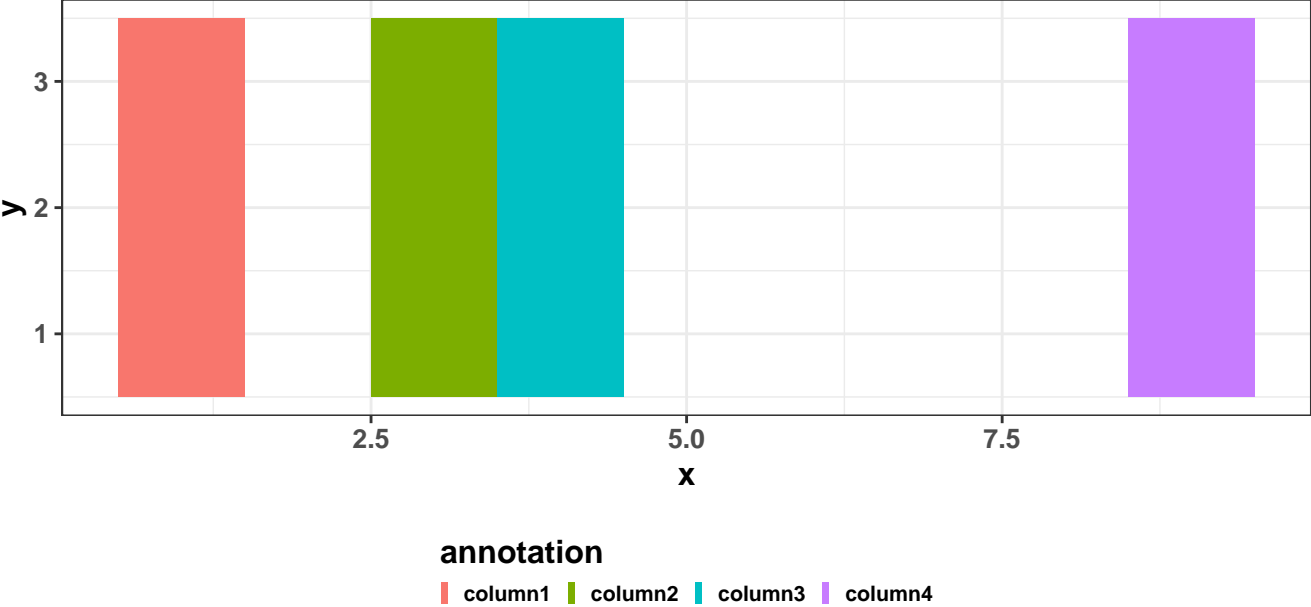


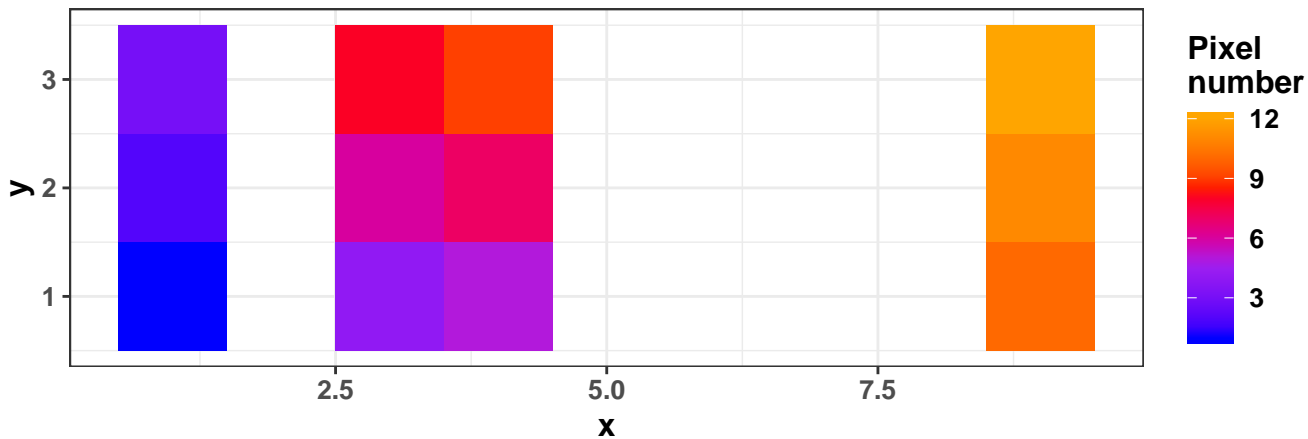
Testfile_rdata

properties	values
Number of m/z features	8399
Range of m/z values	100.08 – 799.92
Number of pixels	12
Range of x coordinates	1 – 9
Range of y coordinates	1 – 3
Range of intensities	0 – 9.24
Number of NA intensities	0
Number of Inf intensities	0
Median of intensities	0
Intensities > 0	31.29 %
Number of empty spectra	0
Median TIC ± sd	161.8 ± 47
Median # peaks per spectrum ± sd	2811 ± 424
Centroided	FALSE
calibrants (#valid/#input) in inputcalibrantfile1.tabular	3 / 3

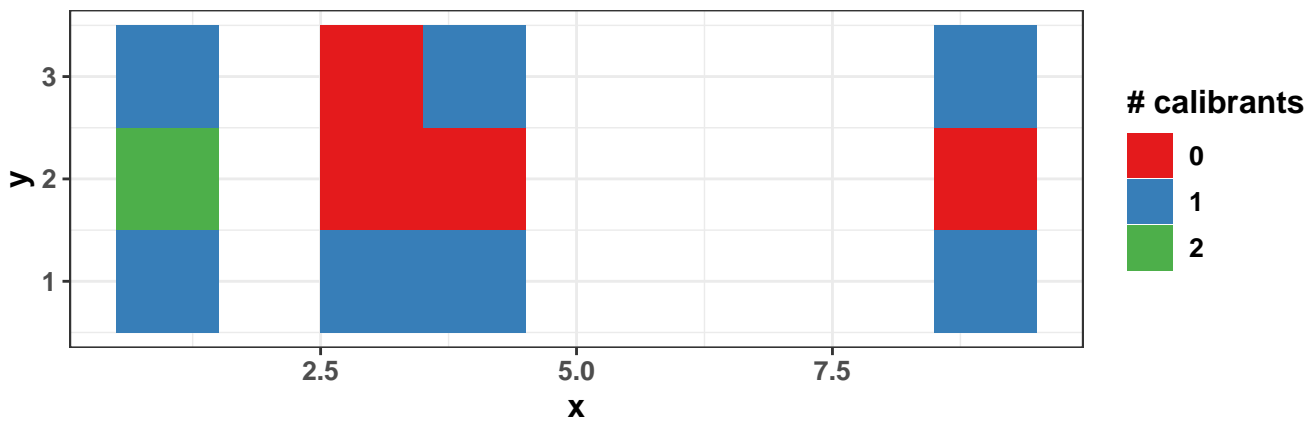
Spatial orientation of pixel annotations



