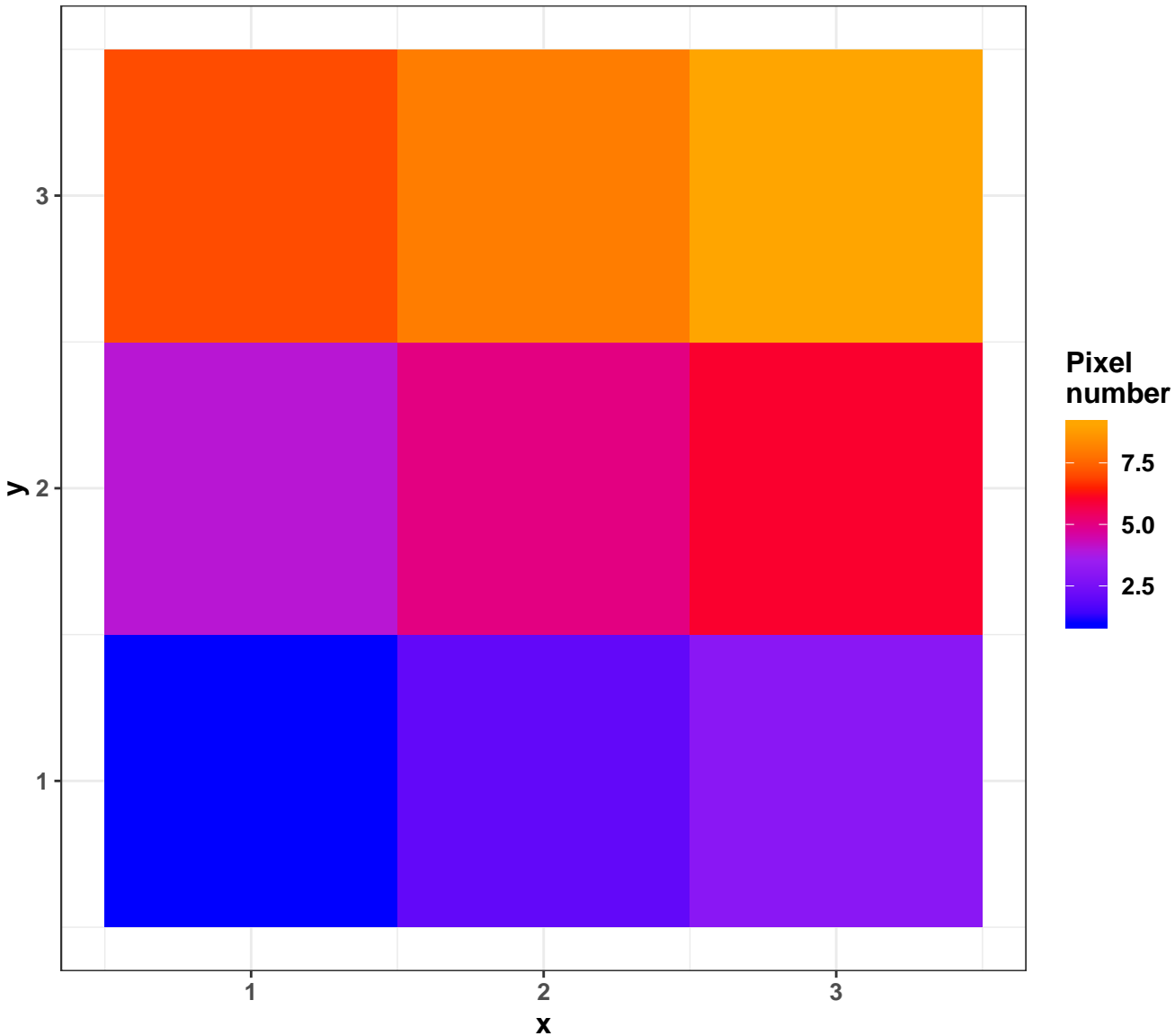


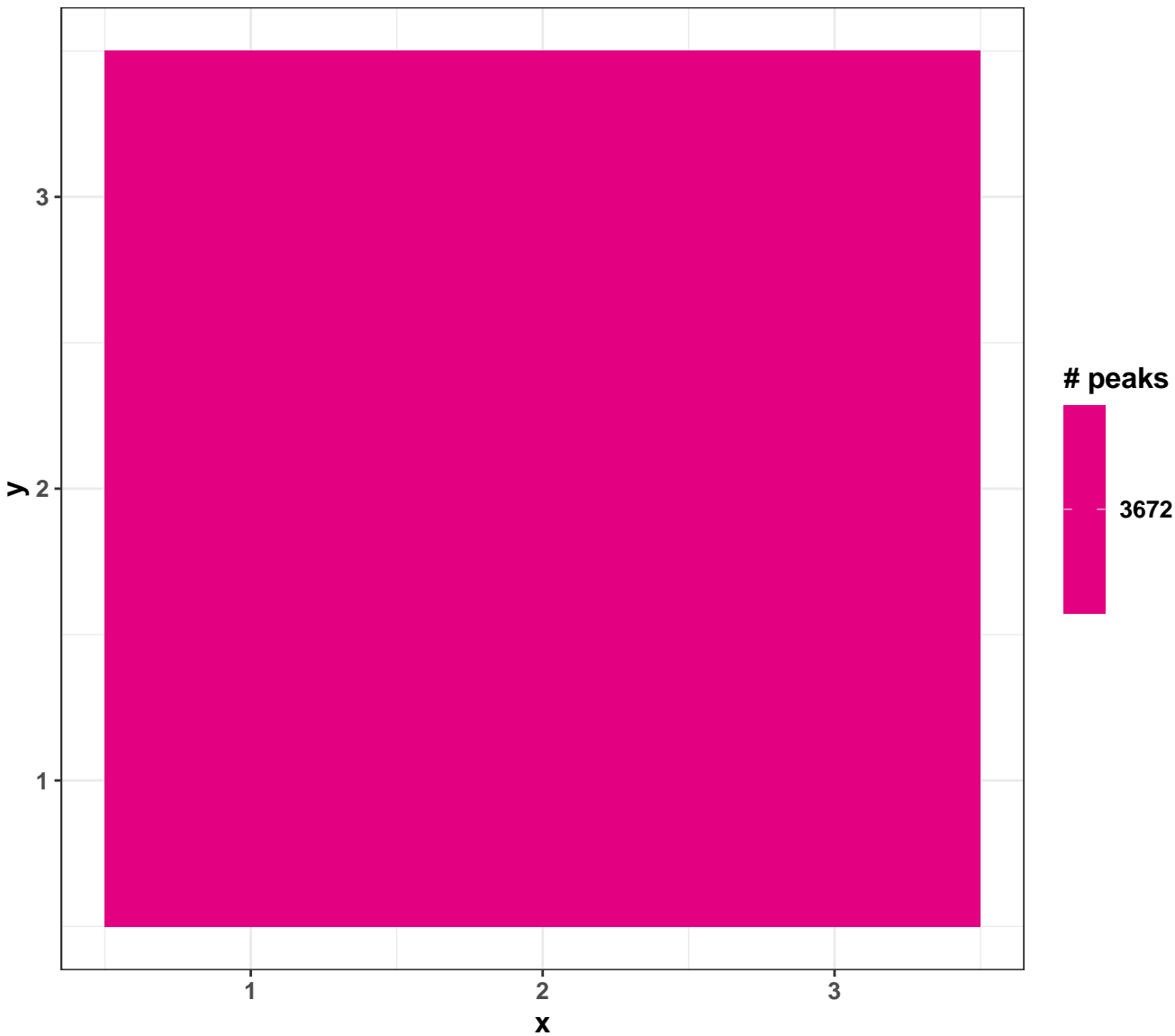
## Testfile\_analyze75

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
Median of intensities	9
Intensities > 0	100 %
Number of empty spectra	0
Median TIC $\pm$ sd	37005 $\pm$ 5329
Median # peaks per spectrum $\pm$ sd	3672 $\pm$ 0
Centroided	FALSE
calibrants (#valid/#input) in None	0 / 0

