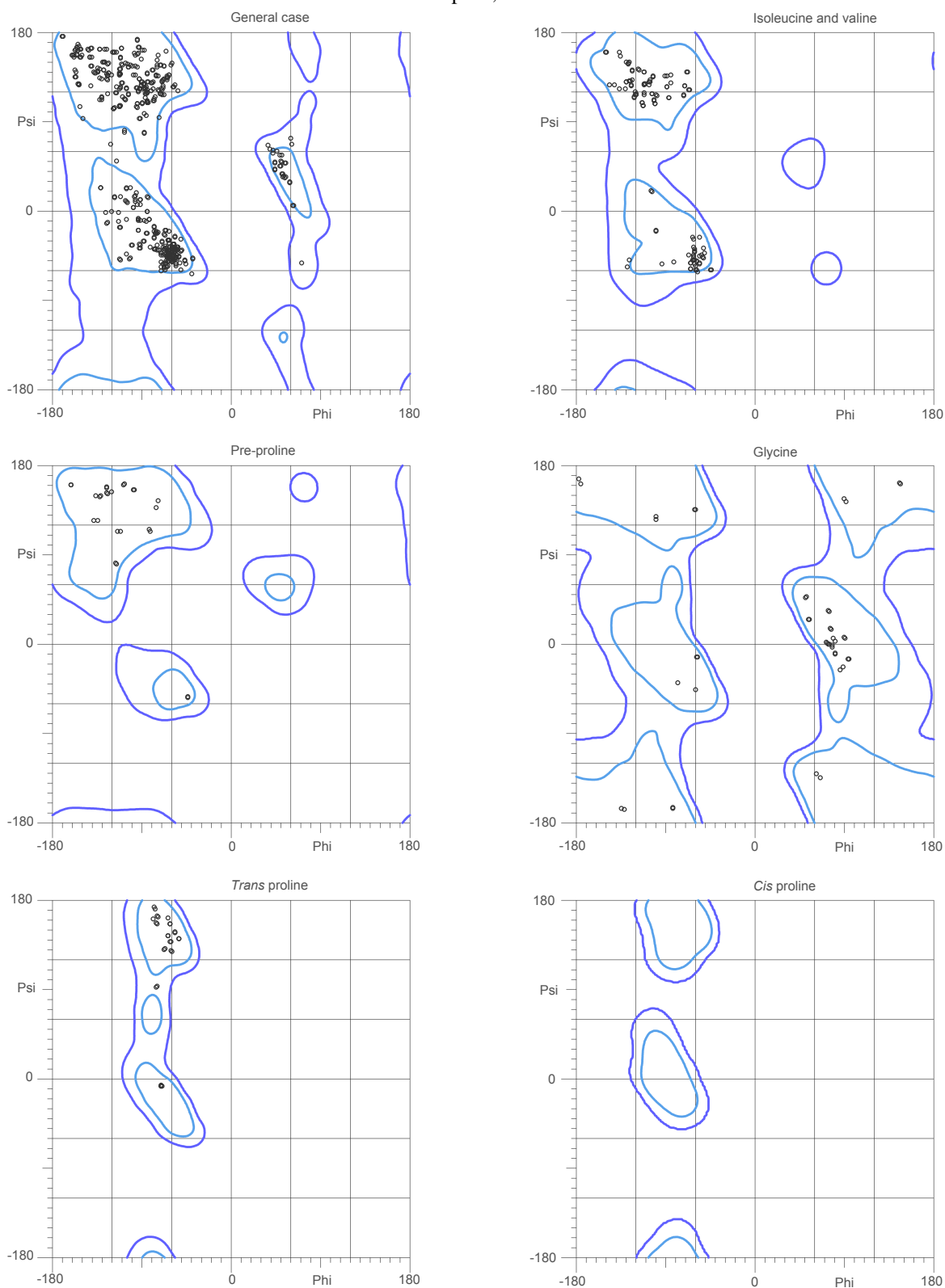


# MolProbity Ramachandran analysis

7keh.H.pdb, model 1



97.3% (674/693) of all residues were in favored (98%) regions.

100.0% (693/693) of all residues were in allowed (>99.8%) regions.

There were no outliers.