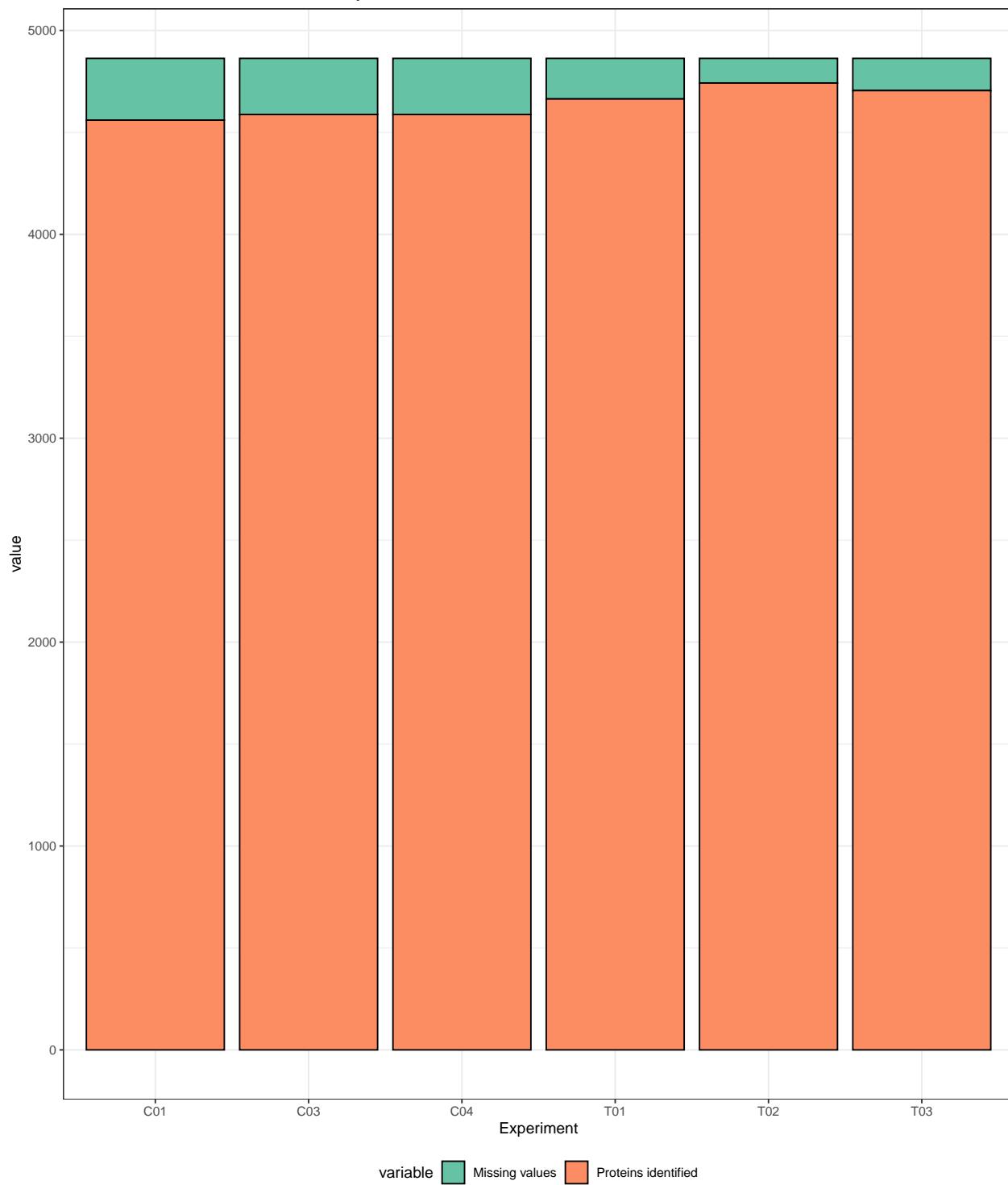


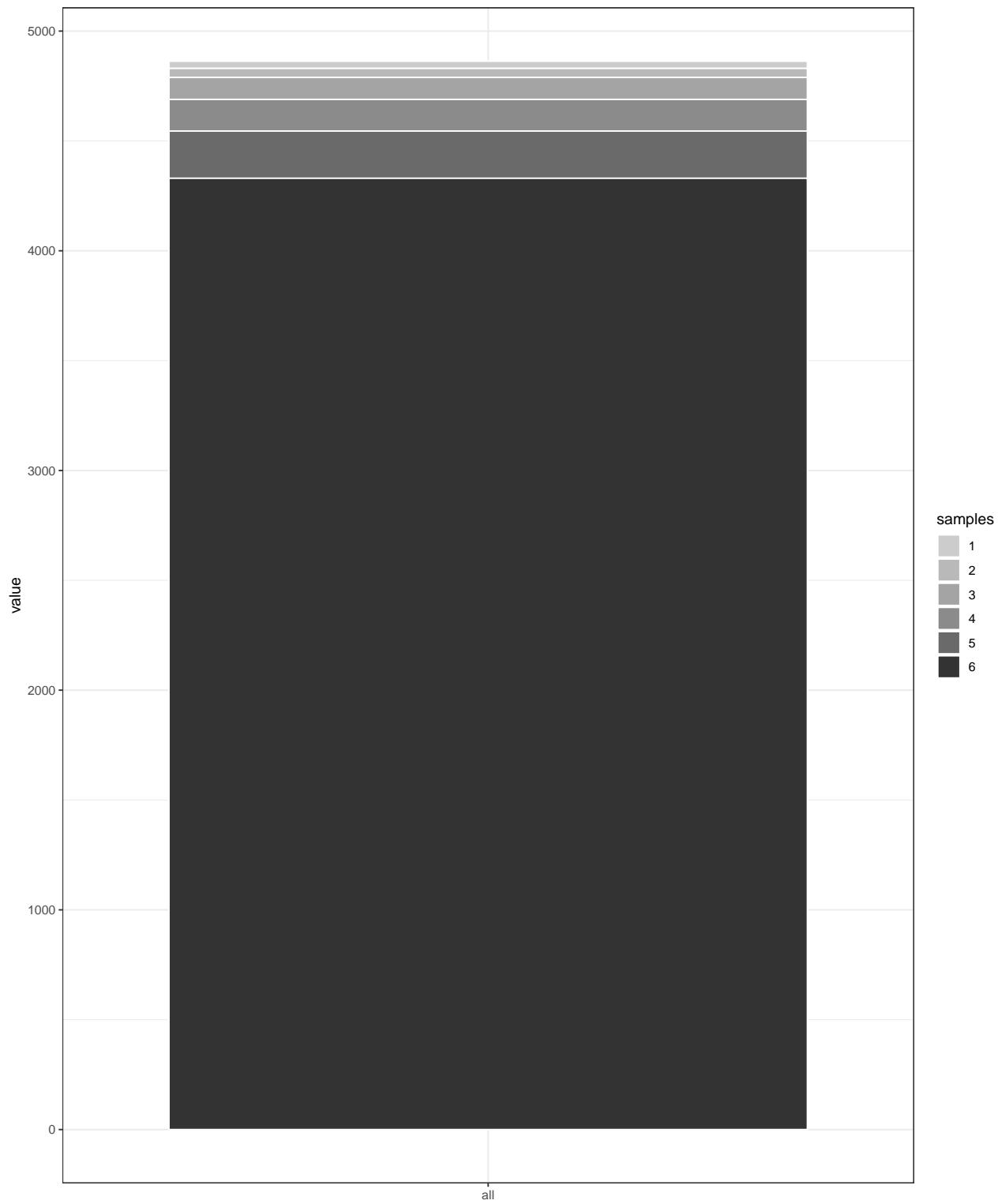
MQmetrics Report

31 March, 2021

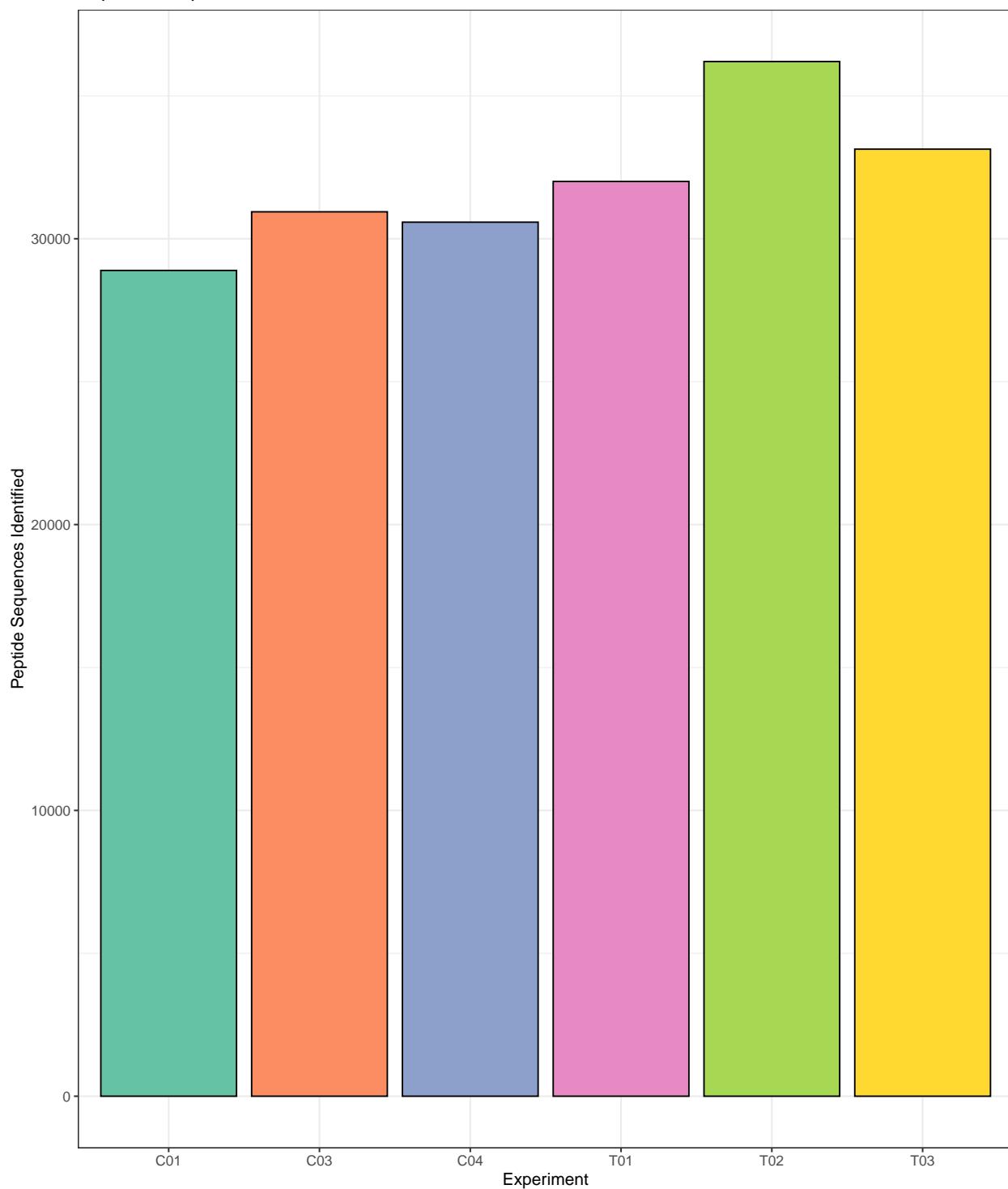
```
## [1] "The experiment started the day: 17/12/2020 at the time: 01:18:03."
## [1] "The whole experiment lasted: 01:19 (hours:minutes)."
## [1] "The MaxQuant version used was: 1.6.12.0"
## [1] "The user was: marek.vrbacky"
## [1] "The machine name was: FGU045PC004"
## [1] "The protein FDR was: 0.01"
## [1] "The match between runs was: True"
## [1] "The fasta file used was: C:\\MaxQuant_Databases\\\\UP000005640_9606.fasta"
```

