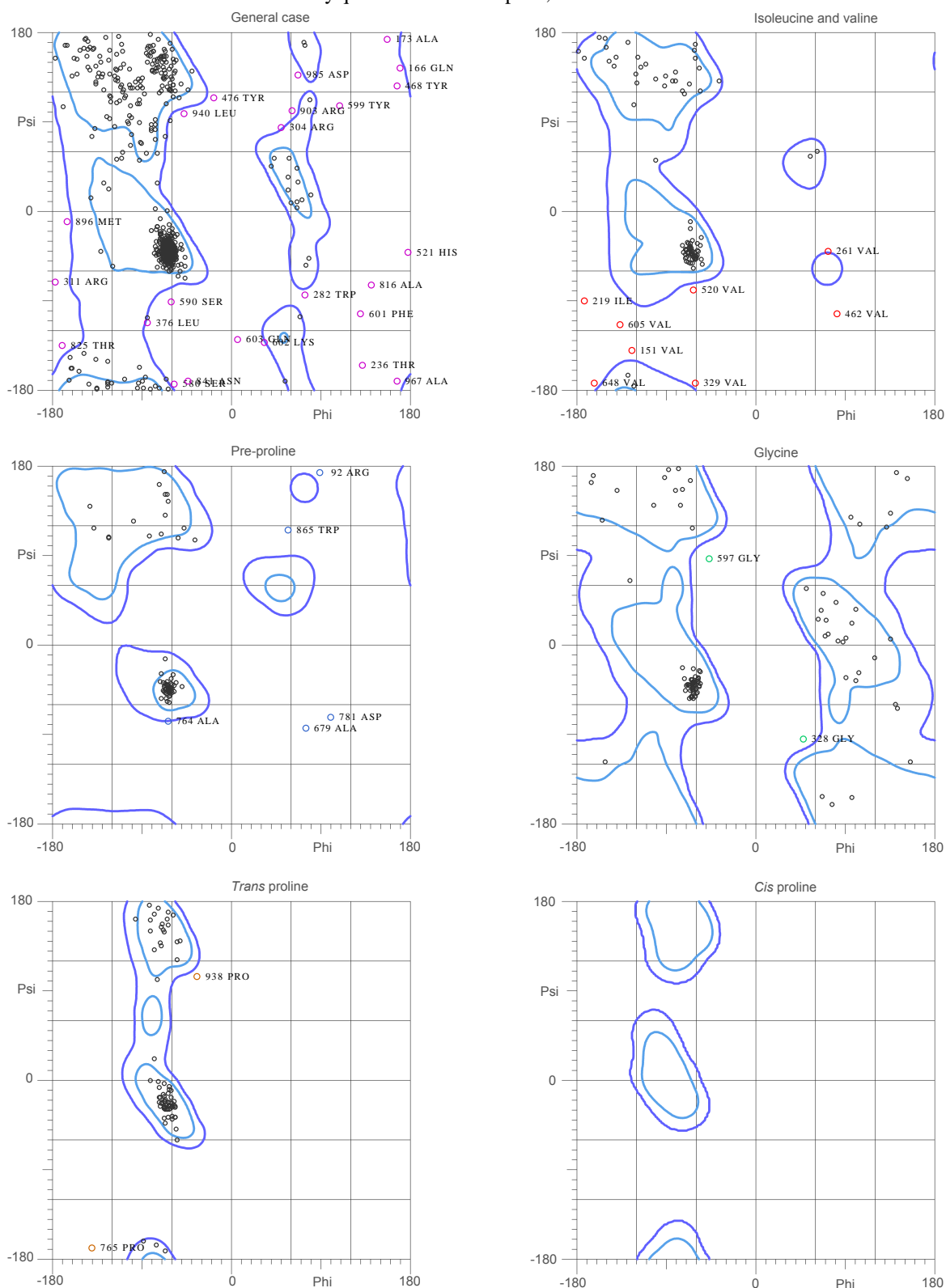


1yq2.B99990042.pdb, model 1



96.9% (95% CI 96.0-97.8) of all isolates were in *Escherichia* (95% CI 96.0-97.8) and 3.1% (95% CI 2.2-4.0) of all isolates were in *Shigella* (95% CI 2.2-4.0).