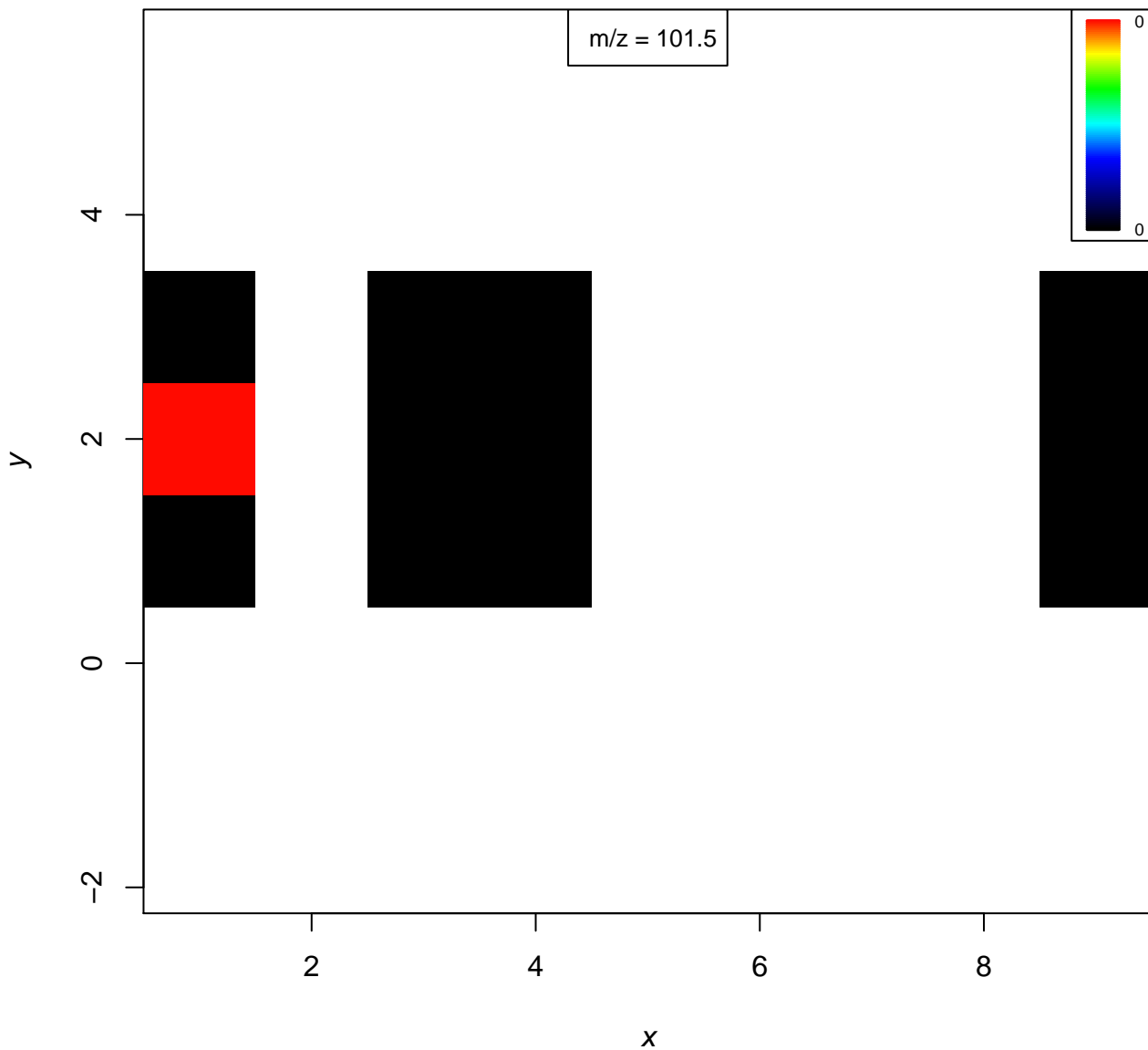
Pixel order



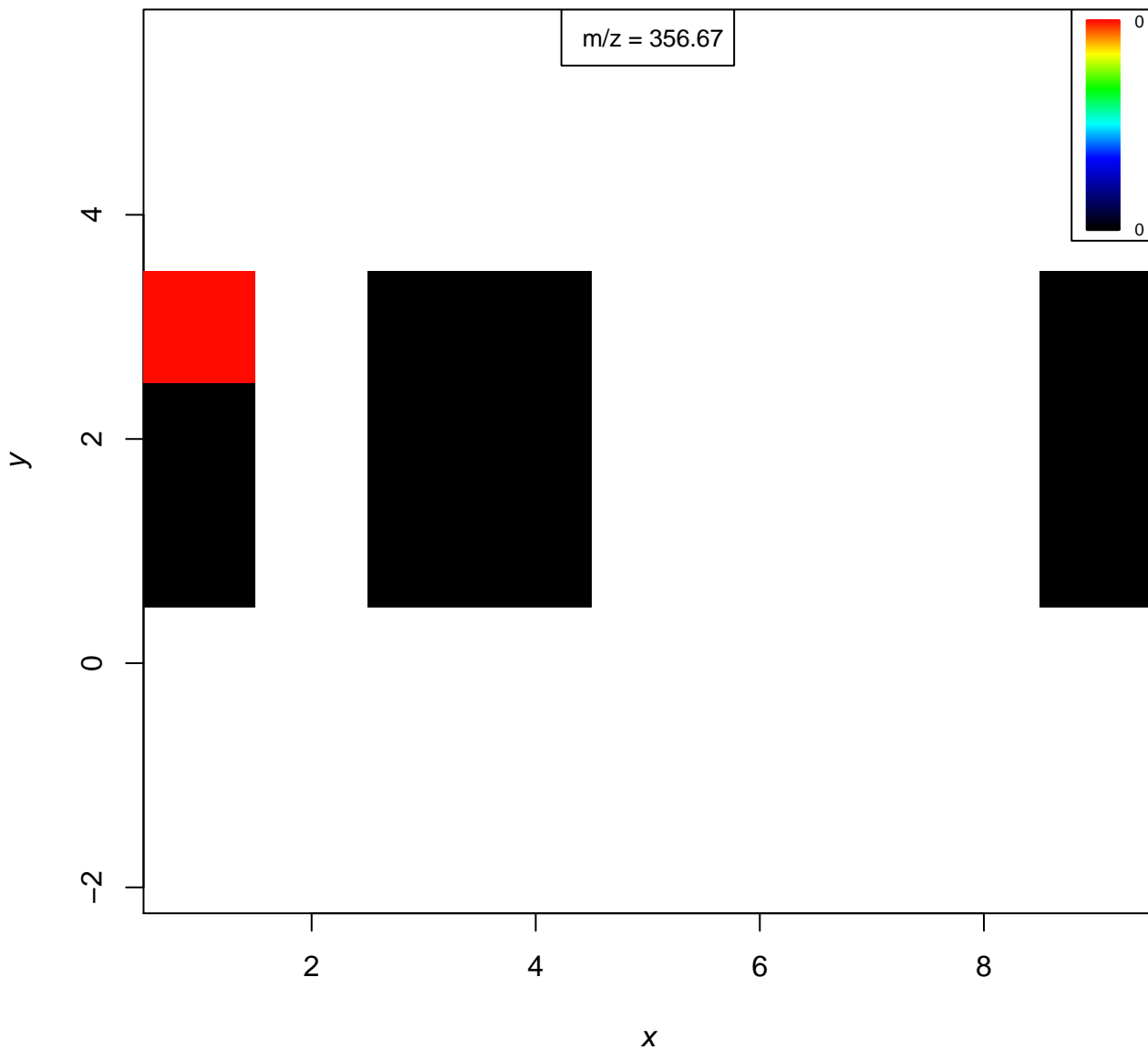
Number of calibrants per pixel (± 100 ppm)



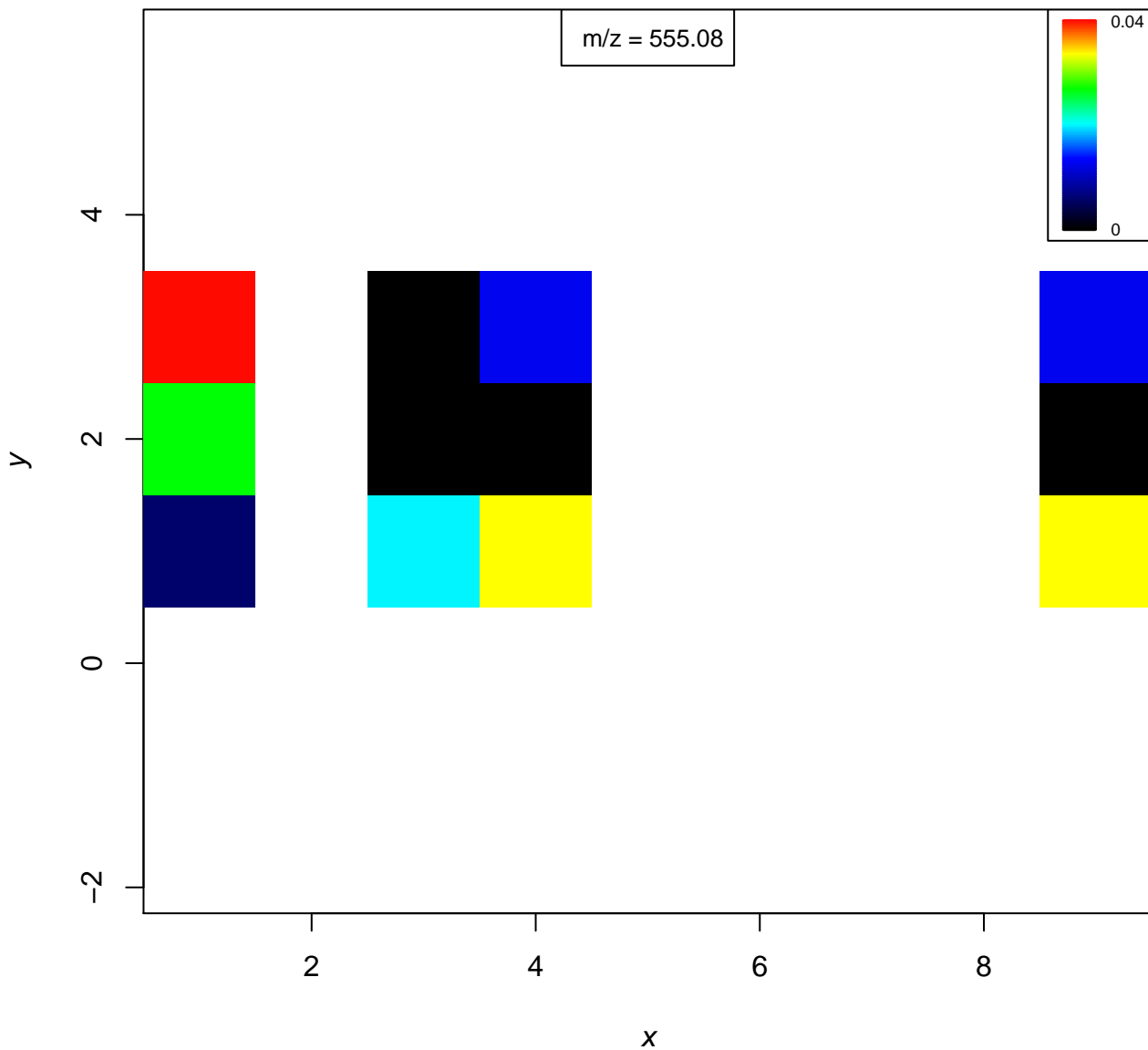
101.5: 101.5 (± 100 ppm)



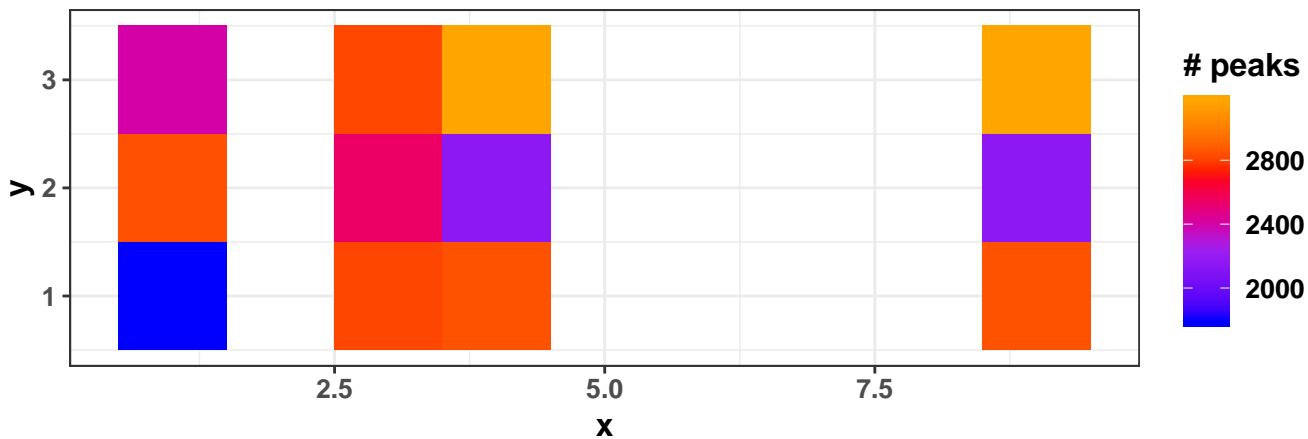
356.7: 356.7 (± 100 ppm)



