

Summarize Peptide Level Measurements

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2018-09-15

Summarize levels

Table 1: summary

NR.Isotope.Label	NR.protein_Id	NR.peptide_Id	NR.precursor_Id	NR.fragment_Id
heavy	12	34	34	163
light	13	45	45	218

Table 2: Protein Summaries

protein_Id	peptide_Id_n_distinct	precursor_Id_n_distinct	fragment_Id_n_distinct	nrpeptide
iRT_Protein	11	11	55	1
sp O95477 ABCA1_HUMAN	3	3	14	3
sp P02786 TFR1_HUMAN	3	3	15	3
sp P04406 G3P_HUMAN	3	3	15	3
sp P08195 4F2_HUMAN	3	3	15	3
sp P49281 NRAM2_HUMAN	3	3	13	3
sp P63104 1433Z_HUMAN	2	2	10	3
sp Q01650 LAT1_HUMAN	3	3	15	3
sp Q15043 S39AE_HUMAN	3	3	15	3
sp Q15365 PCBP1_HUMAN	3	3	15	3
sp Q9C0K1 S39A8_HUMAN	2	2	10	3
sp Q9NP59 S40A1_HUMAN	3	3	11	3
sp Q9UHI5 LAT2_HUMAN	3	3	15	3

```
## Warning in self$trans$transform(x): NaNs produced
```

```
## Warning: Transformation introduced infinite values in continuous x-axis
```

```
## Warning: Removed 106 rows containing non-finite values (stat_bin).
```

