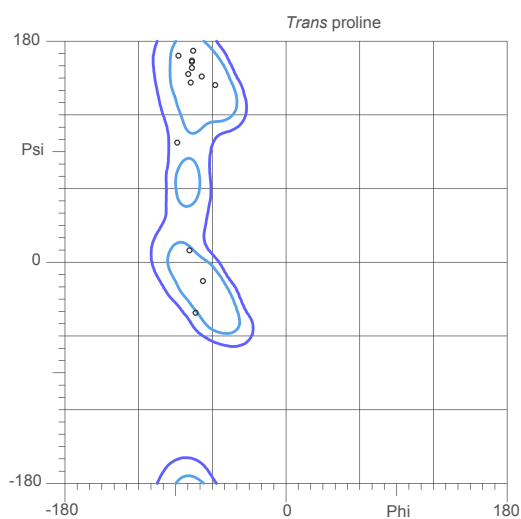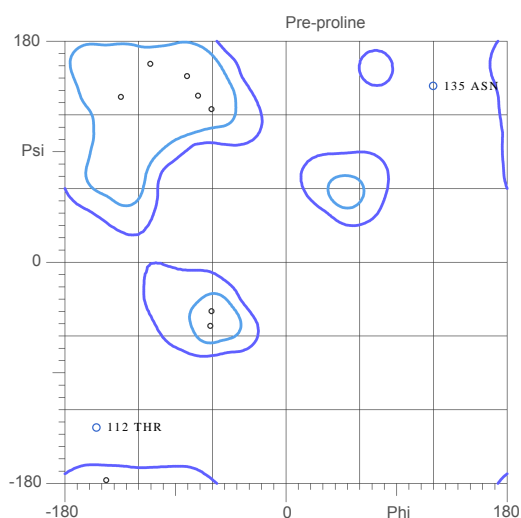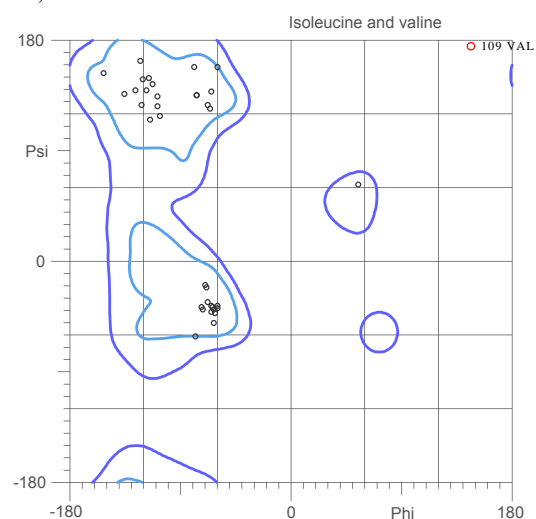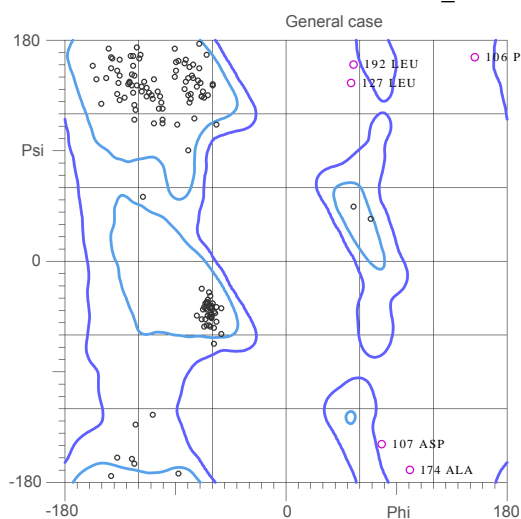
89,6% (199/222) of all residues were in favored (98%) regions.
96,4% (214/222) of all residues were in allowed (>99.8%) regions.

There were 8 outliers (phi, psi):
    106 PHE (154,8, 167,1)
    107 ASP (78,5, -149,5)
    109 VAL (147,5, 176,9)
    112 THR (-155,8, -135,9)
    127 LEU (53,3, 146,4)
    135 ASN (120,7, 144,0)
    174 ALA (101,9, -170,7)
    192 LEU (55,2, 161,0)