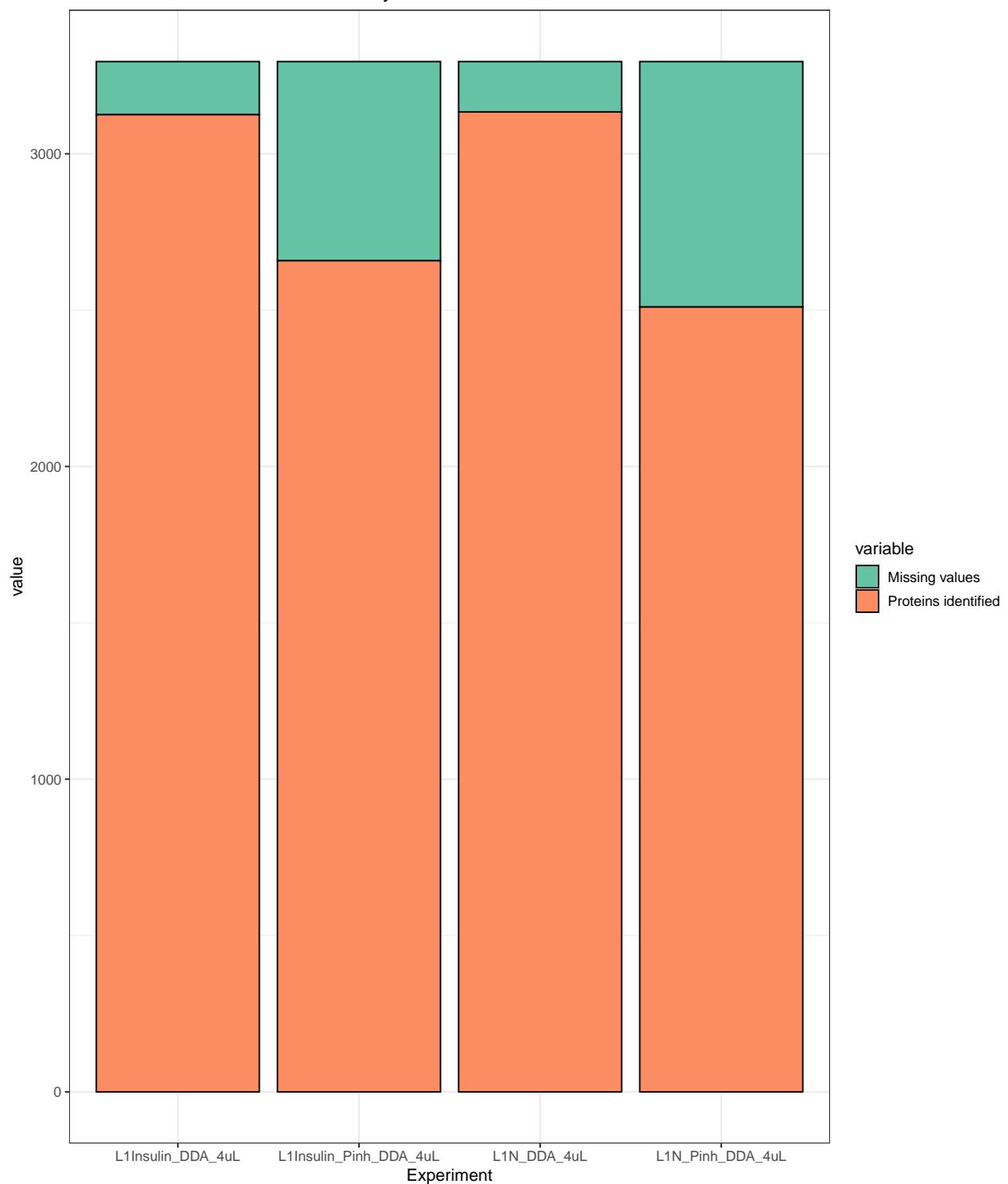


MQmetrics Report

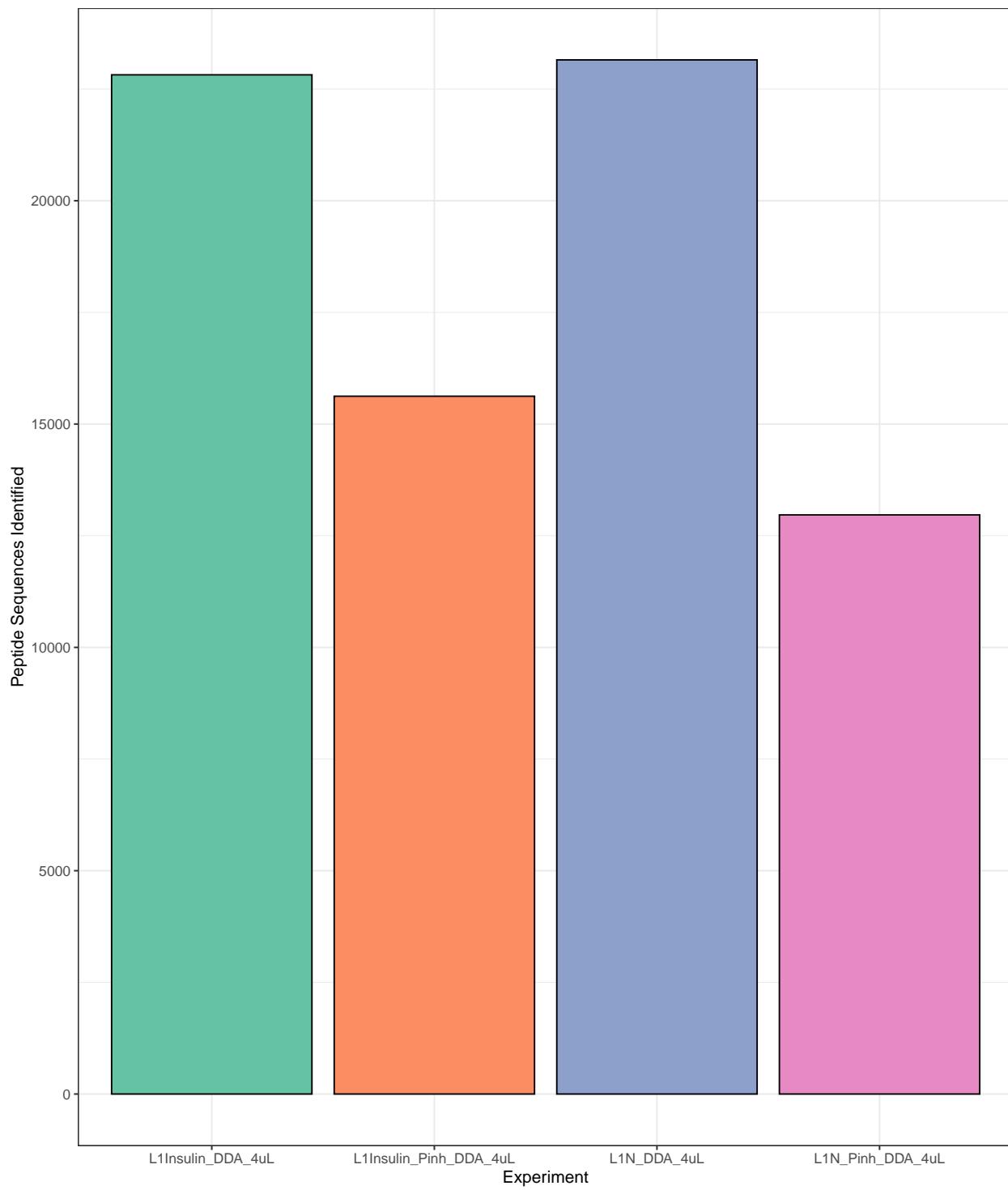
30 March, 2021

```
## [1] "The experiment started the day: 17/02/2021 at the time: 18:57:27."
## [1] "The whole experiment lasted: 05:48 (hours:minutes)."
## [1] "The MaxQuant version used was: 1.6.12.0"
## [1] "The user was: marek.vrbacky"
## [1] "The machine name was: FGU013PC029"
## [1] "The protein FDR was: 0.01"
## [1] "The match between runs was: True"
## [1] "The fasta file used was: C:\\MaxQuant_Databases\\\\UP000000589_10090.fasta"
```

