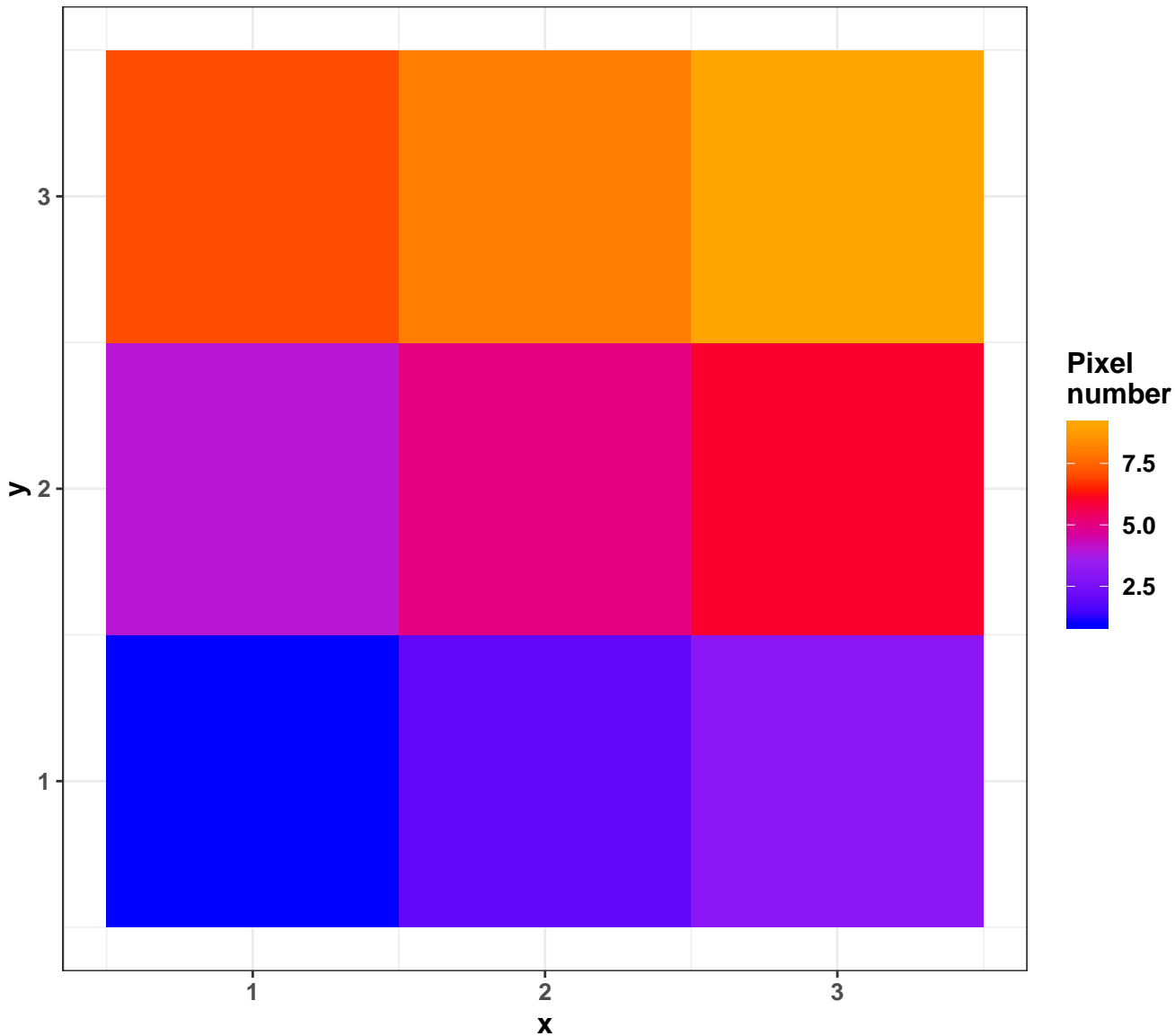


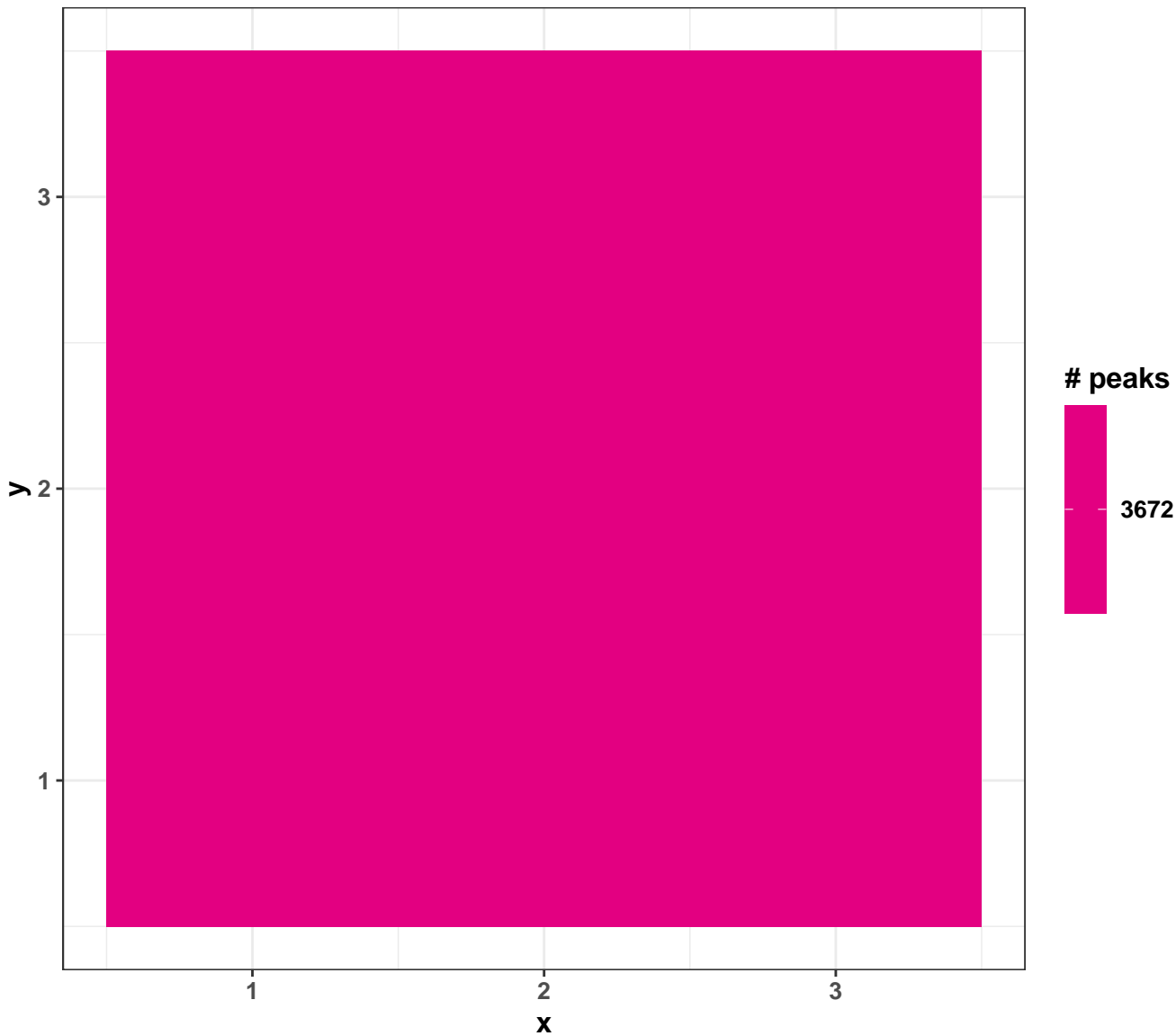
## Testfile\_analyze75

properties	values
Number of m/z features	3672
Range of m/z values	1199.47 – 1356.08
Number of pixels	9
Range of x coordinates	1 – 3
Range of y coordinates	1 – 3
Range of intensities	3 – 84
Number of NA intensities	0
Number of Inf intensities	0
Median of intensities	9
Intensities > 0	100 %
Number of empty spectra	0
Median TIC ± sd	37005 ± 5329
Median # peaks per spectrum ± sd	3672 ± 0
Centroided	FALSE
calibrants (#valid/#input) in None	0 / 0