Proteins Identified based on Intensity

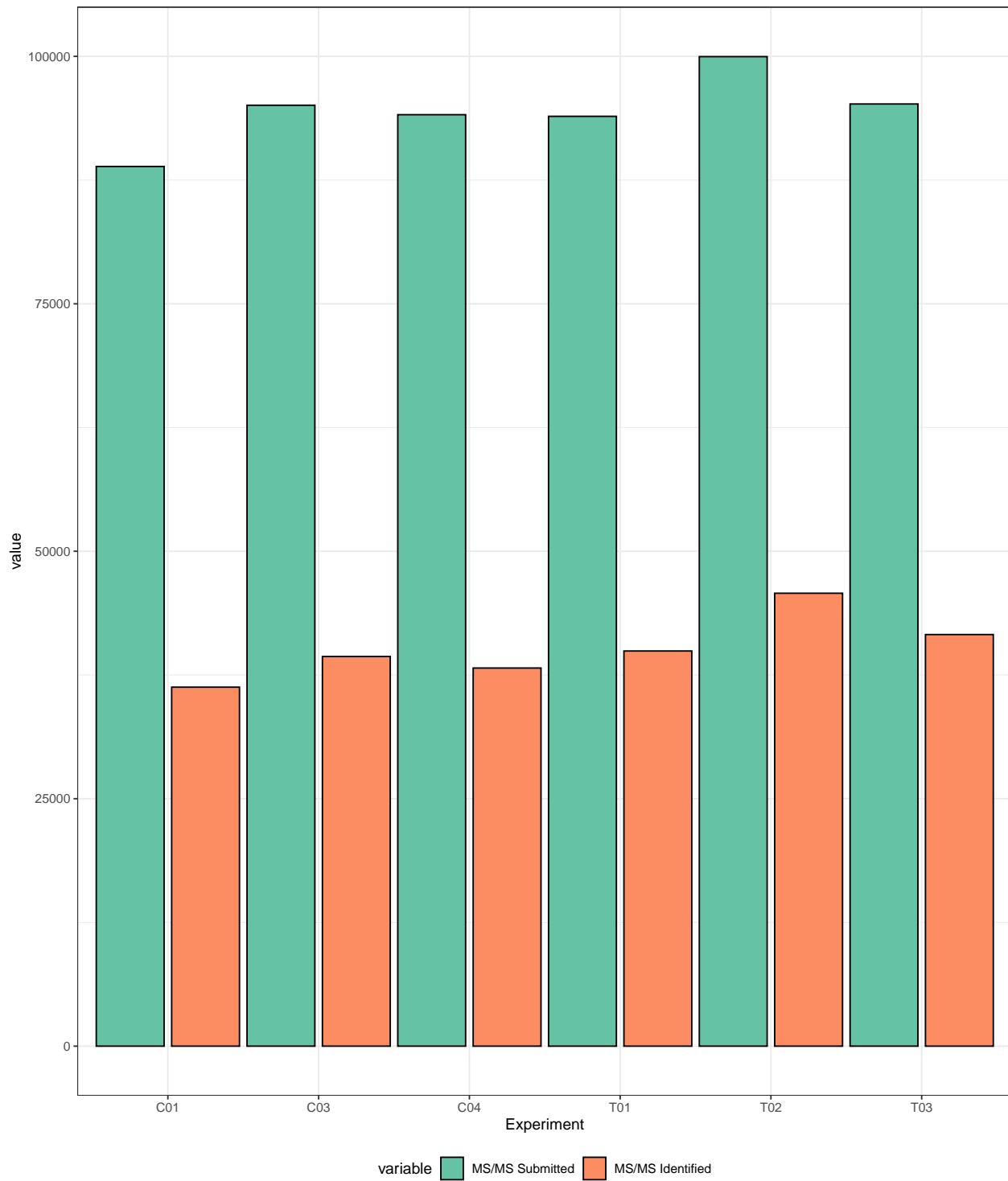




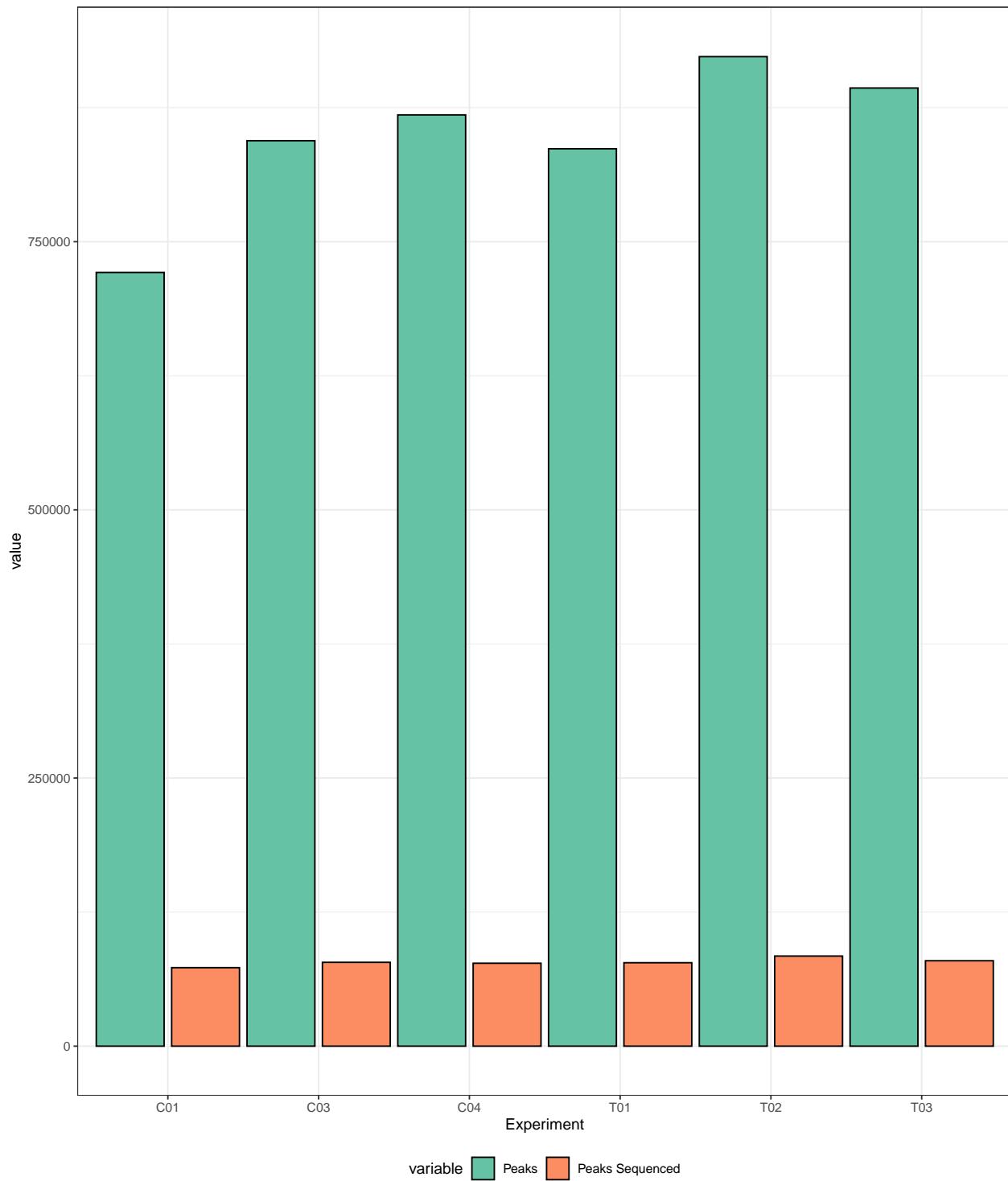
Peptides Sequences Identified

