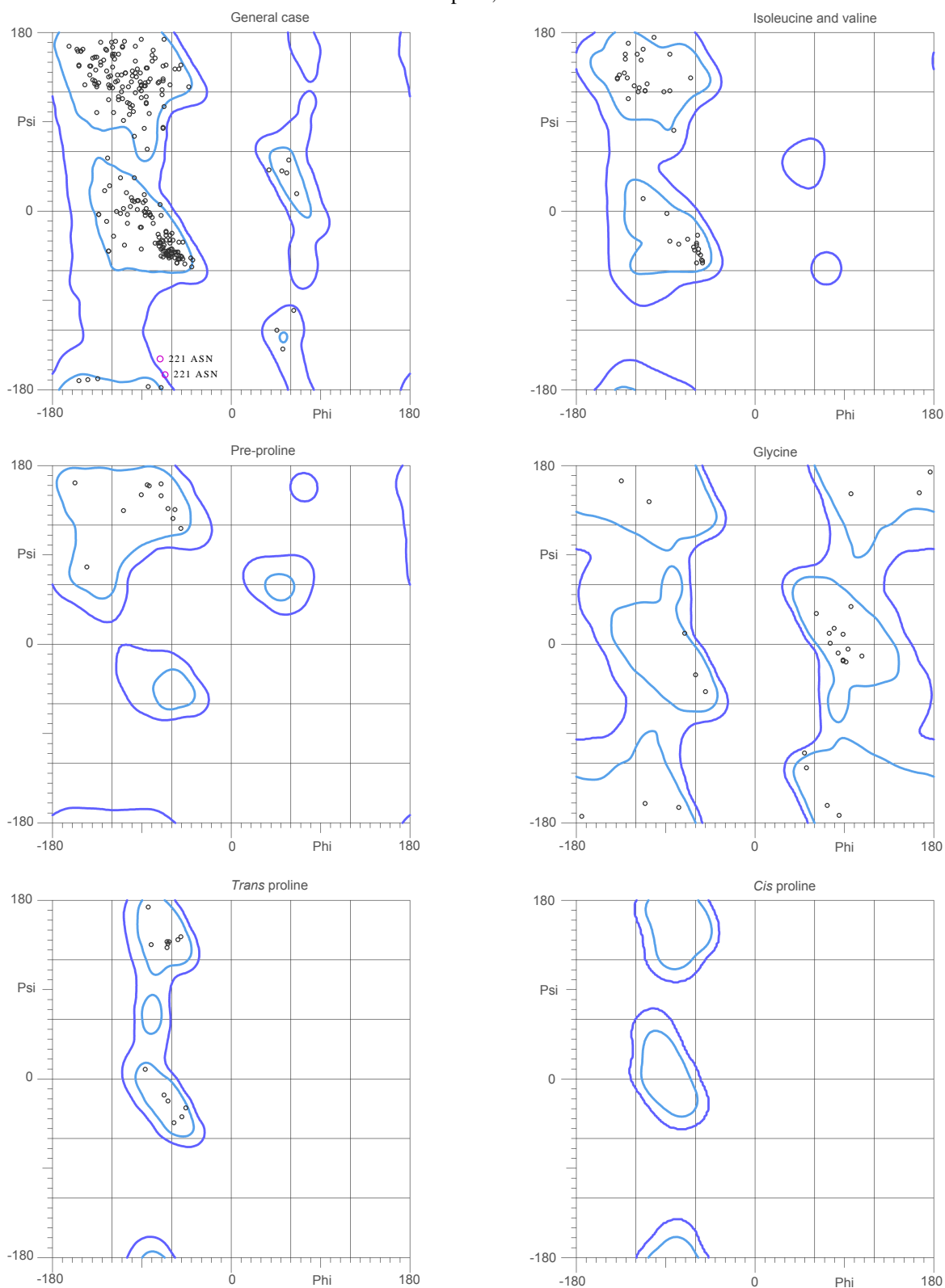


# MolProbity Ramachandran analysis

7ax6.H.pdb, model 1



95.6% (323/338) of all residues were in favored (98%) regions.

99.4% (336/338) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

221 ASN (-72.1, -149.2)

221 ASN (-67.6, -165.2)