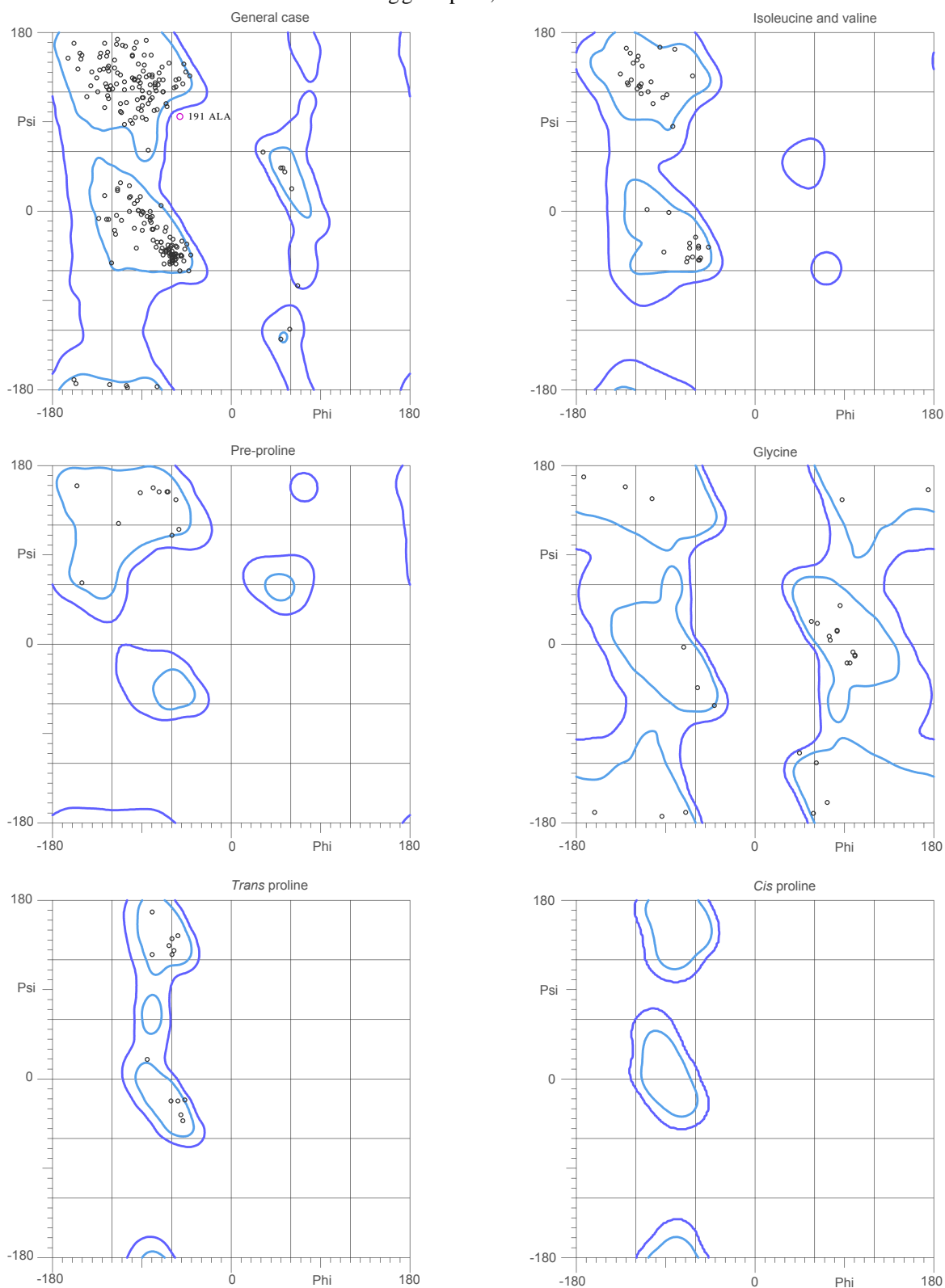


# MolProbity Ramachandran analysis

5rgg.H.pdb, model 1



96.1% (299/311) 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 ( $\phi$ ,  $\psi$ ):

191 ALA (-52.5, 96.2)