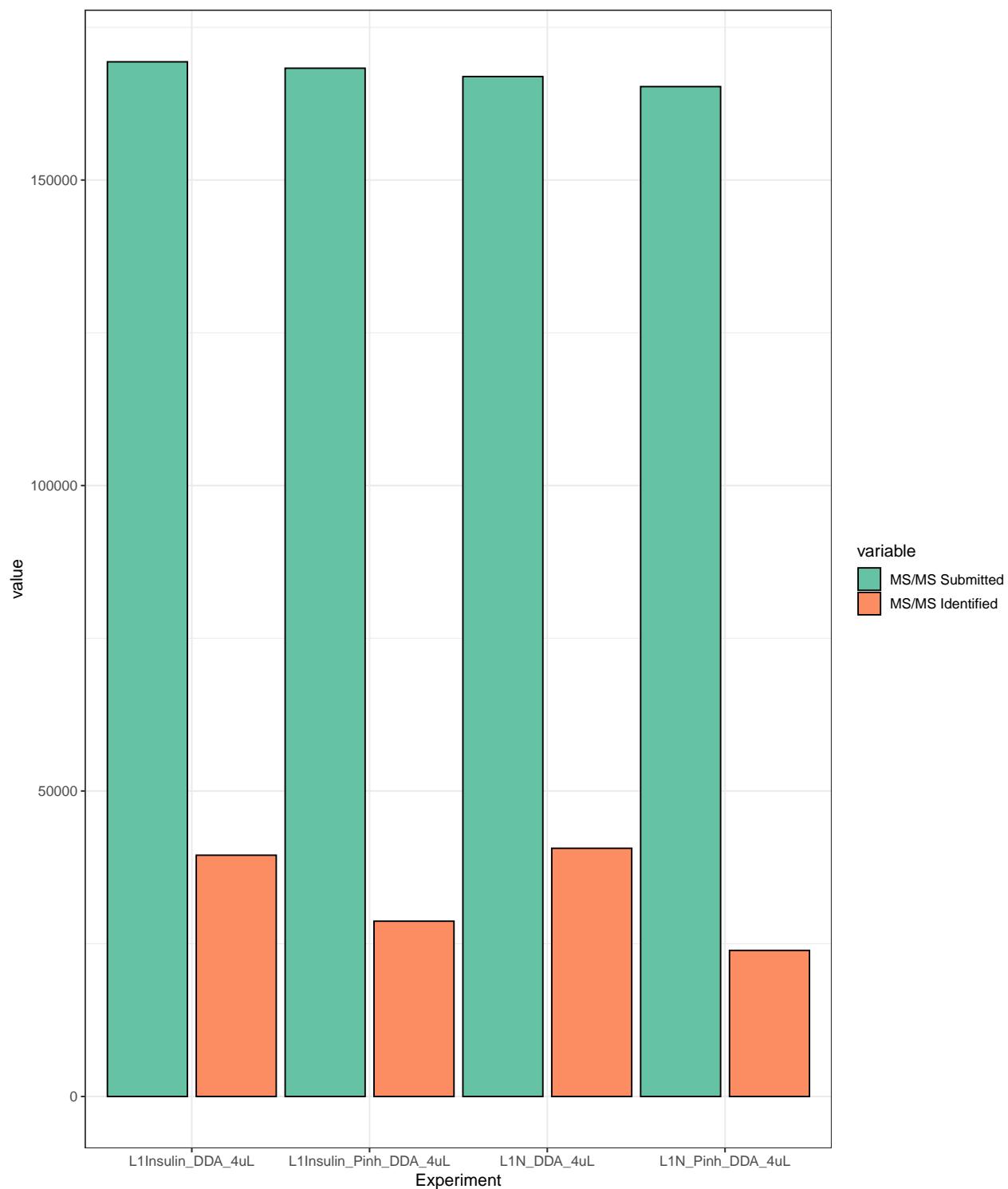
Proteins Identified based on Intensity



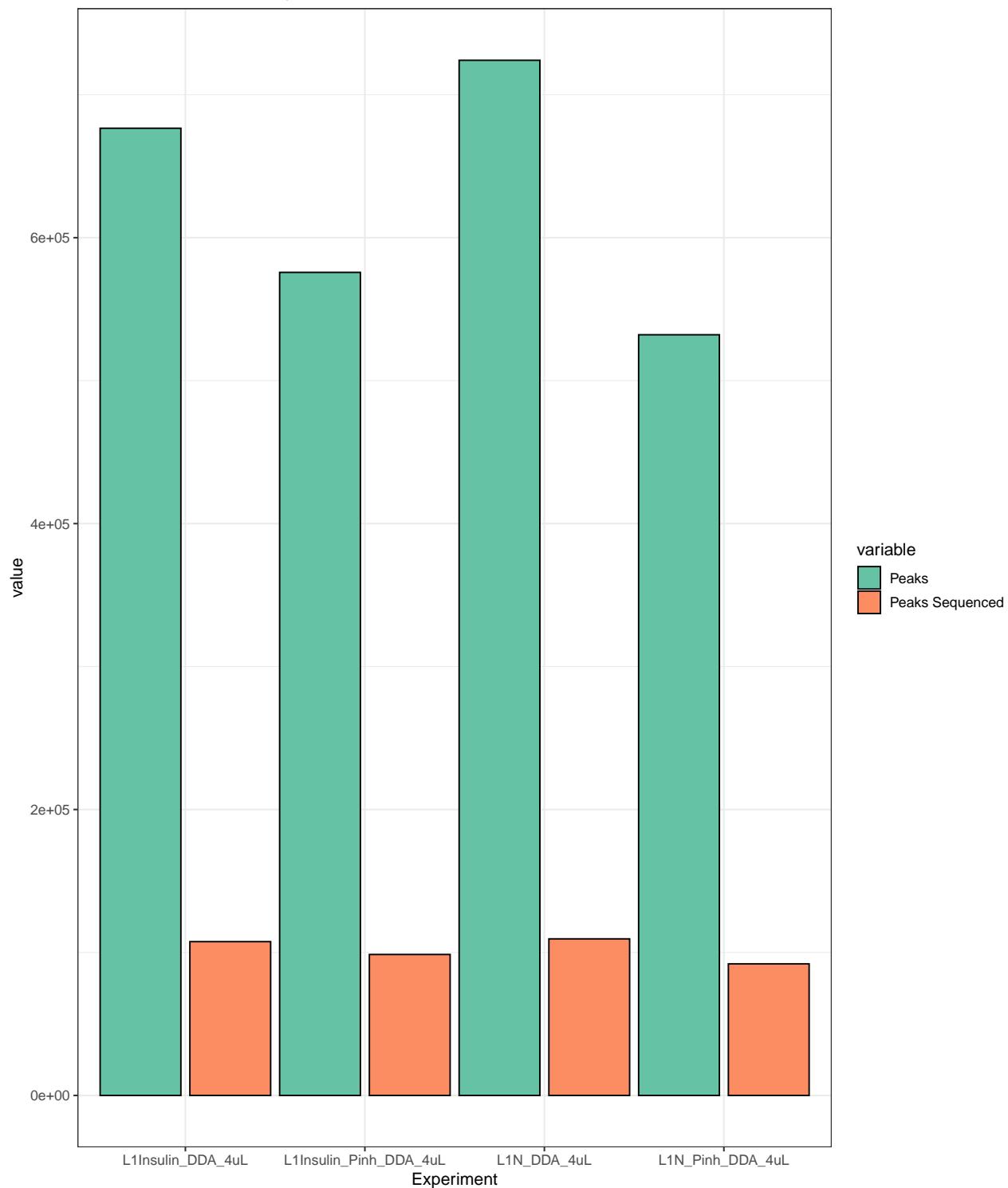
Peptides Sequences Identified



