Summarize Peptide Level Measurements

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Summarize levels

Table 1: summary

NR.Isotope.Label	NR.protein_Id	$NR.peptide_Id$	NR.precursor_Id	$NR.fragment_Id$
L/H	13	45	45	218

[1] 4

Table 2: Protein Summaries

protein_Id	peptide_Id	precursor_Id	fragment_Id	nrpeptides	Precursors	Fragments
iRT_Protein	11	11	55	11	1-1	5-5
sp O95477 ABCA1_HUMAN	3	3	14	3	1-1	4-5
sp P02786 TFR1_HUMAN	3	3	15	3	1-1	5-5
$sp P04406 G3P_HUMAN$	3	3	15	3	1-1	5-5
$sp P08195 4F2_HUMAN$	3	3	15	3	1-1	5-5
sp P49281 NRAM2_HUMAN	3	3	13	3	1-1	4-5
sp P63104 1433Z_HUMAN	2	2	10	2	1-1	5-5
$sp Q01650 LAT1_HUMAN$	3	3	15	3	1-1	5-5
$sp Q15043 S39AE_HUMAN$	3	3	15	3	1-1	5-5
sp Q15365 PCBP1_HUMAN	3	3	15	3	1-1	5-5
sp Q9C0K1 S39A8_HUMAN	2	2	10	2	1-1	5-5
$sp Q9NP59 S40A1_HUMAN$	3	3	11	3	1-1	3-5
${ m sp} { m Q9UHI5} { m LAT2_HUMAN}$	3	3	15	3	1-1	5-5

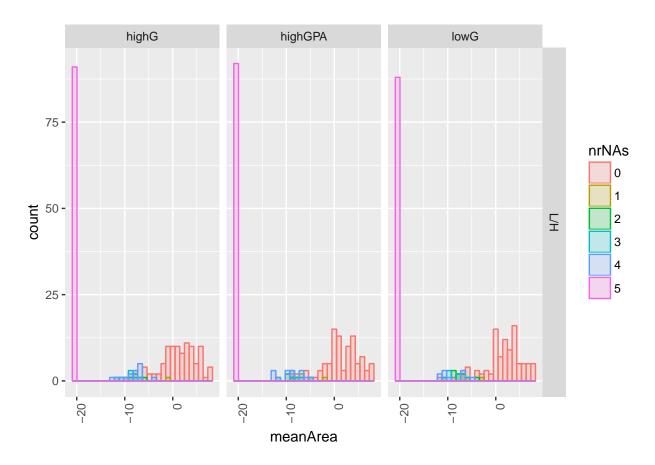


Figure 1: histogram of mean condition intensities per transition

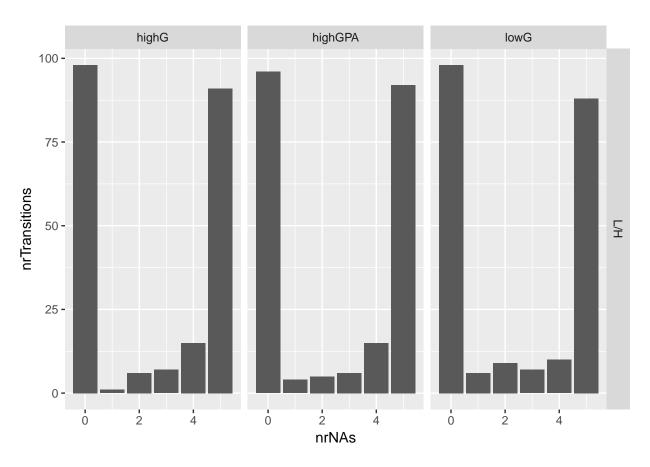


Figure 2: nr of missing

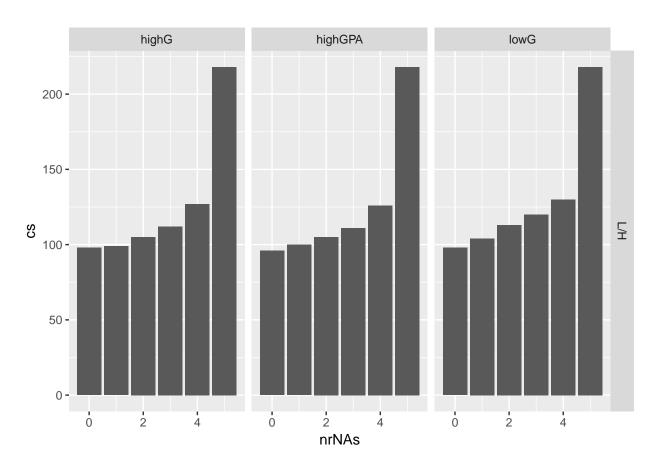


Figure 3: cumulative sum of missing