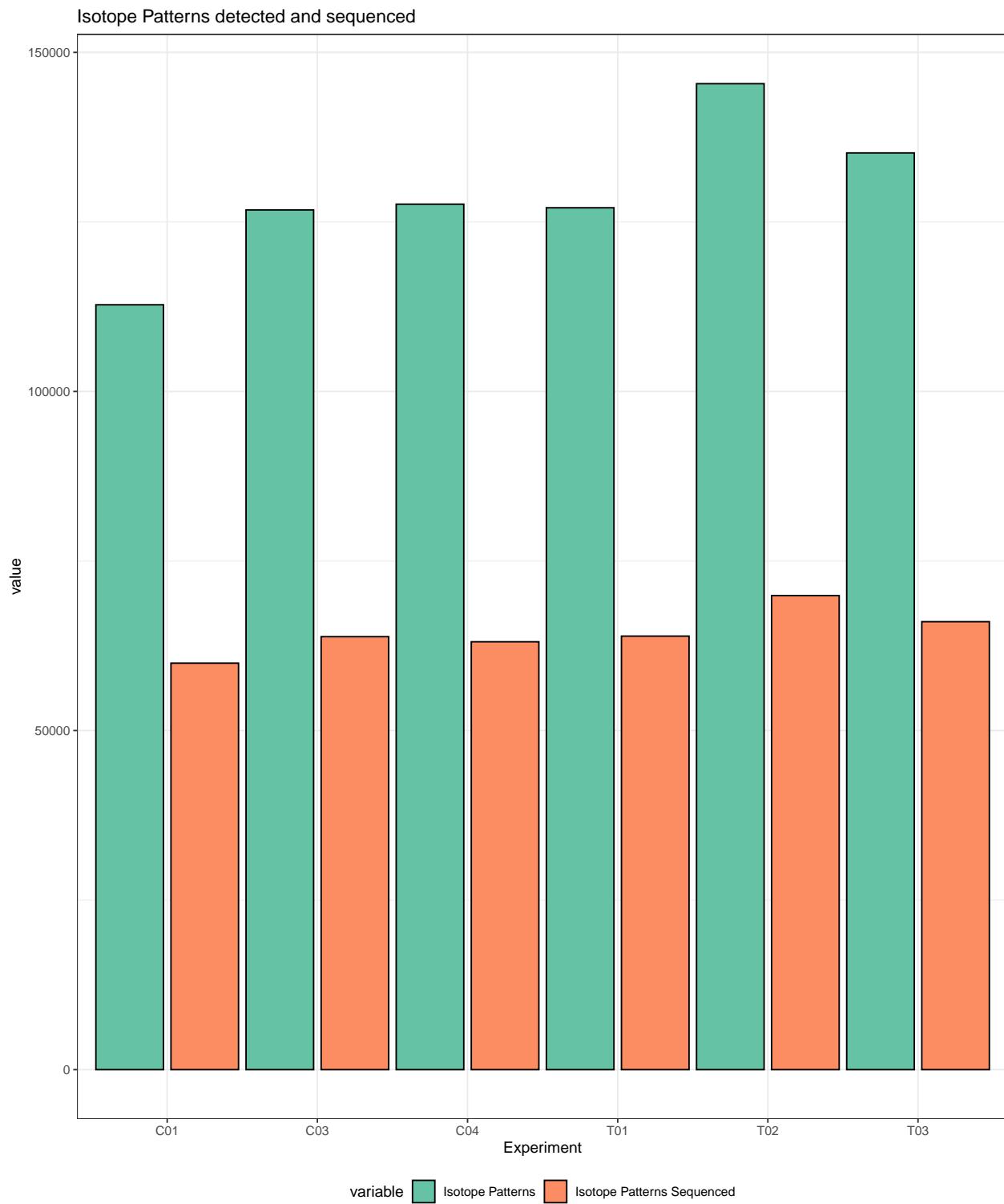


MS/MS Submitted and Identified

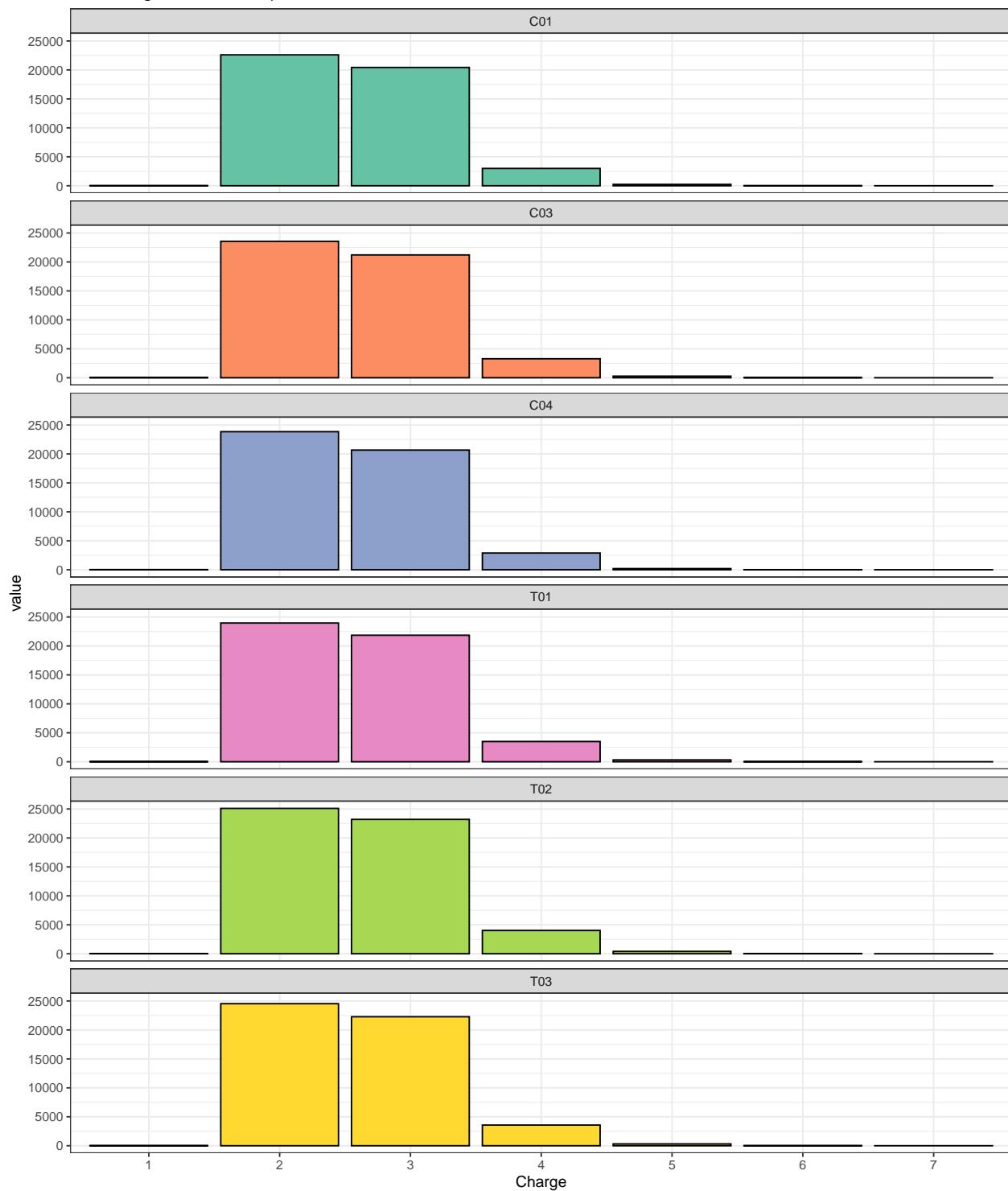


Peaks detected and sequenced in the full scans

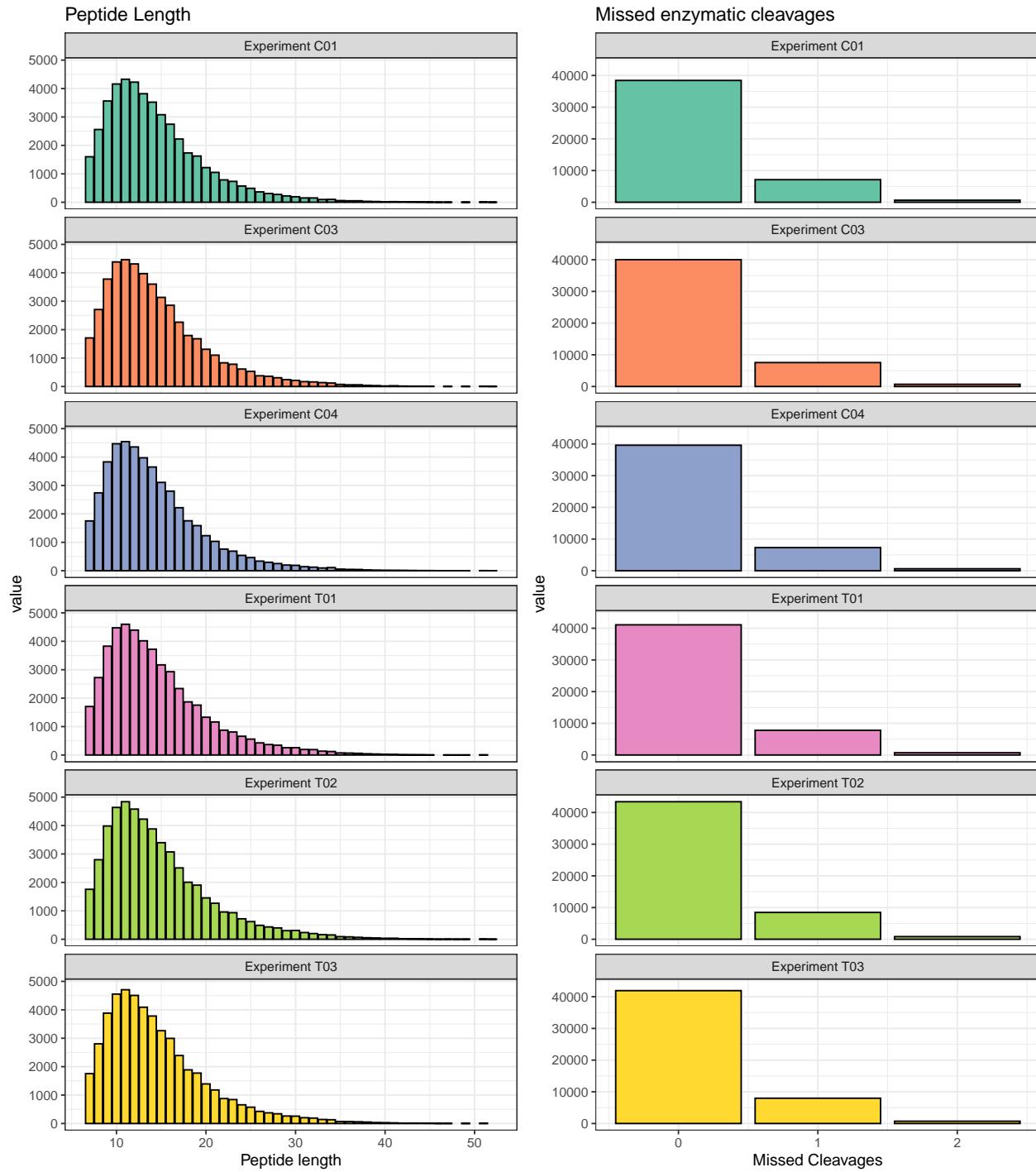