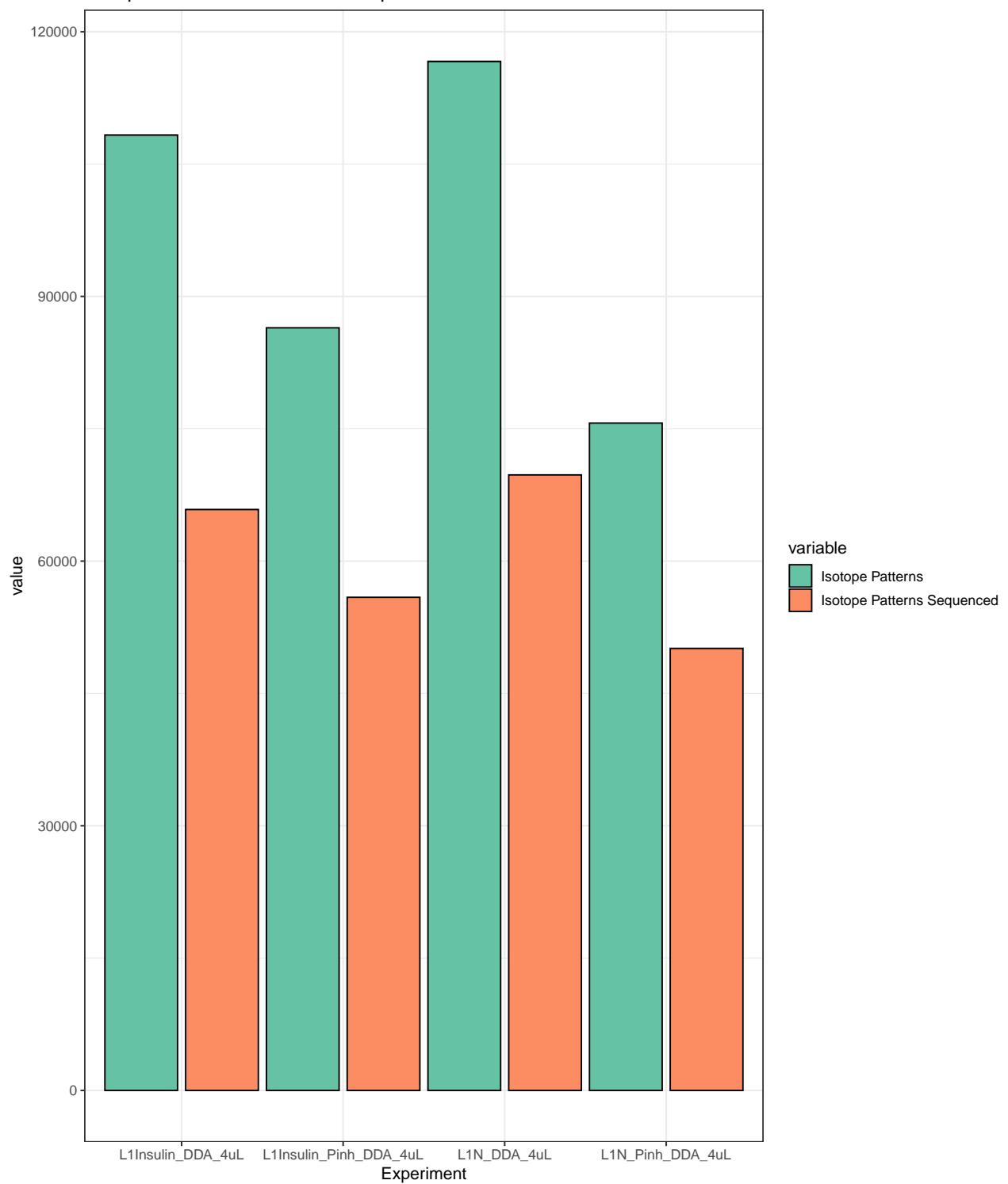
MS/MS Submitted and Identified



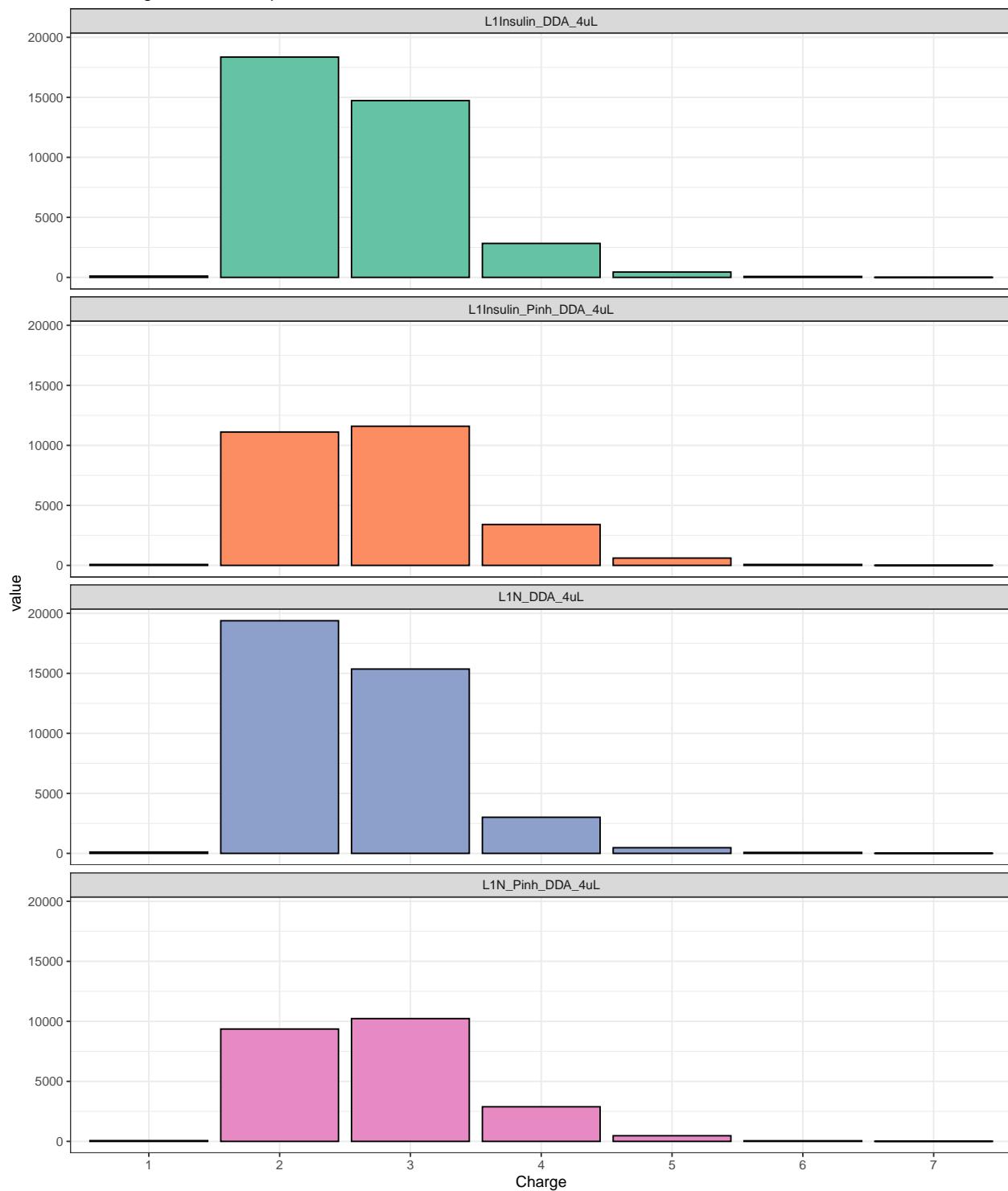
Peaks detected and sequenced in the full scans