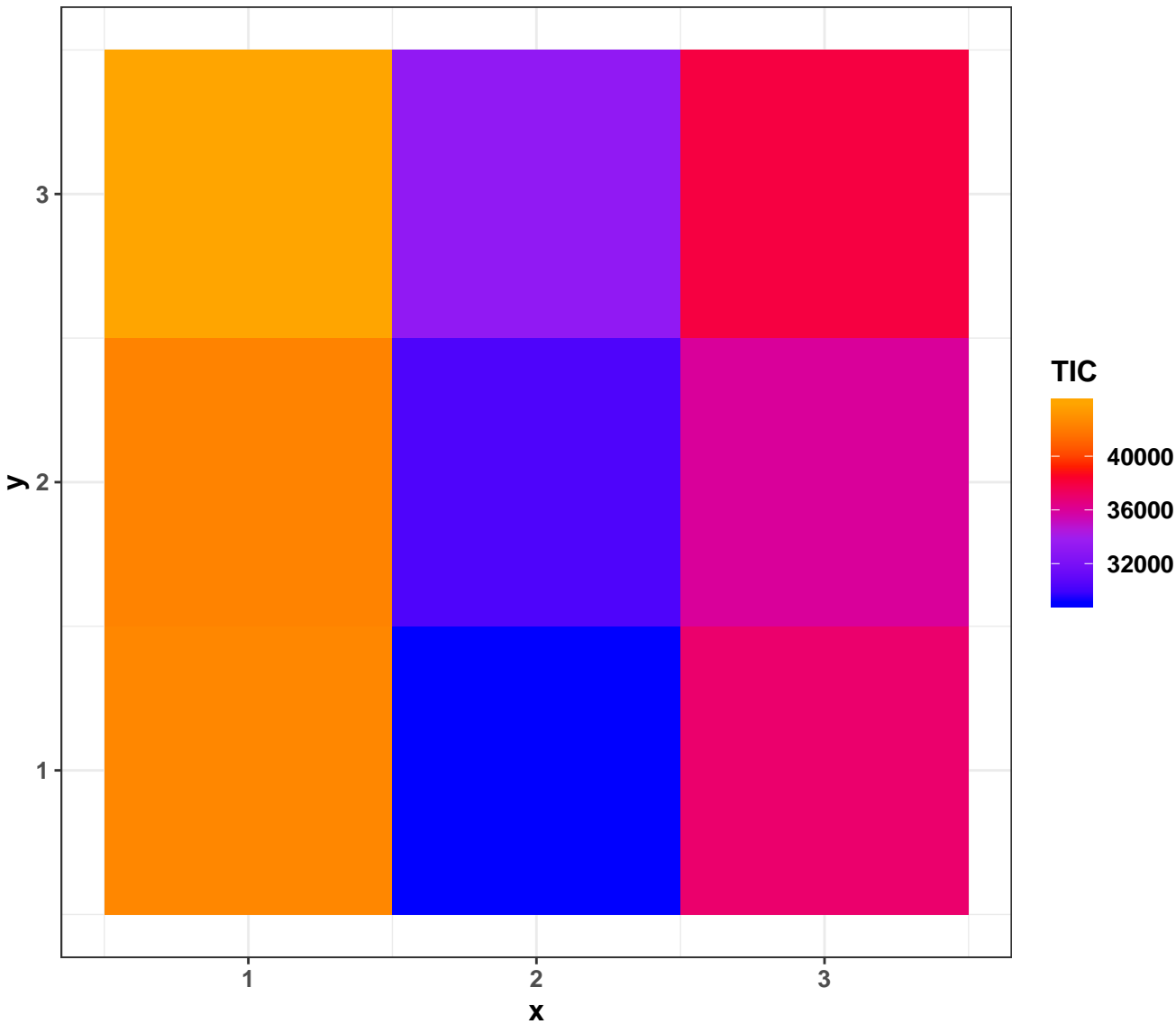
# Pixel order



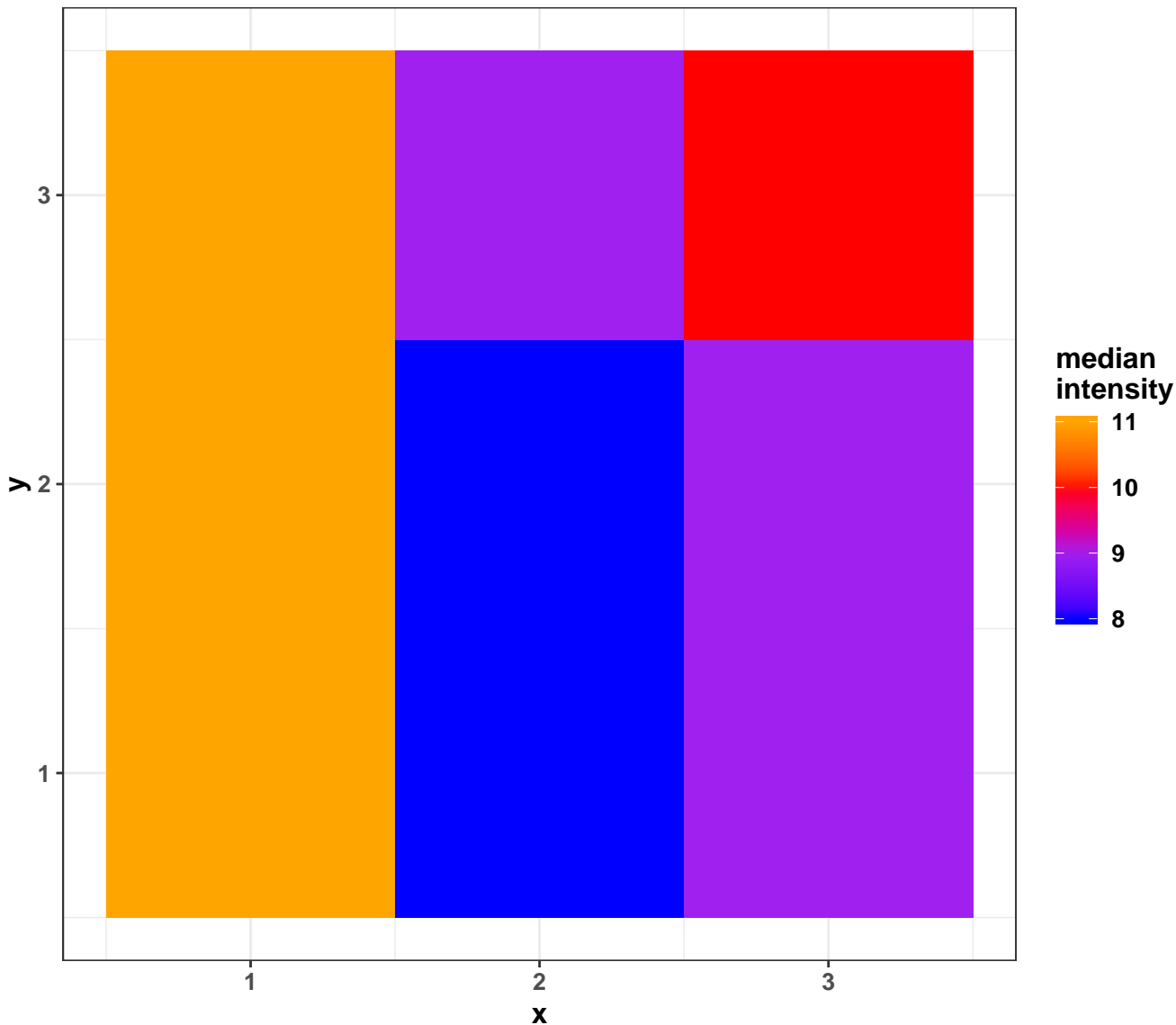