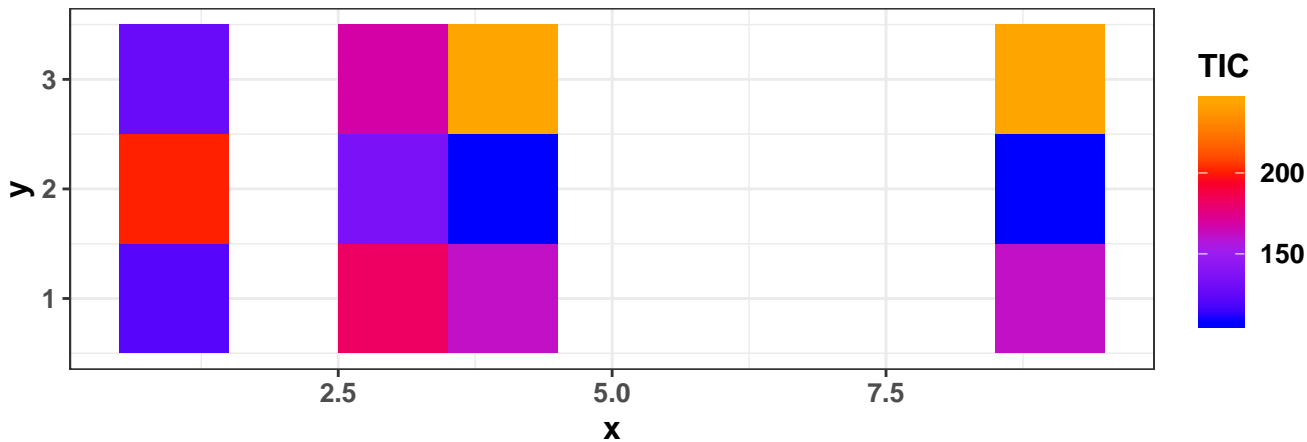
555.1: 555.1 (± 100 ppm)



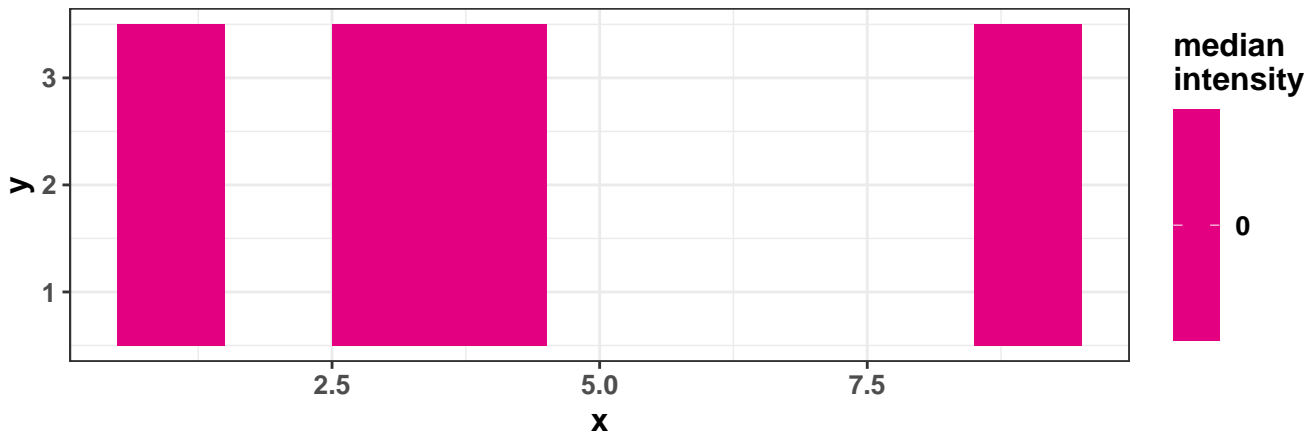
Number of peaks per spectrum



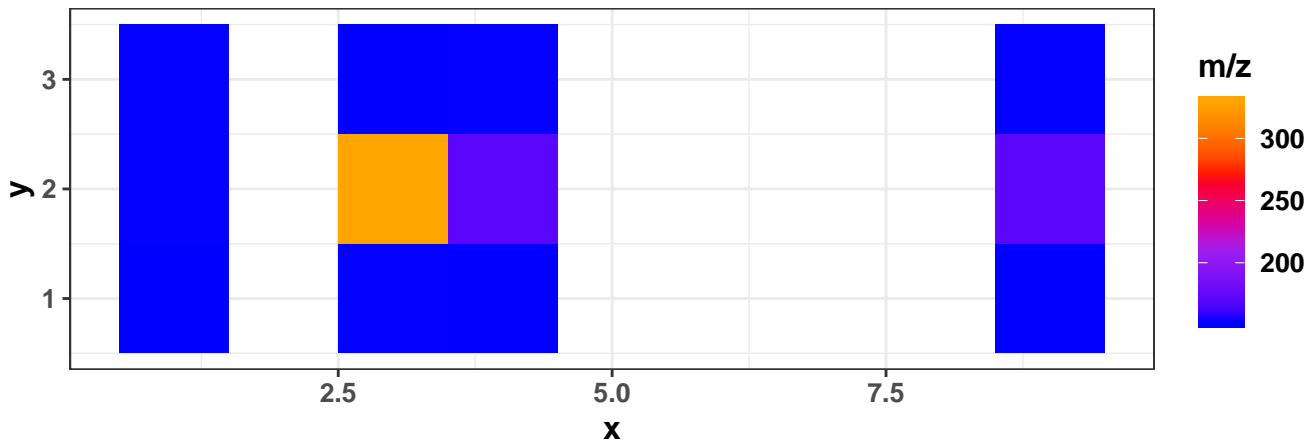
Total Ion Chromatogram



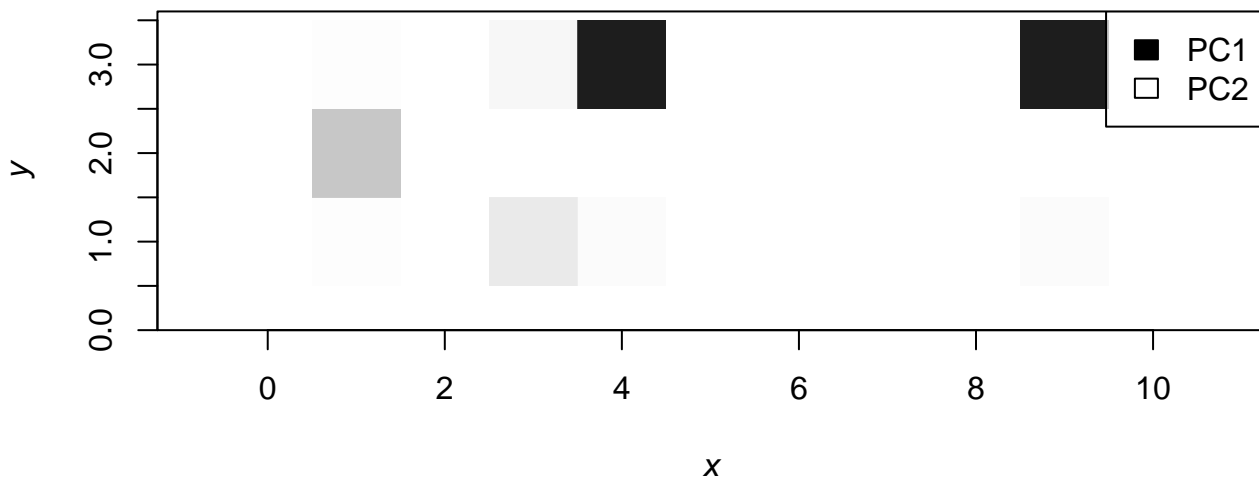
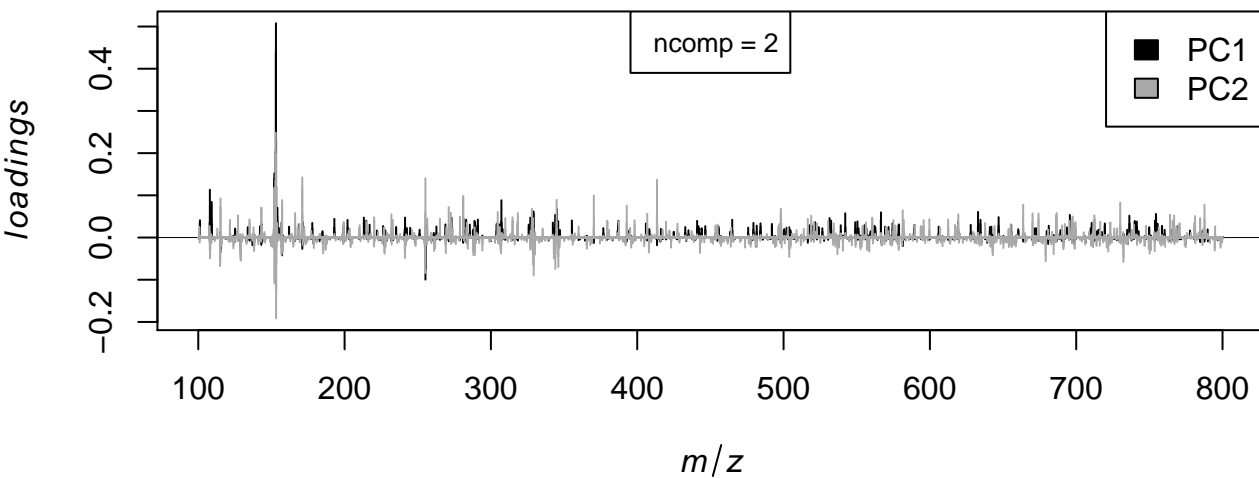
Median intensity per pixel