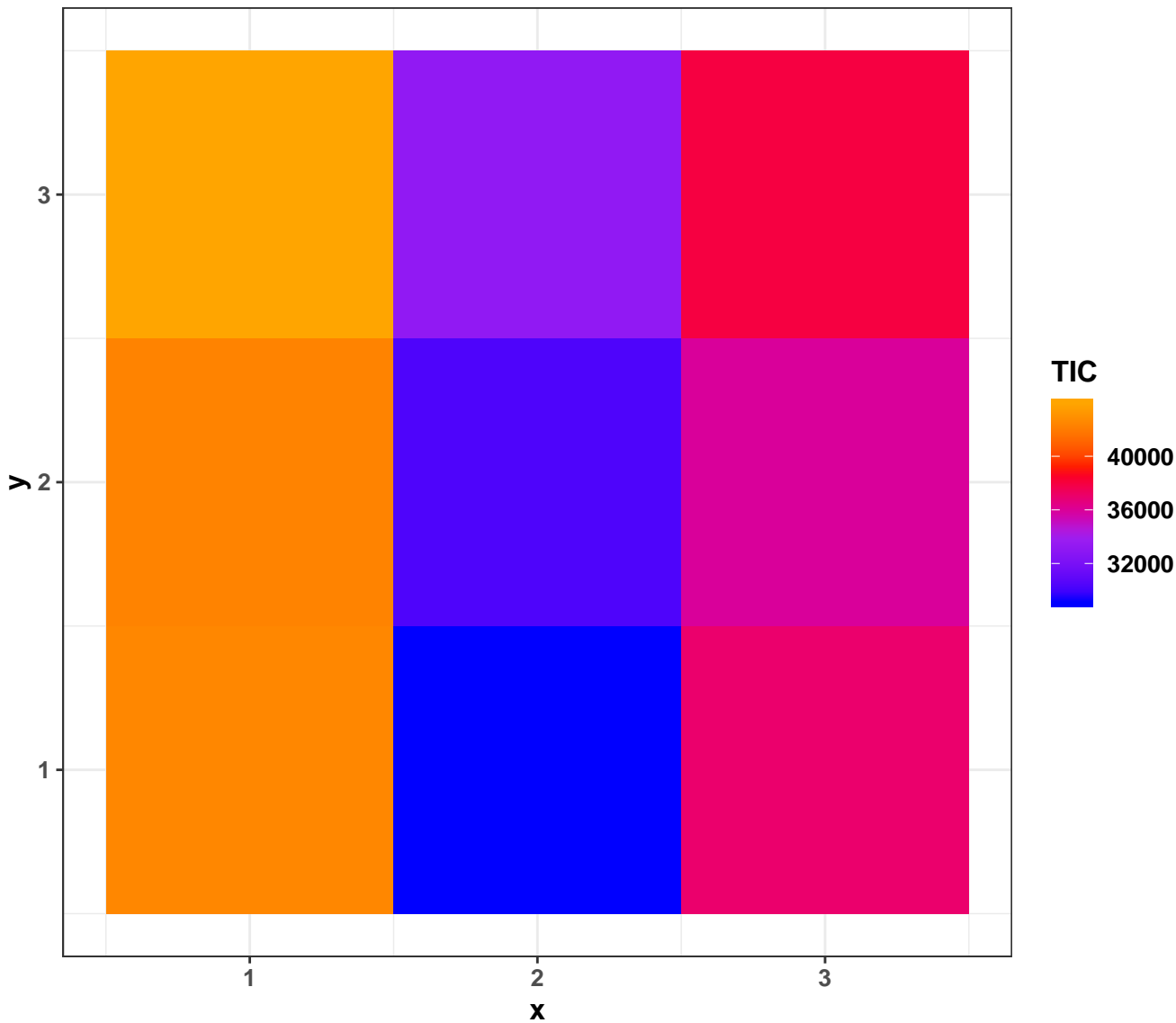
**Pixel order**



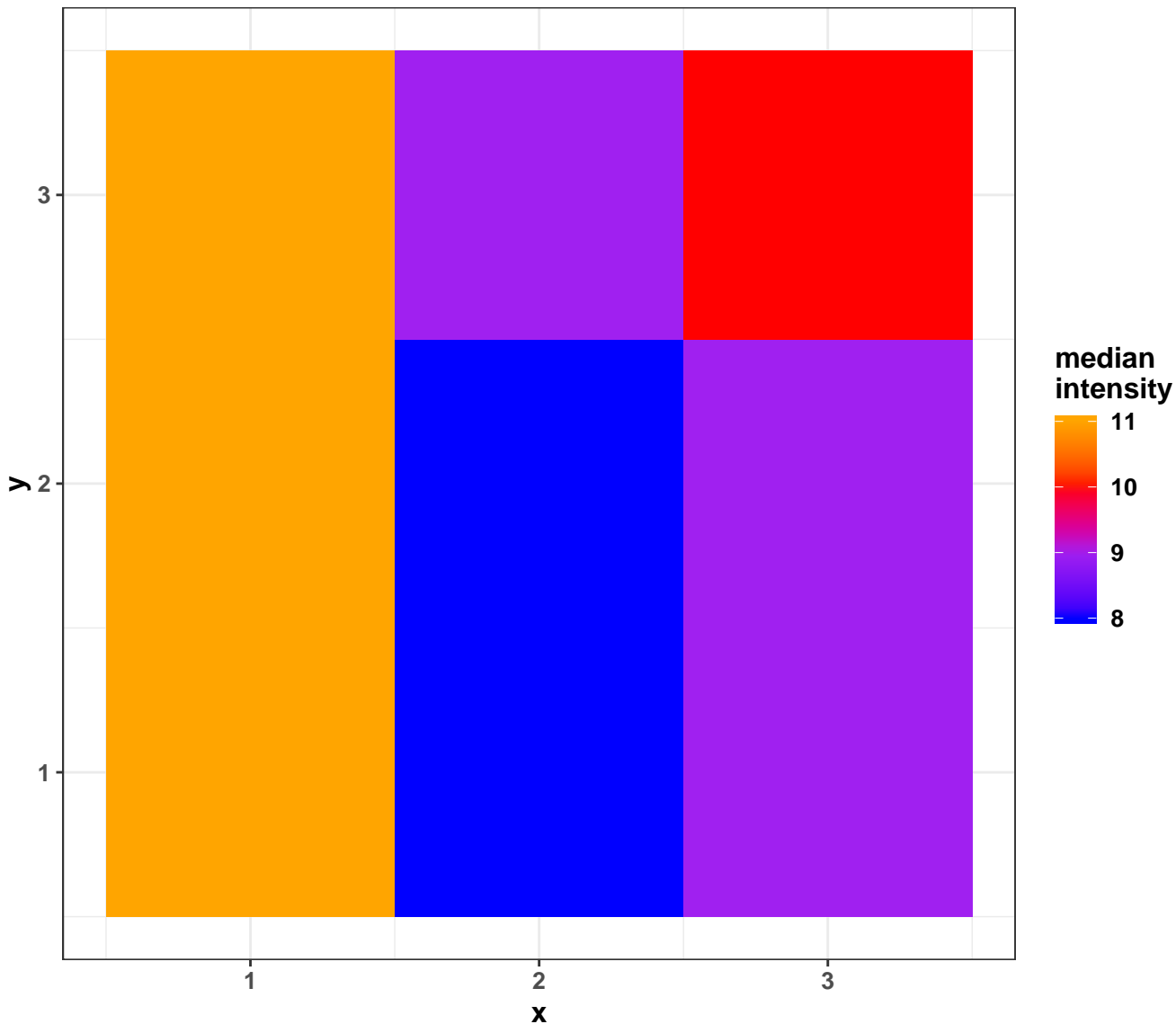
Number of peaks per spectrum



