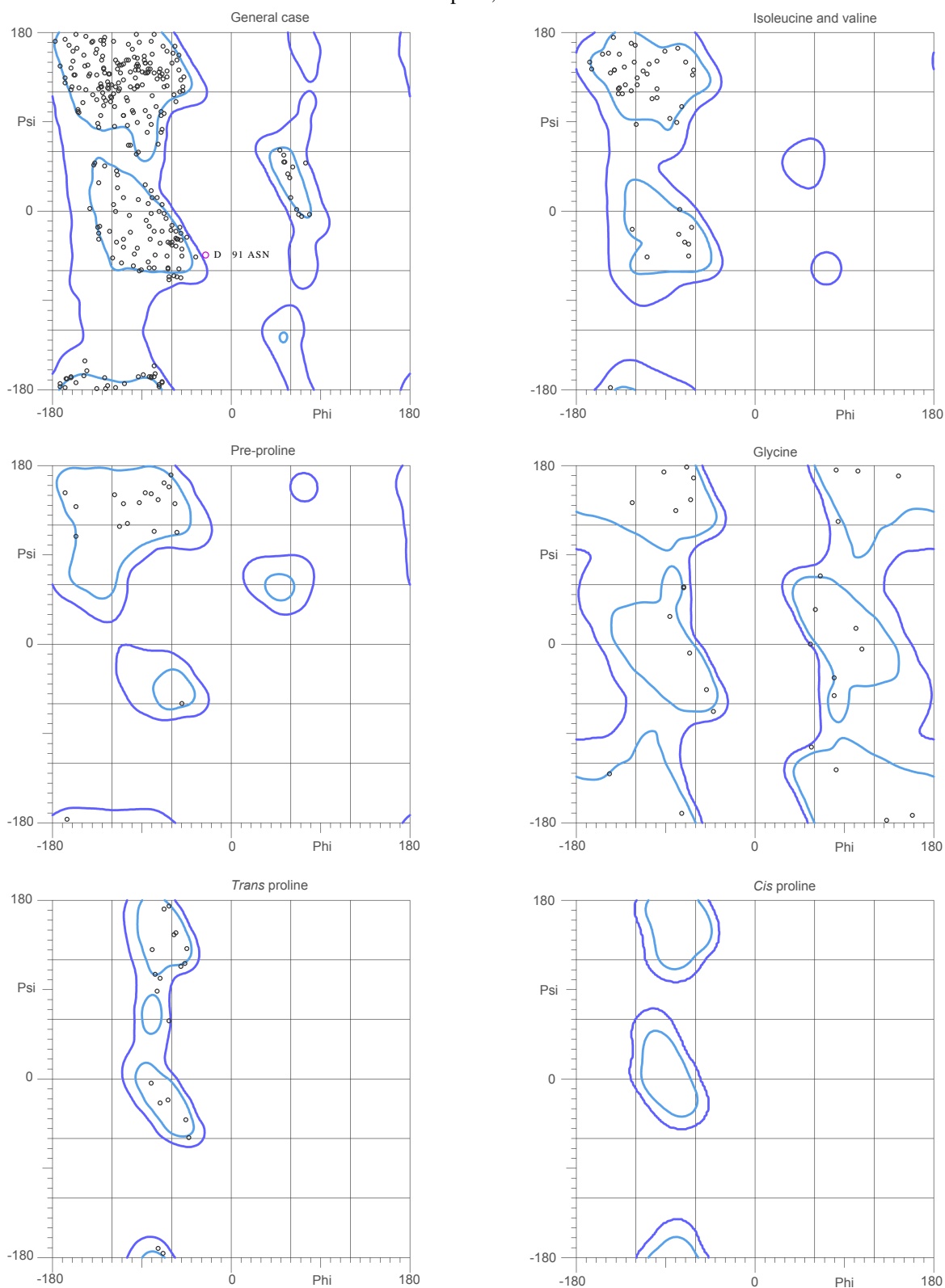


# MolProbity Ramachandran analysis

7cah.H.pdb, model 1



79.8% (316/396) of all residues were in favored (98%) regions.

99.7% (395/396) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

D 91 ASN (-27.0, -44.6)