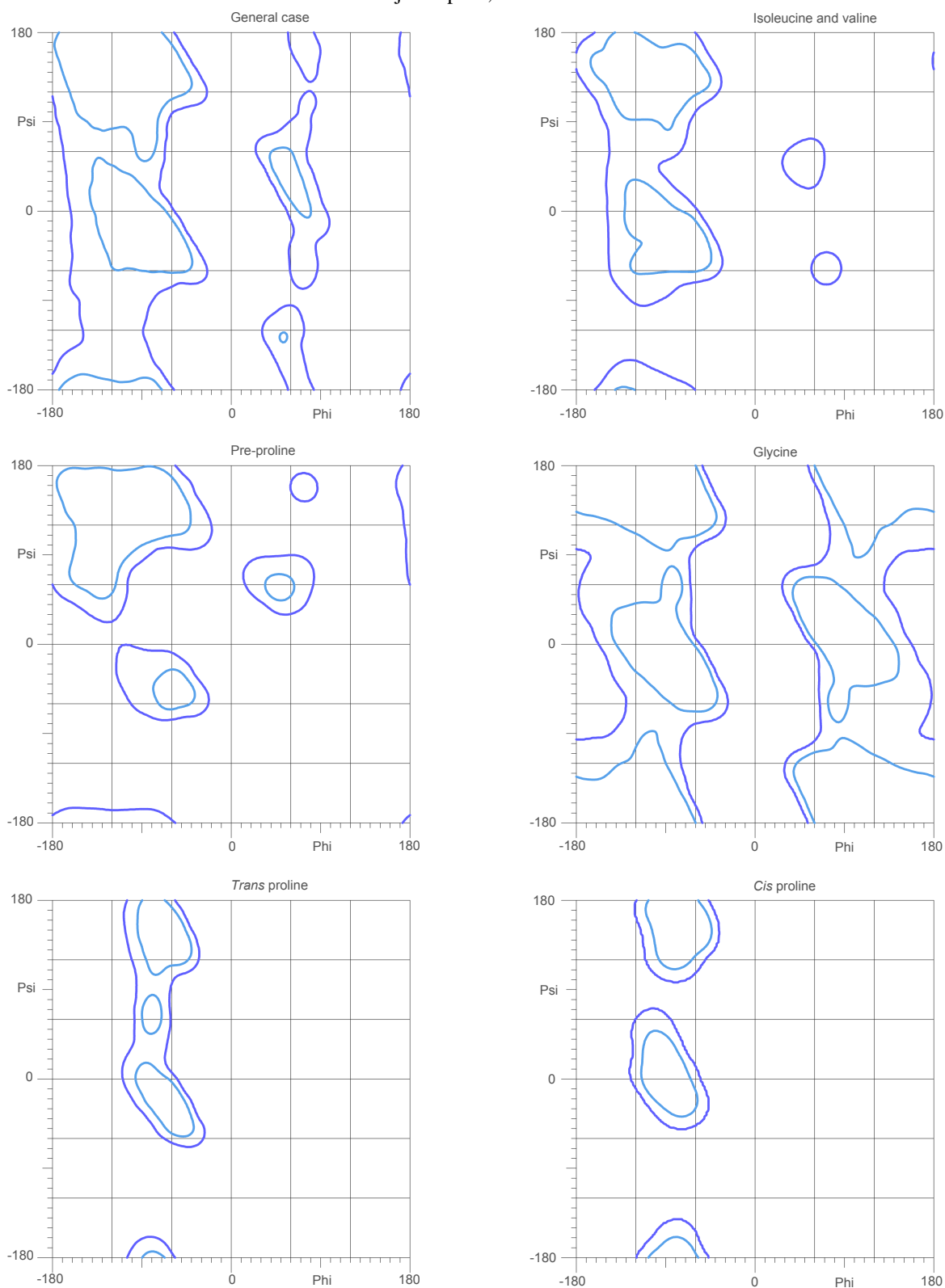


MolProbity Ramachandran analysis

1xjr.H.pdb, model 1



% (0/0) of all residues were in favored (98%) regions.

% (0/0) of all residues were in allowed (>99.8%) regions.

There were no outliers.