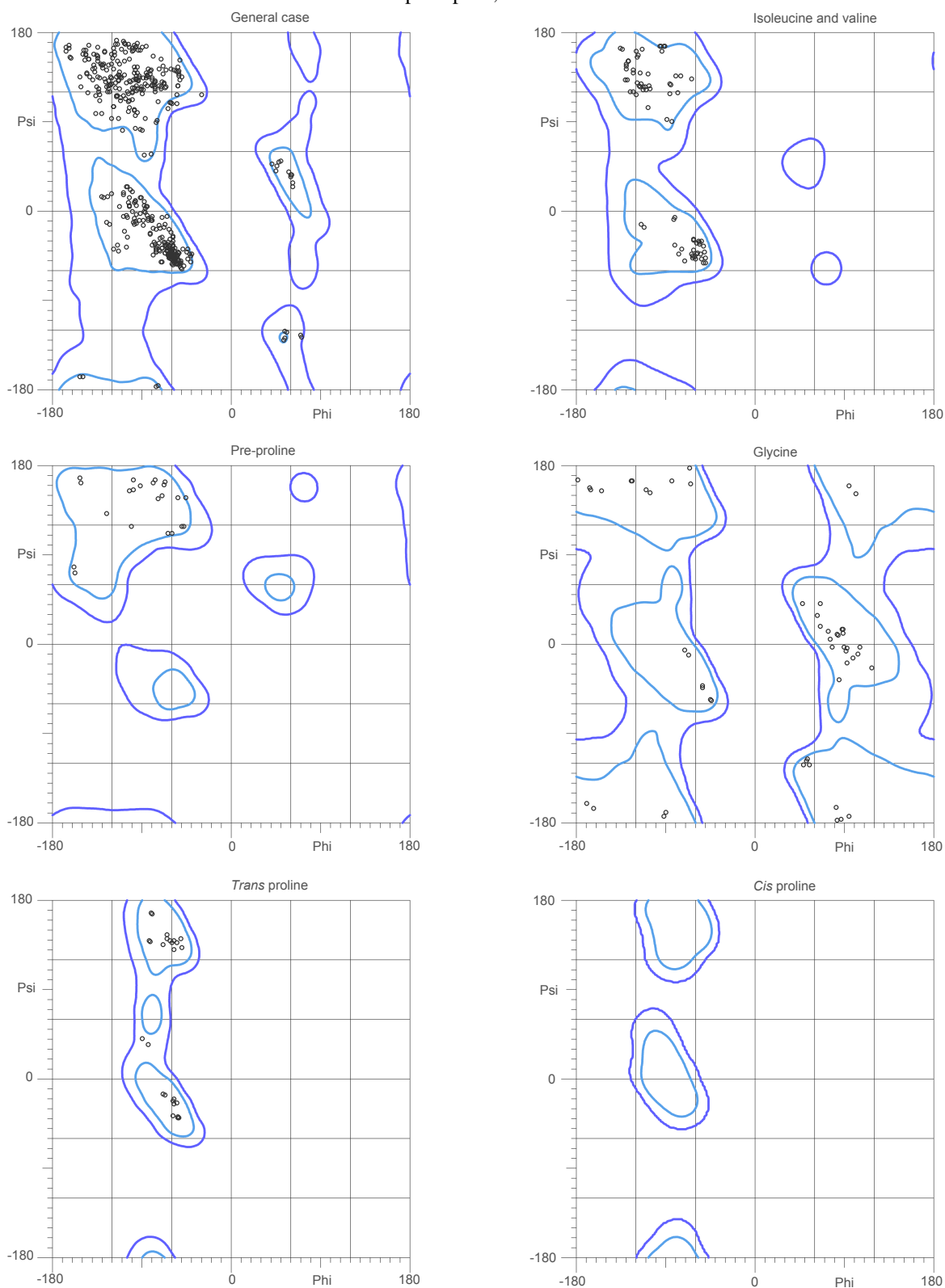


# MolProbity Ramachandran analysis

7khp.H.pdb, model 1



96.9% (589/608) of all residues were in favored (98%) regions.

100.0% (608/608) of all residues were in allowed (>99.8%) regions.

There were no outliers.