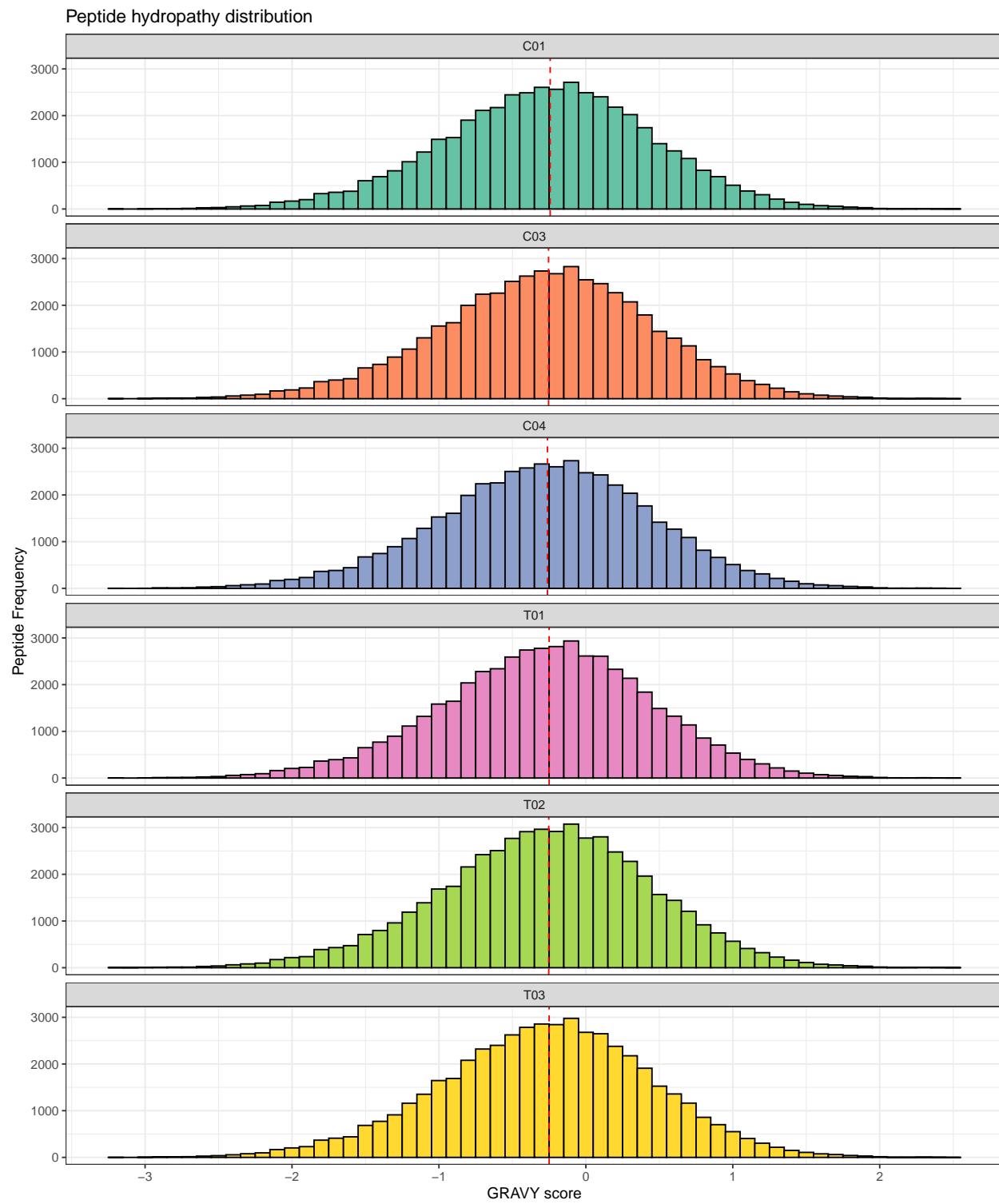


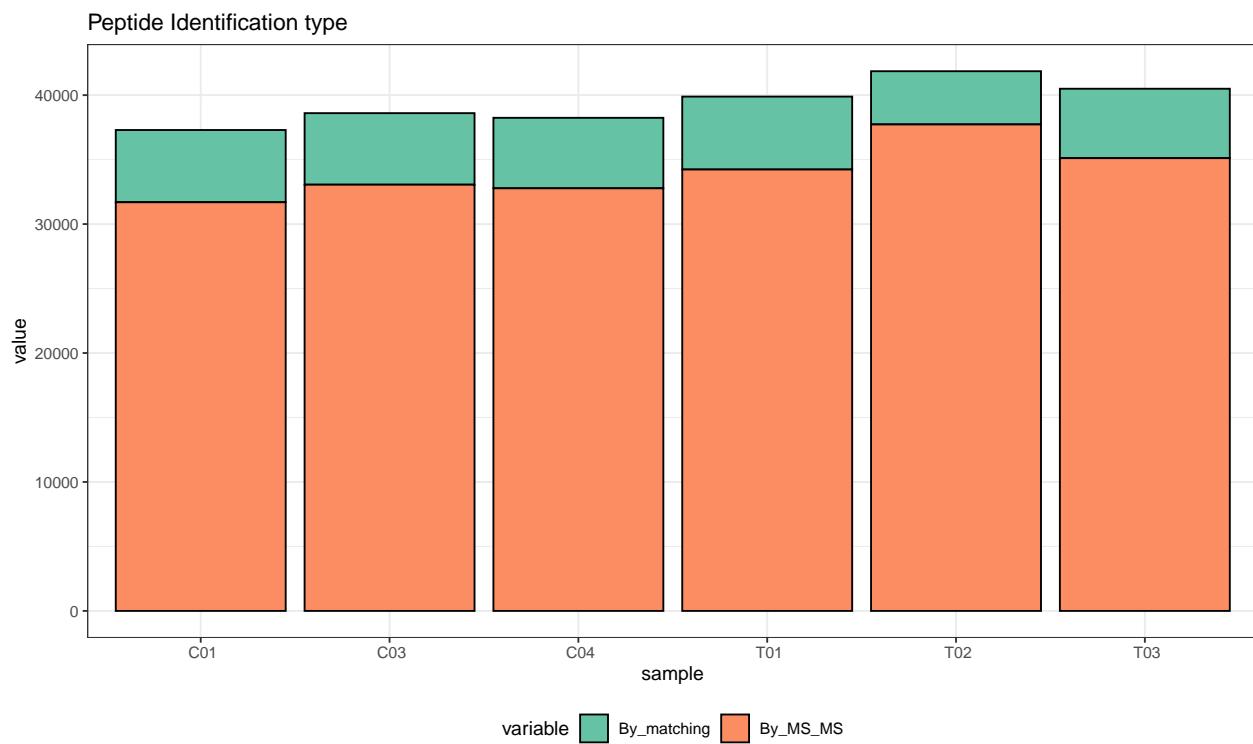
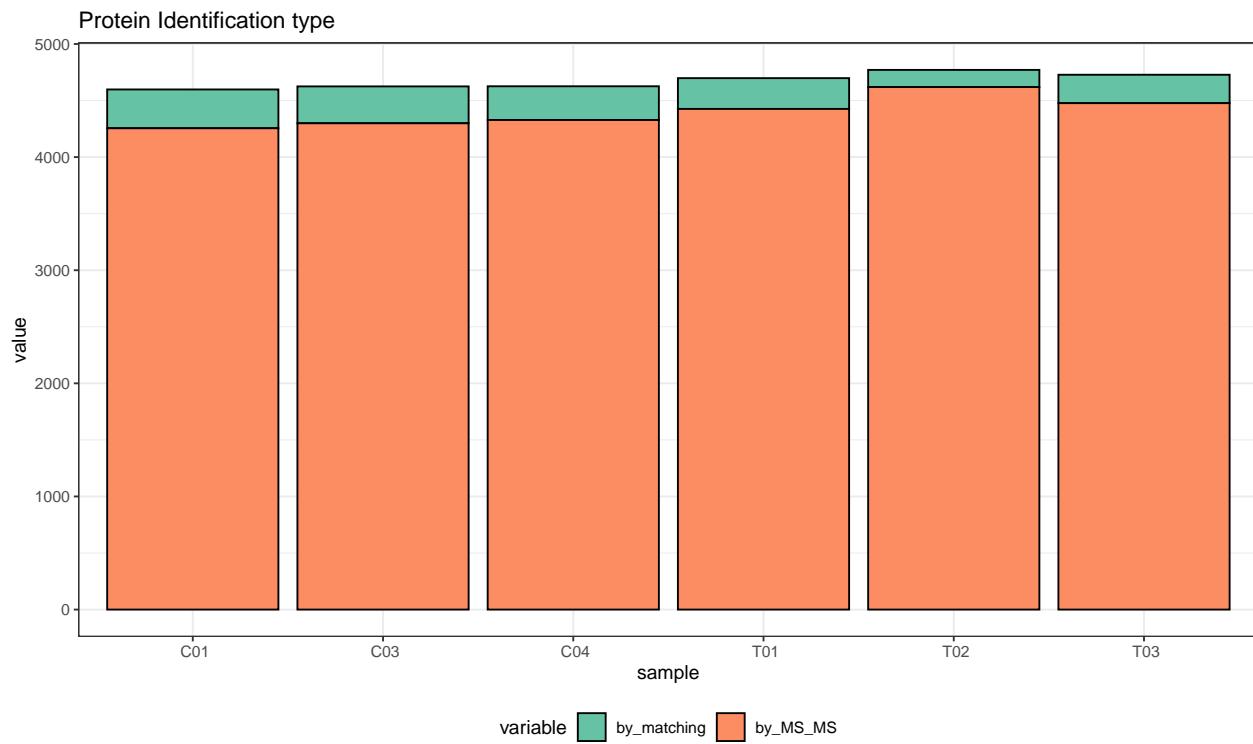
The charge-state of the precursor ion.



