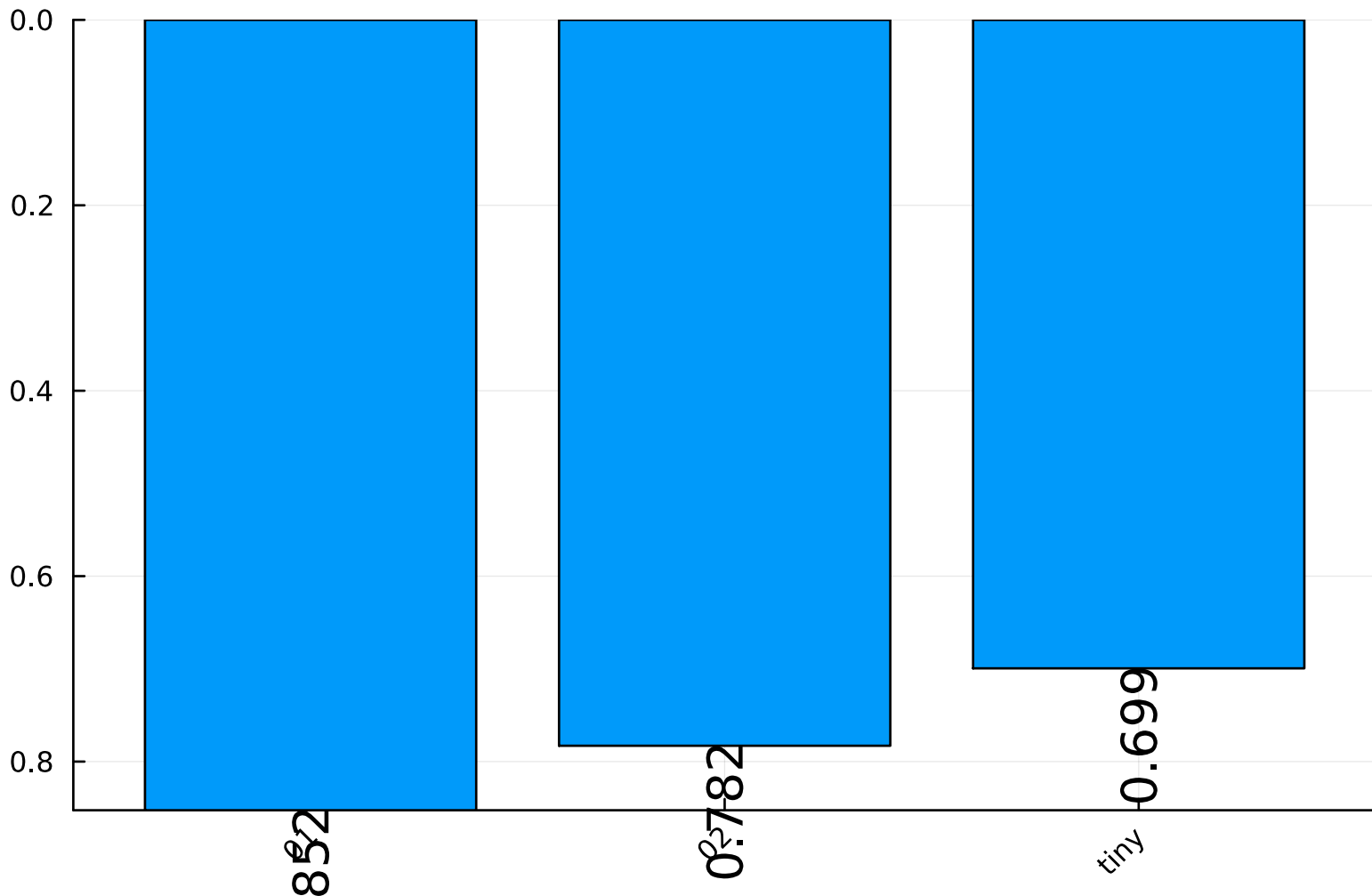
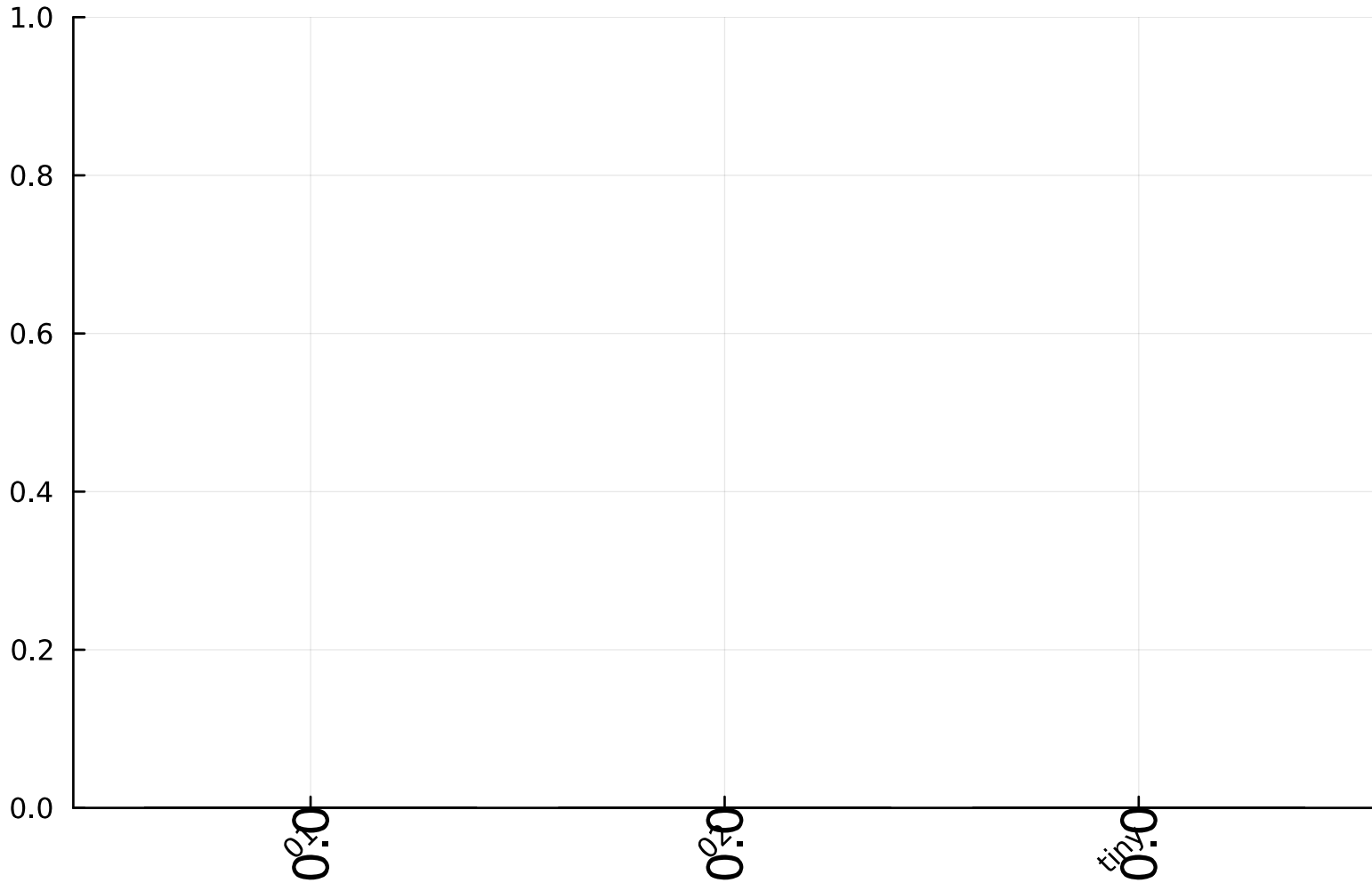


