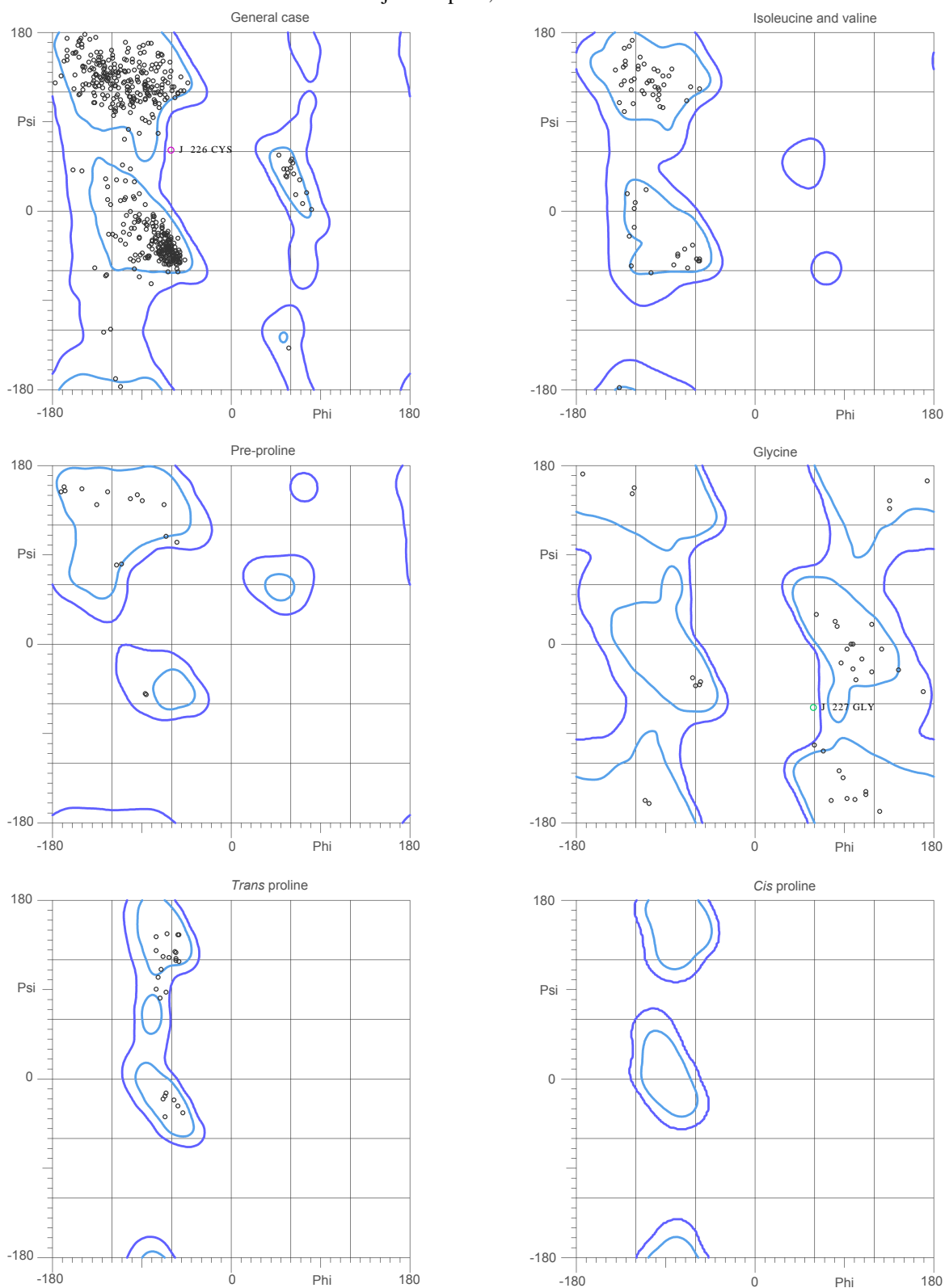


# MolProbity Ramachandran analysis

7jrn.H.pdb, model 1



94.4% (591/626) of all residues were in favored (98%) regions.

99.7% (624/626) of all residues were in allowed (>99.8%) regions.

There were 2 outliers (phi, psi):

J 226 CYS (-61.2, 62.7)

J 227 GLY (59.4, -64.7)