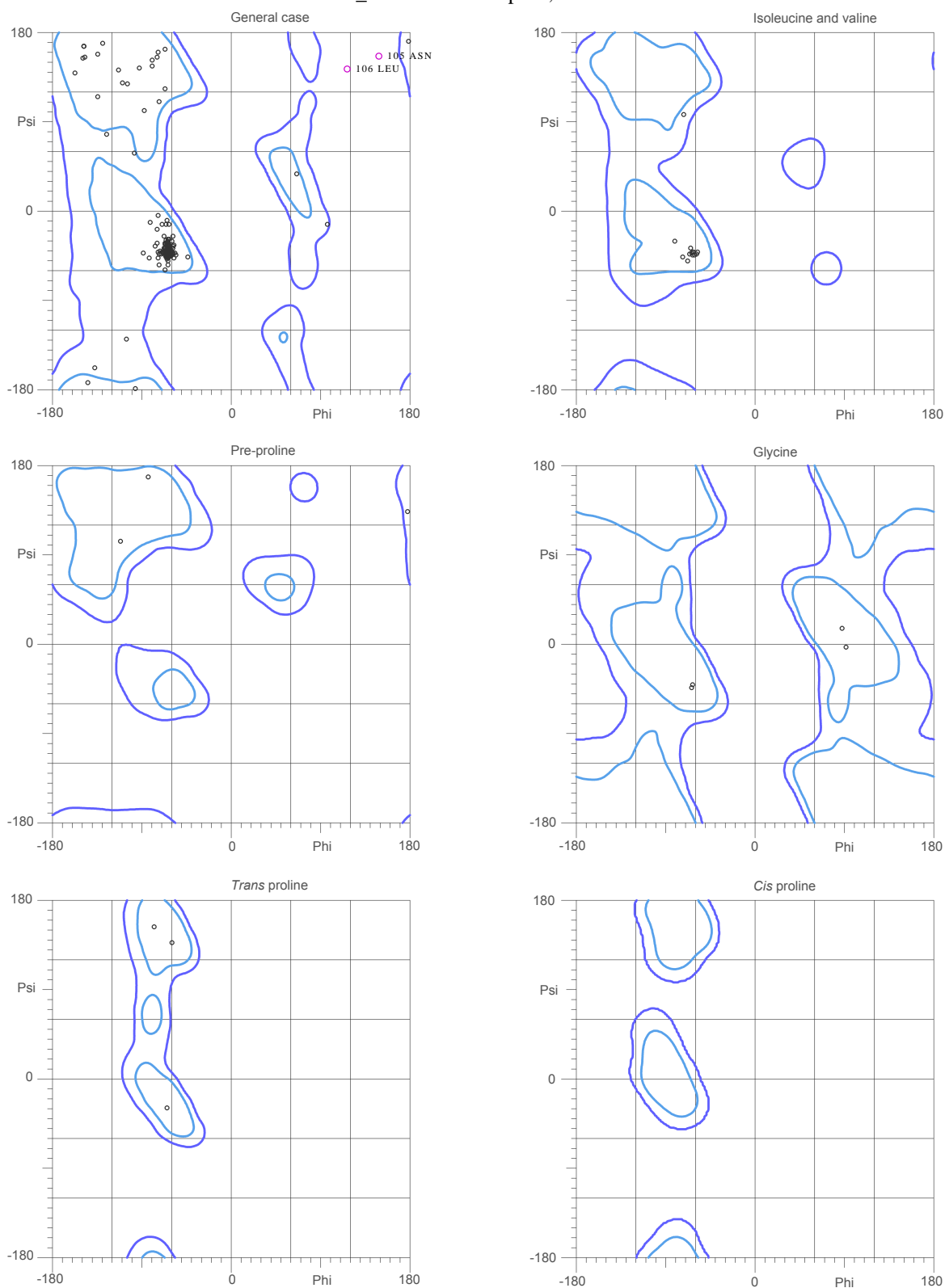# MolProbity Ramachandran analysis

1alu_.B99990032.pdb, model 1

93,5% (145/155) of all residues were in favored (98%) regions.

98,7% (153/155) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):
    105 ASN (149,8, 157,9)
    106 LEU (117,8, 144,1)

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