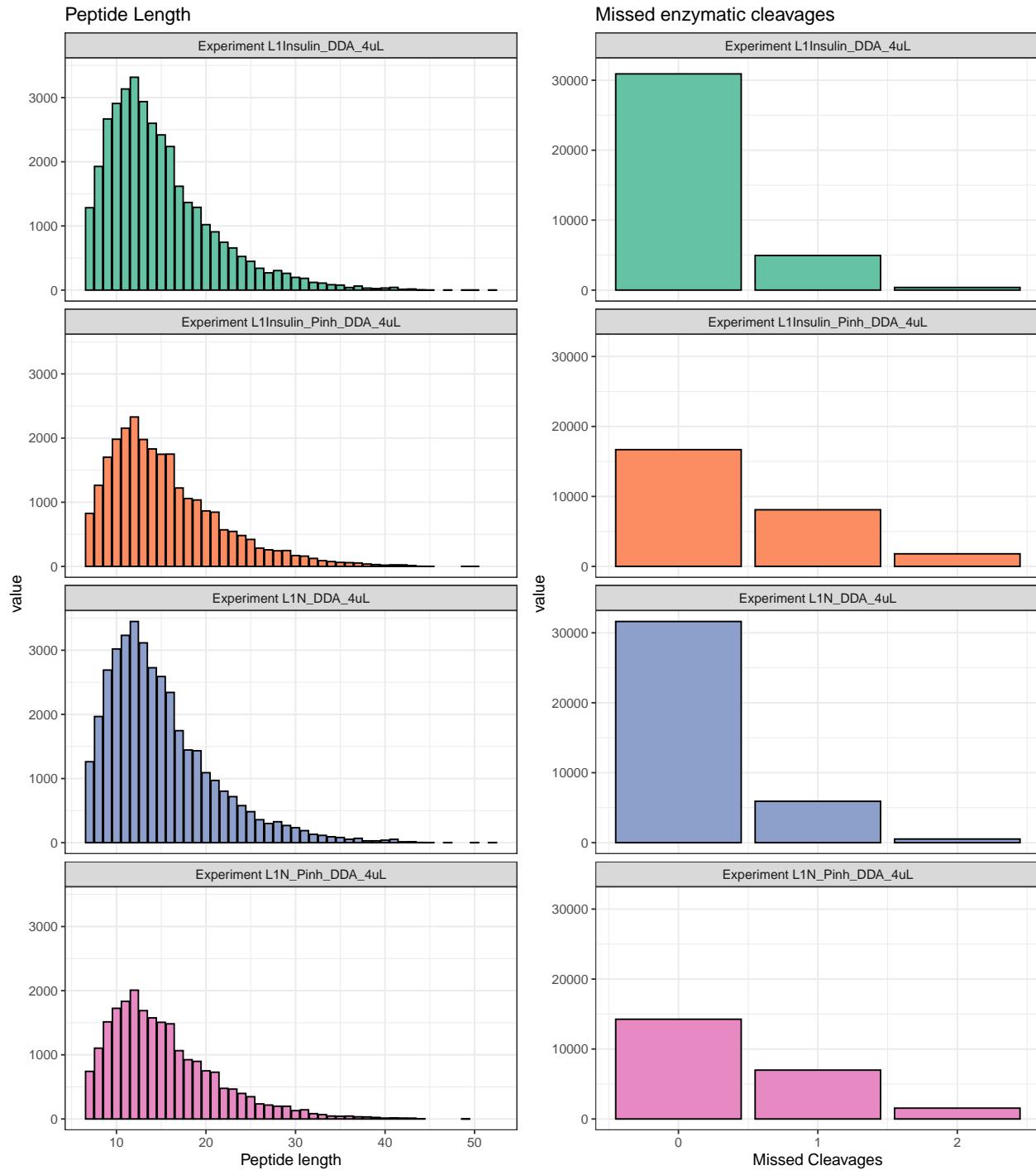
Isotope Patterns detected and sequenced



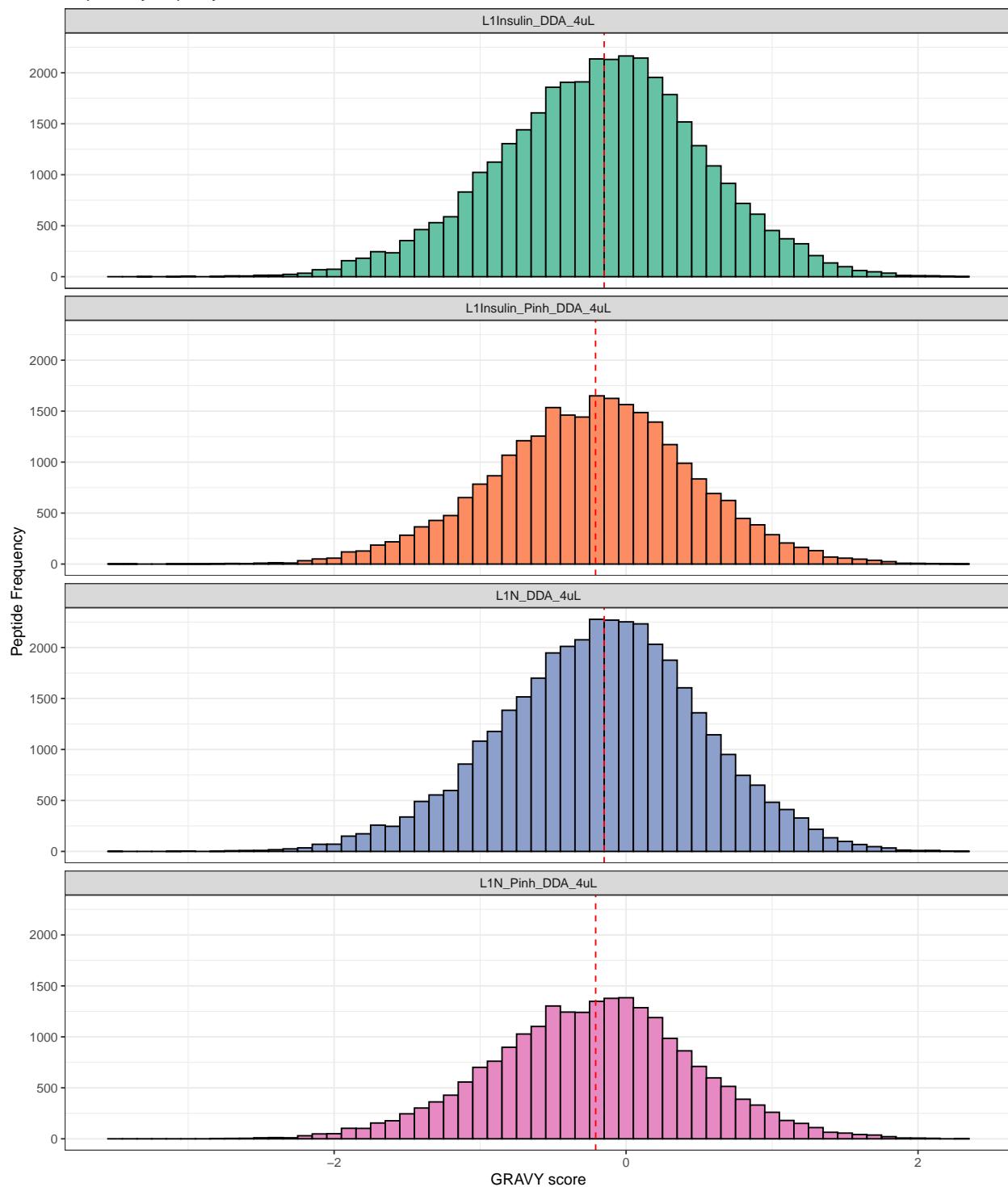
The charge-state of the precursor ion.