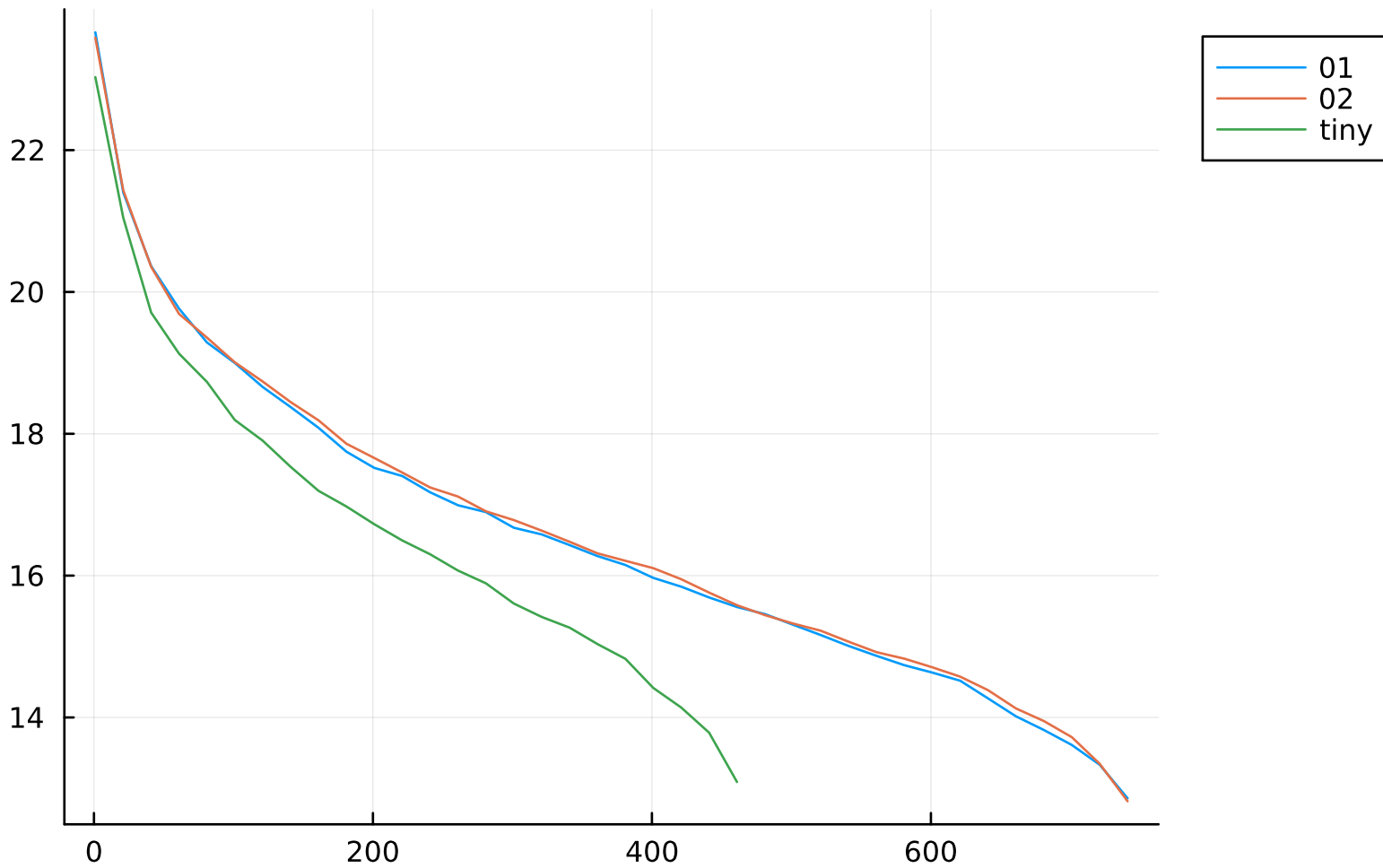
MS2 Mass Error Correction



