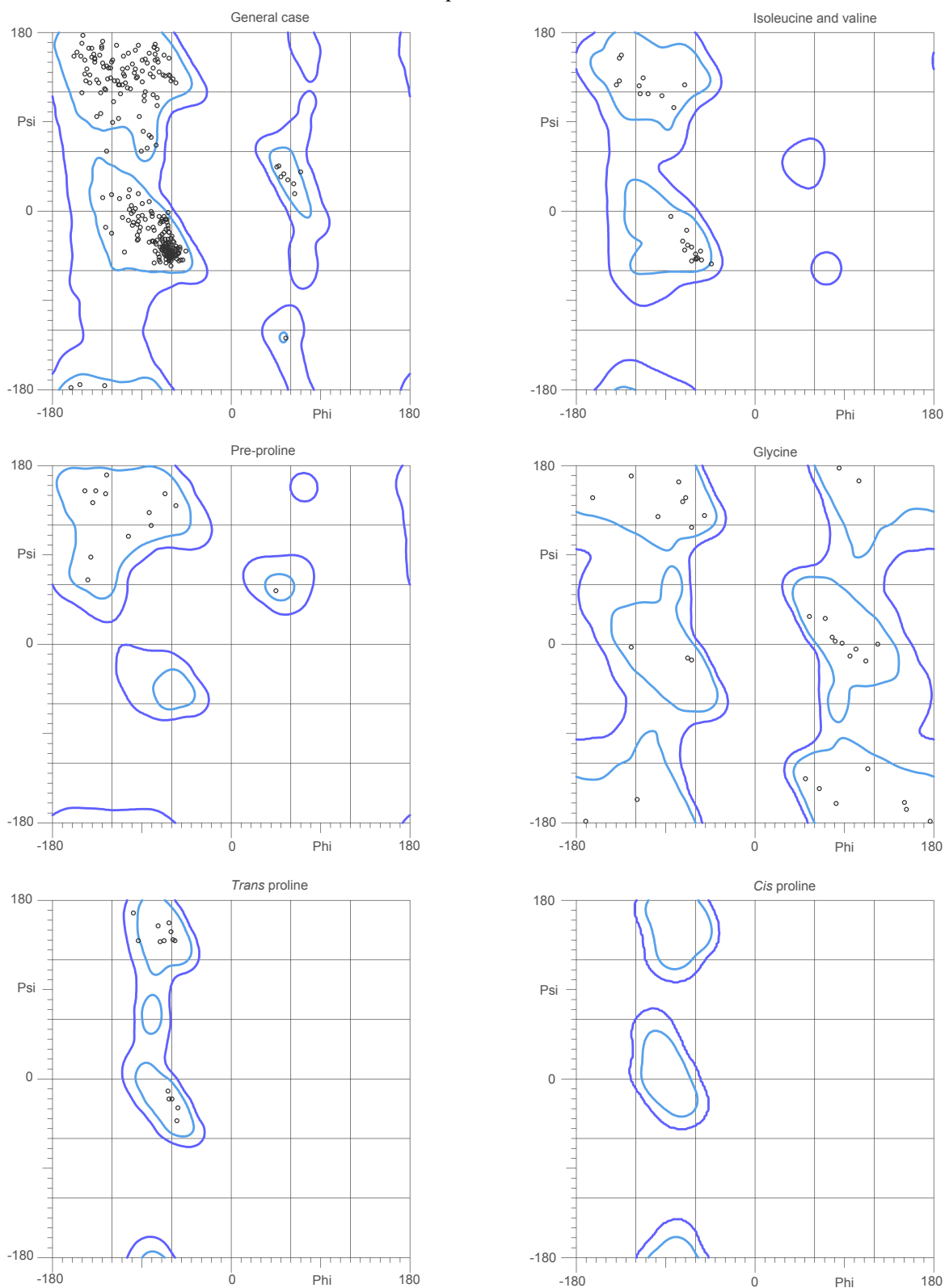


# MolProbity Ramachandran analysis

1hkk.pdb, model 1



98.3% (356/362) of all residues were in favored (98%) regions.  
100.0% (362/362) of all residues were in allowed (>99.8%) regions.

There were no outliers.