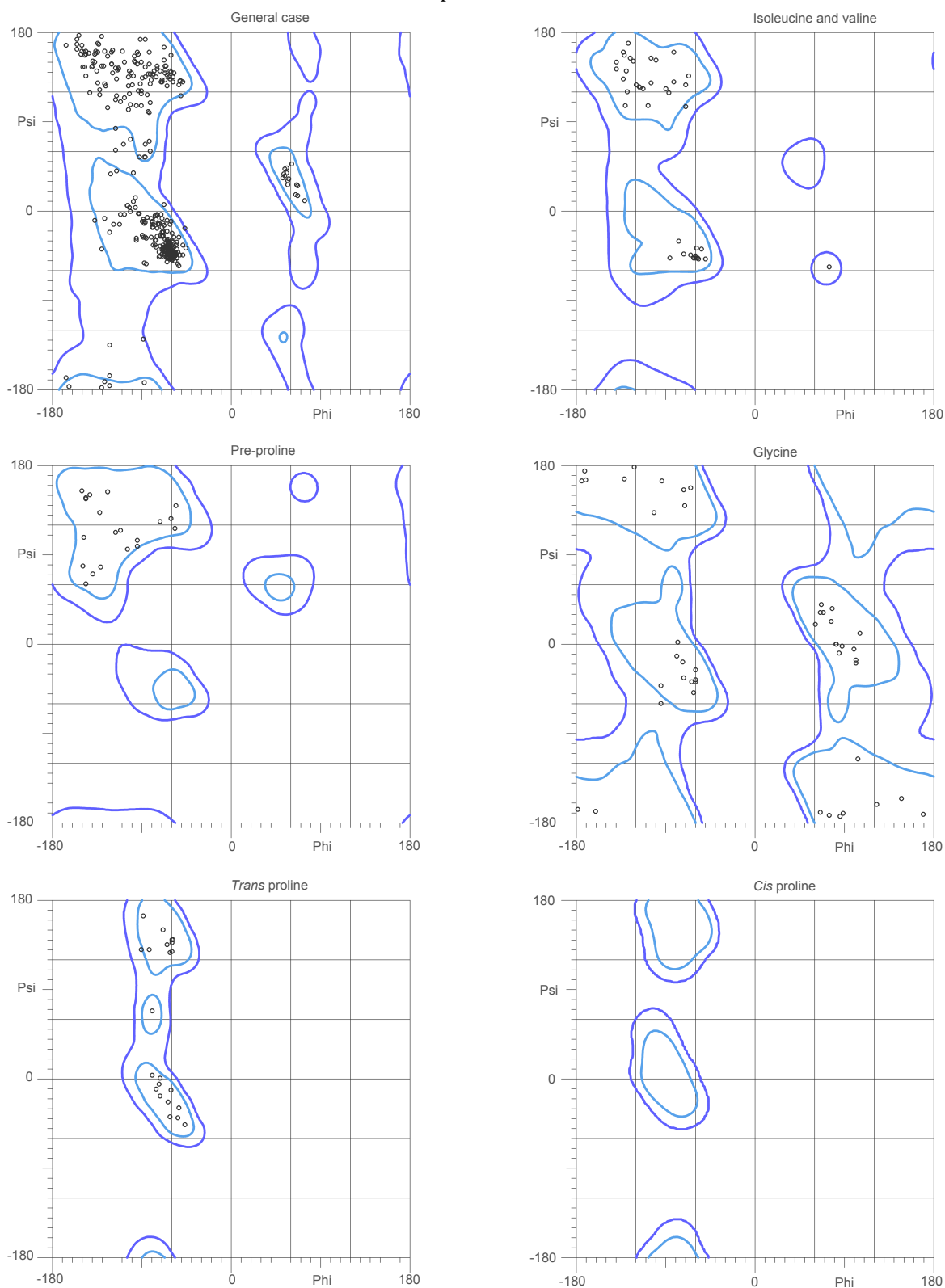


MolProbity Ramachandran analysis

litx.pdb, model 1



96.6% (456/472) of all residues were in favored (98%) regions.
100.0% (472/472) of all residues were in allowed (>99.8%) regions.

There were no outliers.