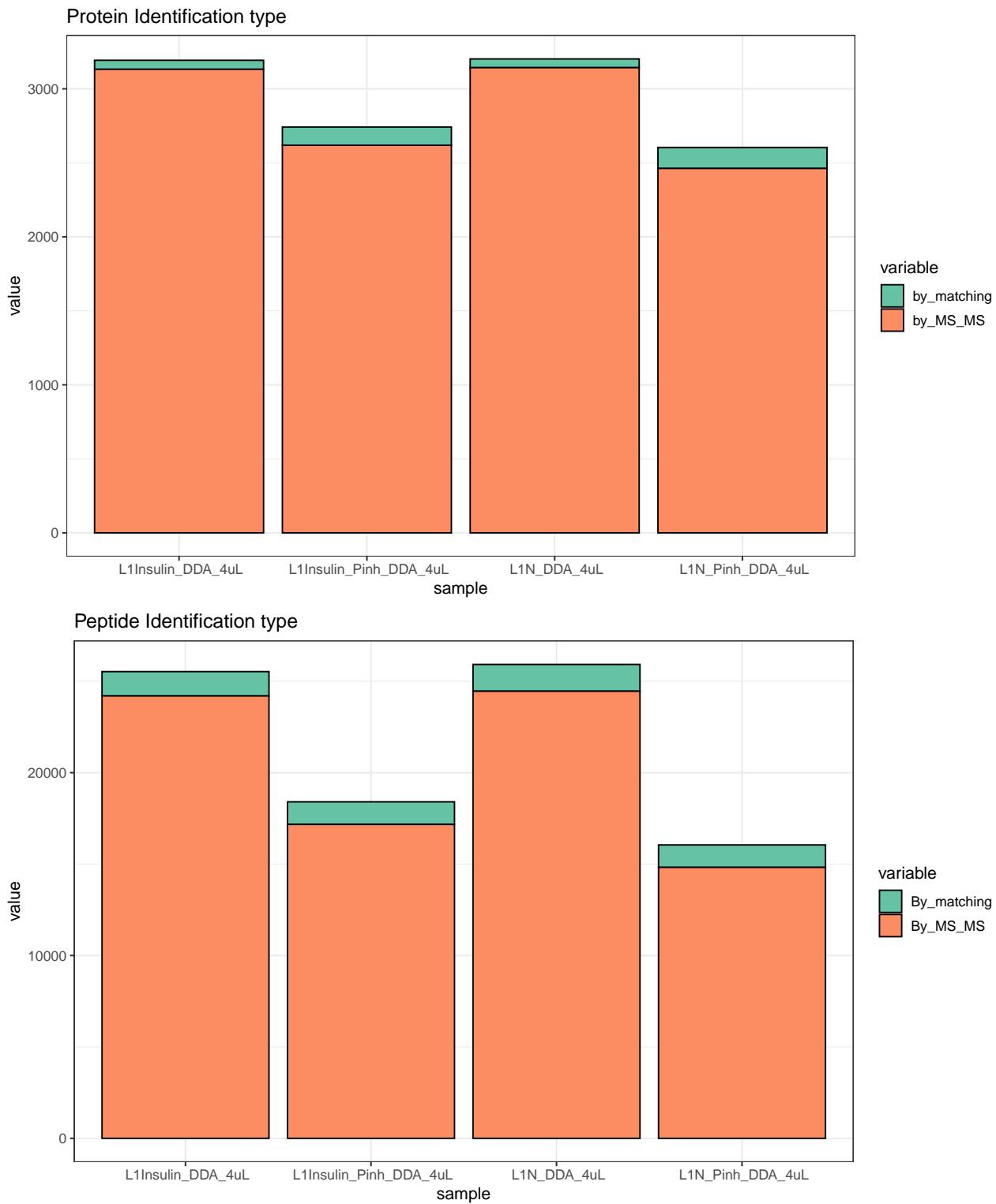


Protease Specificity

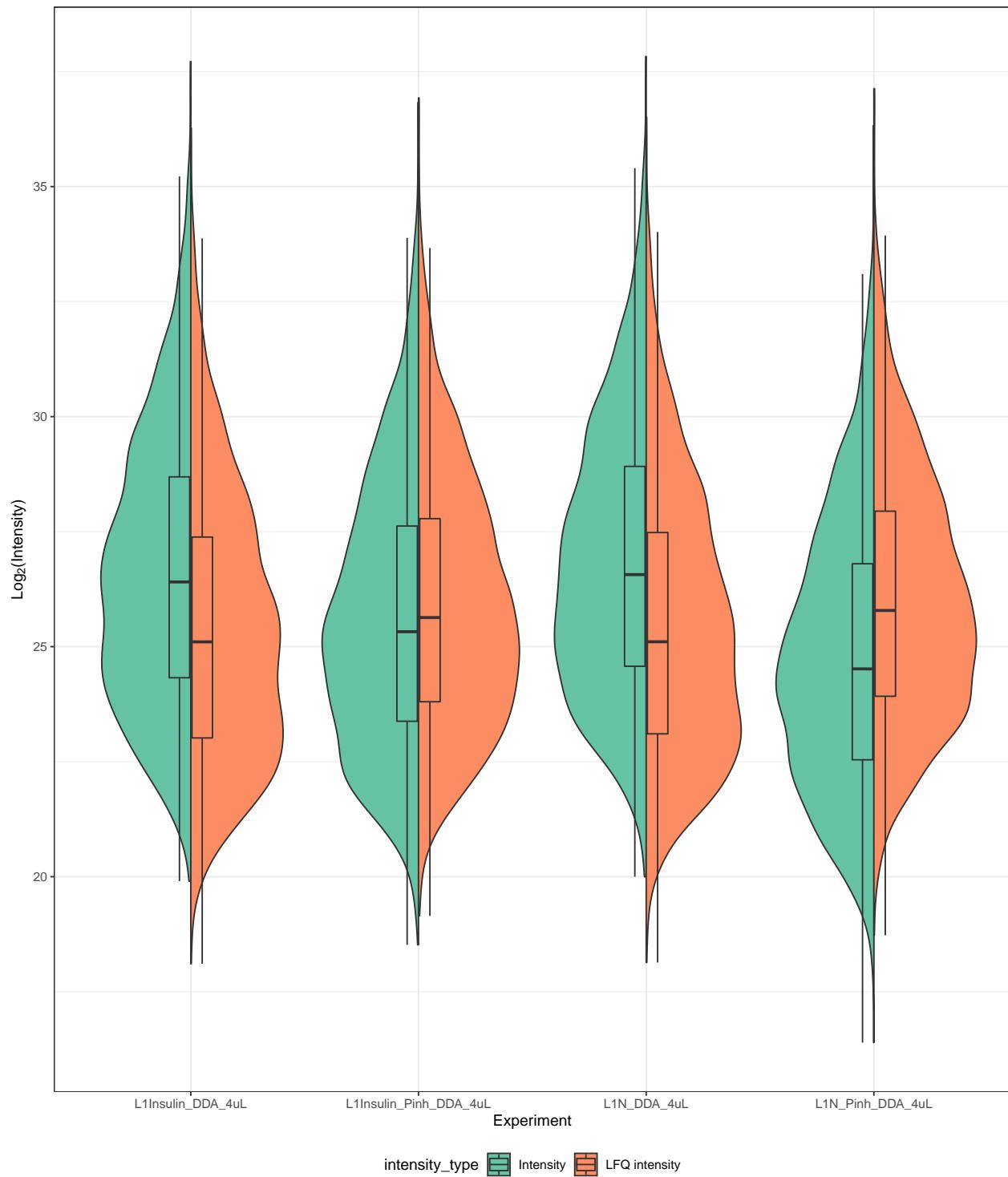


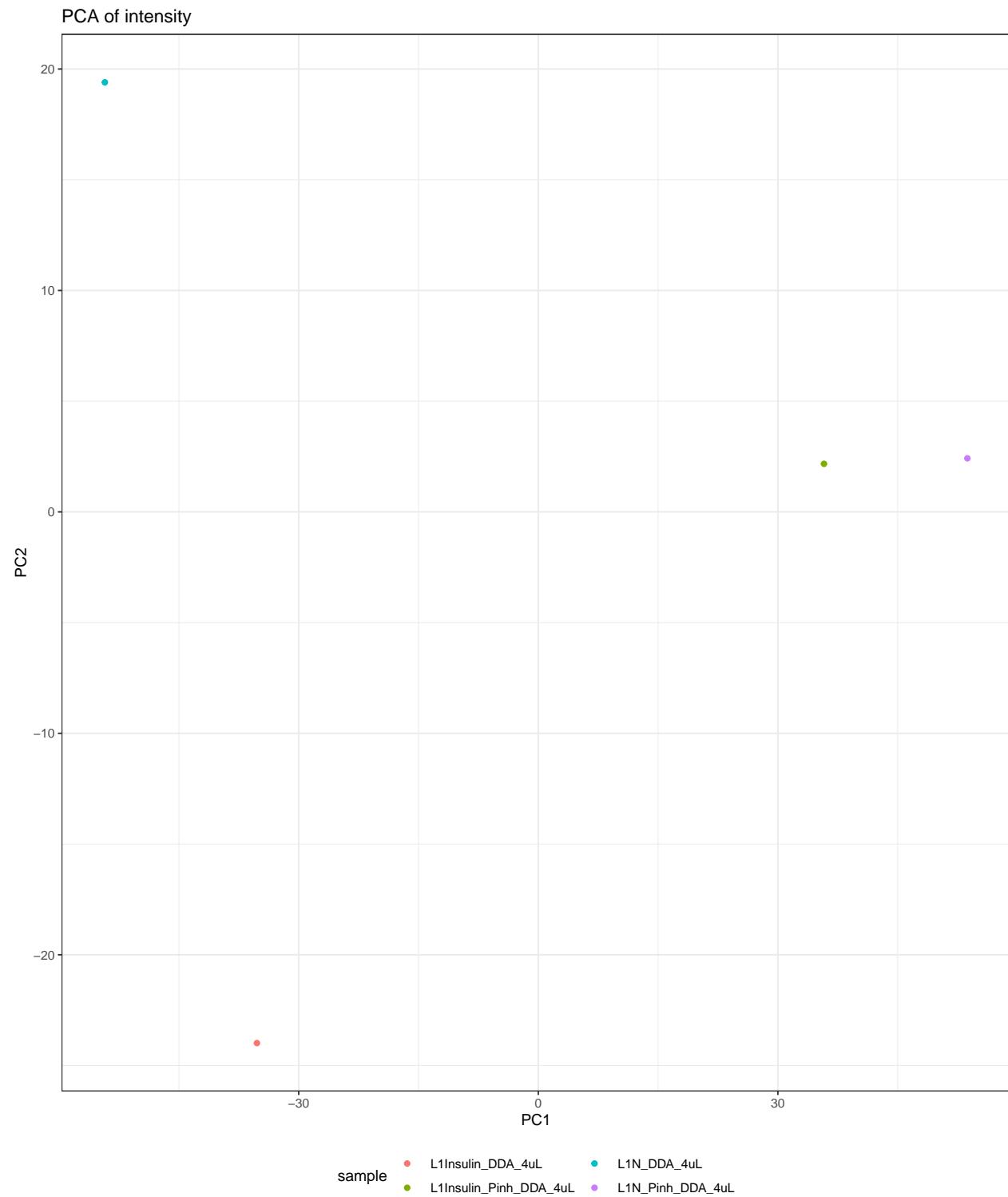
