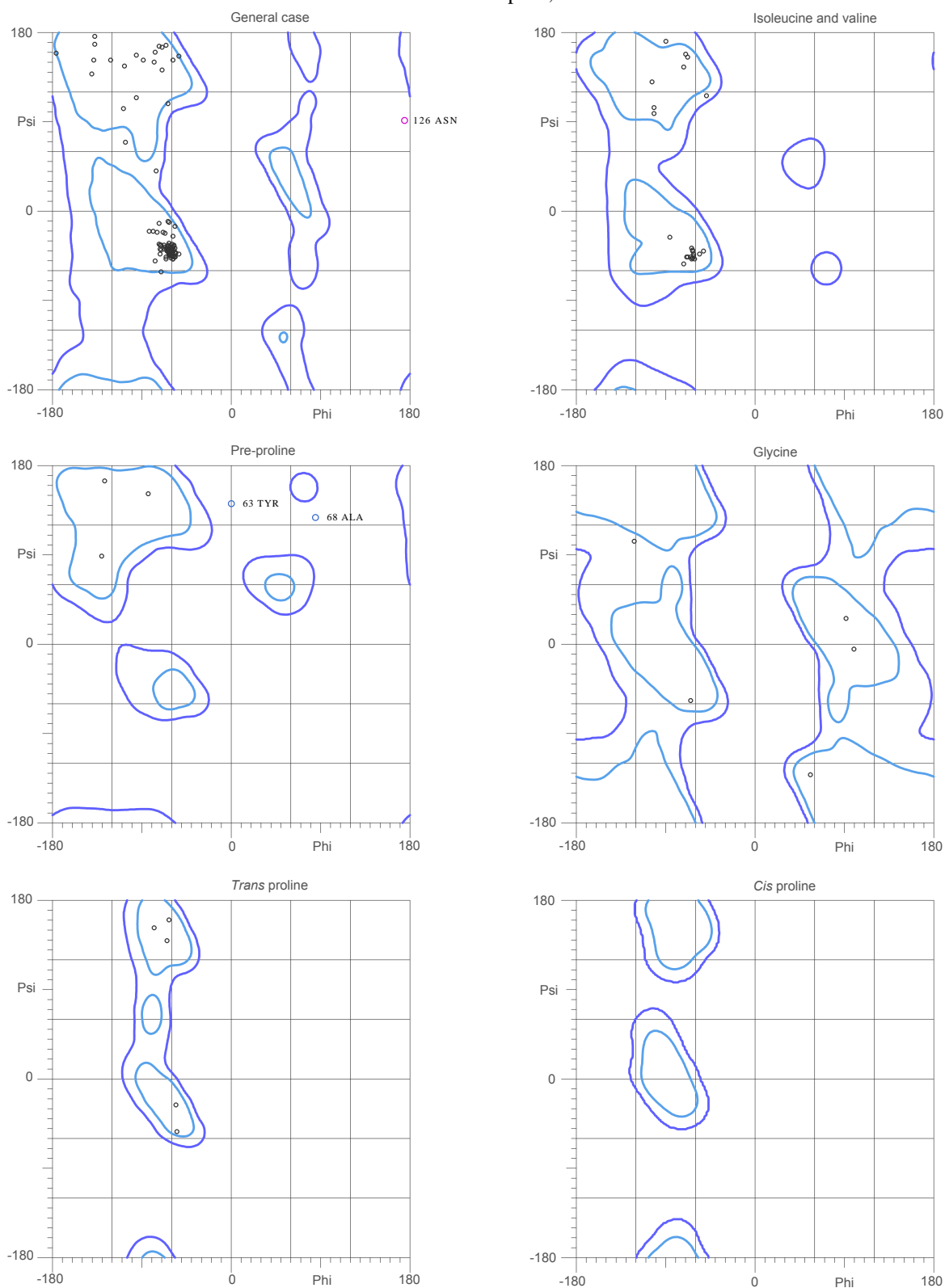# MolProbity Ramachandran analysis

1al01.B99990047.pdb, model 1

91,5% (129/141) of all residues were in favored (98%) regions.
97,9% (138/141) of all residues were in allowed (>99.8%) regions.

There were 3 outliers (phi, psi):
    63 TYR (-0,7, 142,2)
    68 ALA (85,8, 128,8)
    126 ASN (175,6, 92,0)