# Number of peaks per spectrum



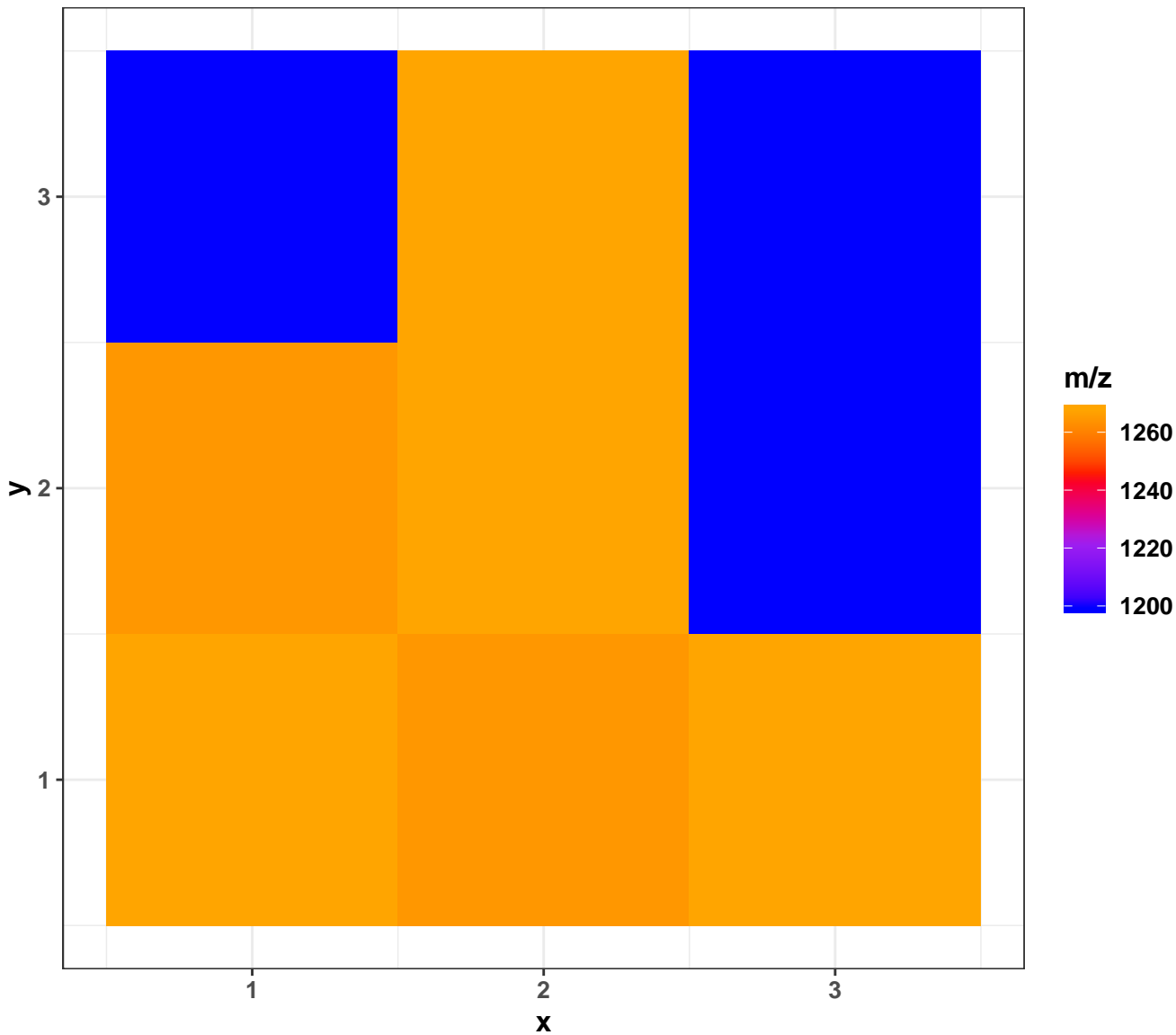
# Total Ion Chromatogram