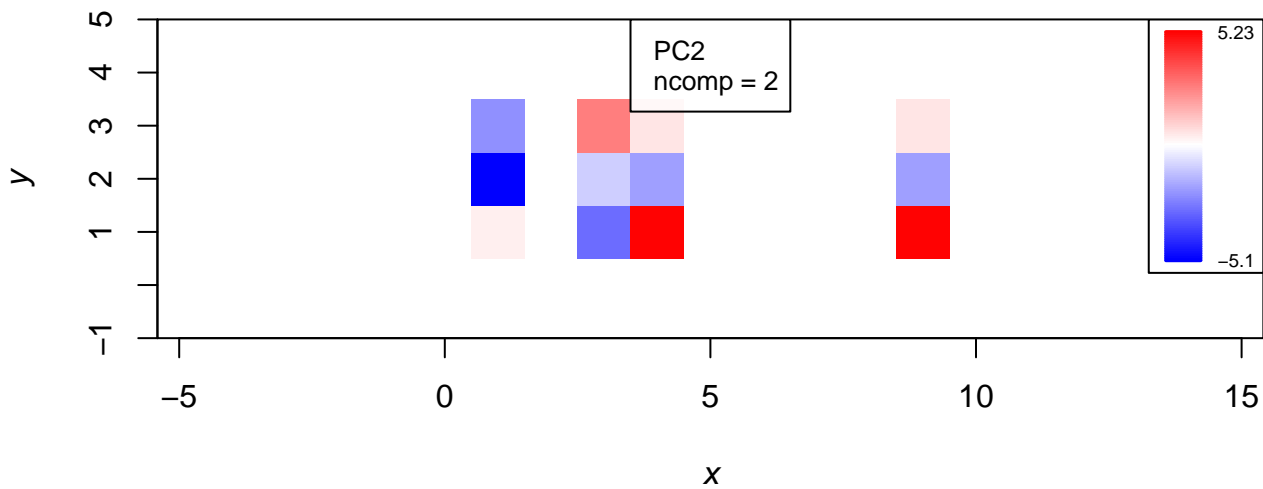
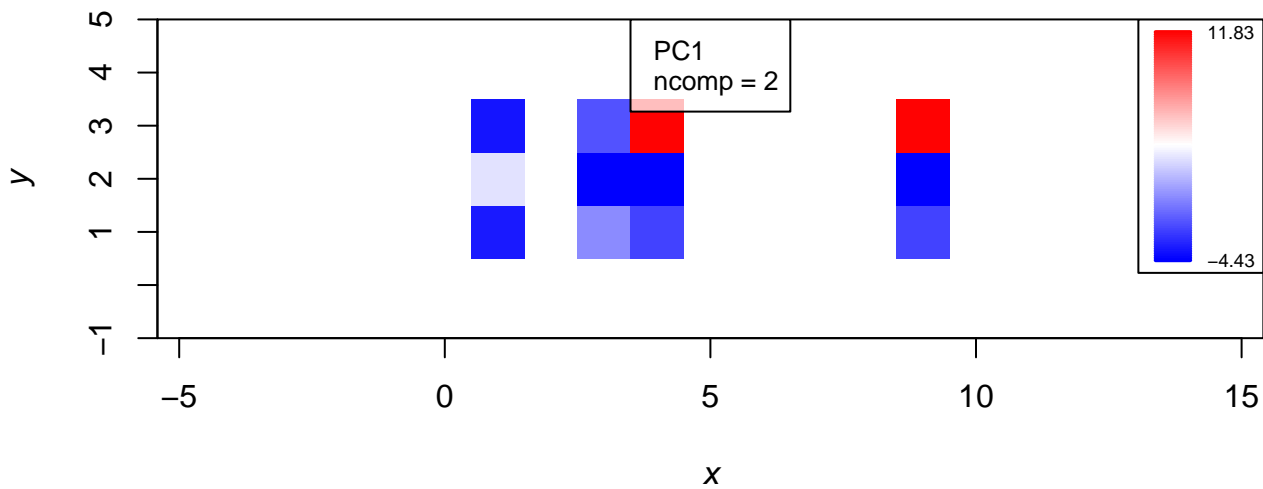


