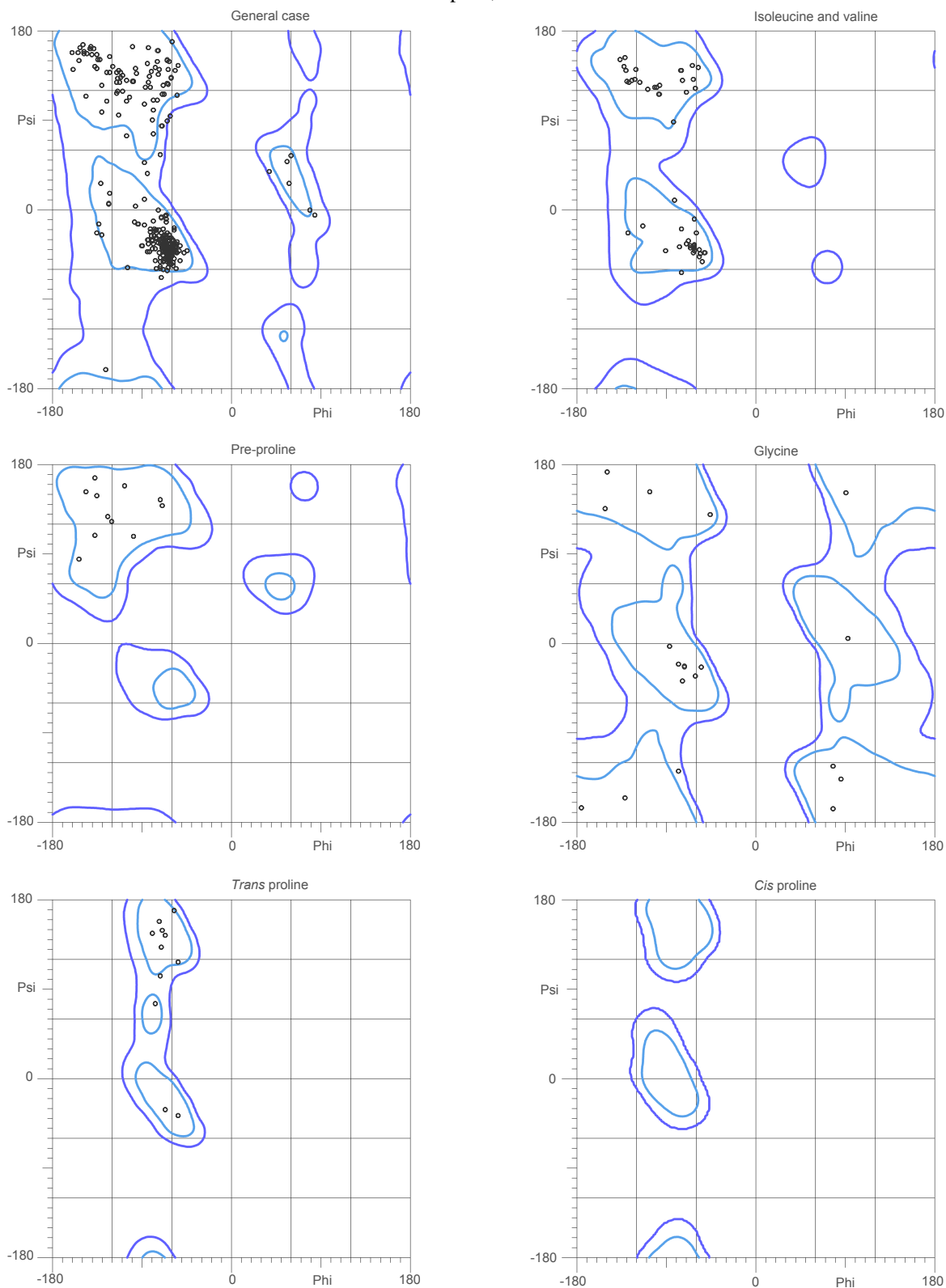


# MolProbity Ramachandran analysis

6xra.H.pdb, model 1



94.4% (969/1026) of all residues were in favored (98%) regions.

100.0% (1026/1026) of all residues were in allowed (>99.8%) regions.

There were no outliers.