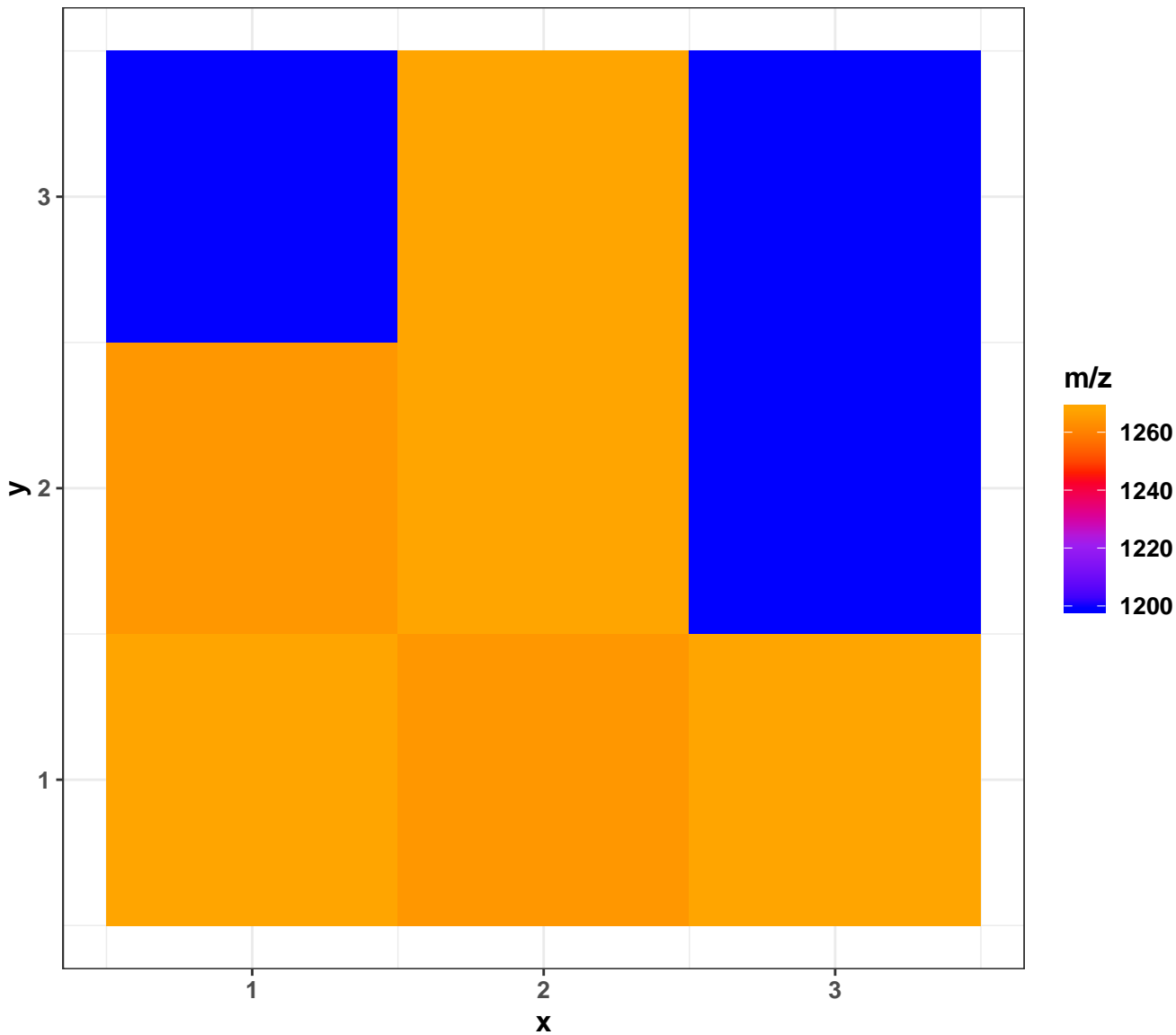
Total Ion Chromatogram



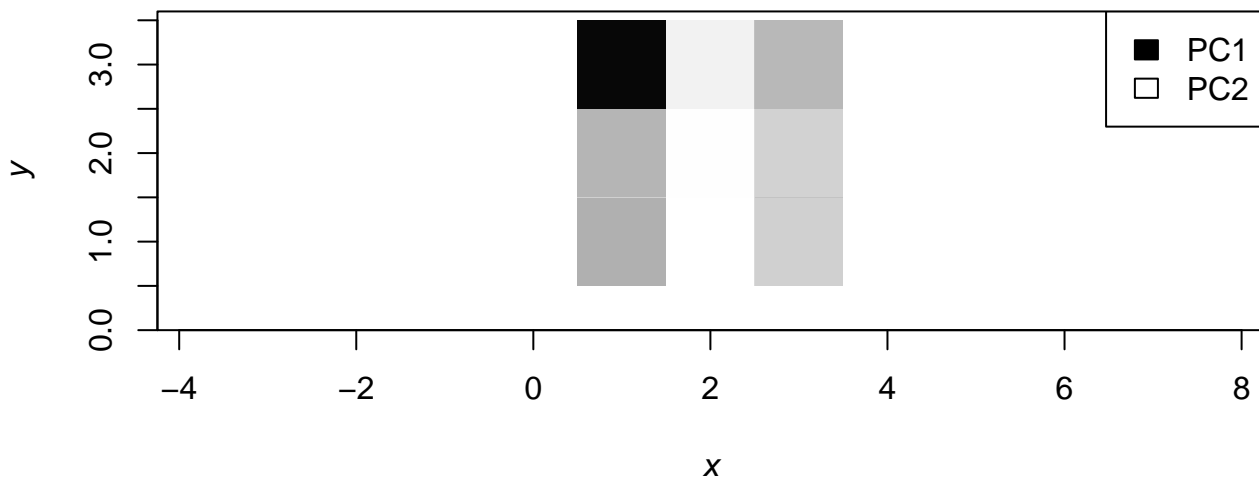
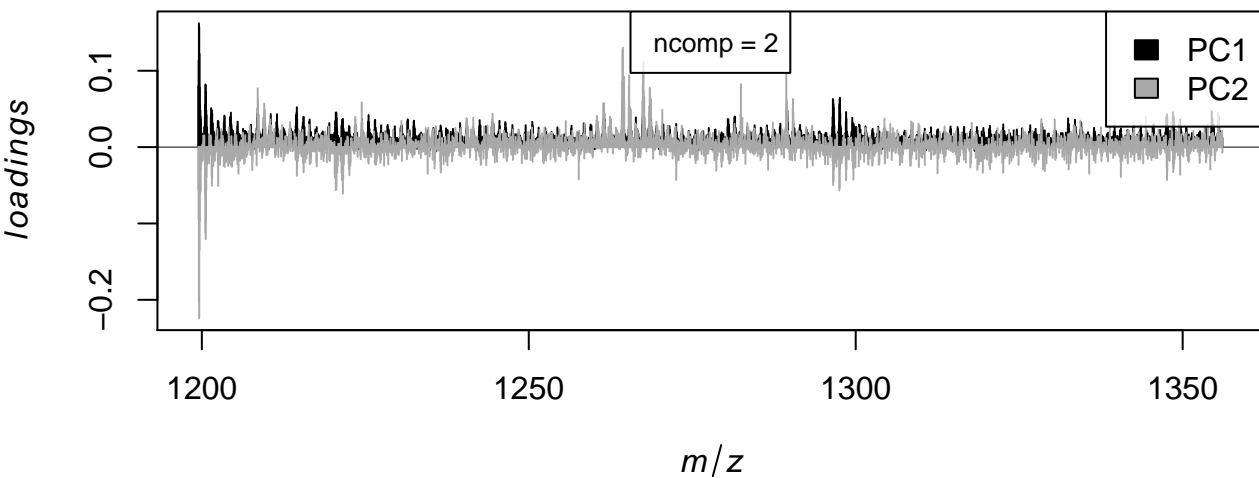