Most abundant m/z in each spectrum

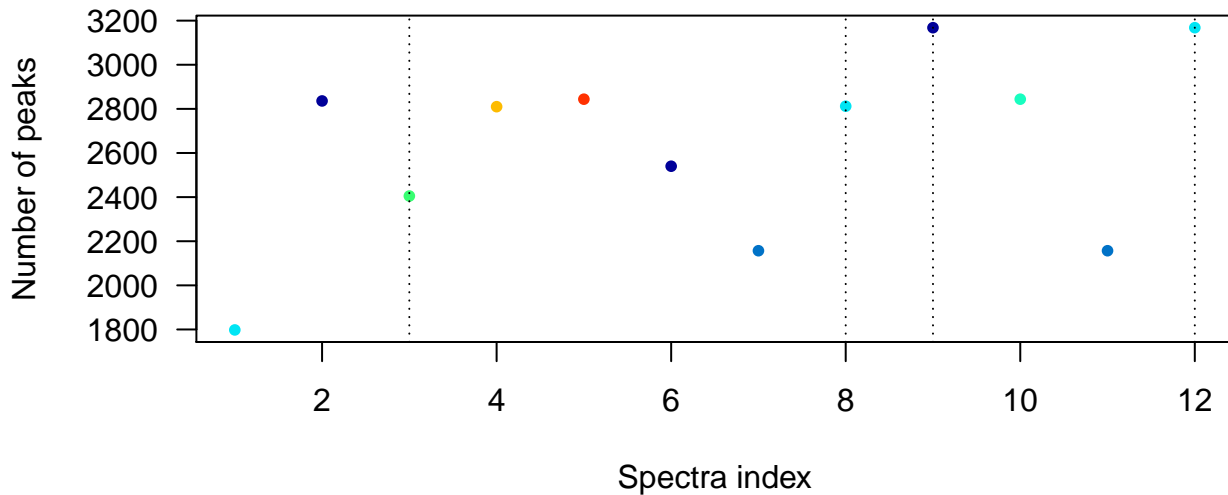


PCA for two components

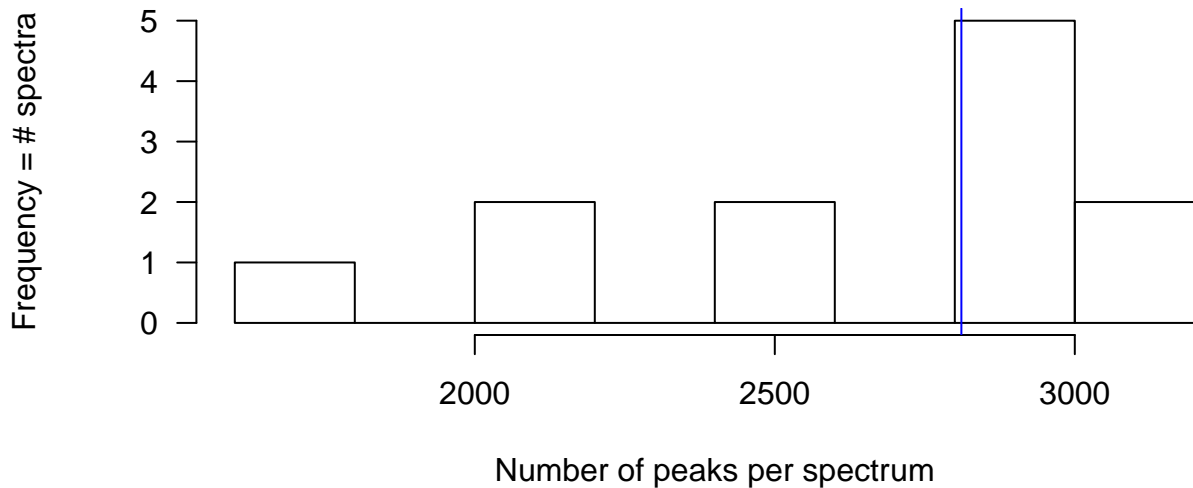




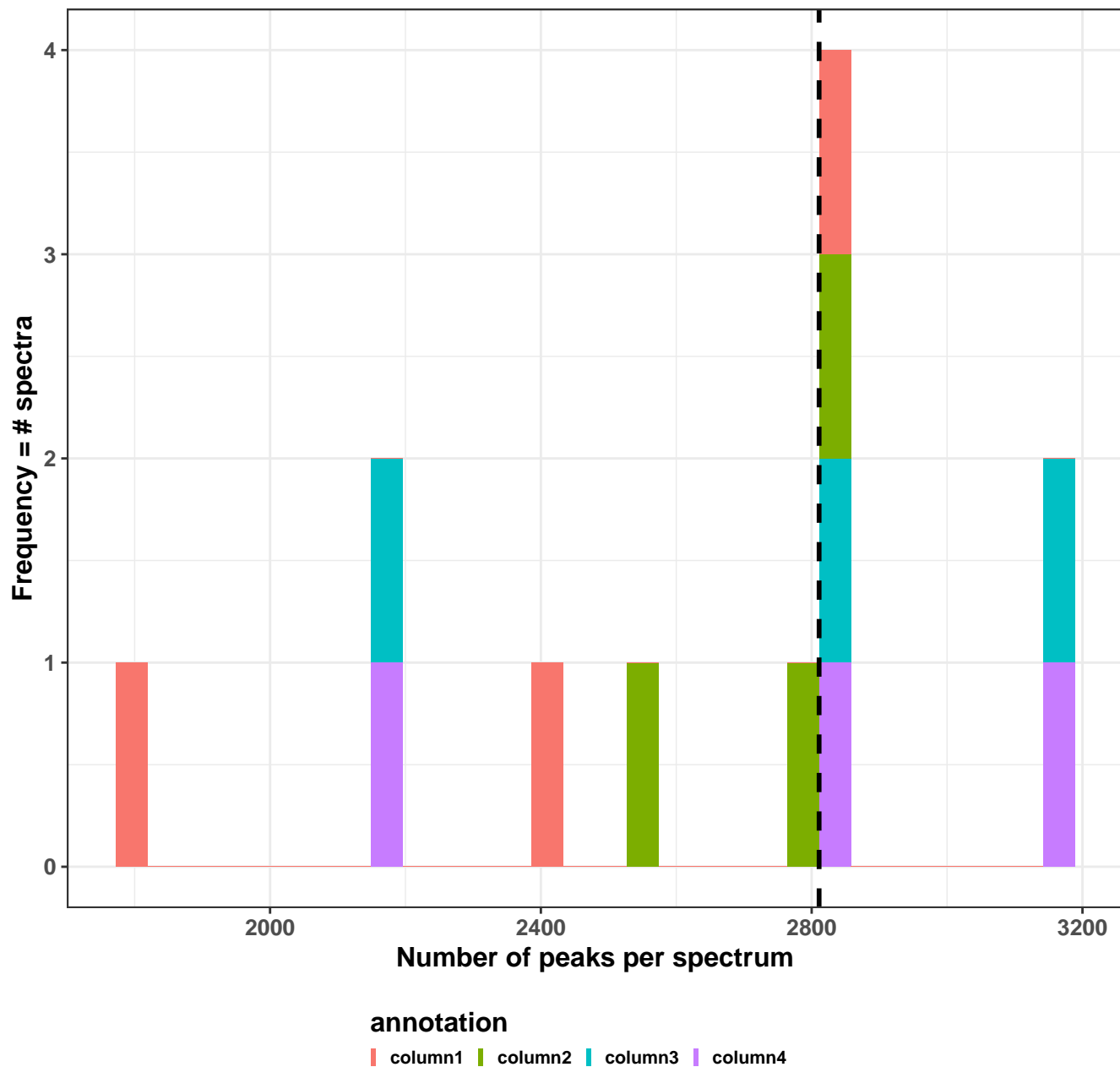
Number of peaks per spectrum

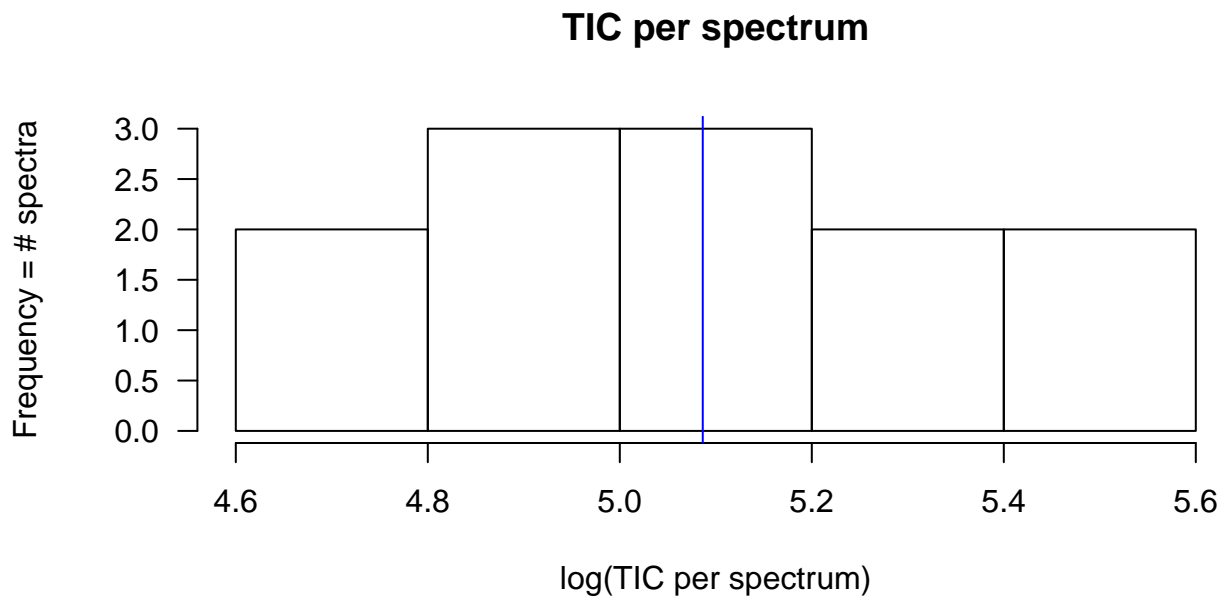
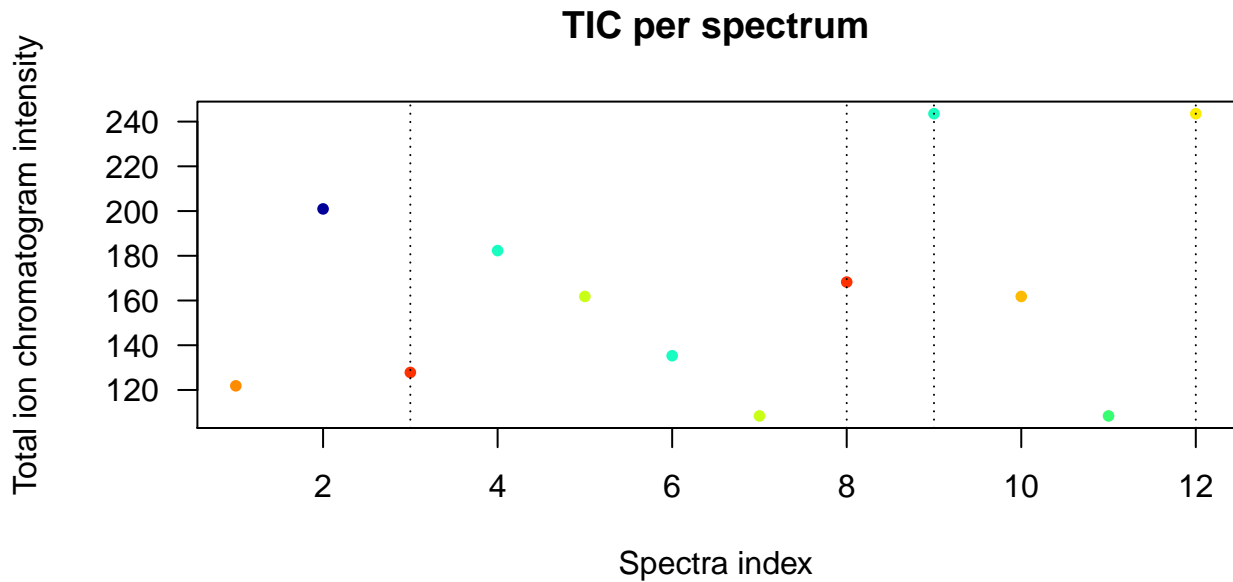