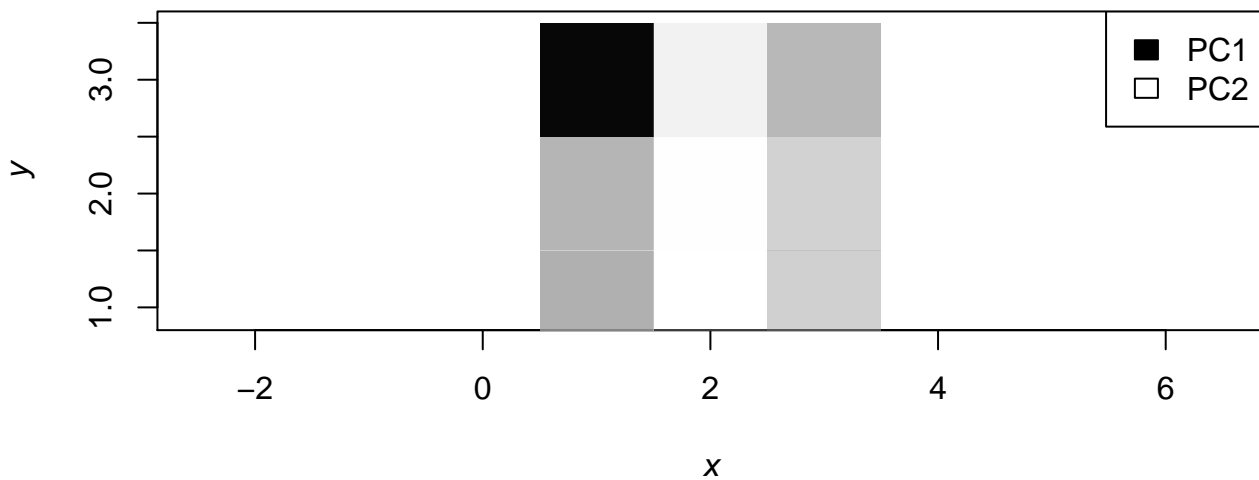
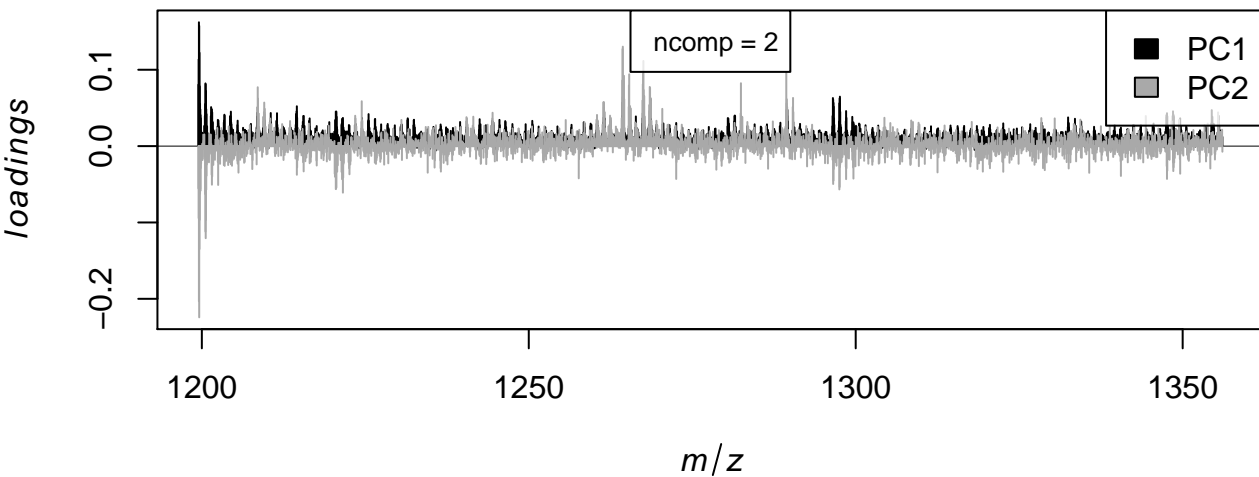
Median intensity per pixel



