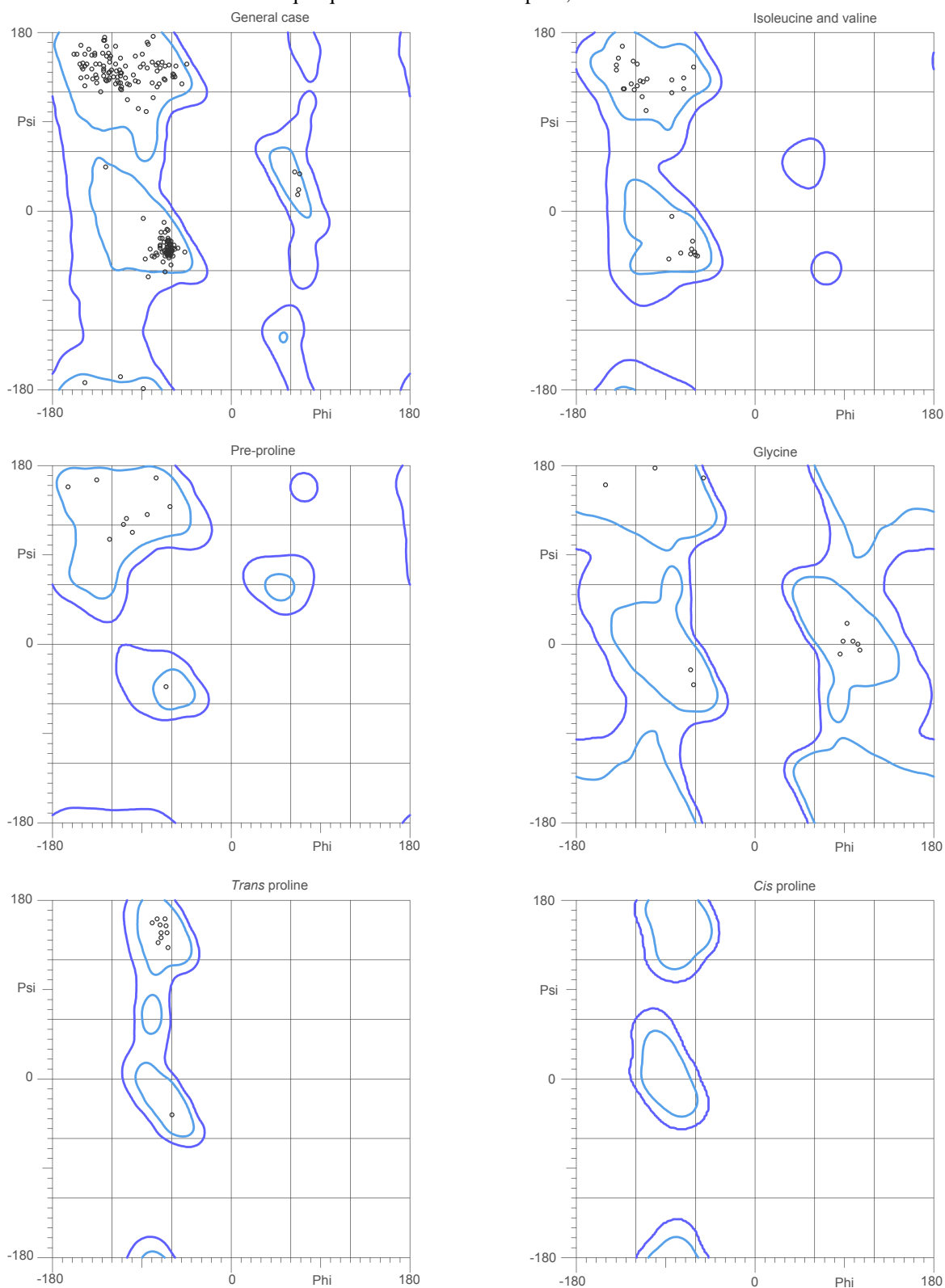# MolProbity Ramachandran analysis

qseq1.B99990002FH.pdb, model 1

97.6% (239/245) of all residues were in favored (98%) regions.

100.0% (245/245) of all residues were in allowed (>99.8%) regions.

There were no outliers.