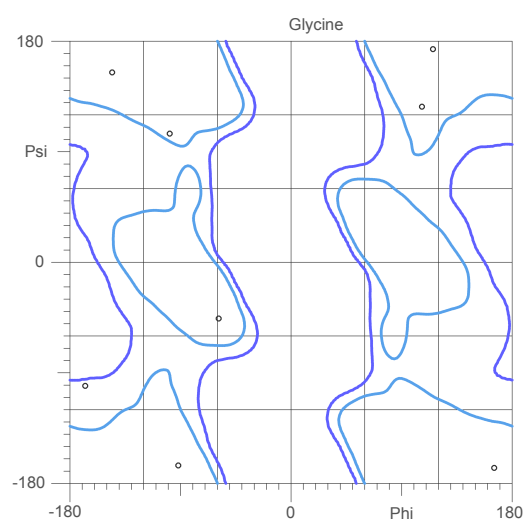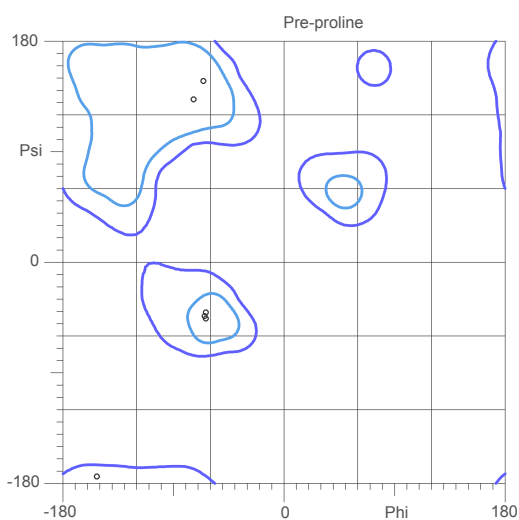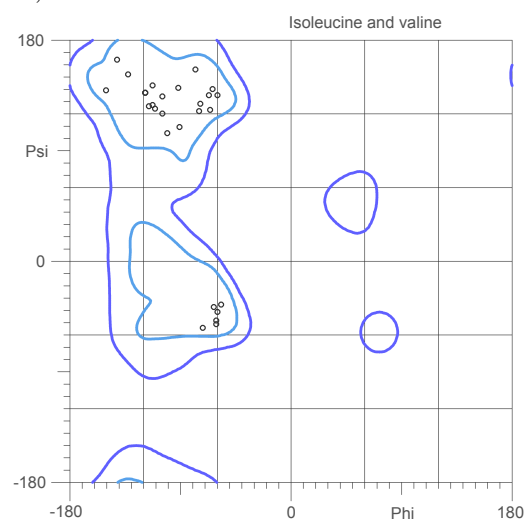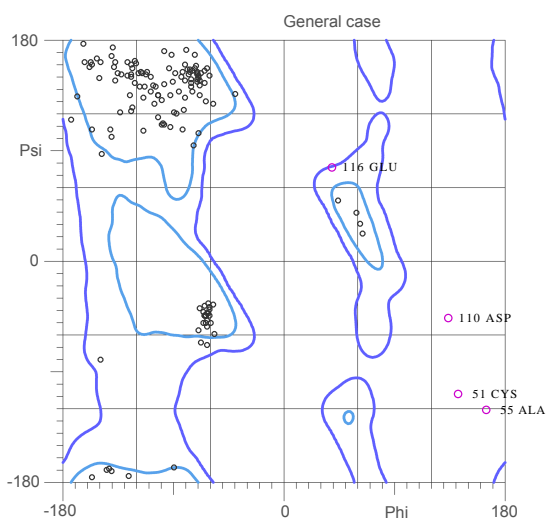# MolProbity Ramachandran analysis

1aisB.B99990027.pdb, model 1
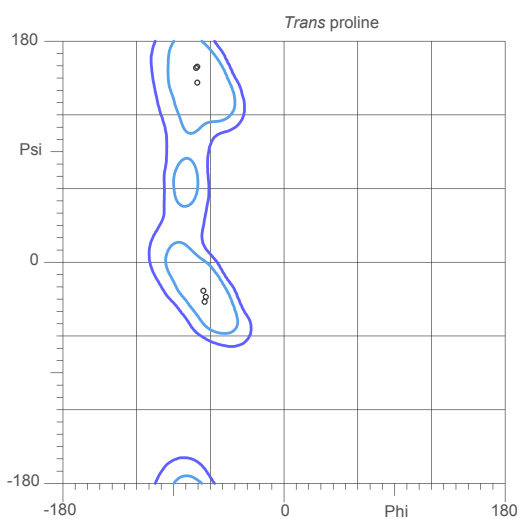
93,2% (178/191) of all residues were in favored (98%) regions.
97,9% (187/191) of all residues were in allowed (>99.8%) regions.

There were 4 outliers (phi, psi):
    51 CYS (142,7, -108,4)
    55 ALA (165,4, -121,1)
    110 ASP (134,4, -46,3)
    116 GLU (39,9, 77,8)

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Lovell, Davis, et al. Proteins 50:437 (2003)