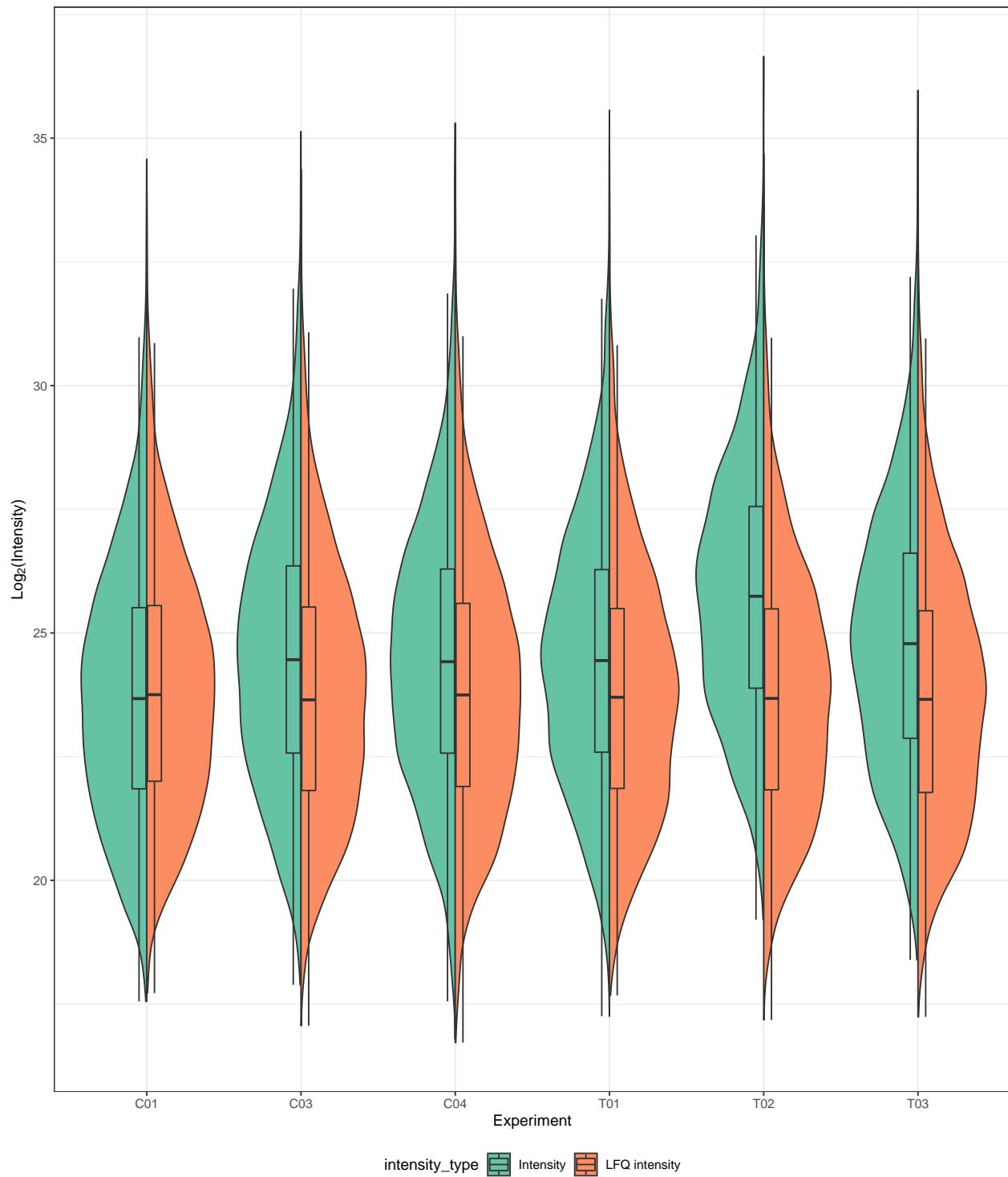
Protease Specificity



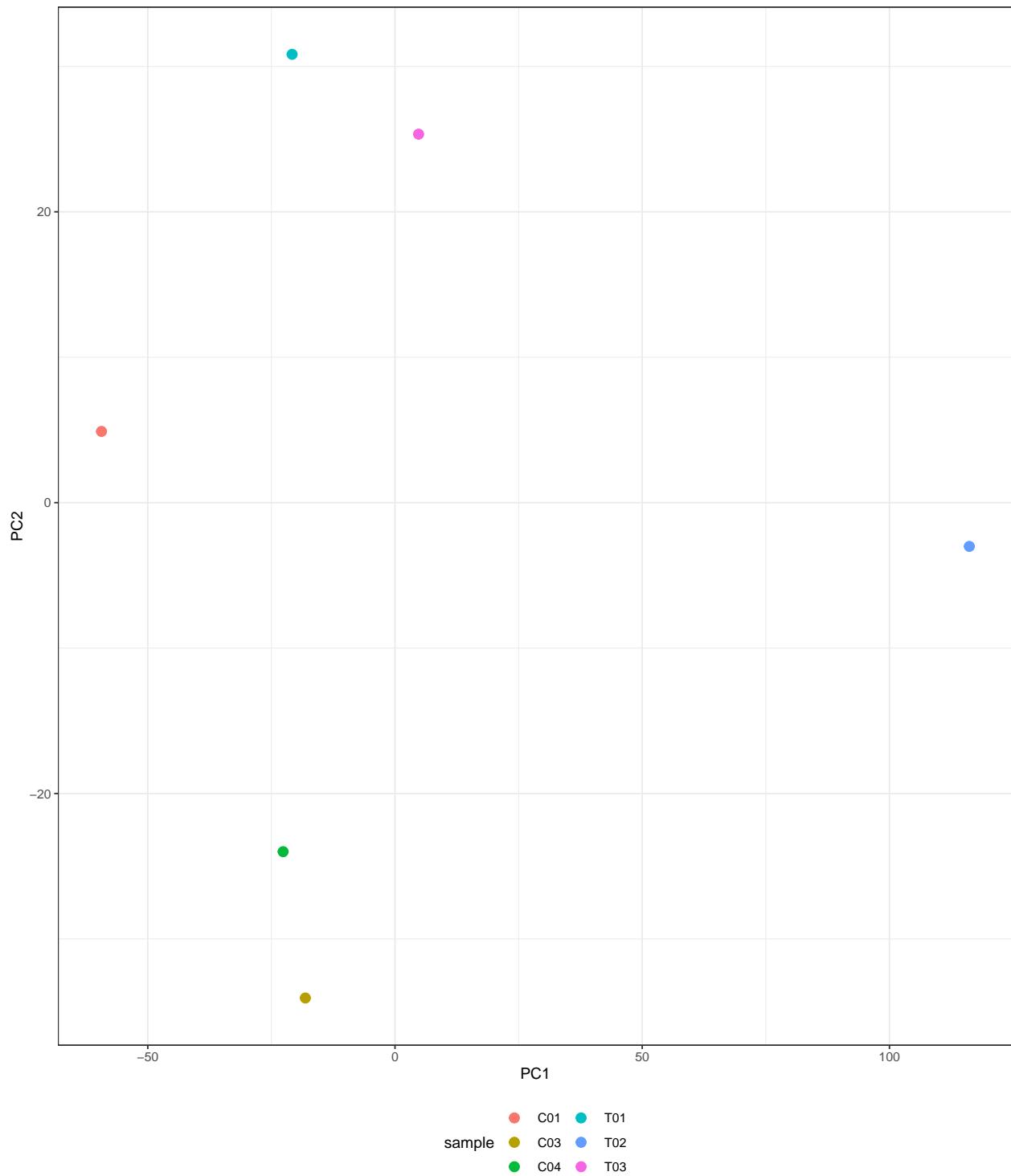




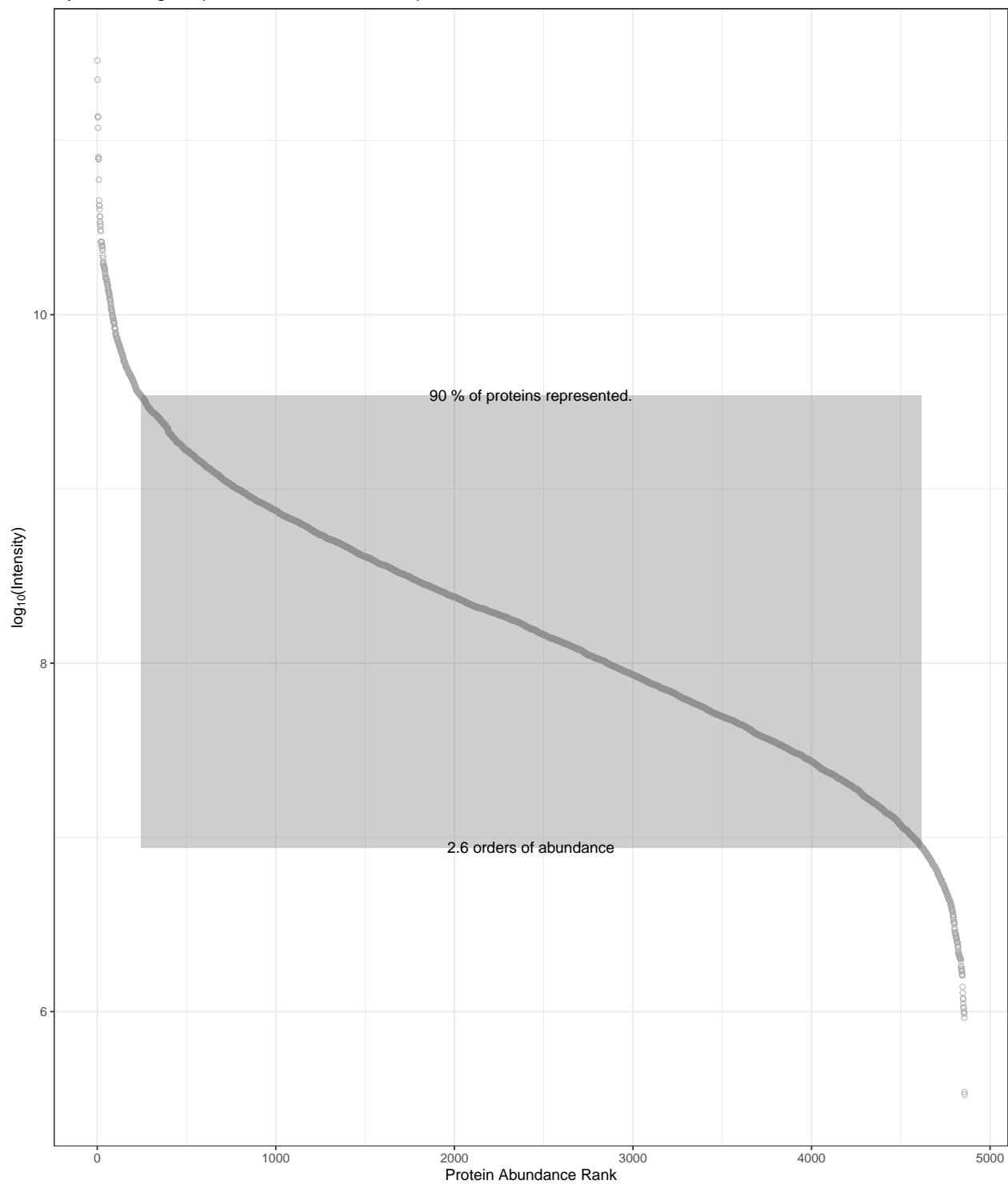
Protein Intensity & LFQ intensity

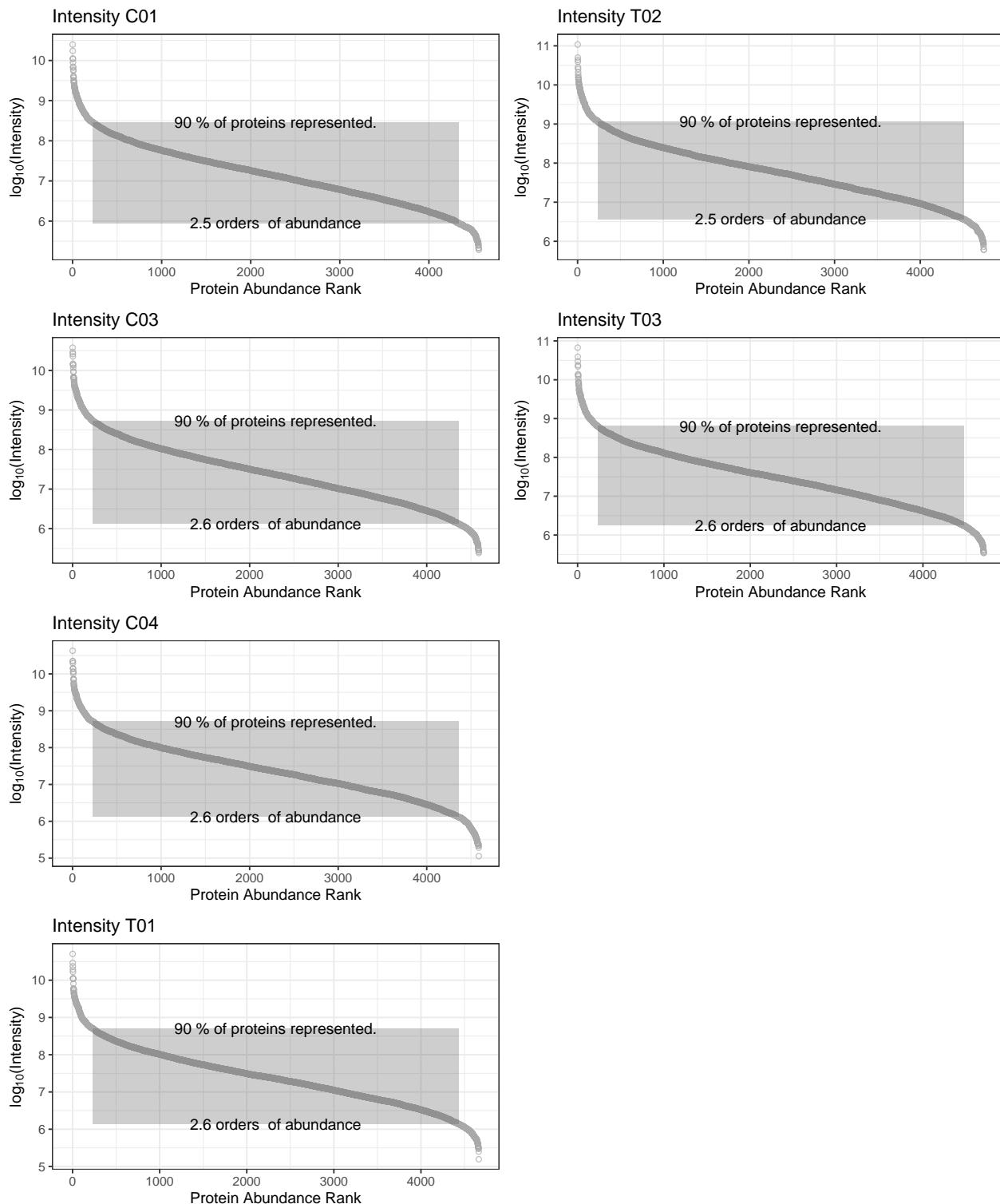


PCA of intensity