Number of peaks per spectrum

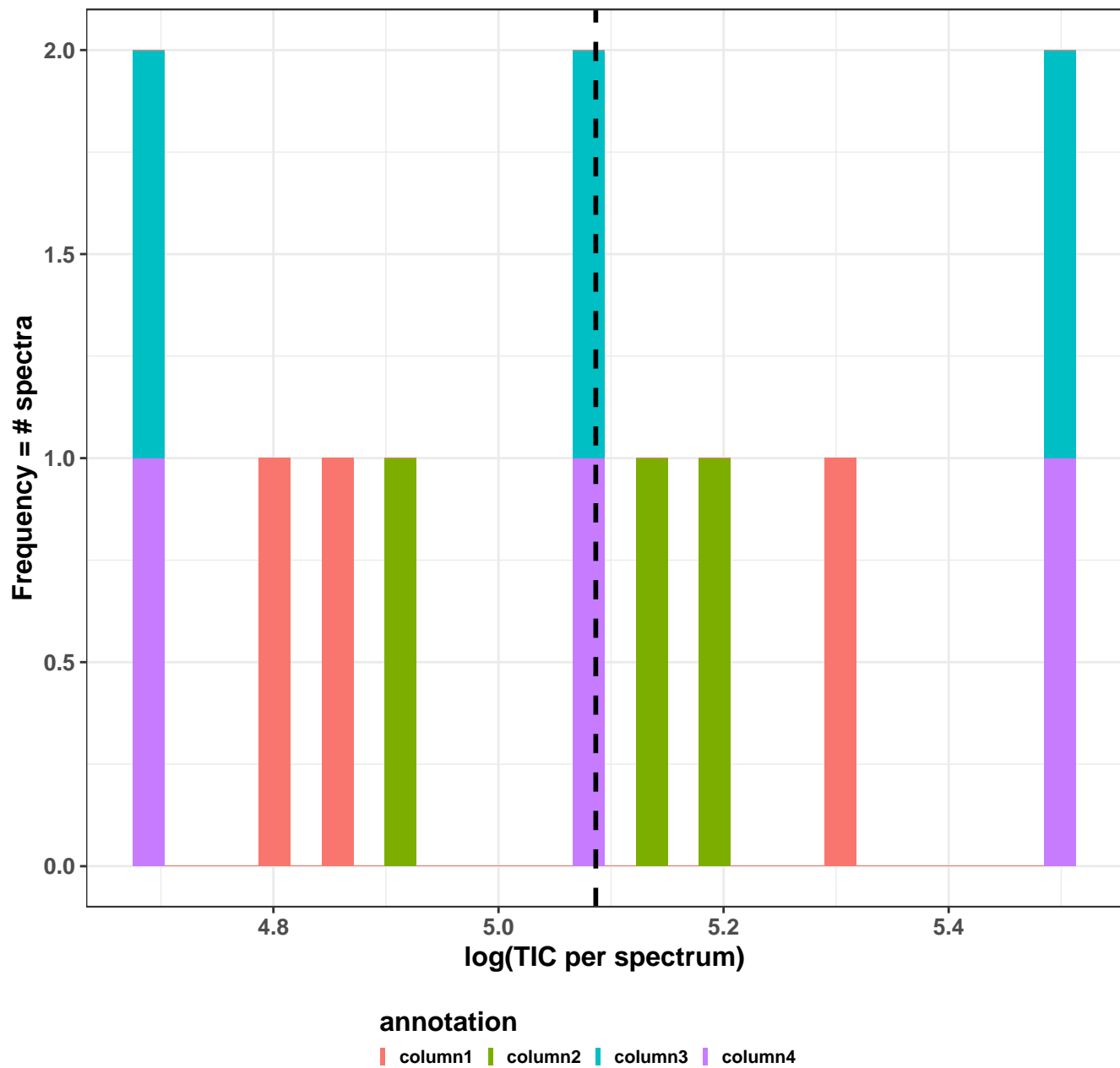


Number of peaks per spectrum and annotation group

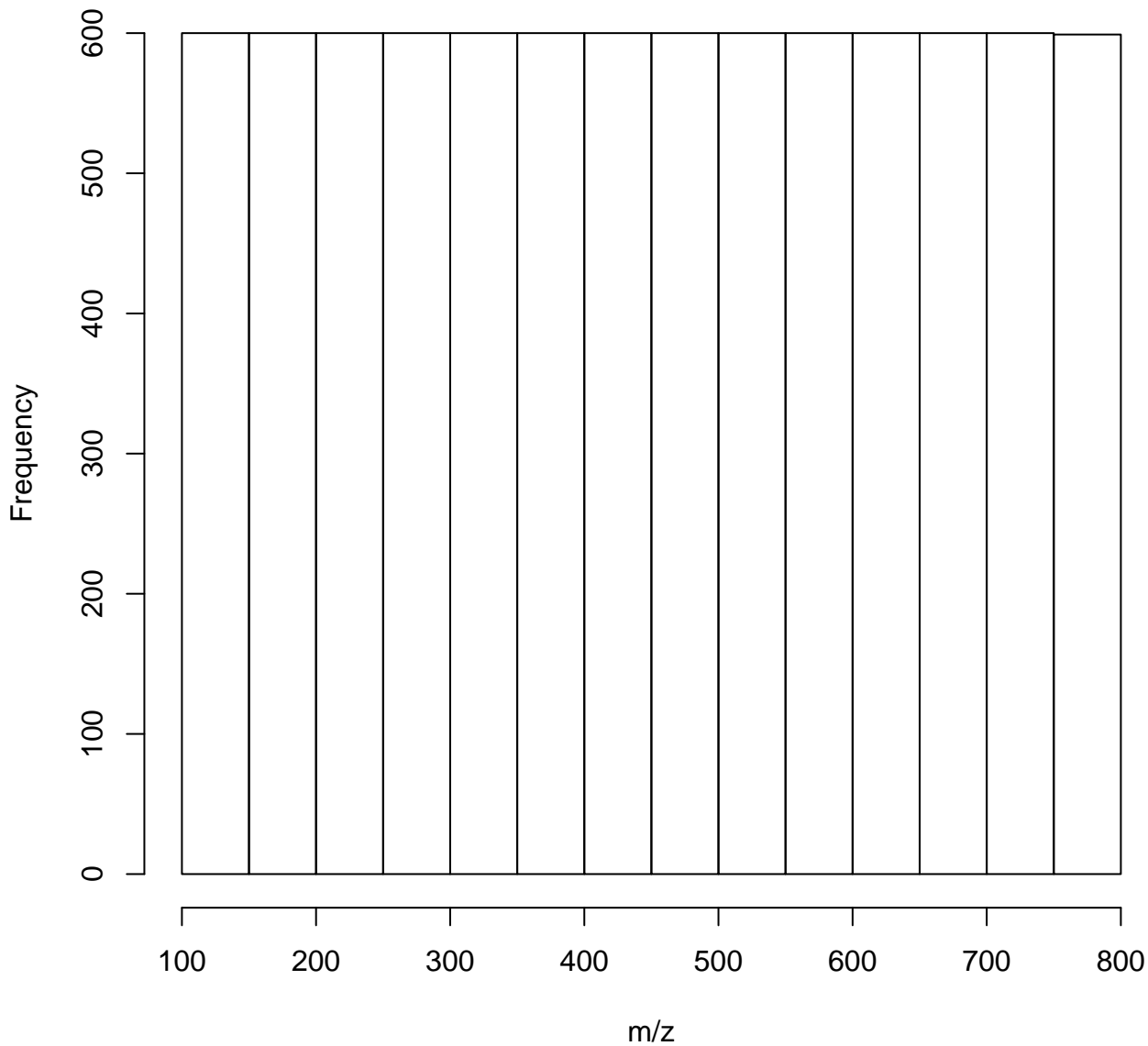




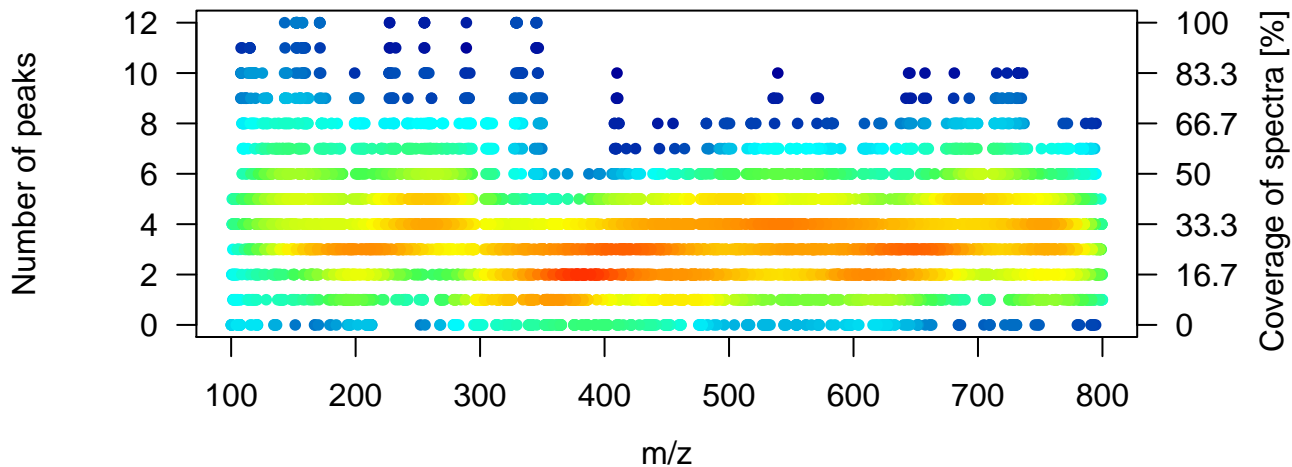
TIC per spectrum and annotation group



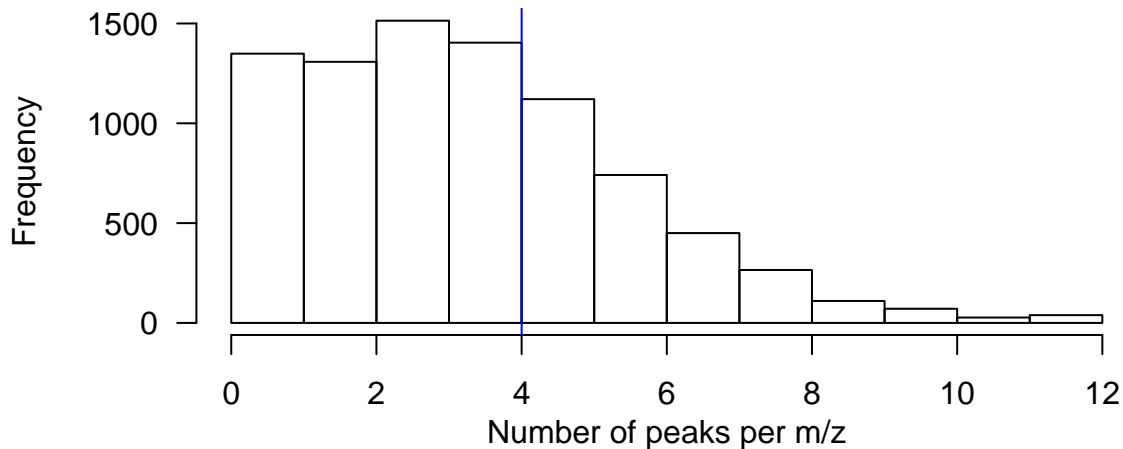
Histogram of m/z values



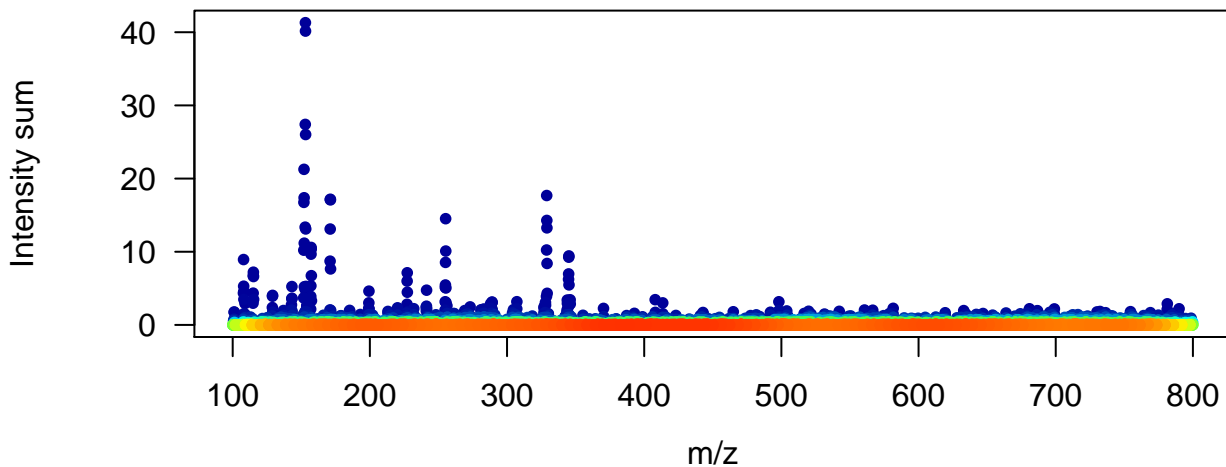
Number of peaks per m/z