Median intensity per pixel

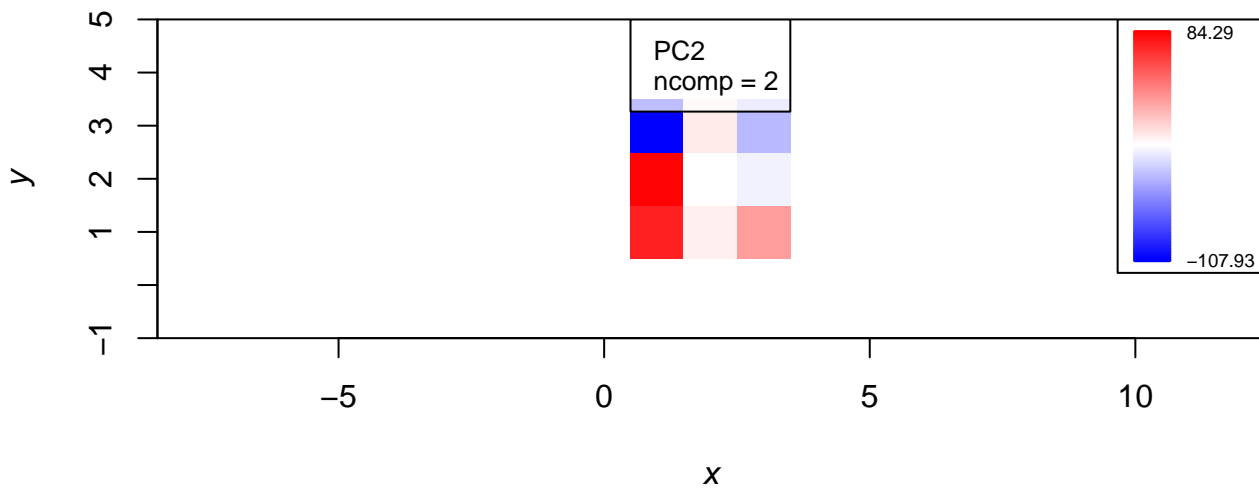
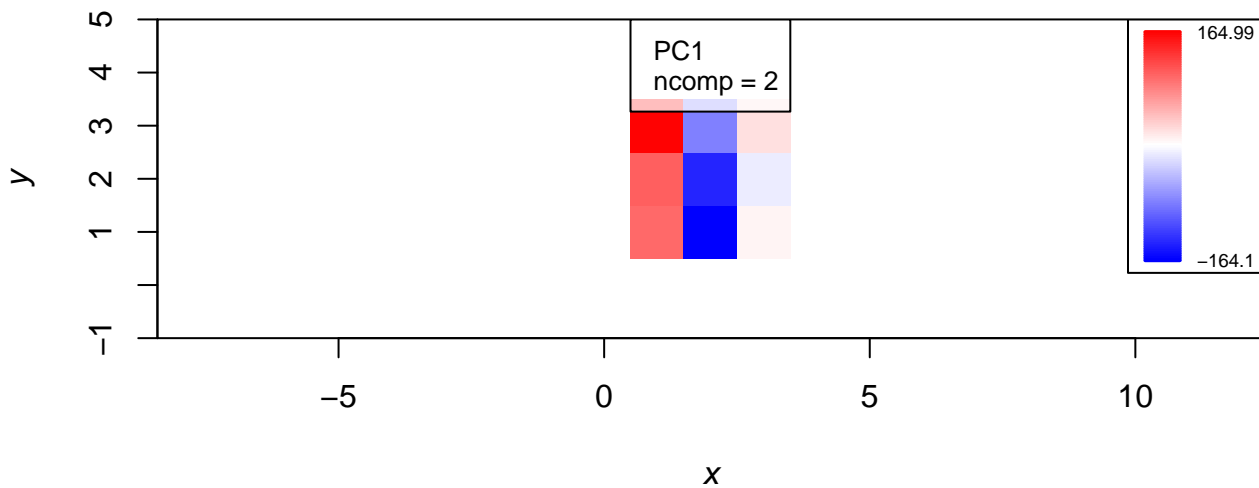


