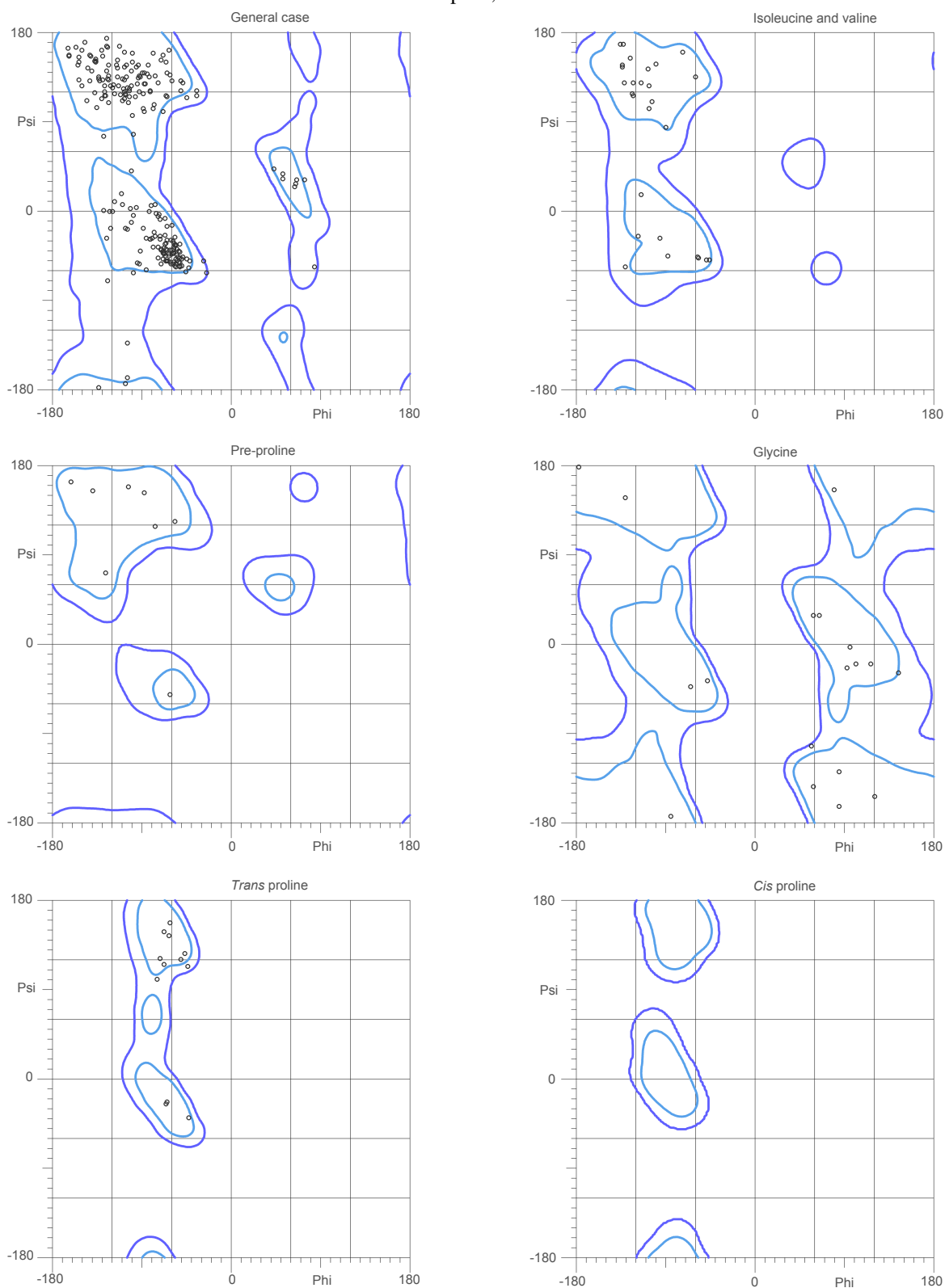


# MolProbity Ramachandran analysis

7krx.H.pdb, model 1



93.4% (285/305) of all residues were in favored (98%) regions.

100.0% (305/305) of all residues were in allowed (>99.8%) regions.

There were no outliers.