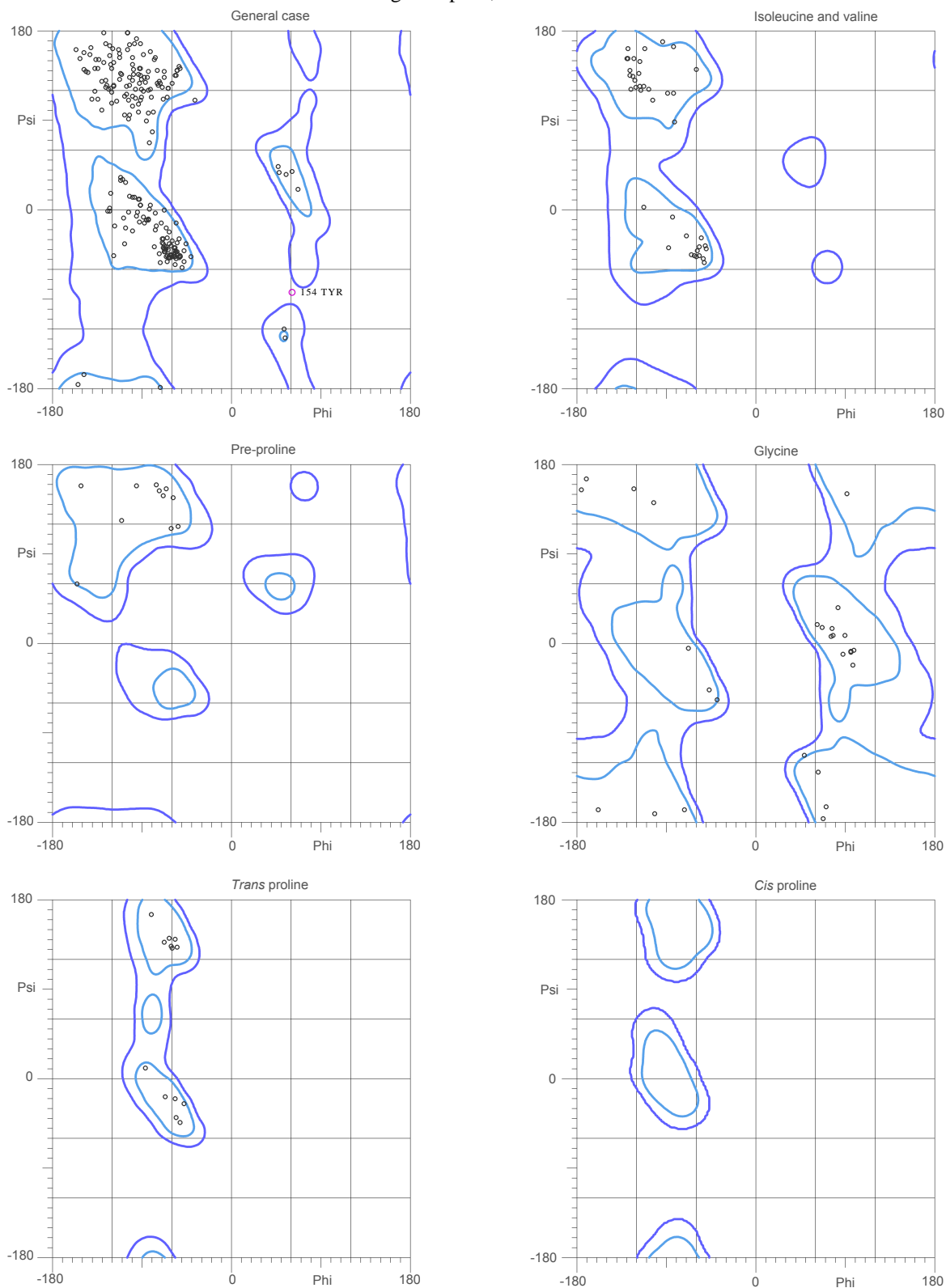


# MolProbity Ramachandran analysis

5rgs.H.pdb, model 1



98.1% (305/311) of all residues were in favored (98%) regions.

99.7% (310/311) of all residues were in allowed (>99.8%) regions.

There were 1 outliers ( $\phi$ ,  $\psi$ ):

154 TYR (61.6, -83.9)