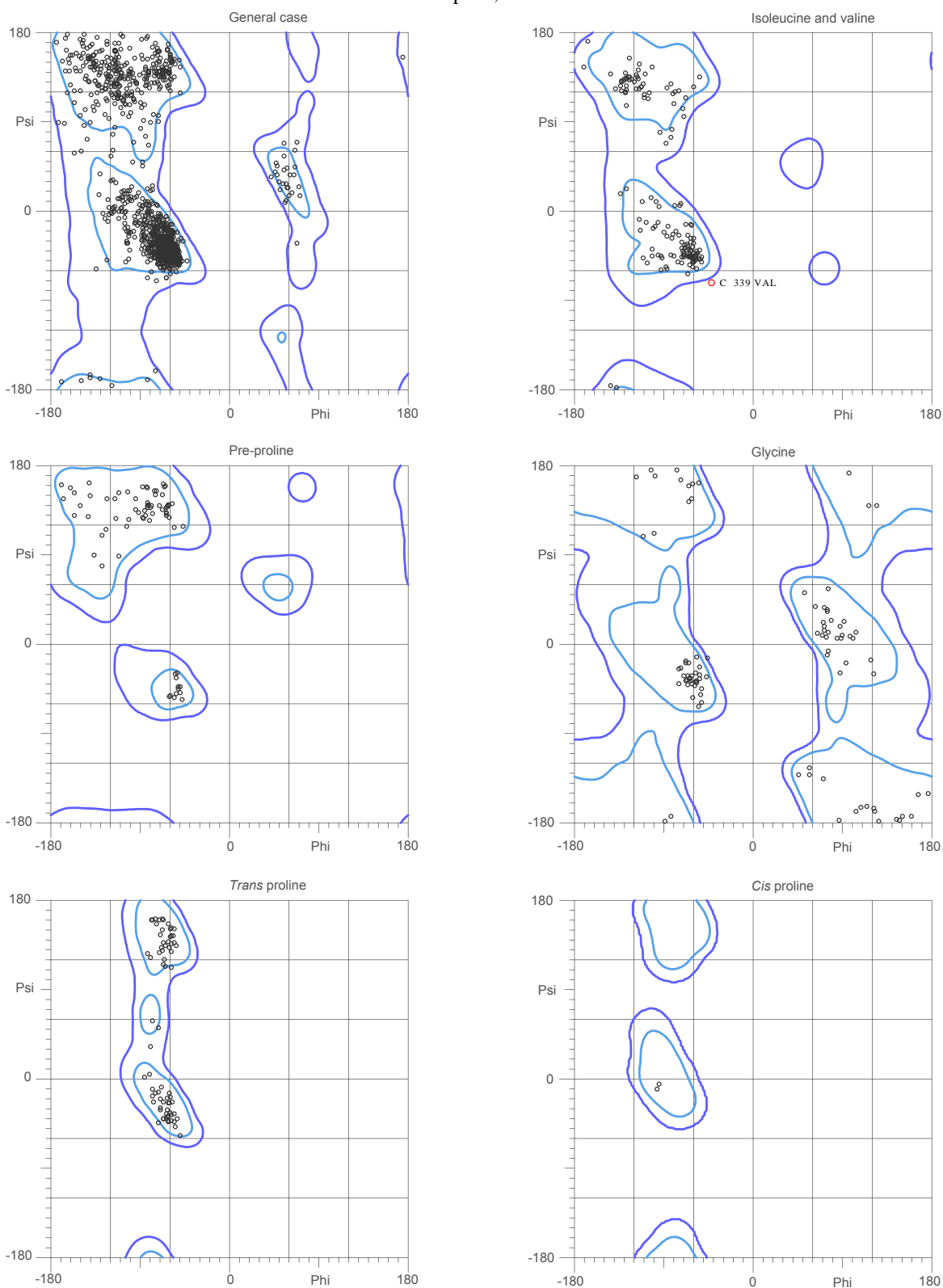


# MolProbity Ramachandran analysis

7dmu.H.pdb, model 1



96.3% (1520/1578) of all residues were in favored (98%) regions.

99.9% (1577/1578) of all residues were in allowed (>99.8%) regions.

There were 1 outliers (phi, psi):

C 339 VAL (-42.6, -72.2)