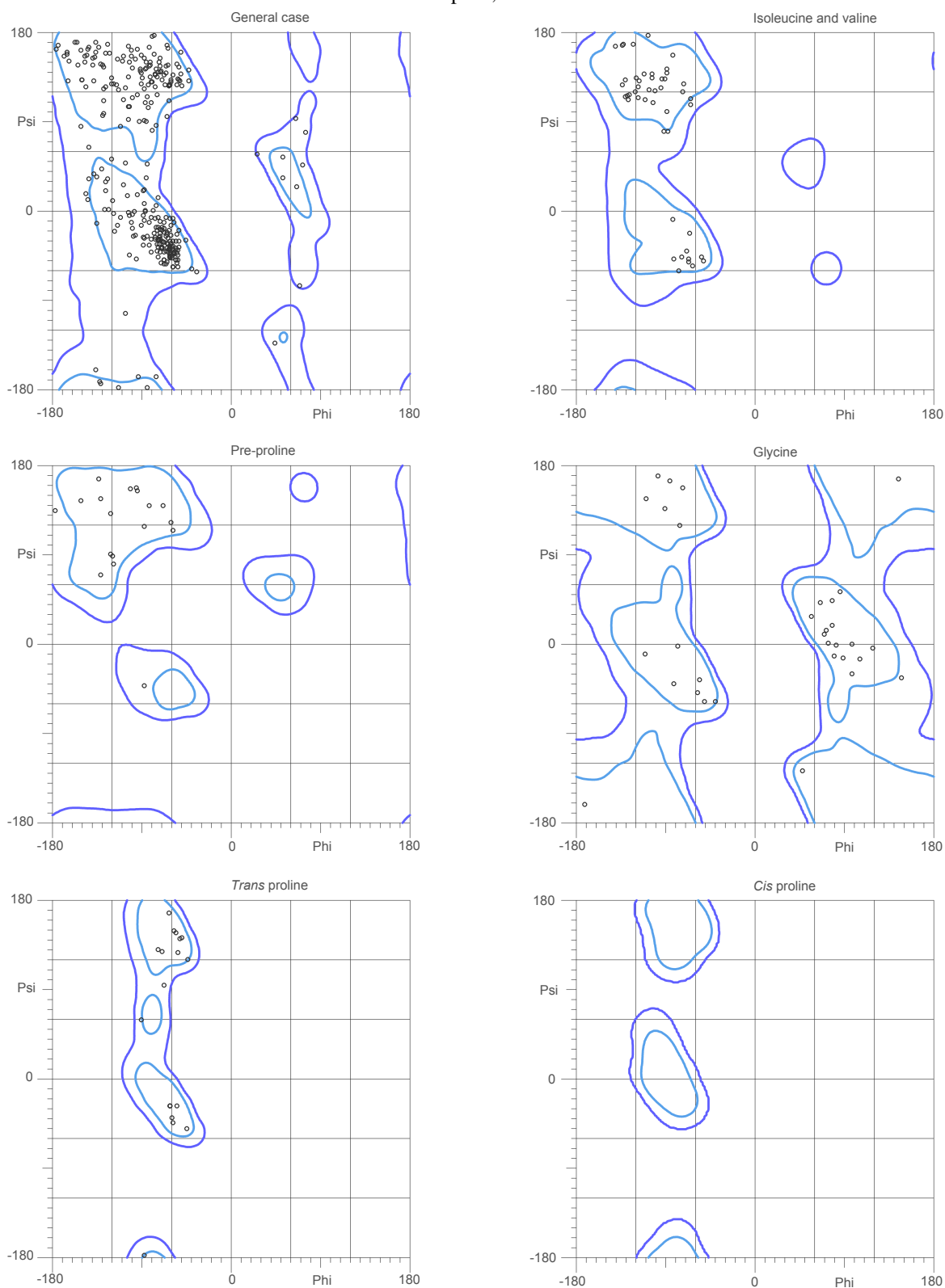


# MolProbity Ramachandran analysis

7koa.H.pdb, model 1



91.0% (383/421) of all residues were in favored (98%) regions.

100.0% (421/421) of all residues were in allowed (>99.8%) regions.

There were no outliers.