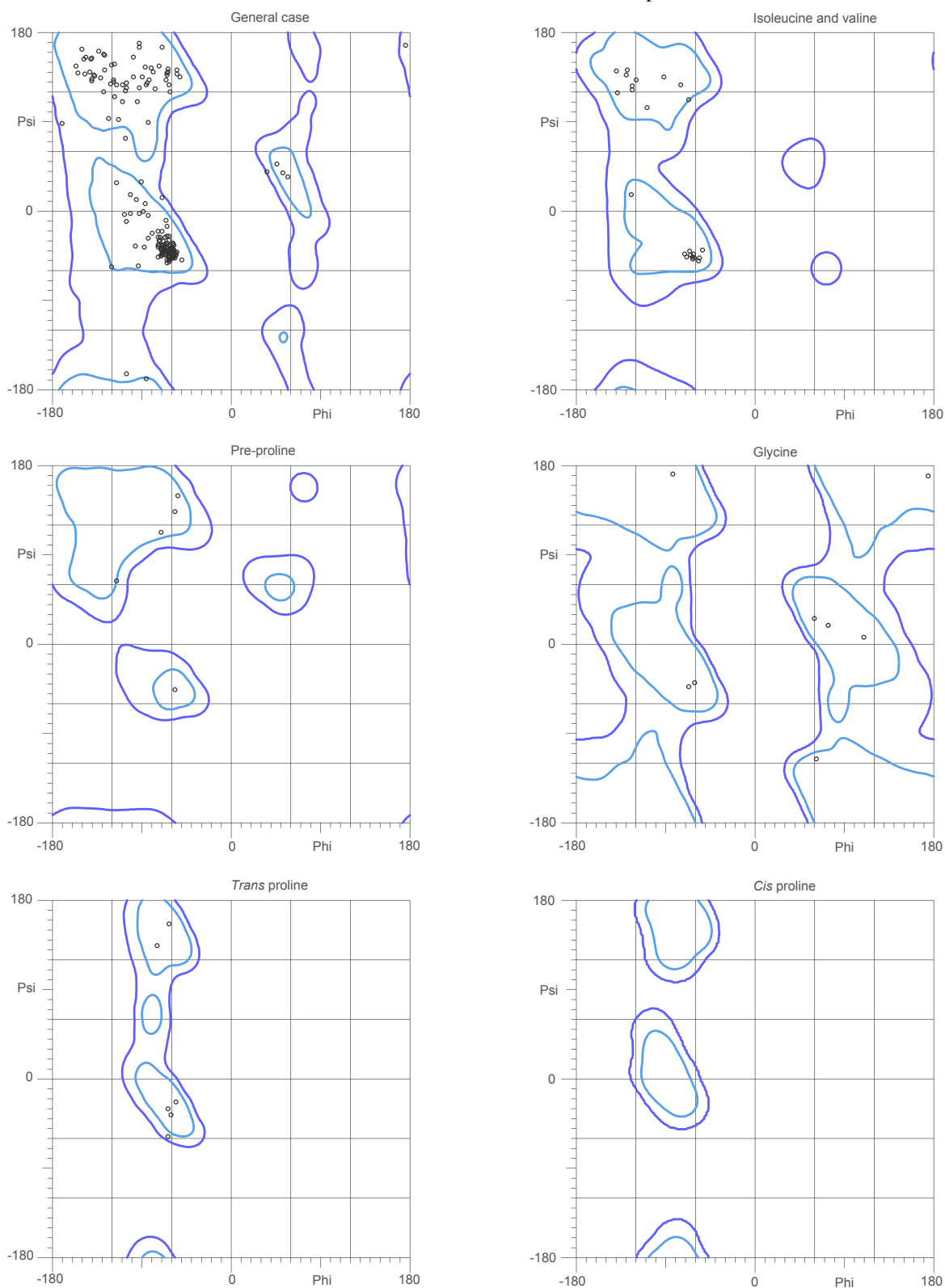


# MolProbity Ramachandran analysis

EntAAA-3-11-xds\_refine\_66-TER\_clean\_trimmed.pdb, model 1



96.1% (219/228) of all residues were in favored (98%) regions.  
100.0% (228/228) of all residues were in allowed (>99.8%) regions.

There were no outliers.