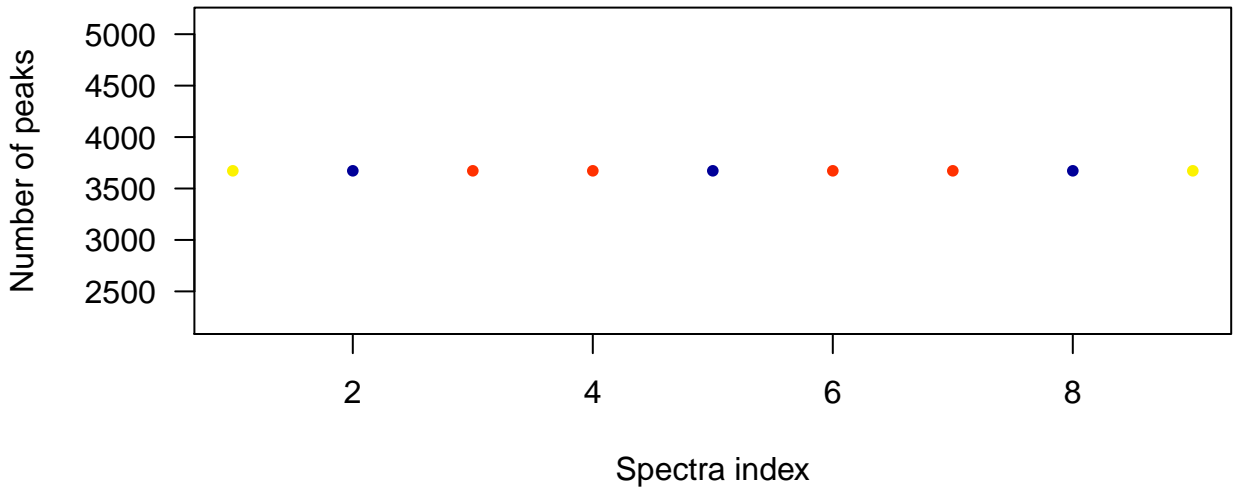
# Most abundant m/z in each spectrum



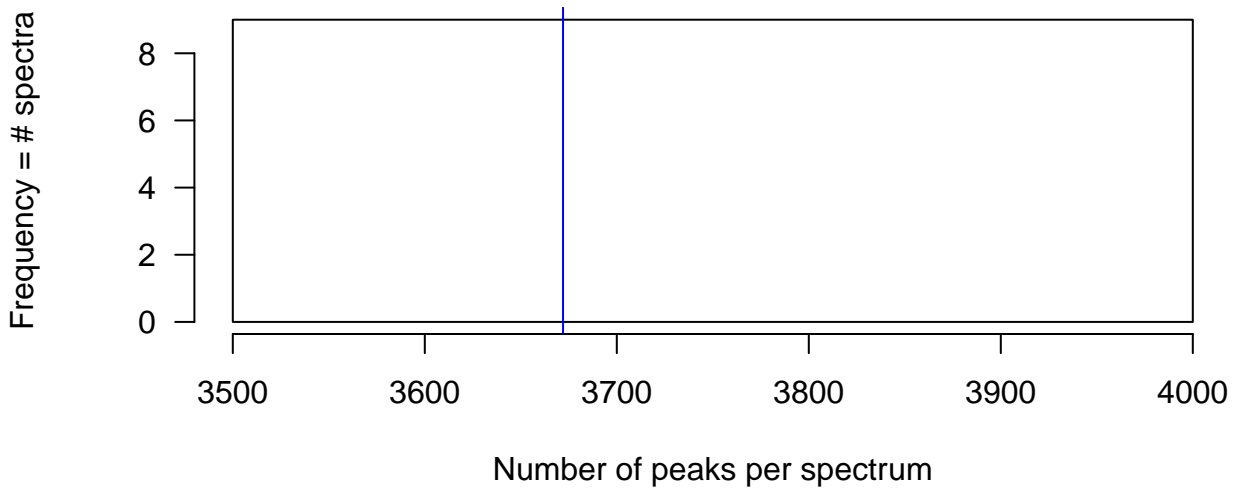
# PCA for two components



### Number of peaks per spectrum

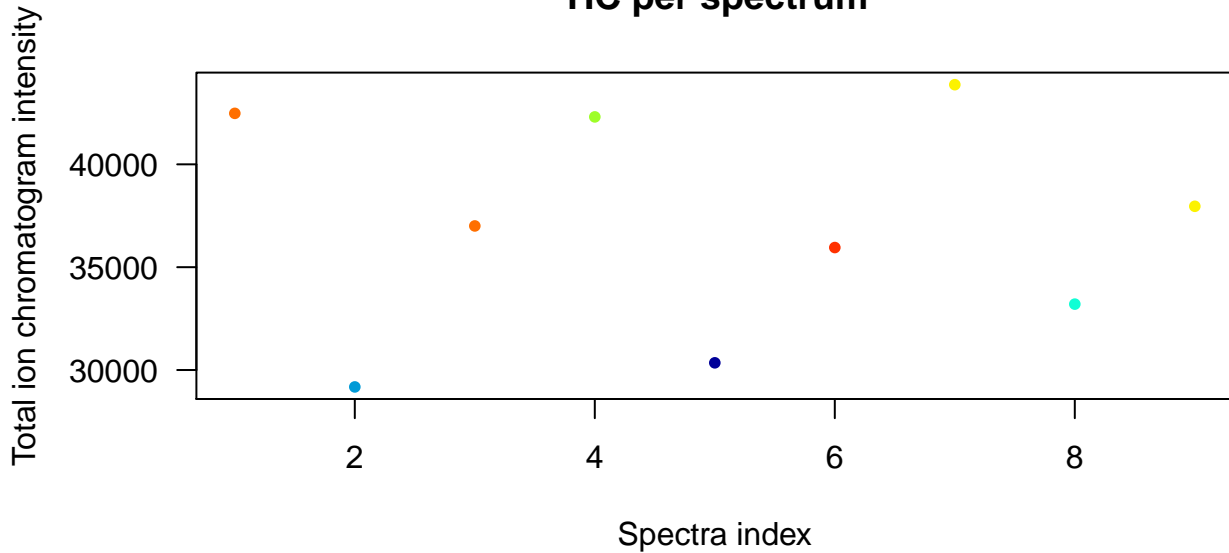


### Number of peaks per spectrum

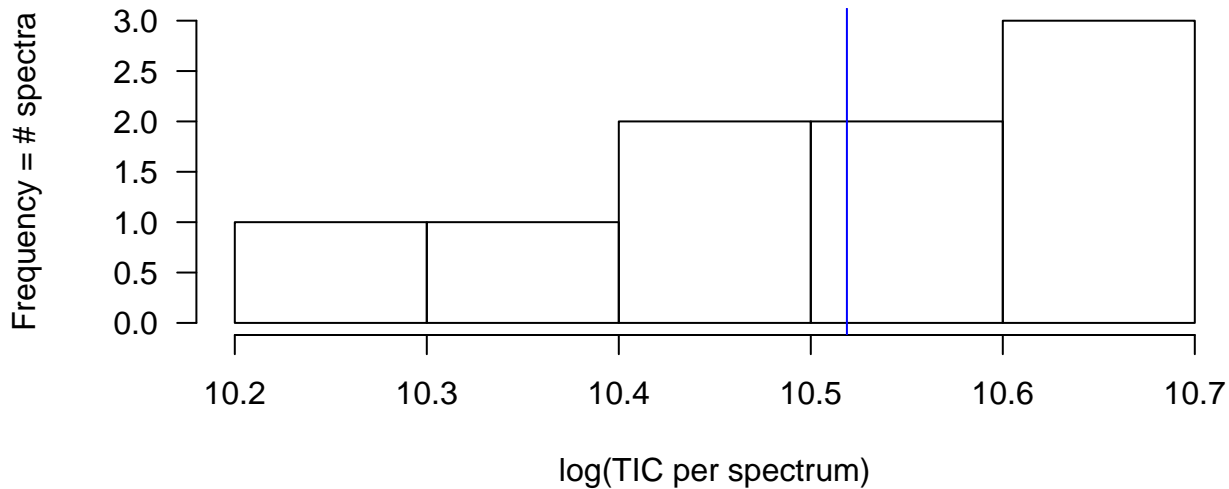