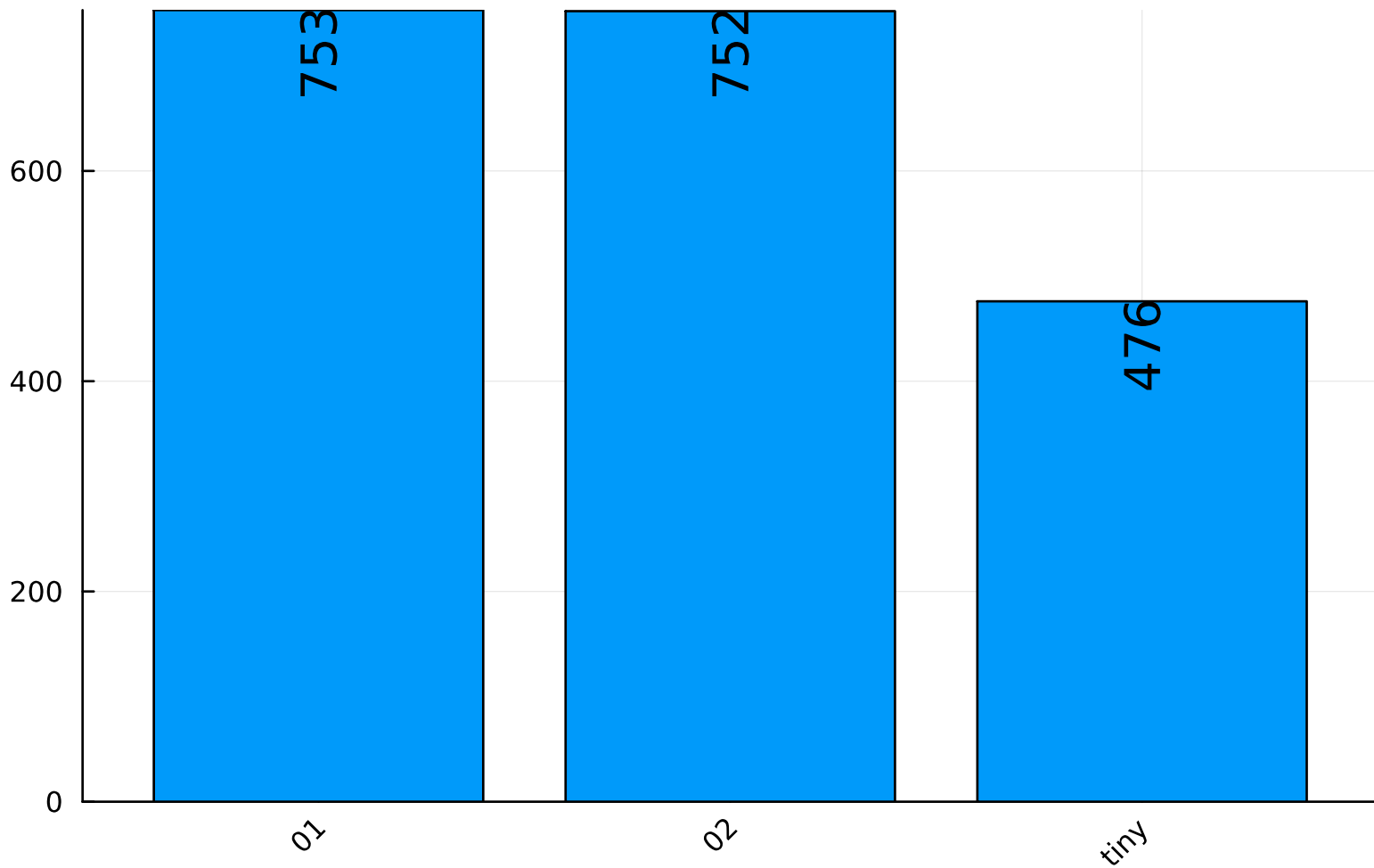
Missed Cleavage Percentage