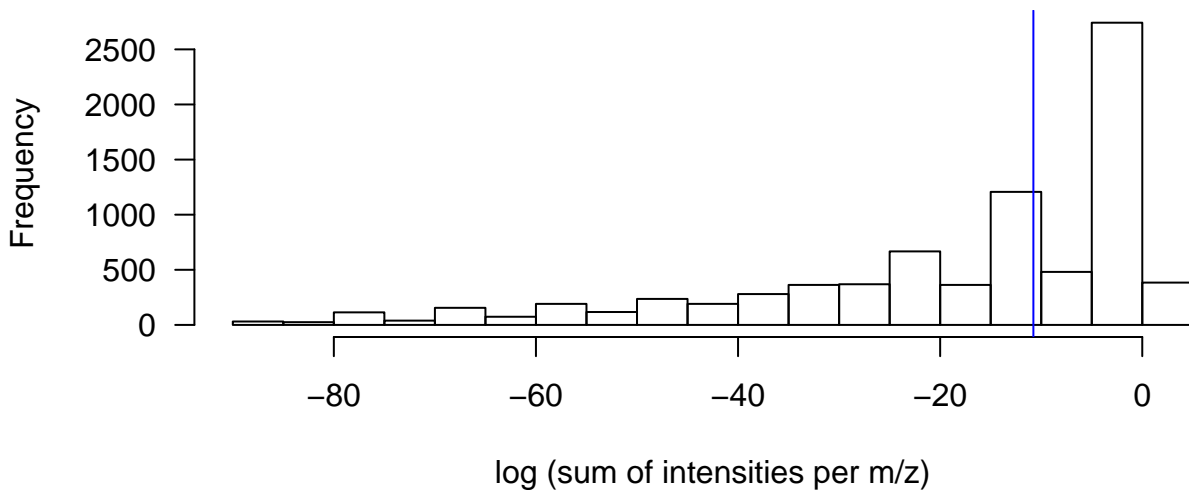
Number of peaks per m/z



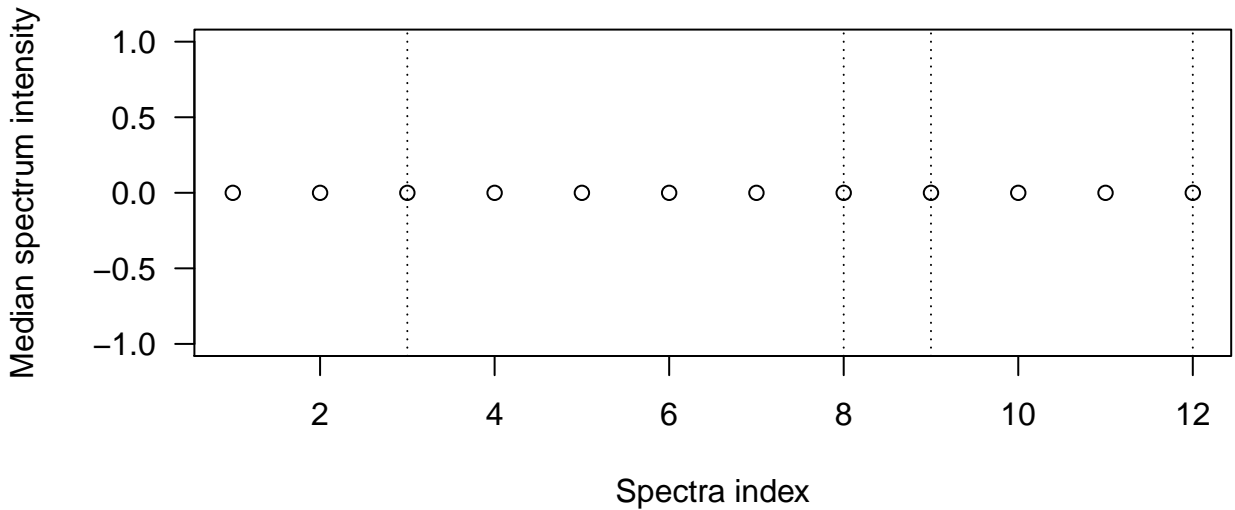
Sum of intensities per m/z



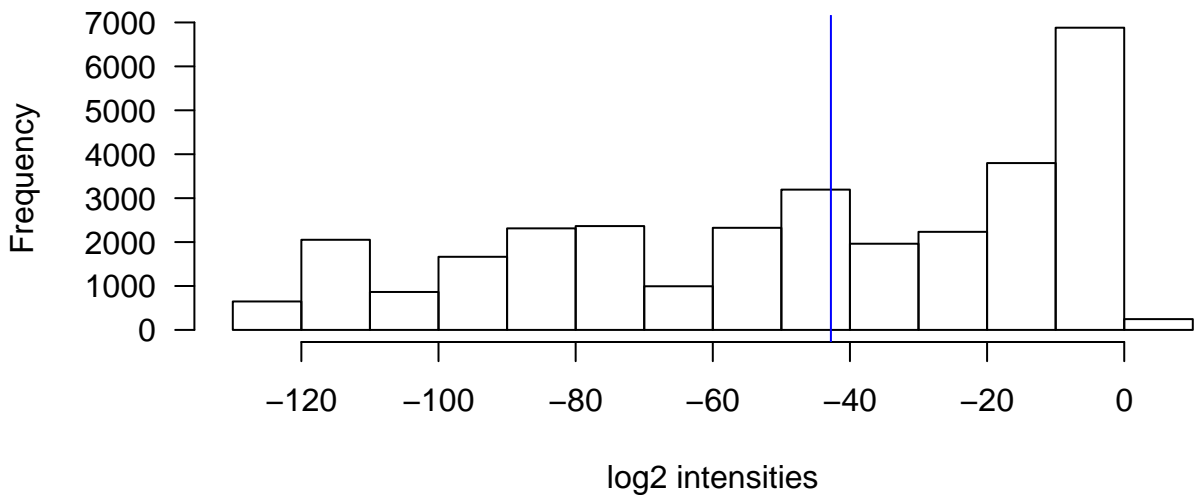
Sum of intensities per m/z



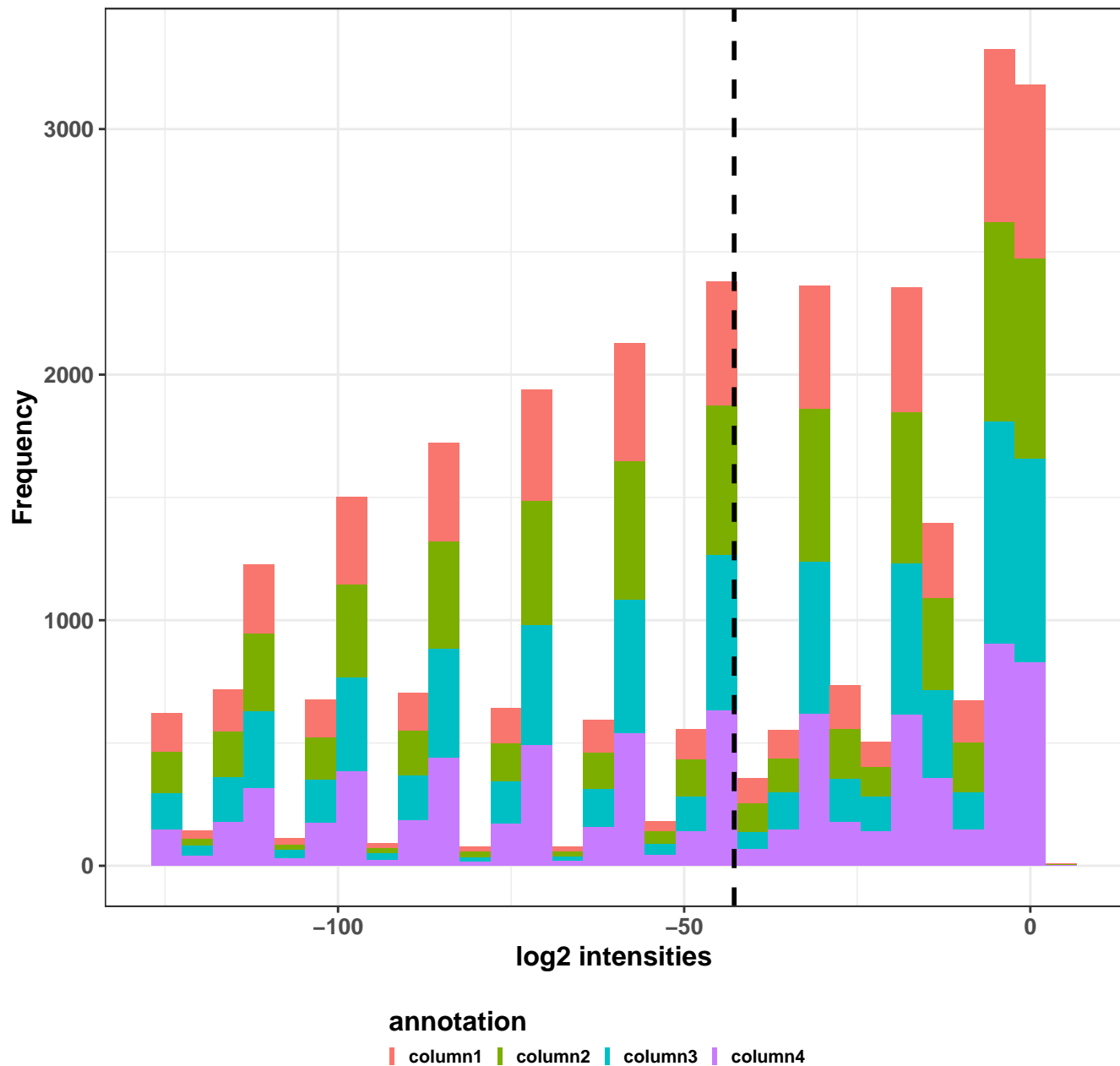
Median intensity per spectrum



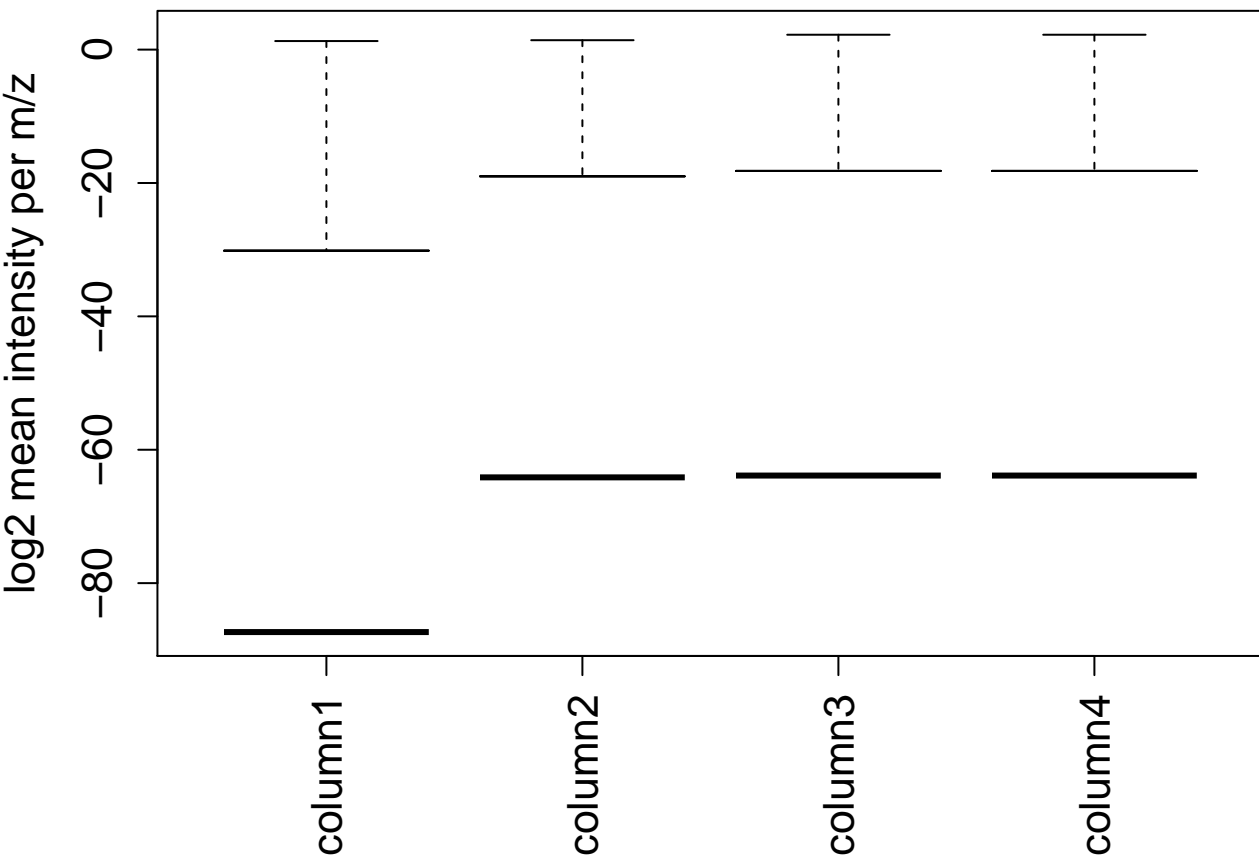
Log2-transformed intensities



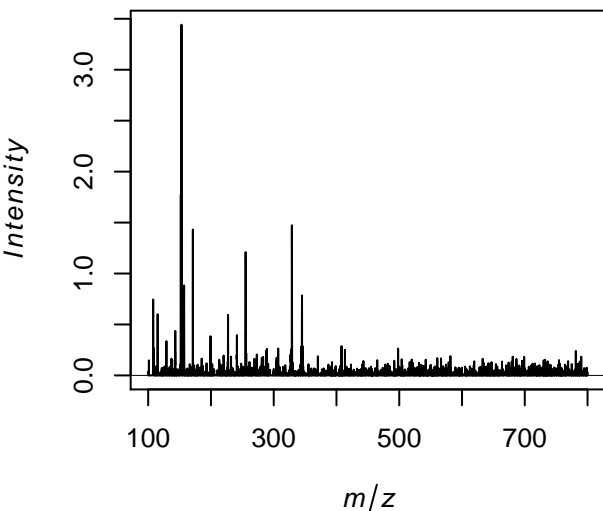