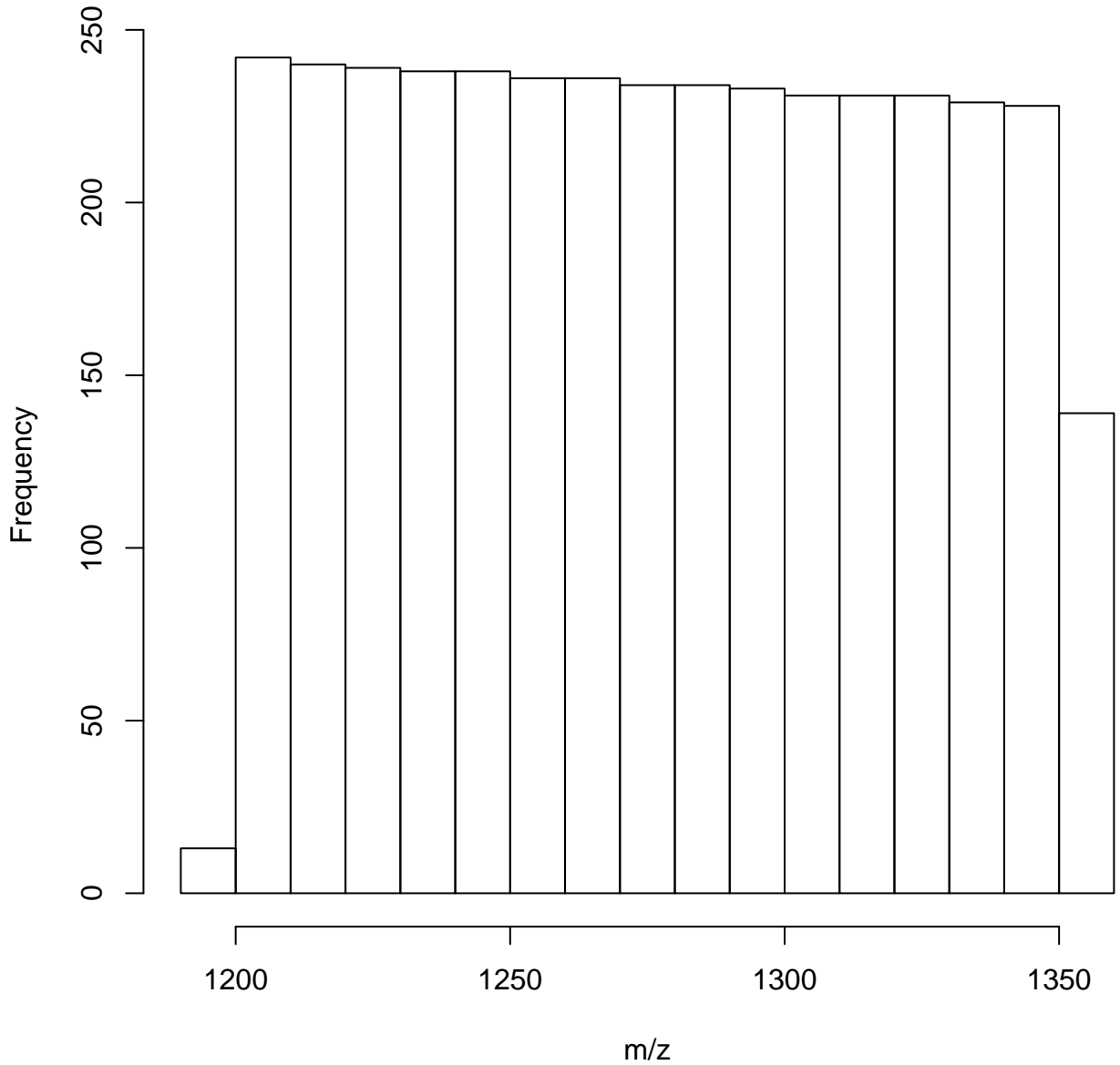
### TIC per spectrum



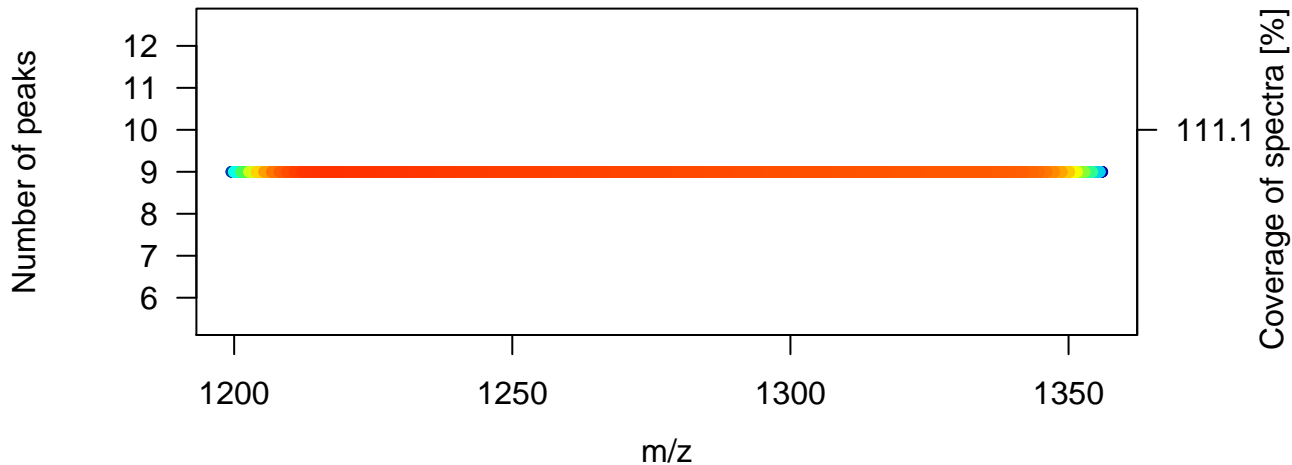
### TIC per spectrum



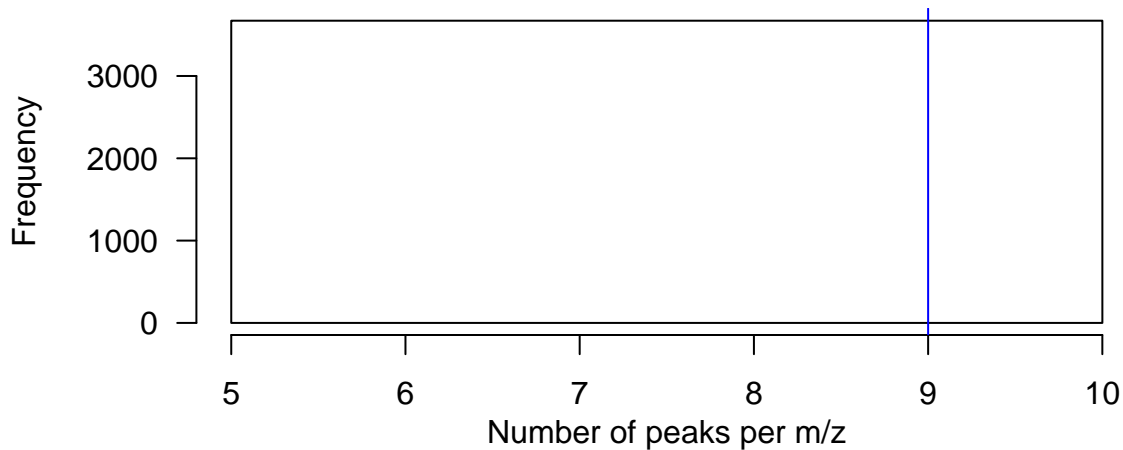
# Histogram of m/z values



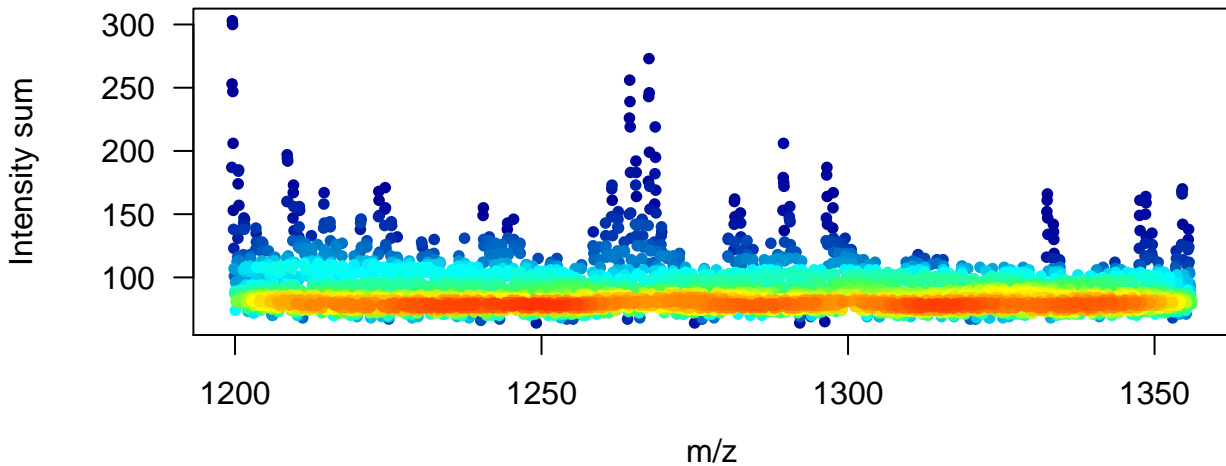