Most abundant m/z in each spectrum



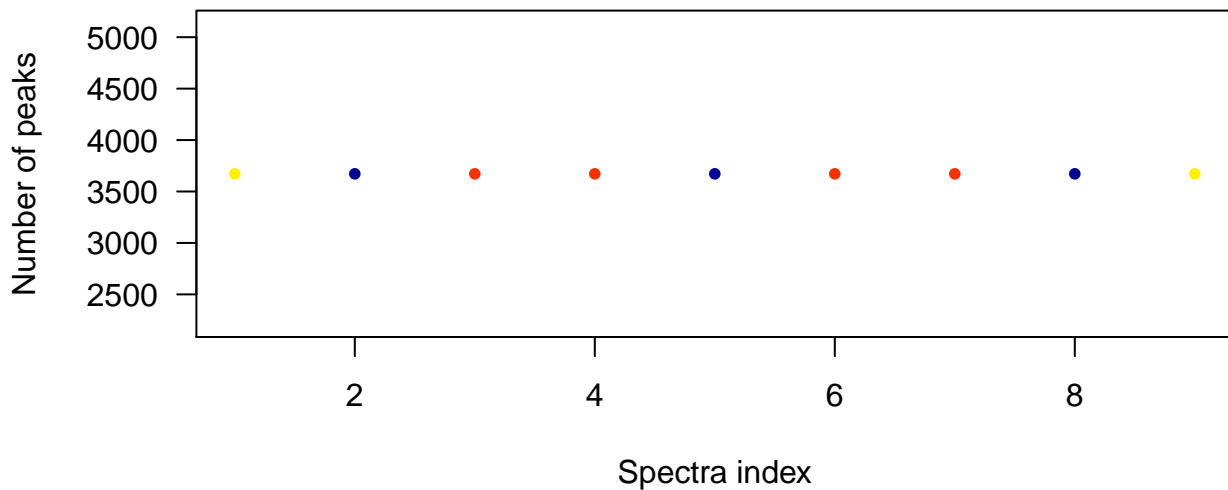
# PCA for two components



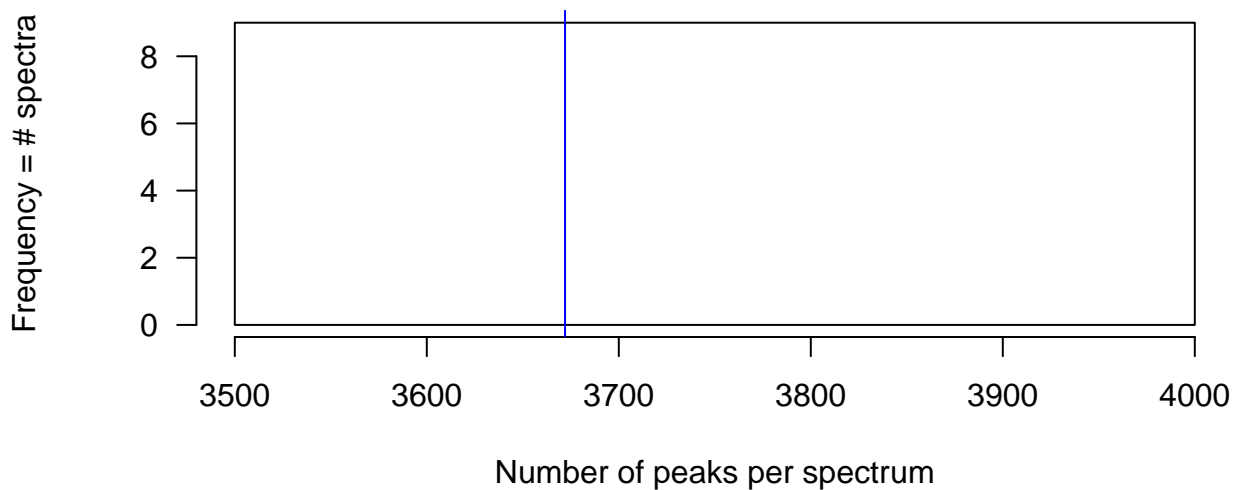


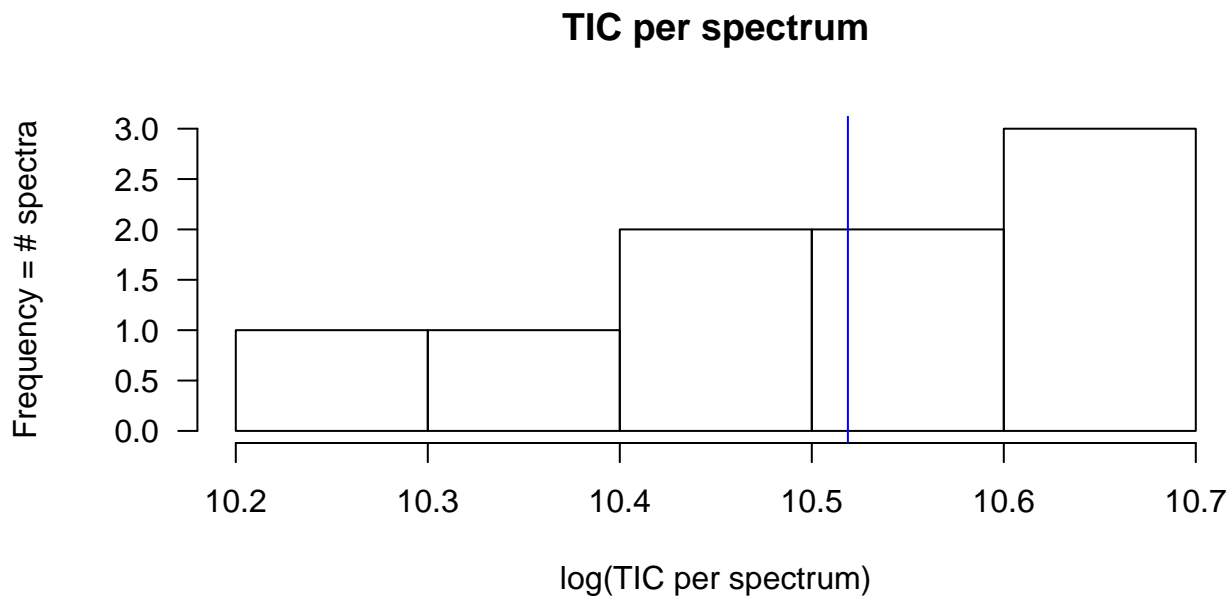
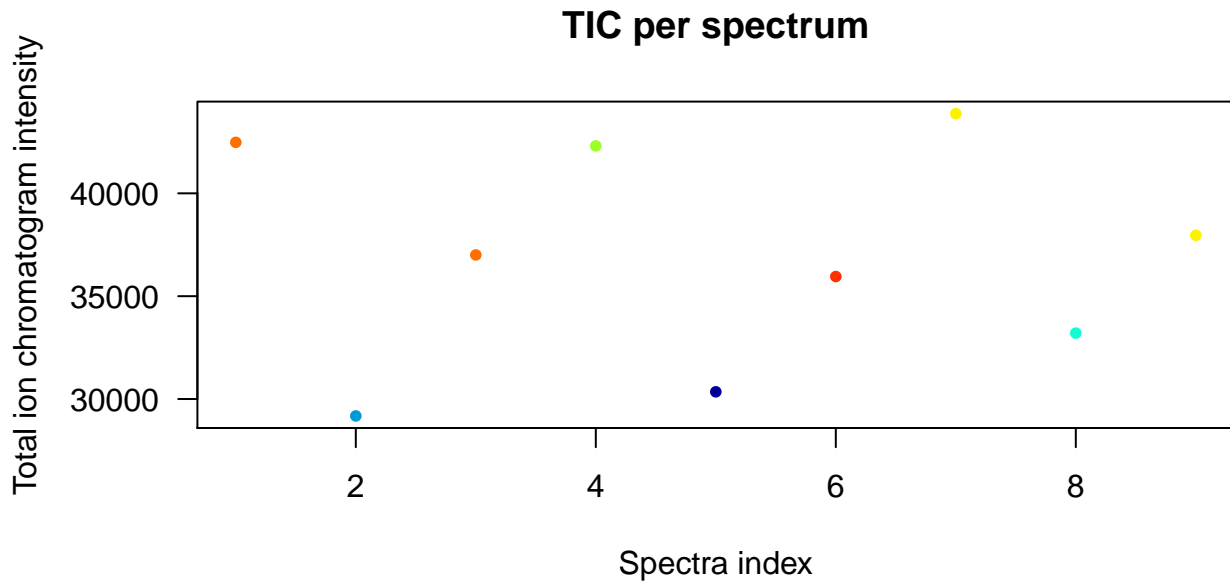


