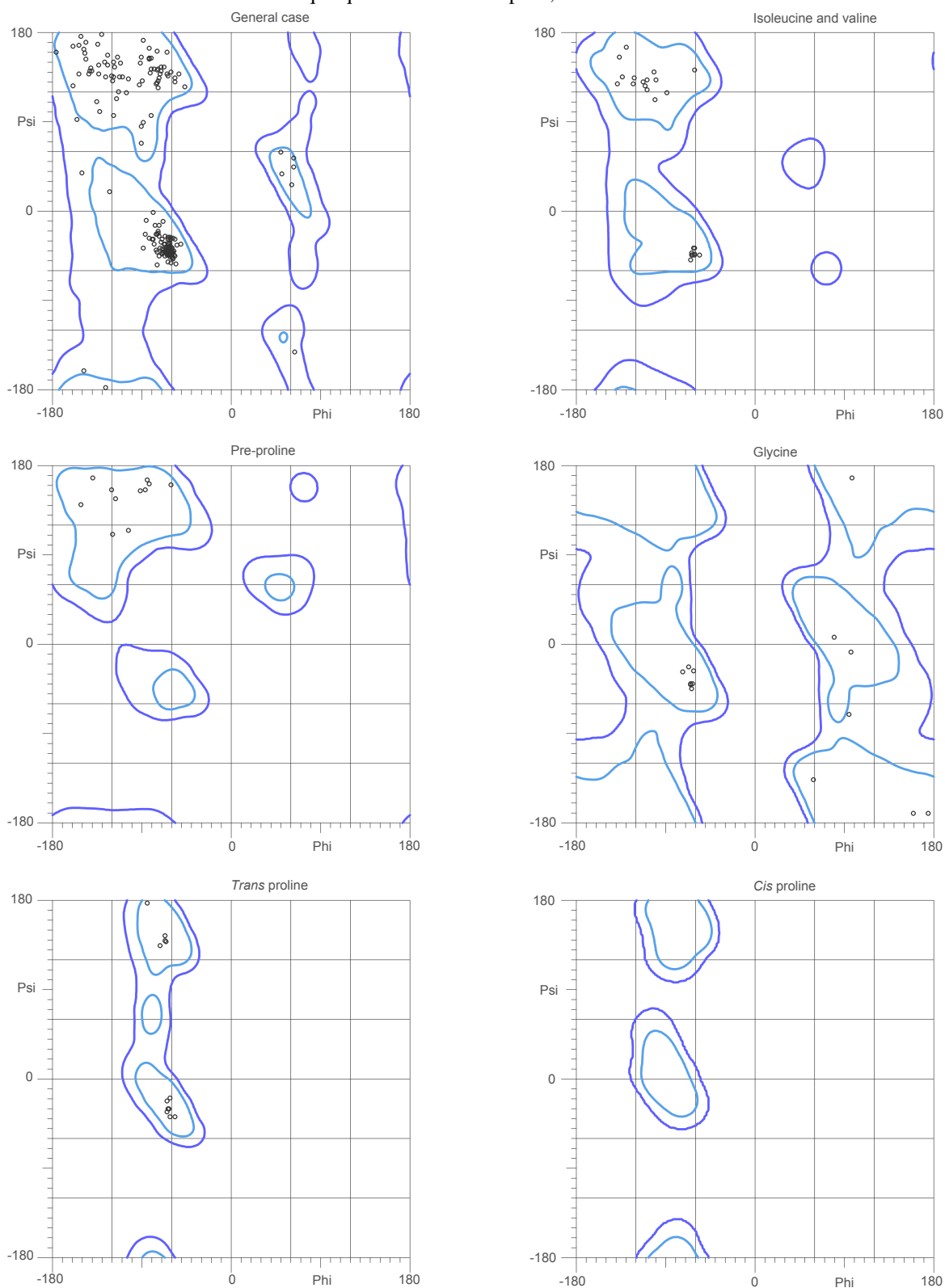# MolProbity Ramachandran analysis

qseq1.B99990001.pdb, model 1

97.7% (250/256) of all residues were in favored (98%) regions.

100.0% (256/256) of all residues were in allowed (>99.8%) regions.

There were no outliers.