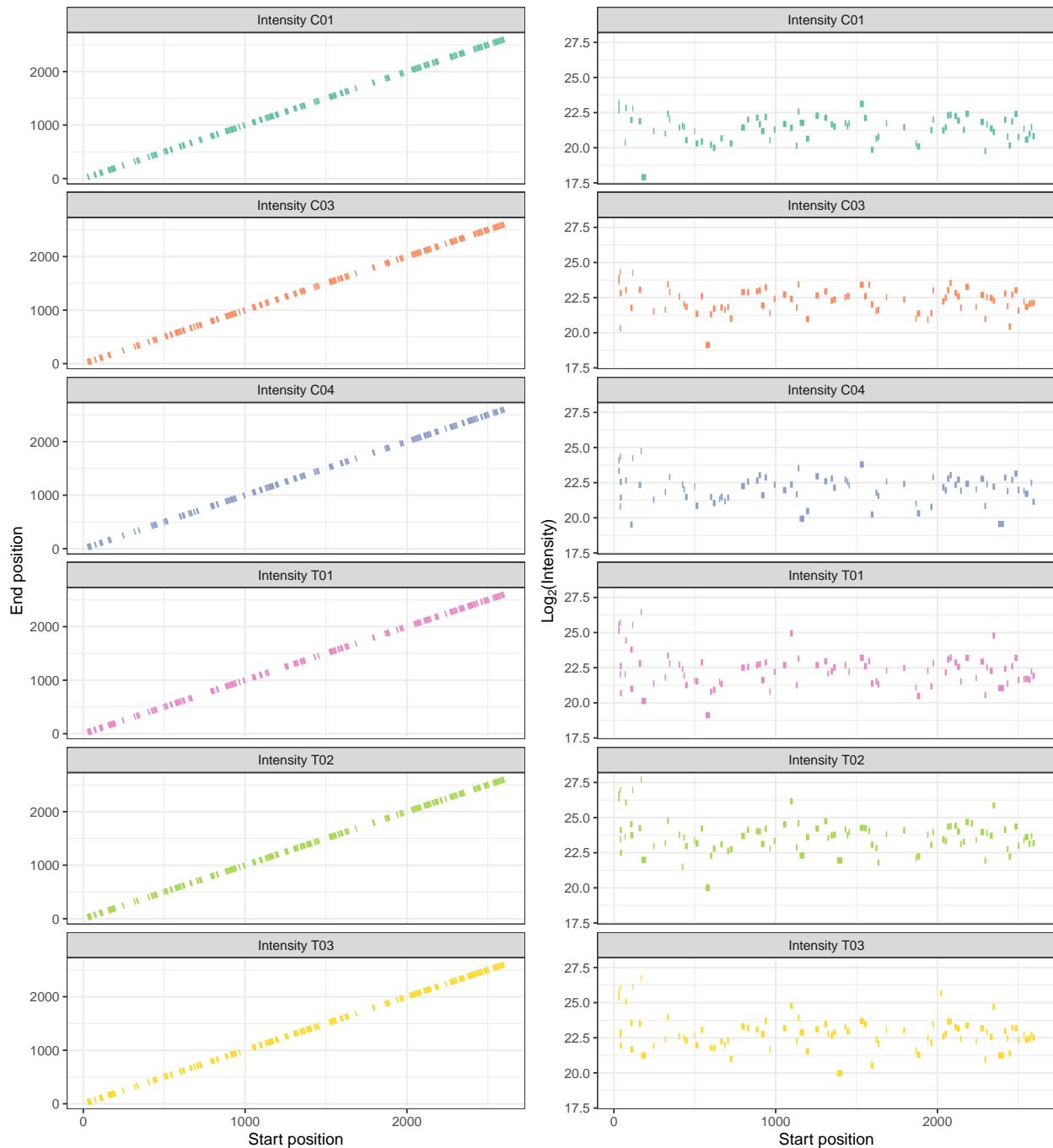


Dynamic range of protein abundance all samples



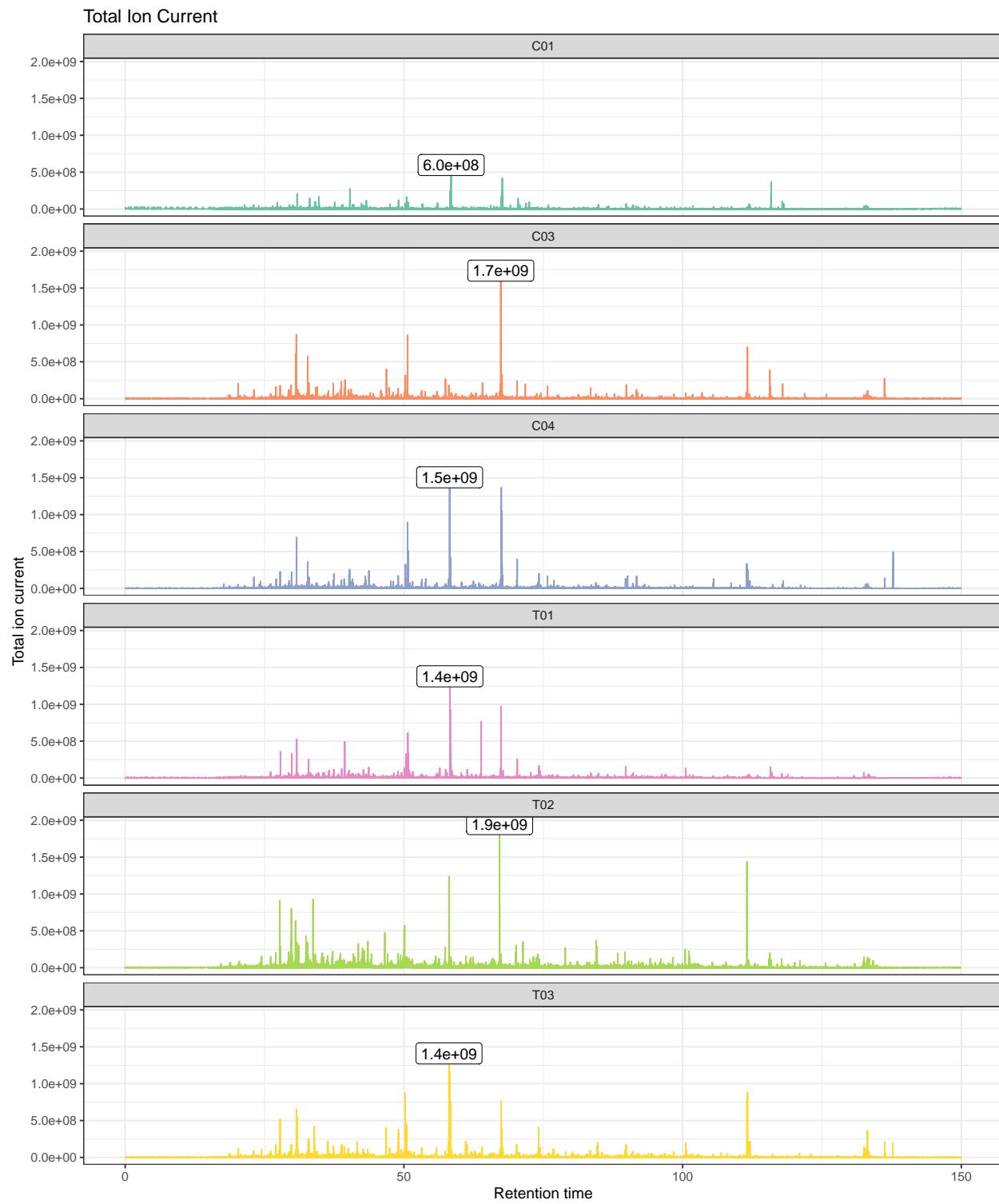


The Protein Coverage of: O75369 (2602 amino acids), Gene: FLNB
is: 54.8%

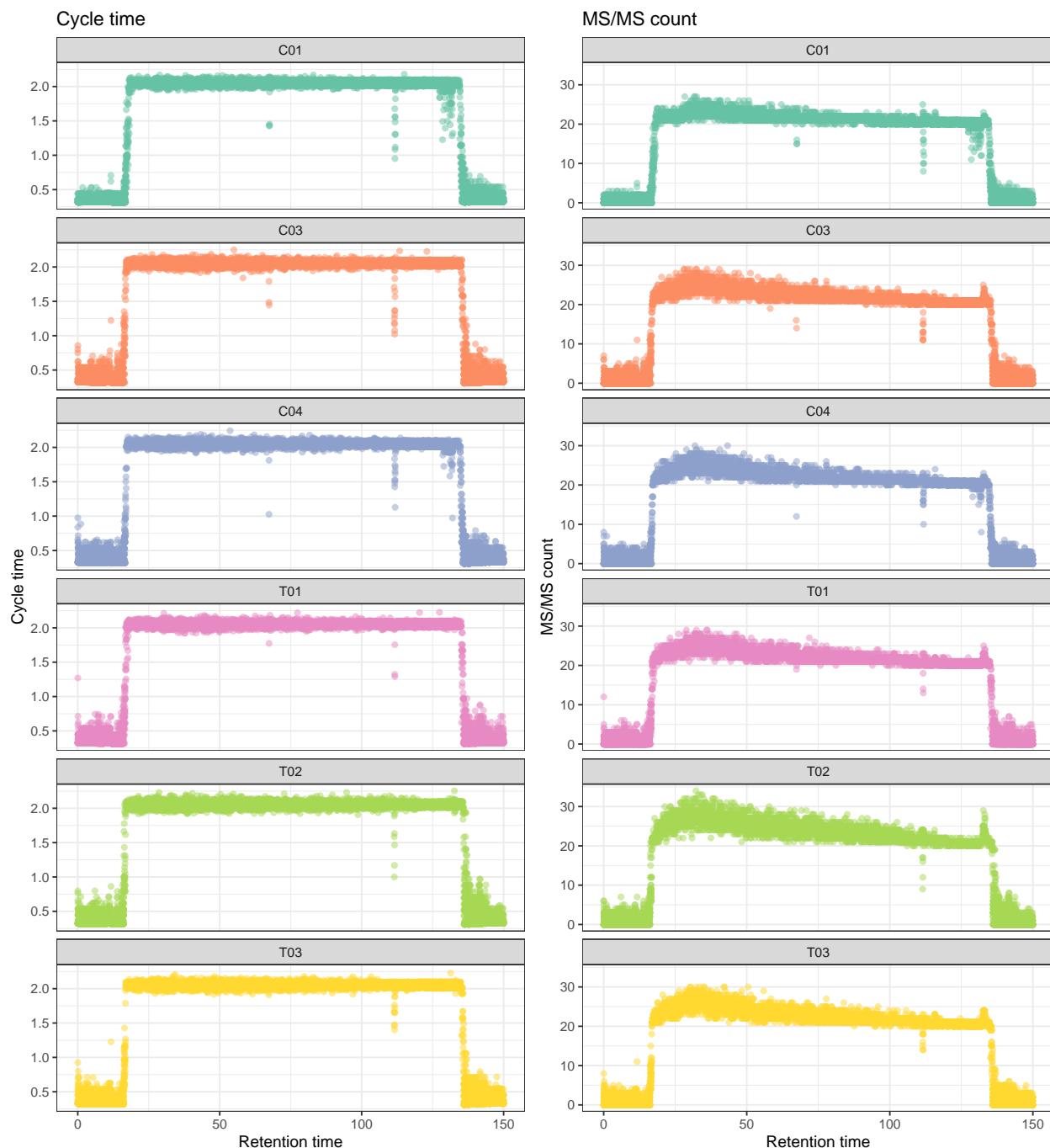


```
## [1] "No iRT peptides found in the MaxQuant output."
```

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



Acquisition Cycle



Post–Translational Modifications



Modifications 2 Oxidation (M) Acetyl (Protein N-term);2 Oxidation (M) Unmodified
 Acetyl (Protein N-term) Oxidation (M)