**Number of peaks per spectrum**

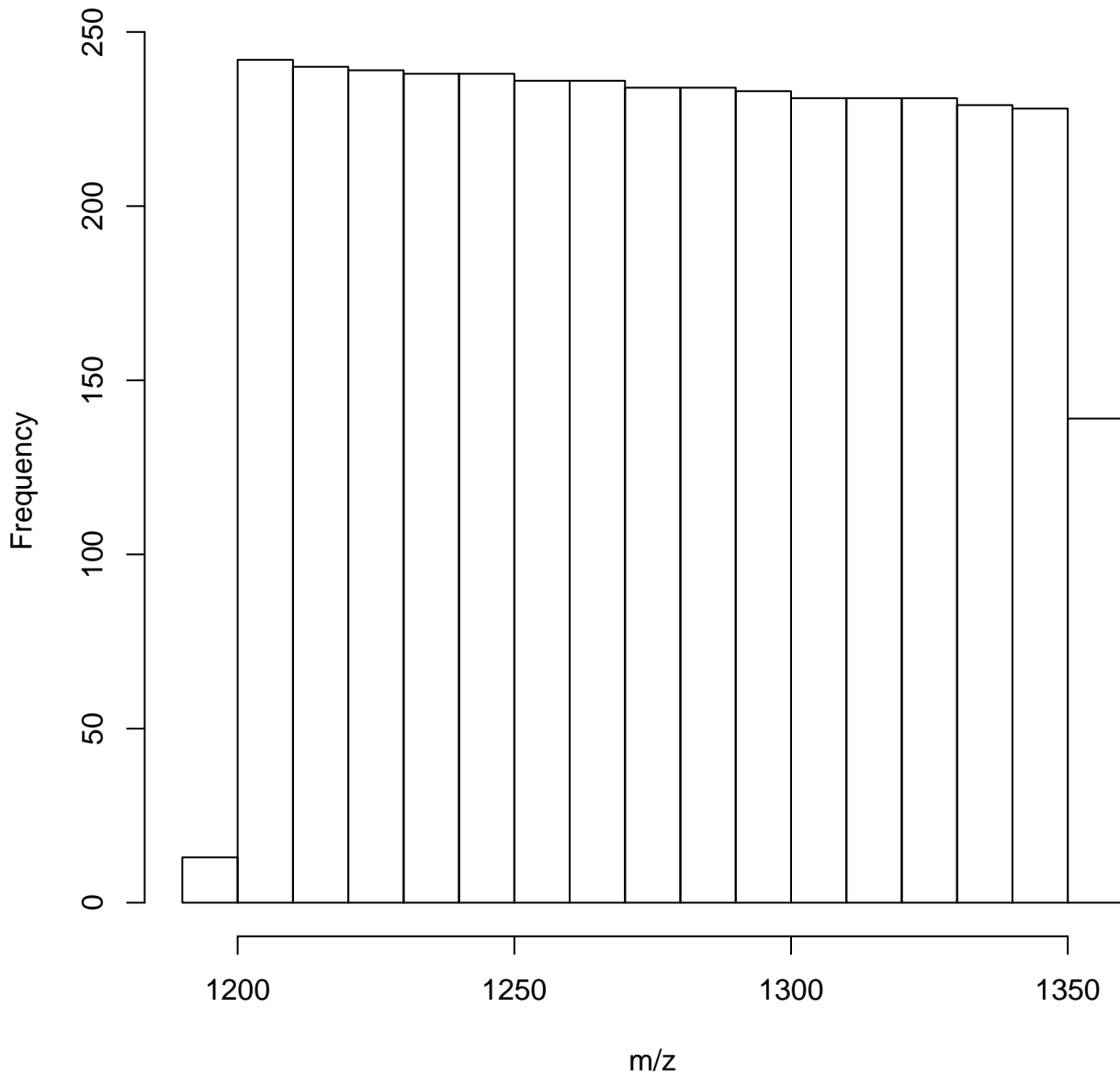


**Number of peaks per spectrum**

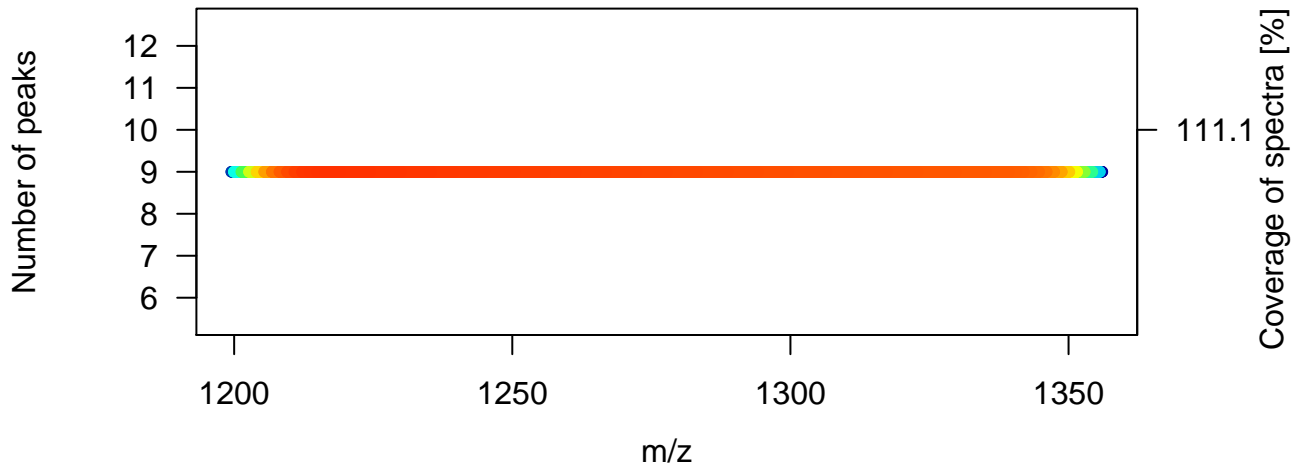




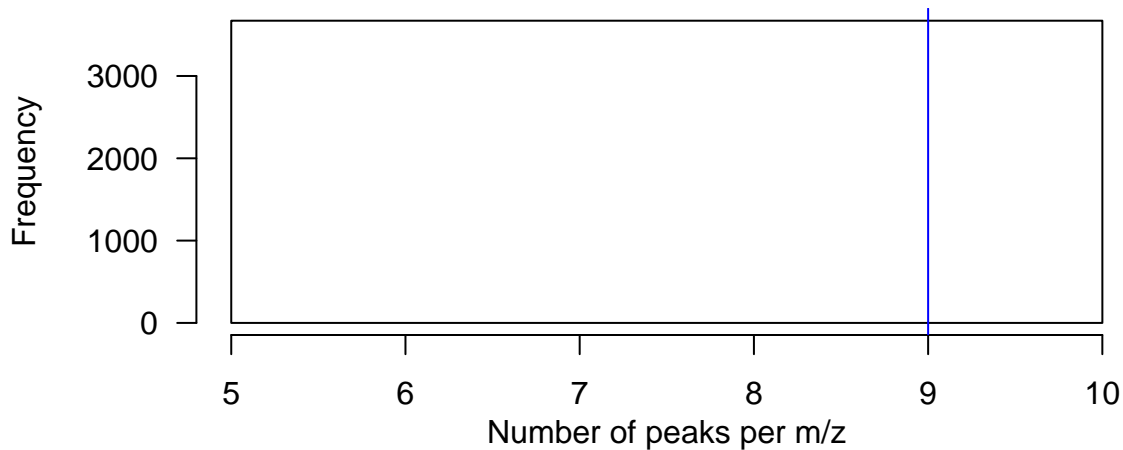
**Histogram of m/z values**



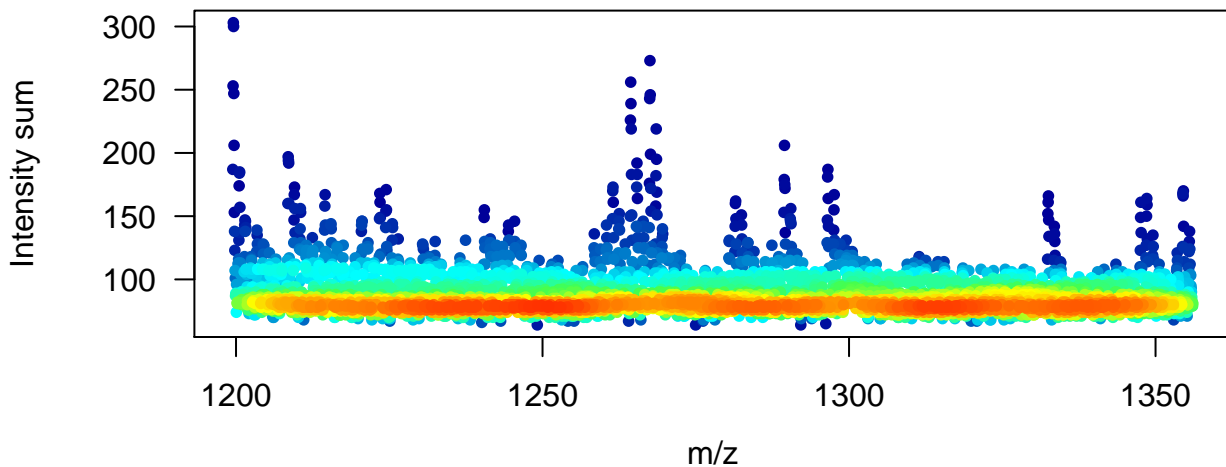