Peptide hydrophathy distribution



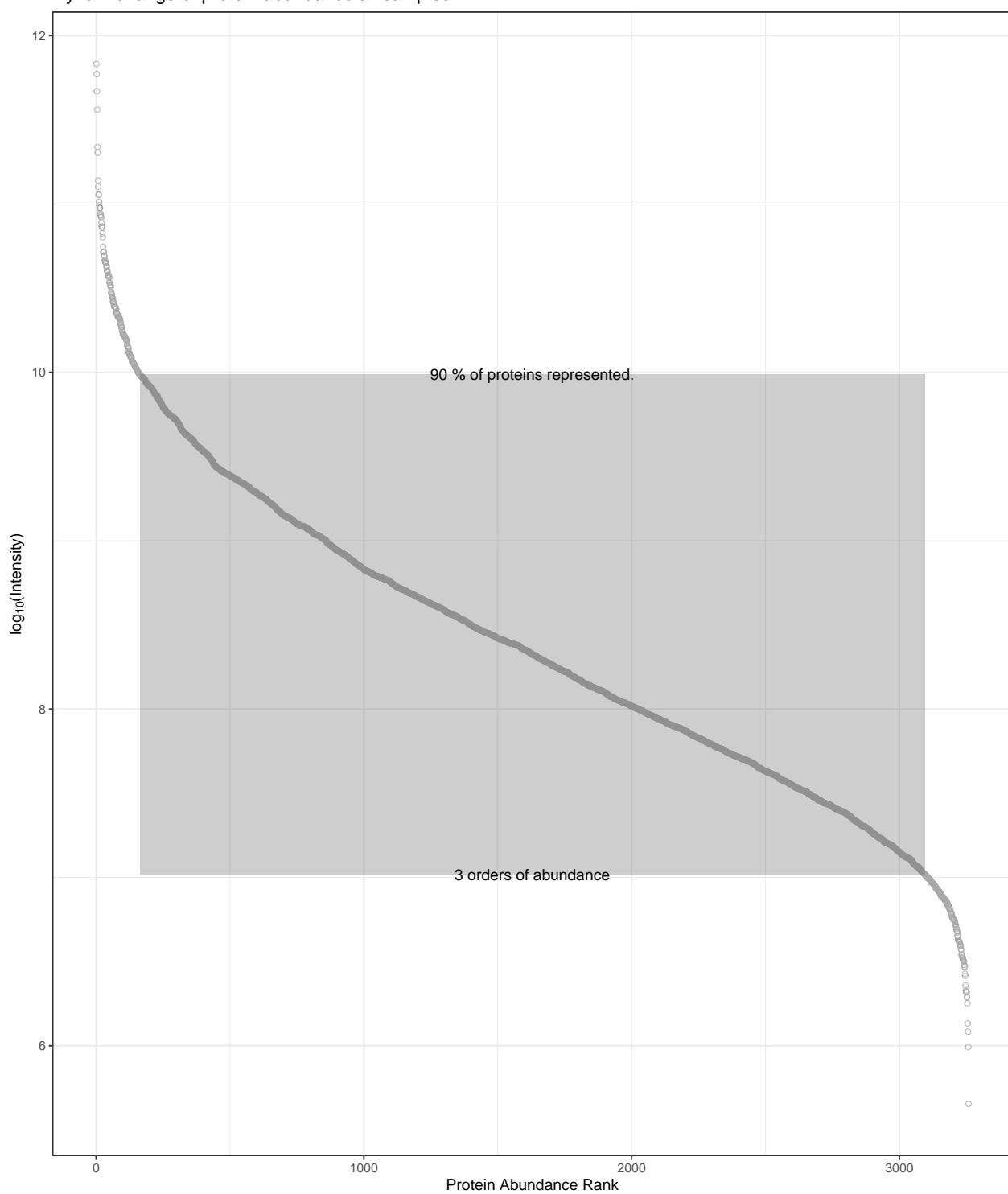


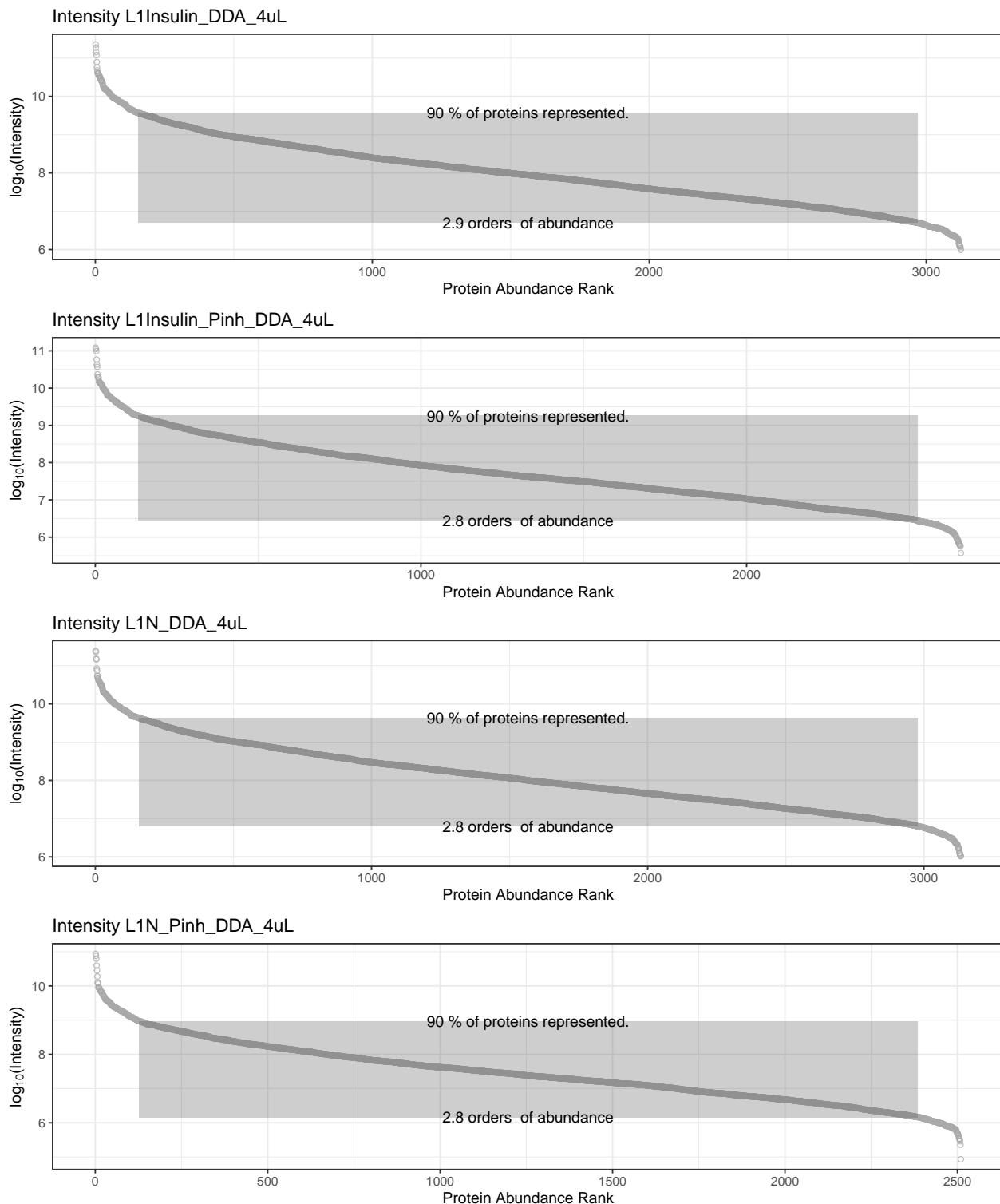
Protein Intensity & LFQ intensity





Dynamic range of protein abundance all samples

