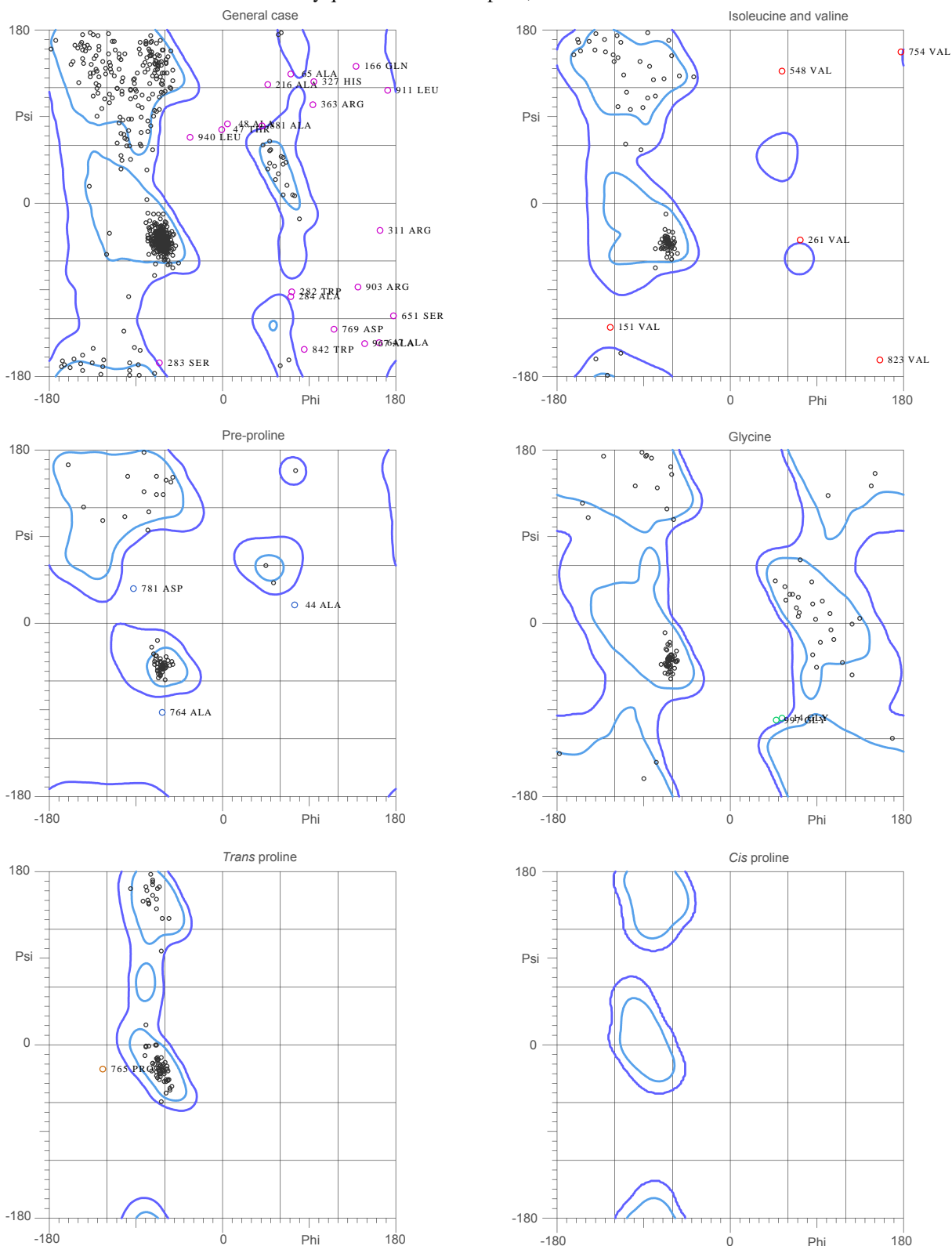


MolProbity Ramachandran analysis

1yq2.B99990002.pdb, model 1



90.0% (157/173) of all residues were in favored (blue) regions.

97.0% (167/173) of all residues were in allowed (blue) regions.

There were 11 outliers (Phi, Psi):

166 GLN (Phi, Psi)
216 ALA (Phi, Psi)
327 HIS (Phi, Psi)
363 ARG (Phi, Psi)
48 ALA (Phi, Psi)
147 THR (Phi, Psi)
181 ALA (Phi, Psi)
940 LEU (Phi, Psi)
283 SER (Phi, Psi)
282 TRP (Phi, Psi)
284 ALA (Phi, Psi)
903 ARG (Phi, Psi)
769 ASP (Phi, Psi)
842 TRP (Phi, Psi)
976 ALA (Phi, Psi)
651 SER (Phi, Psi)
311 ARG (Phi, Psi)
781 ASP (Phi, Psi)
44 ALA (Phi, Psi)
764 ALA (Phi, Psi)
765 PRO (Phi, Psi)
974 GLY (Phi, Psi)
754 VAL (Phi, Psi)
548 VAL (Phi, Psi)
151 VAL (Phi, Psi)
261 VAL (Phi, Psi)
823 VAL (Phi, Psi)