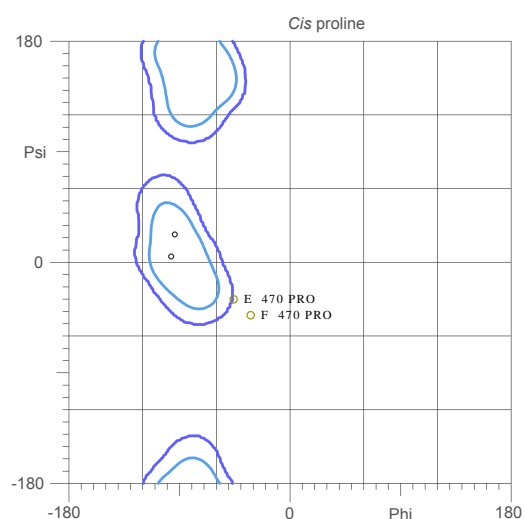
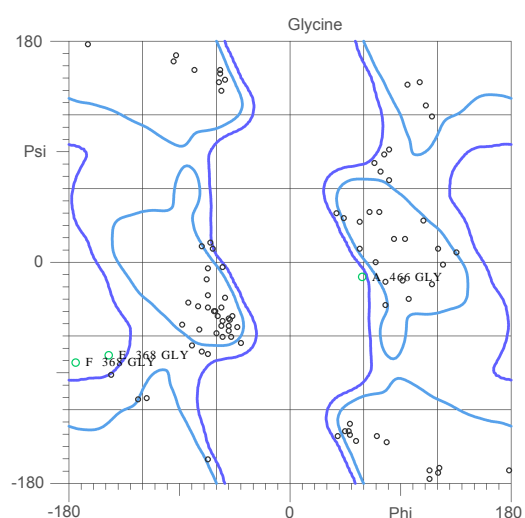
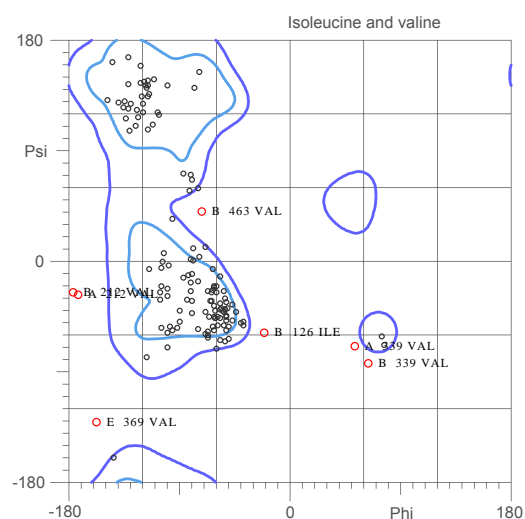


3d0h.H.pdb, model 1



Phylogenetic tree and confidence values (bootstrap values) are shown for the 1000 random samples generated by 10000 iterations of the MCMC analysis. The scale bar represents 0.01 substitutions per site. The scale bar represents 0.01 substitutions per site. The scale bar represents 0.01 substitutions per site.