## Number of peaks per m/z



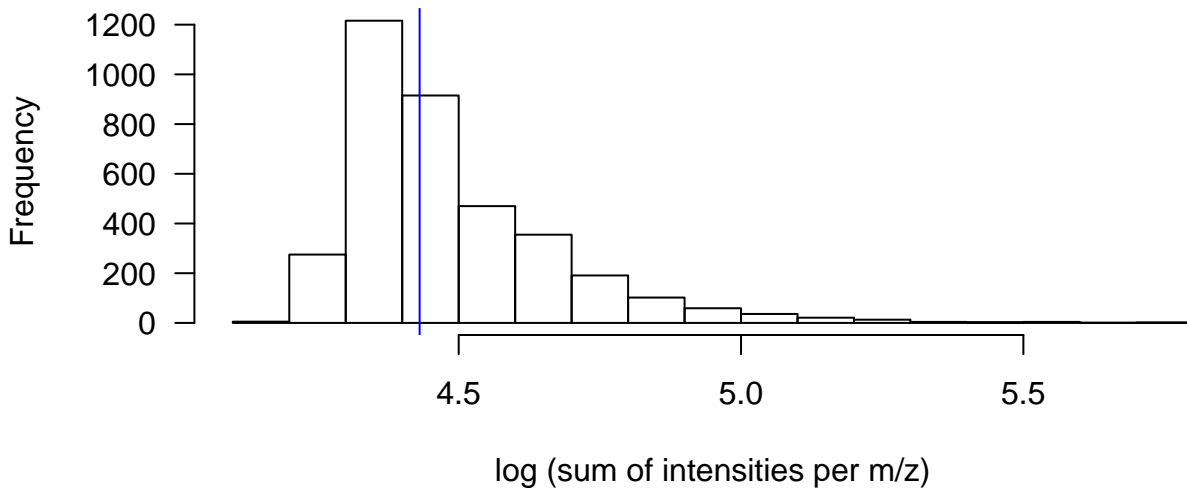
## Number of peaks per m/z



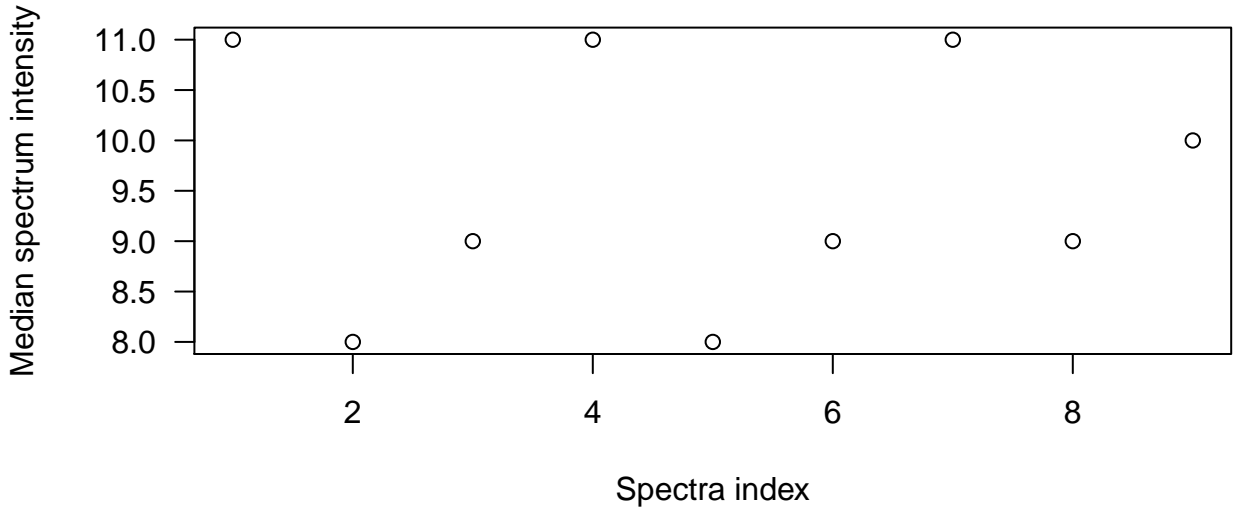
Sum of intensities per m/z



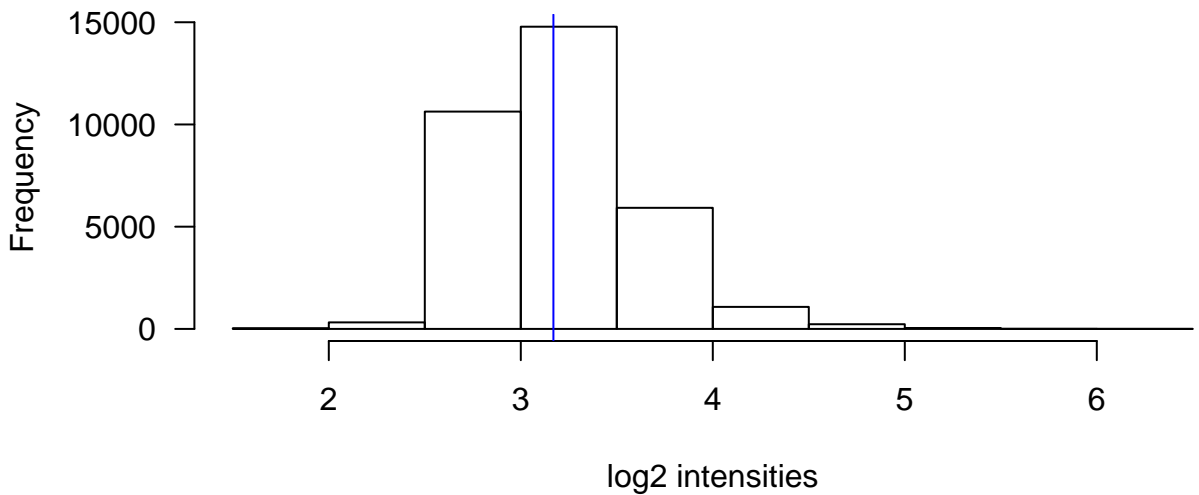
Sum of intensities per m/z



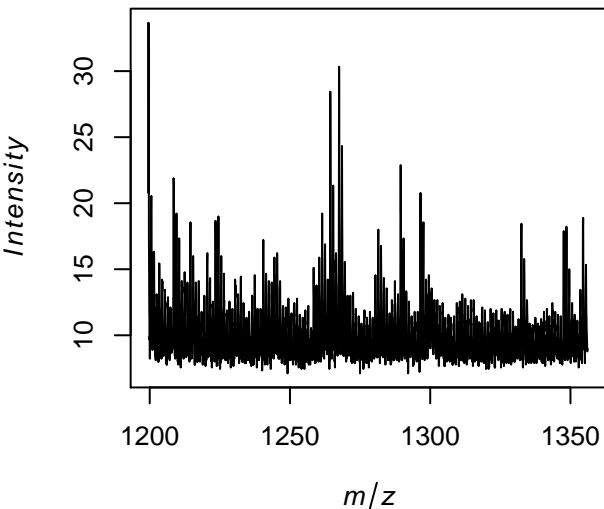