### Number of peaks per m/z



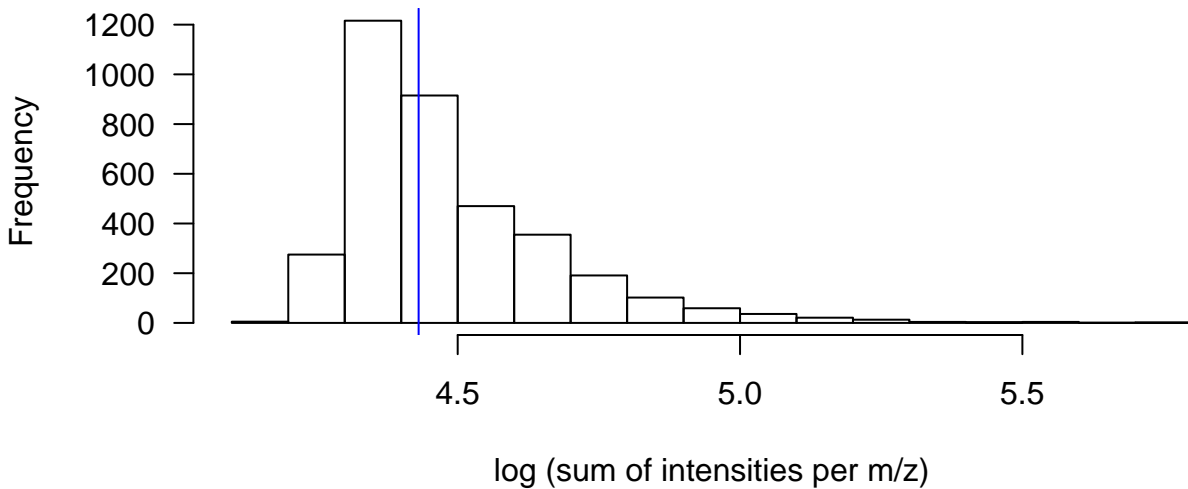
### Number of peaks per m/z



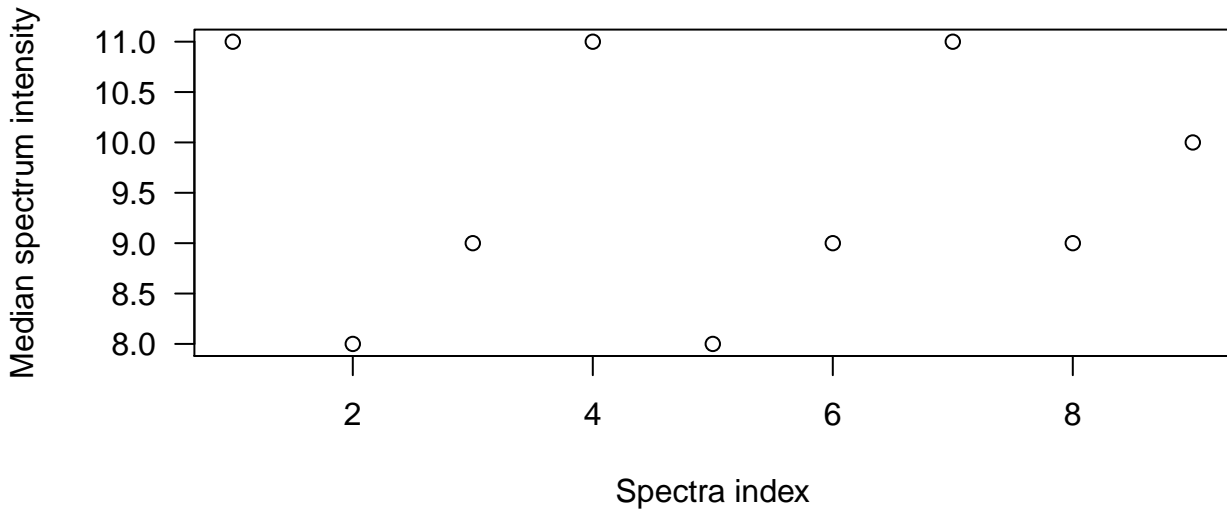
Sum of intensities per m/z



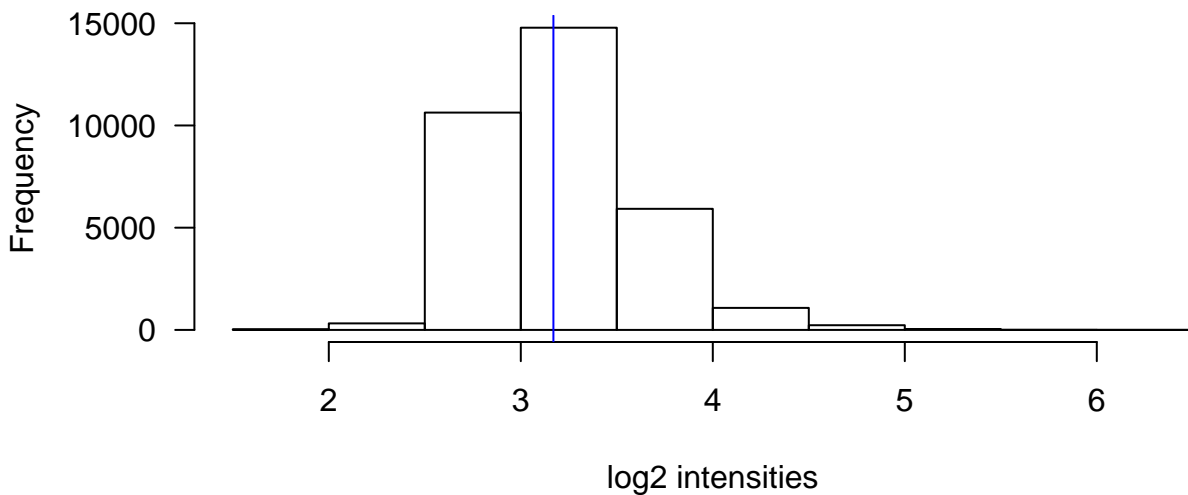
Sum of intensities per m/z



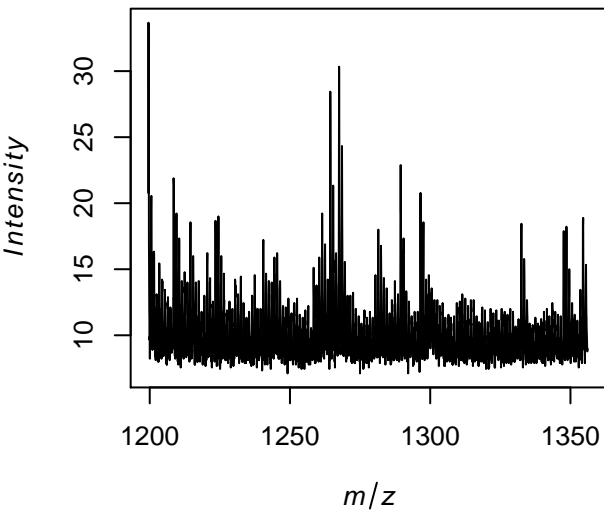
