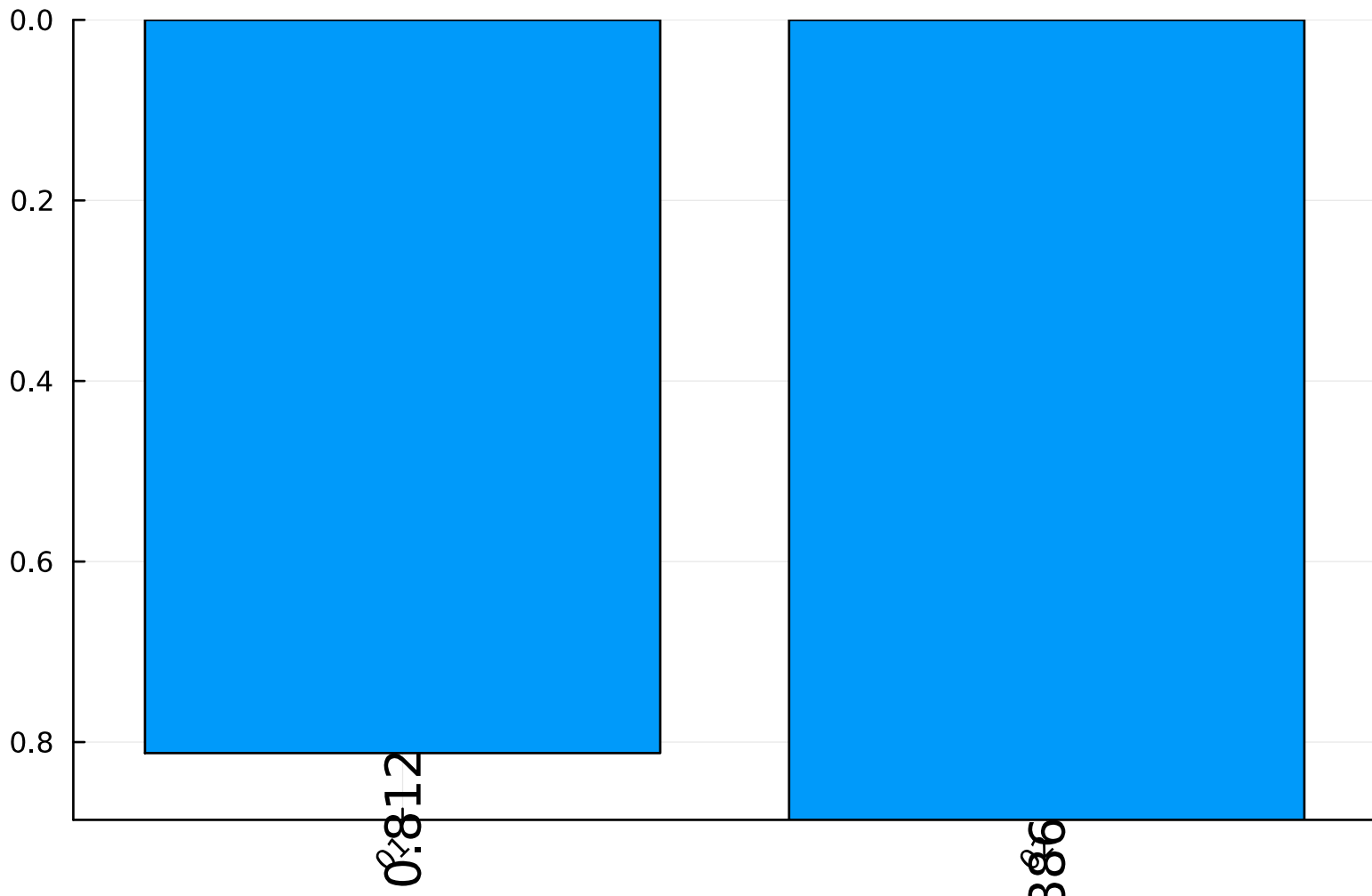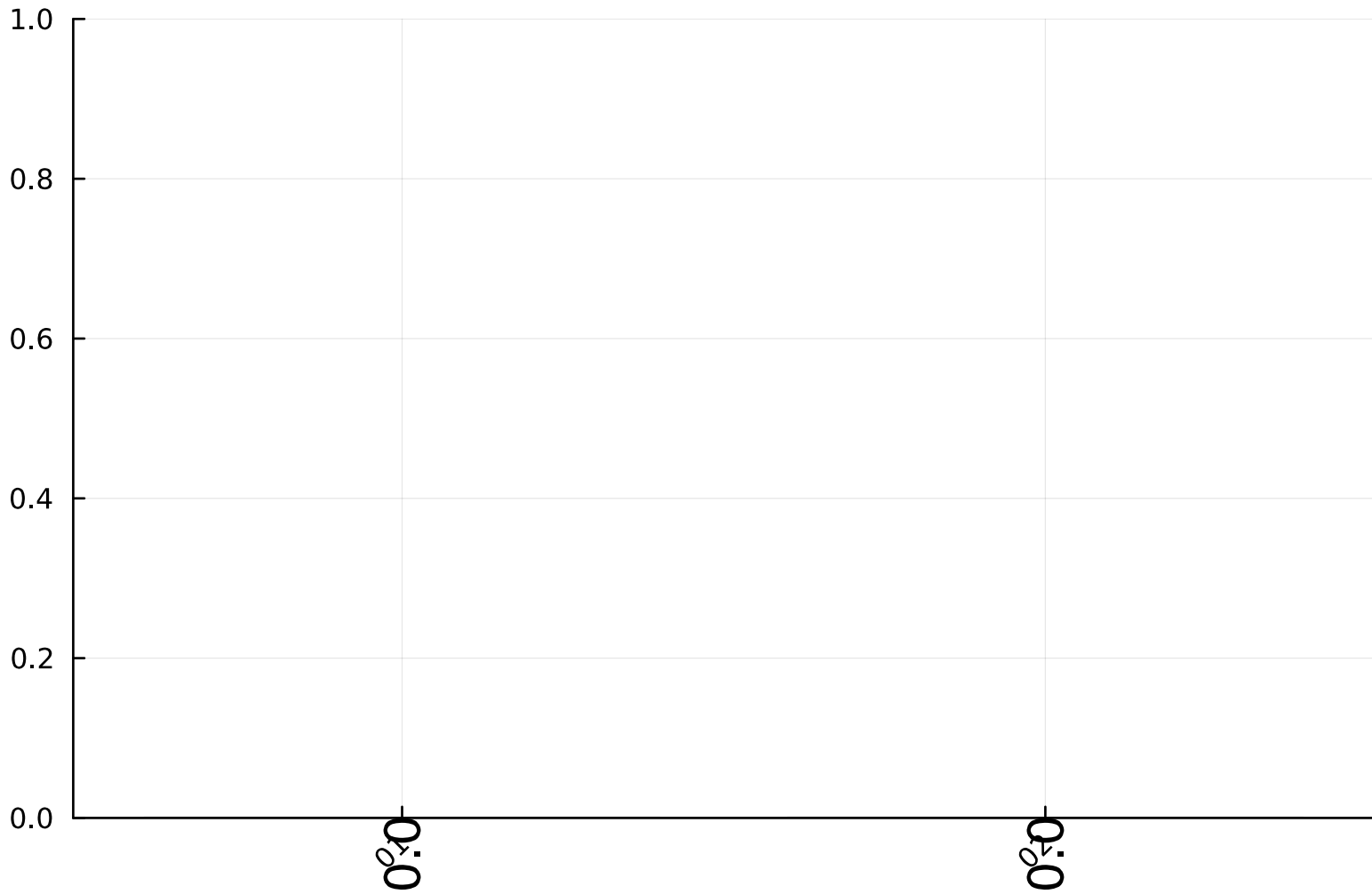