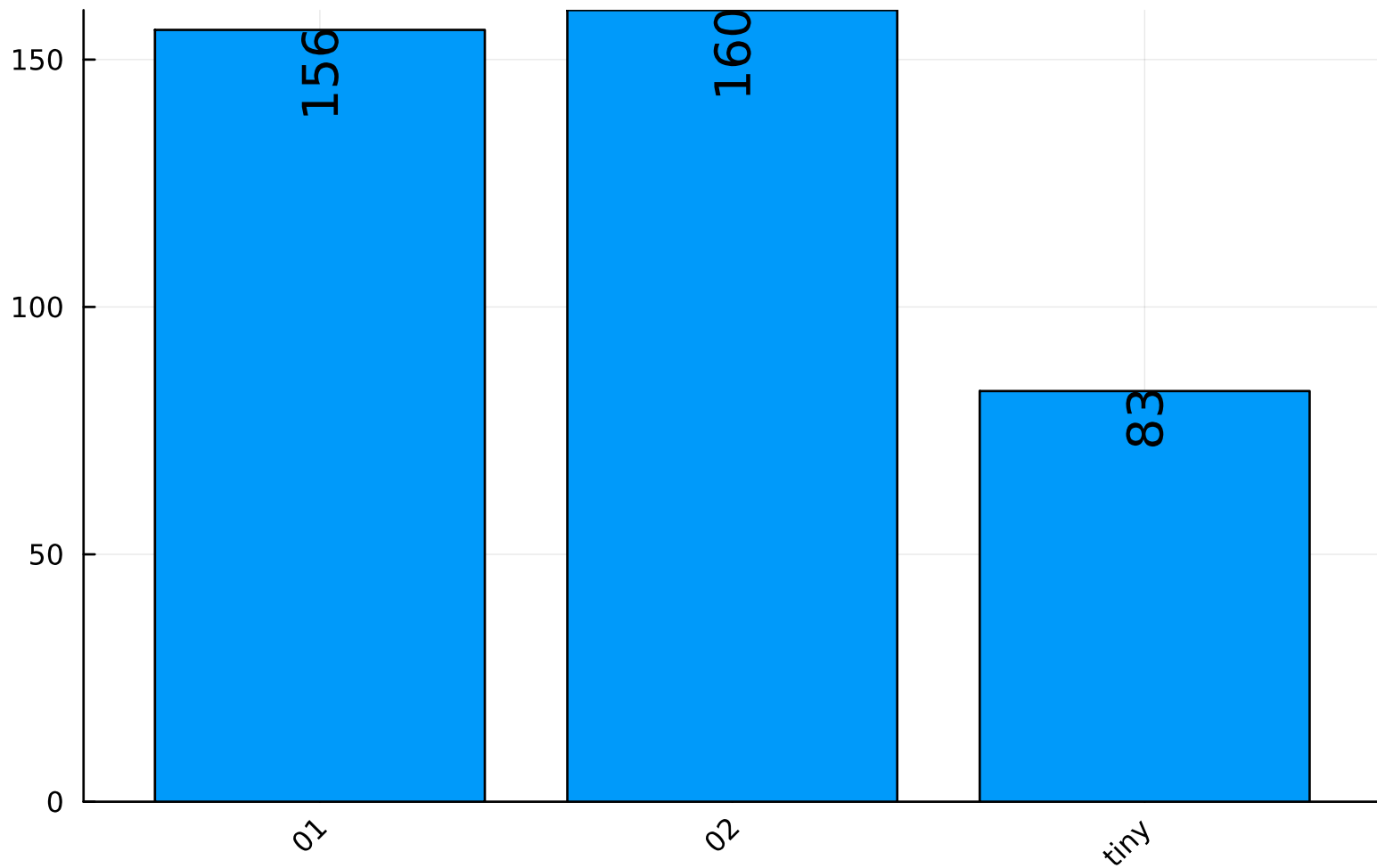
Precursor Abundance by Rank



Precursor ID's per File



Protein Group ID's per File



TIC plot