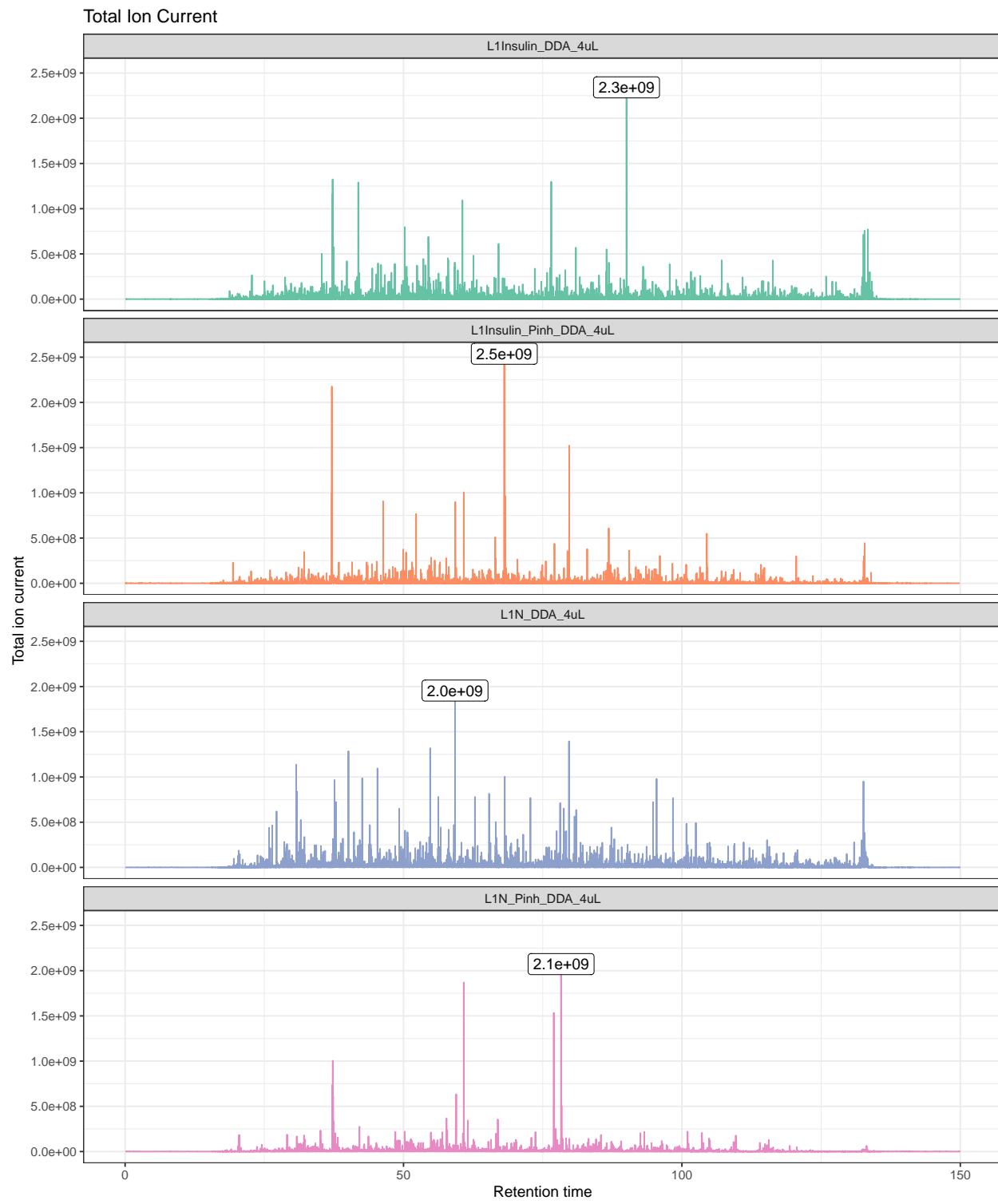




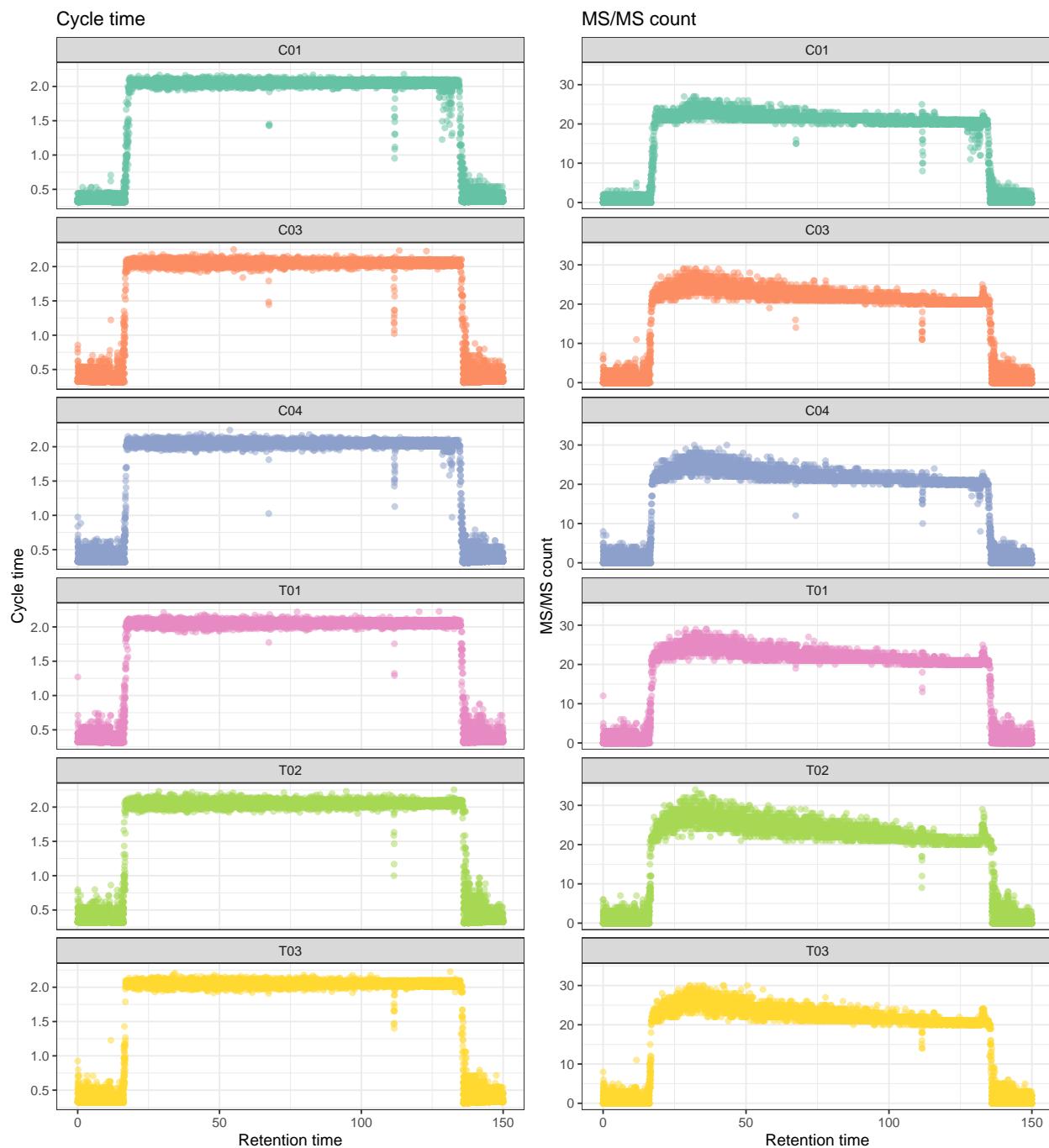
```
## [1] "No UniprotID provided."
```

```
## [1] "No iRT peptides found in the MaxQuant output."
```

```
## [1] "No iRT peptides found in the MaxQuant output."
```



Acquisition Cycle



Post-Translational Modifications