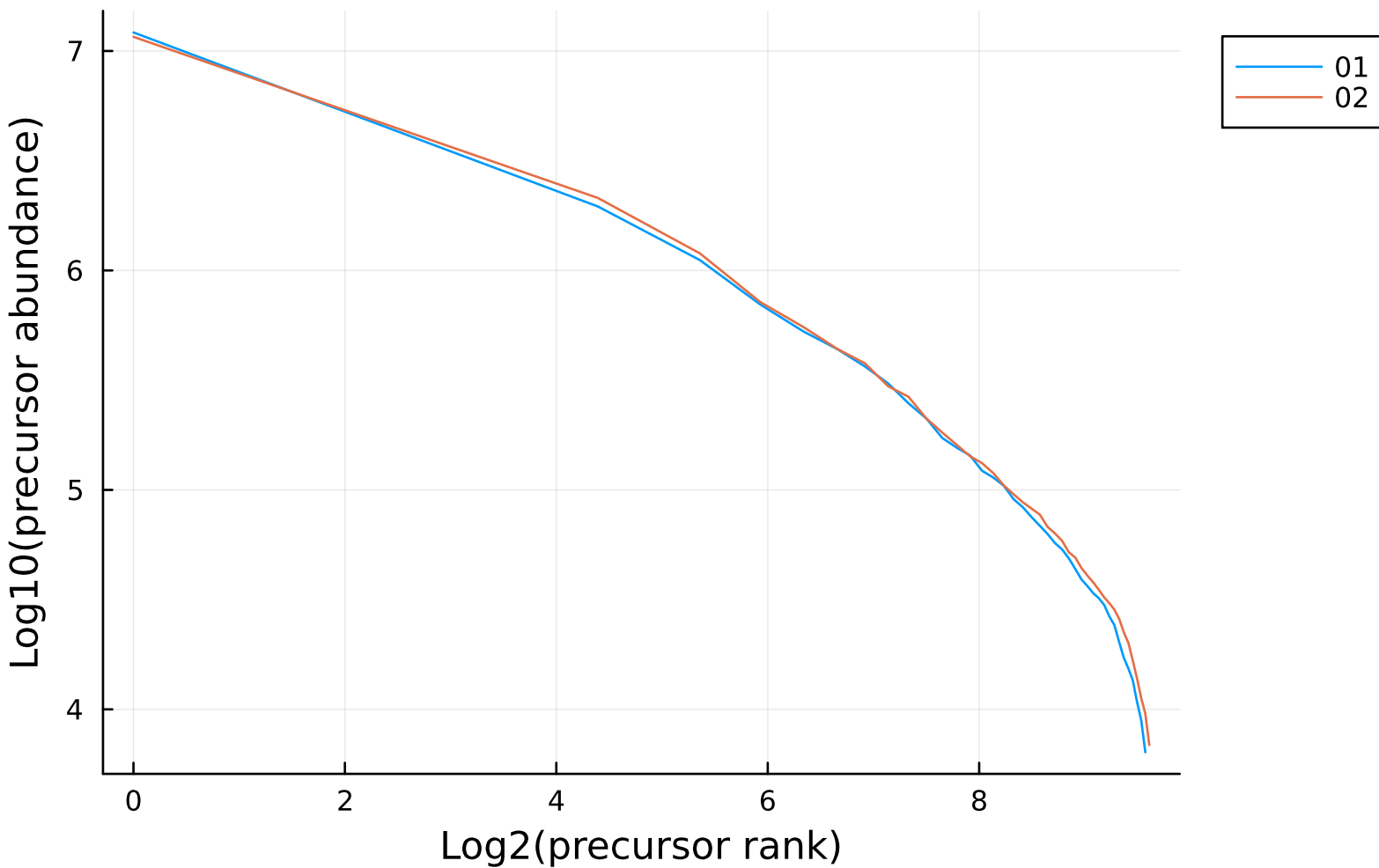# MS2 Mass Error Correction