## Median intensity per spectrum



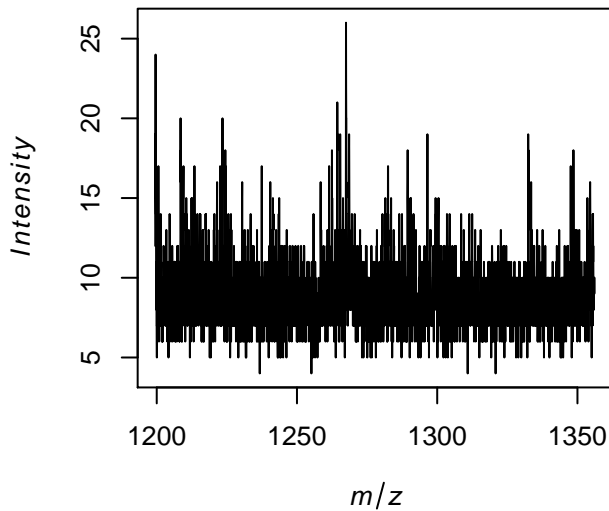
## Log2-transformed intensities



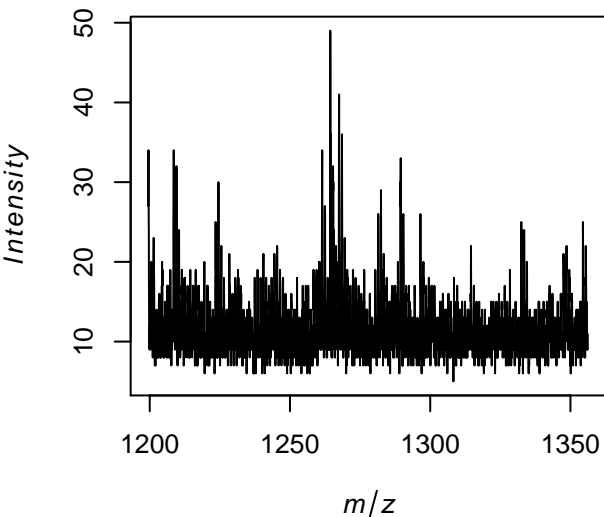
**Average spectrum**



**Spectrum at  $x = 2, y = 3$**



**Spectrum at  $x = 1, y = 2$**



**Spectrum at  $x = 3, y = 3$**