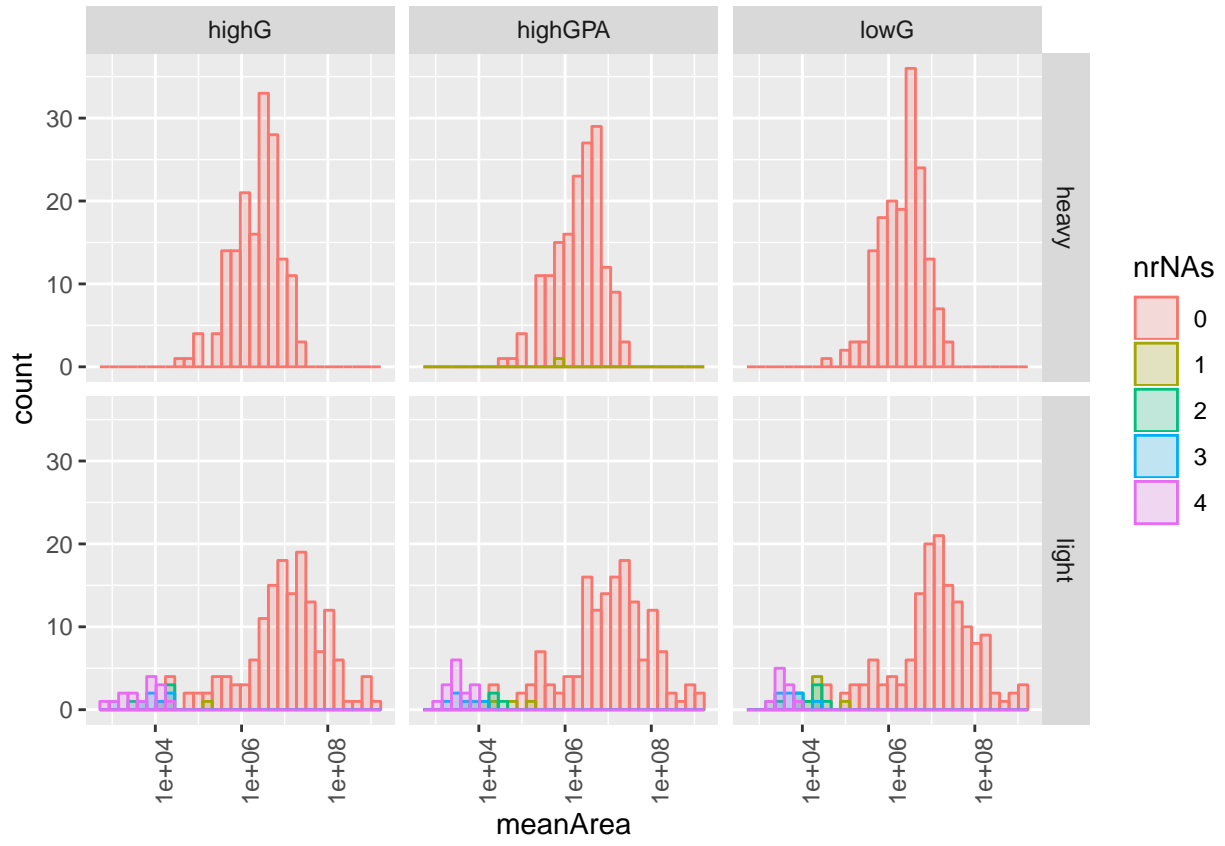


Figure 1: histogram of mean condition intensities per transition