**Median intensity per spectrum**



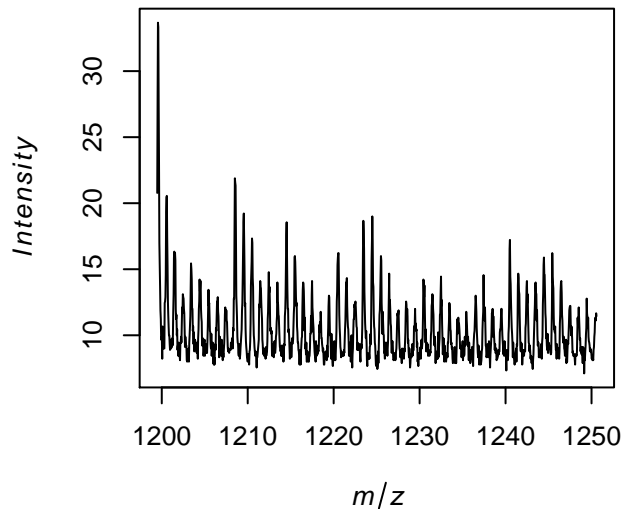
**Log2-transformed intensities**



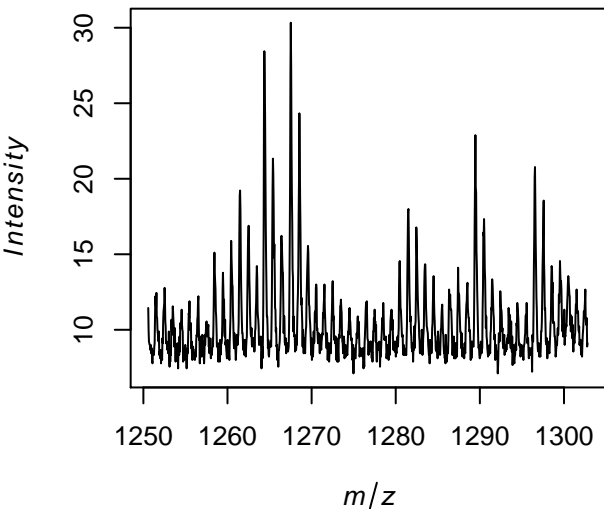
**Average spectrum**



**Zoomed average spectrum**



**Zoomed average spectrum**



**Zoomed average spectrum**

