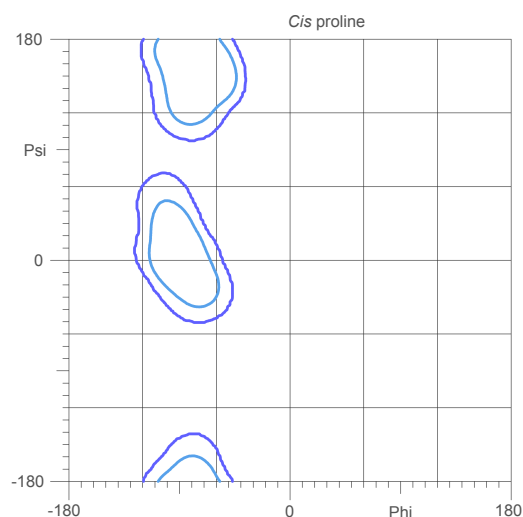
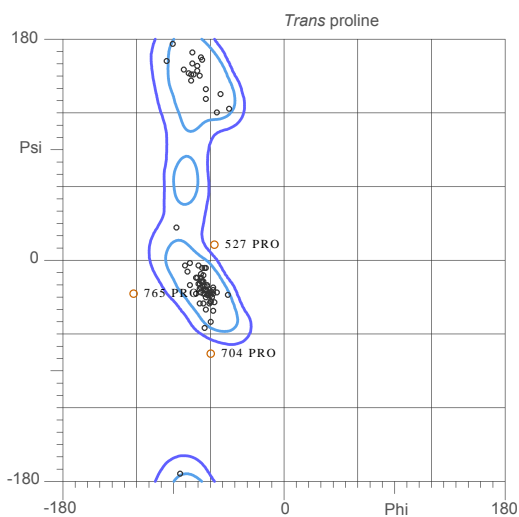
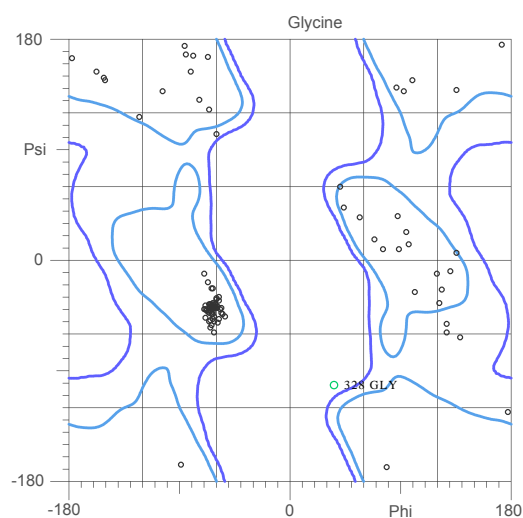
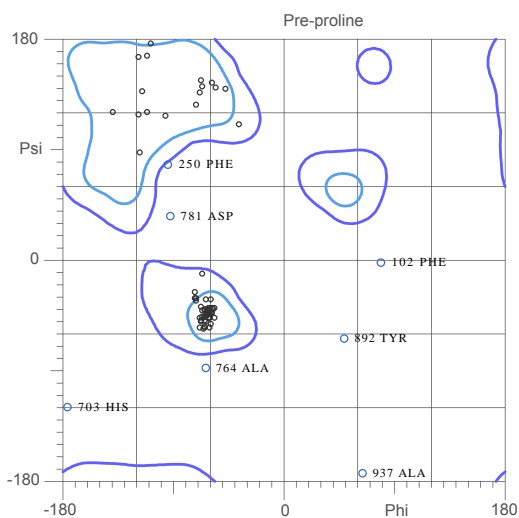
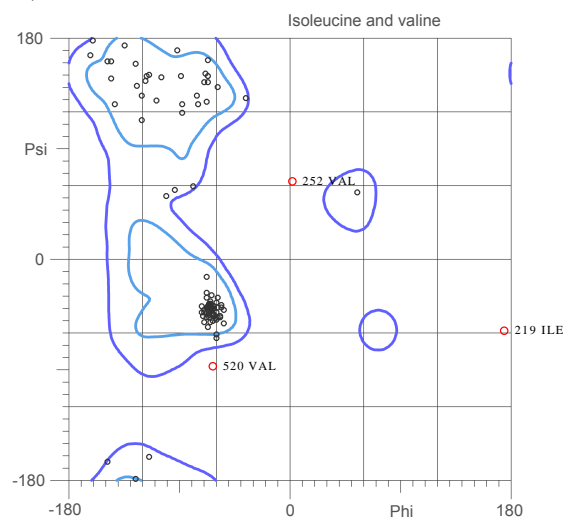
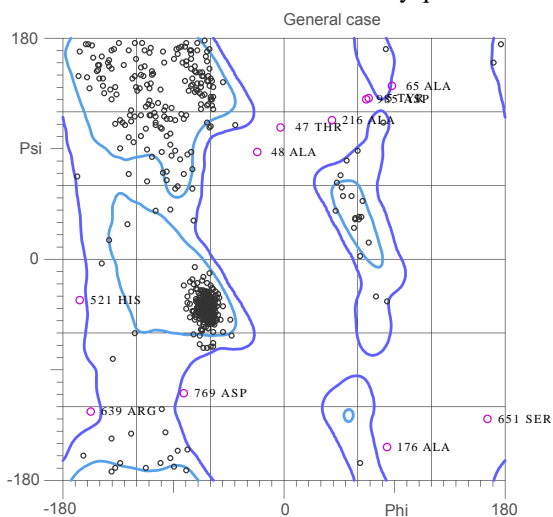


MolProbity Ramachandran analysis

1yq2.B99990036.pdb, model 1



85.9% (918/1058) of all residues were in favored (90%) regions.
10.2% (107/1058) of all residues were in allowed (100%) regions.

There were 25 outliers (phi, psi):

6 TYR (phi, psi)
47 THR (phi, psi)
48 ALA (phi, psi)
49 ALA (phi, psi)
102 PHE (phi, psi)
176 ALA (phi, psi)
216 ALA (phi, psi)
250 PHE (phi, psi)
252 VAL (phi, psi)
328 GLY (phi, psi)
47 THR (phi, psi)
48 ALA (phi, psi)
521 HIS (phi, psi)
639 ARG (phi, psi)
769 ASP (phi, psi)
937 ALA (phi, psi)
965 ASP (phi, psi)
176 ALA (phi, psi)
219 ILE (phi, psi)
250 PHE (phi, psi)
781 ASP (phi, psi)
892 TYR (phi, psi)
102 PHE (phi, psi)
704 PRO (phi, psi)
765 PRO (phi, psi)
527 PRO (phi, psi)