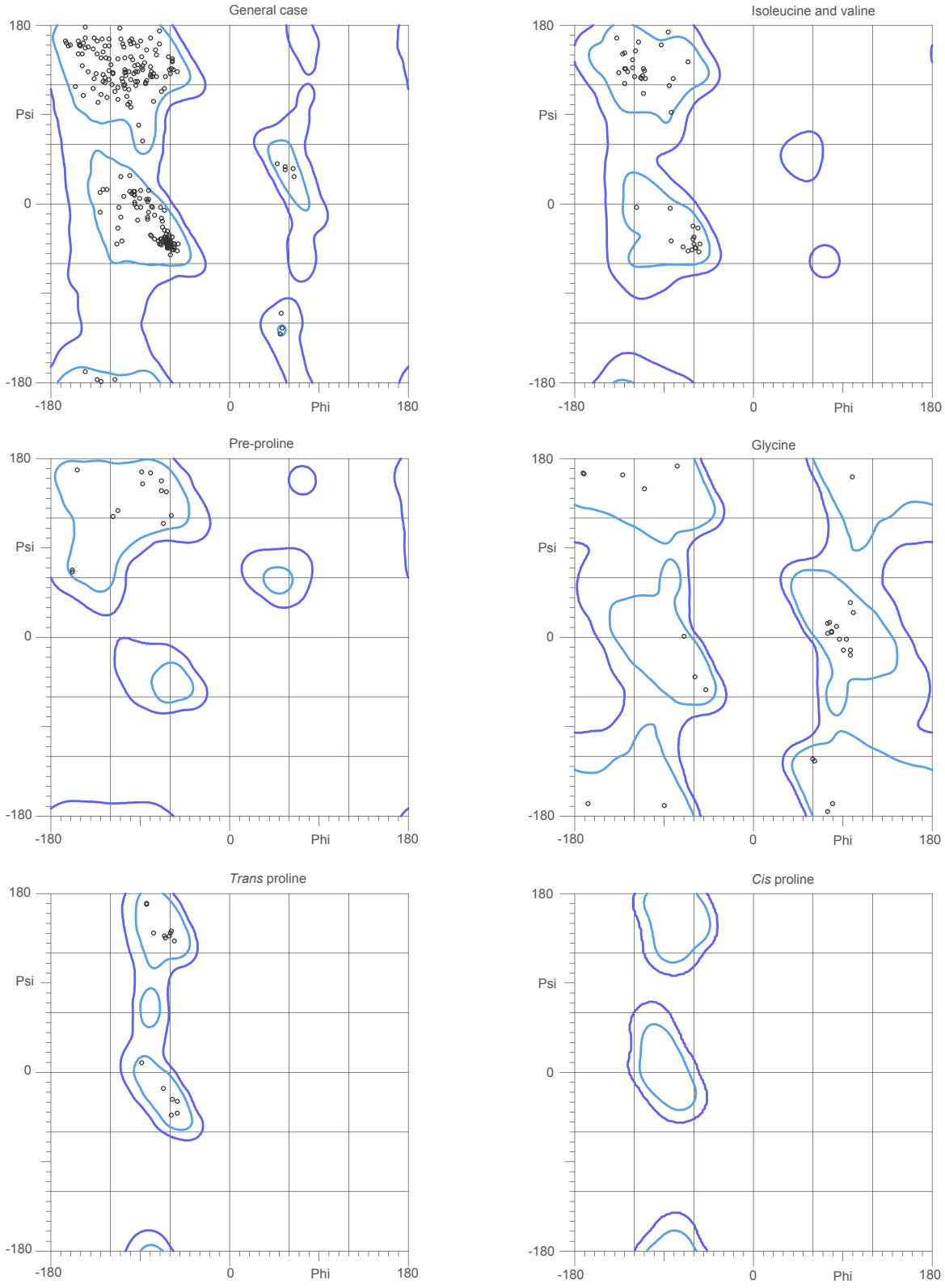
MolProbity Ramachandran analysis

6xkh.H.pdb, model 1



99.4% (329/331) of all residues were in favored (98%) regions.
100.0% (331/331) of all residues were in allowed (>99.8%) regions.

There were no outliers.