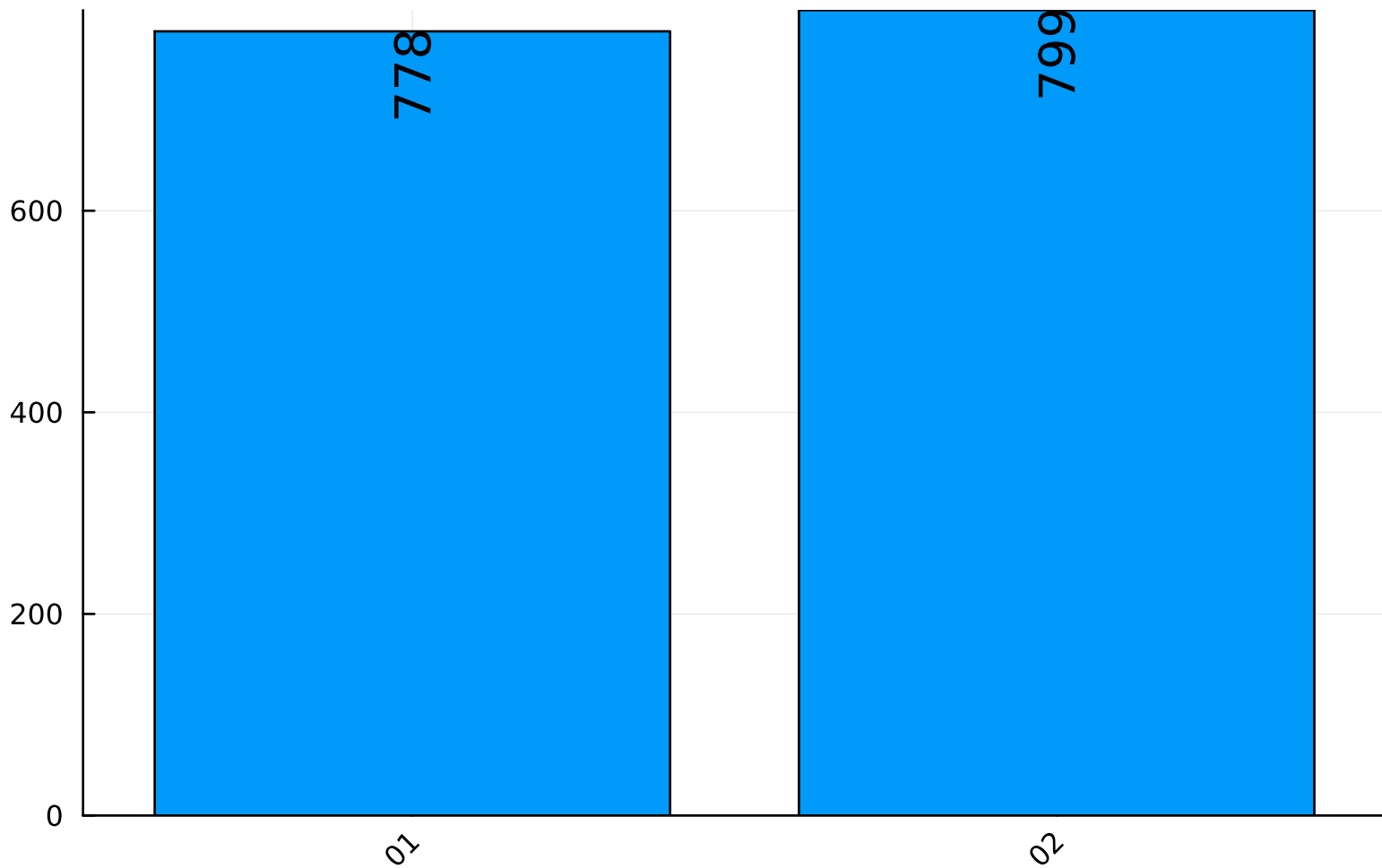
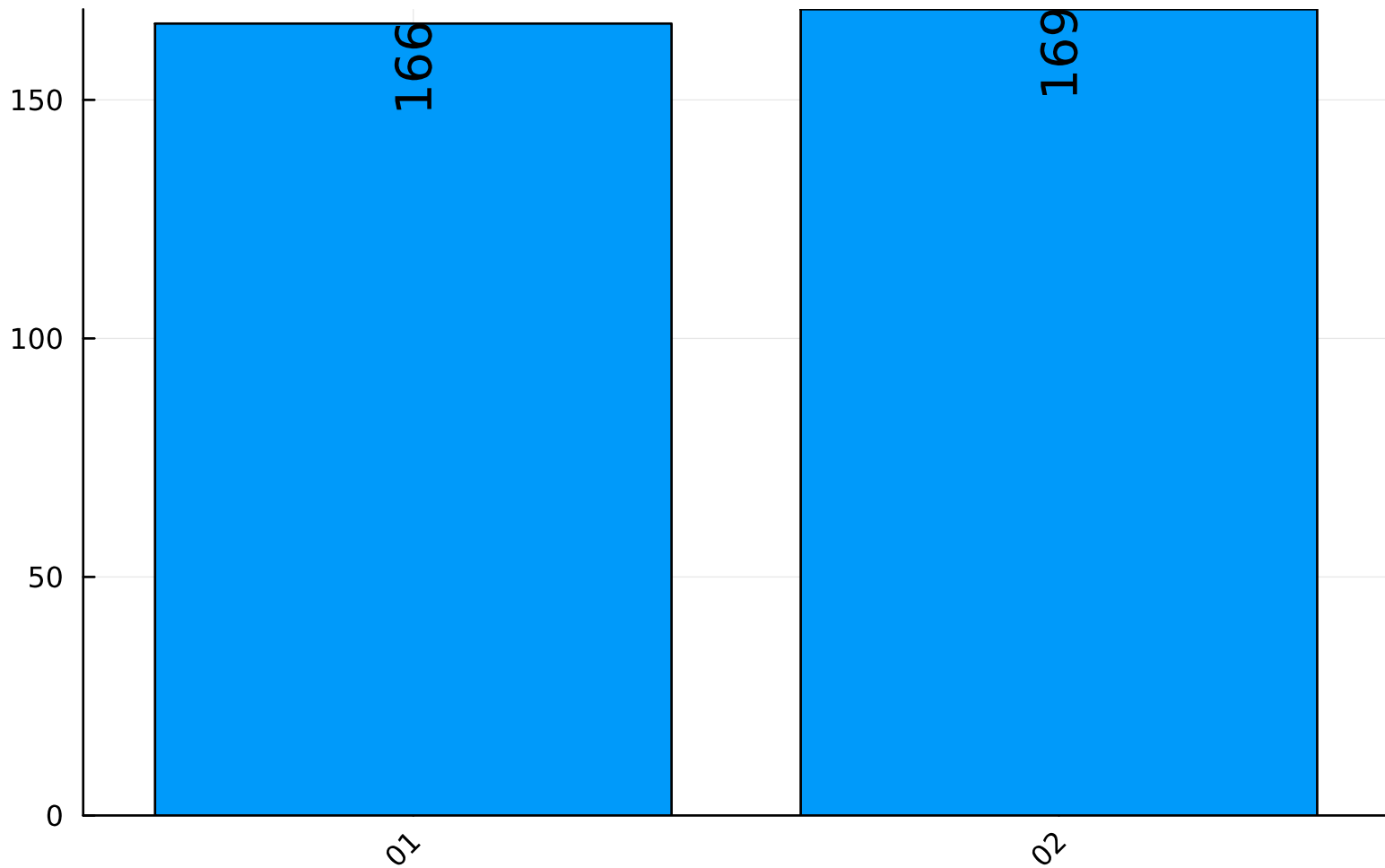# Missed Cleavage Percentage

0.0   0.0

Precursor Abundance by Rank

Precursor ID's per File

Protein Group ID's per File

TIC plot

retention time (min)

01
02