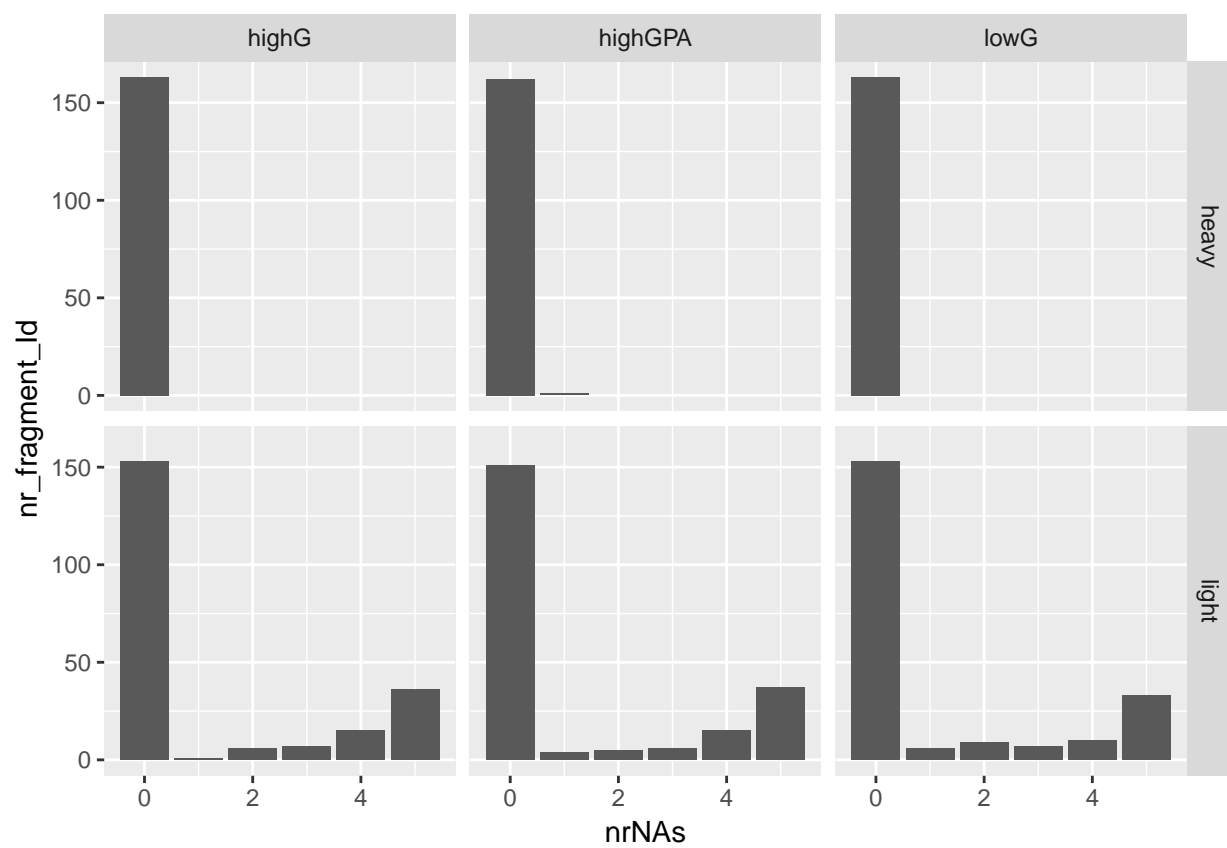


Figure 2: nr of missing

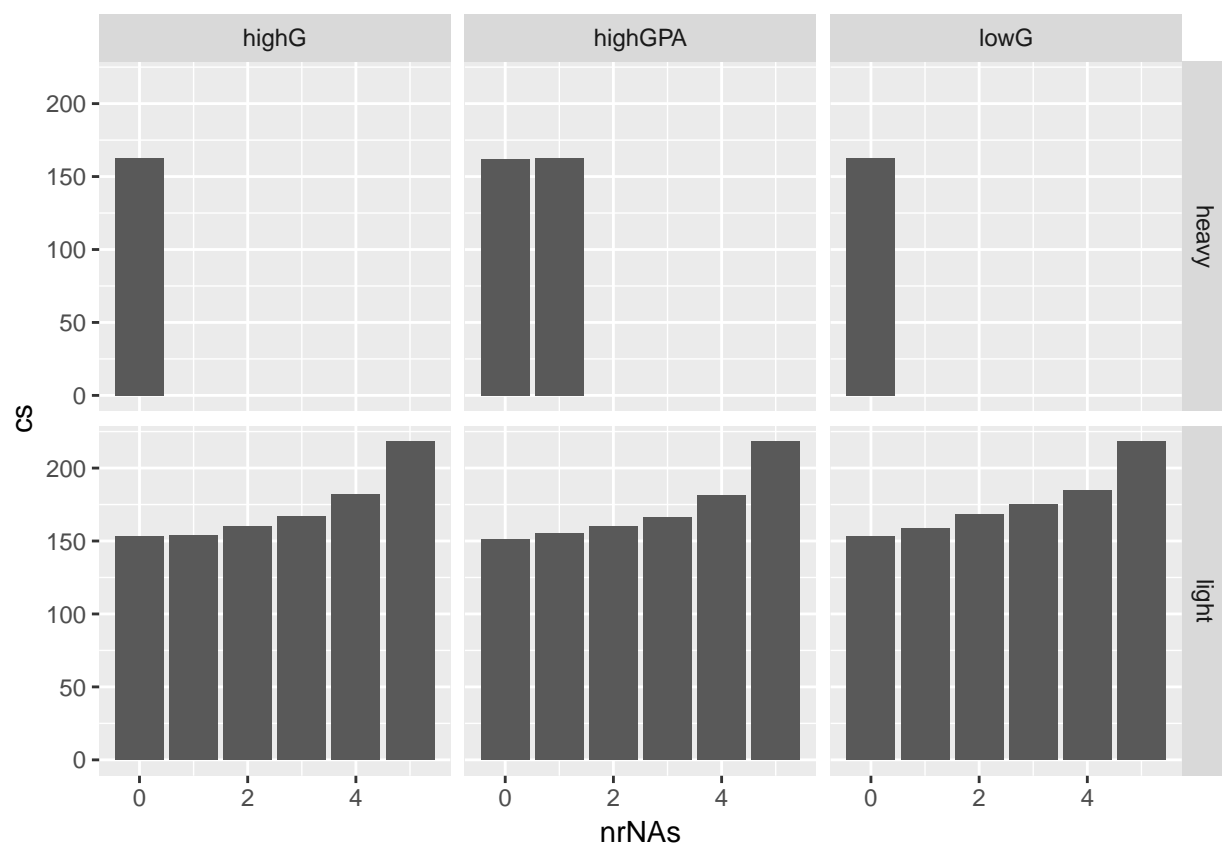


Figure 3: cumulative sum of missing