Log2-transformed intensities per sample



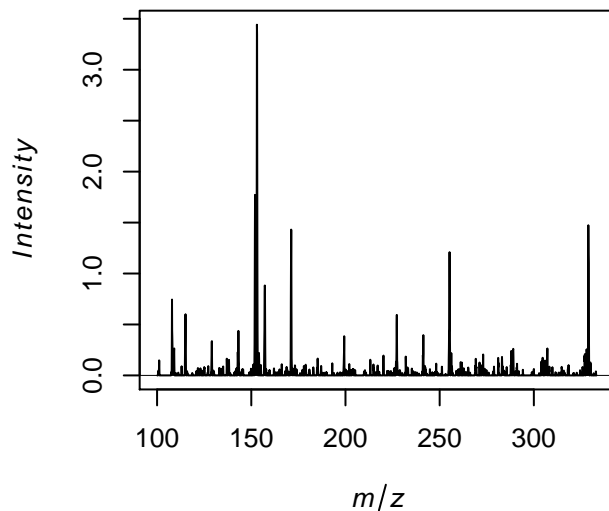
Mean m/z intensities per annotation group



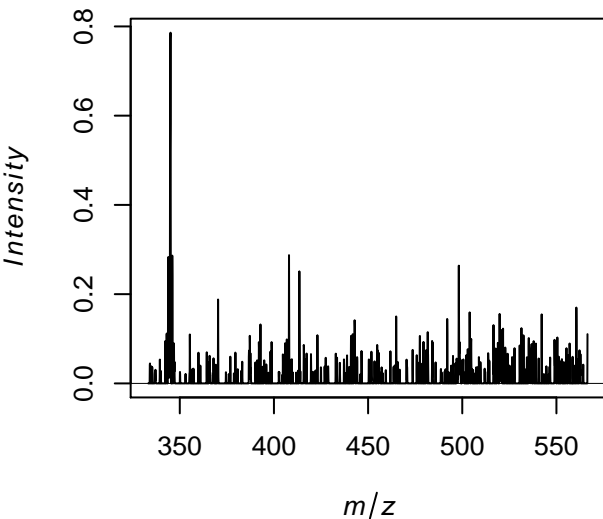
Average spectrum



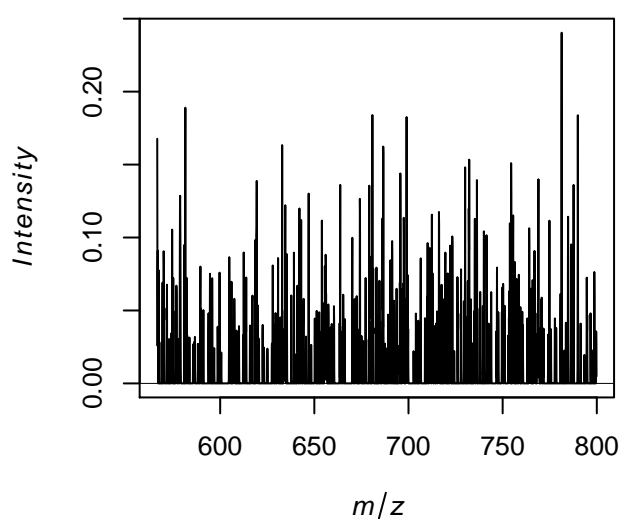
Zoomed average spectrum



Zoomed average spectrum



Zoomed average spectrum

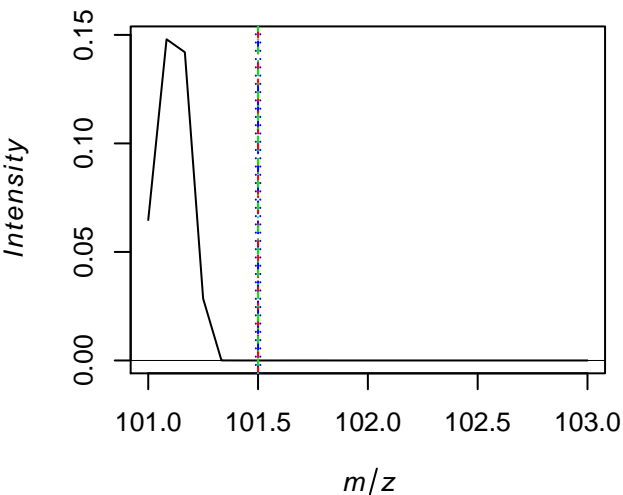


theor. m/z: 101.5

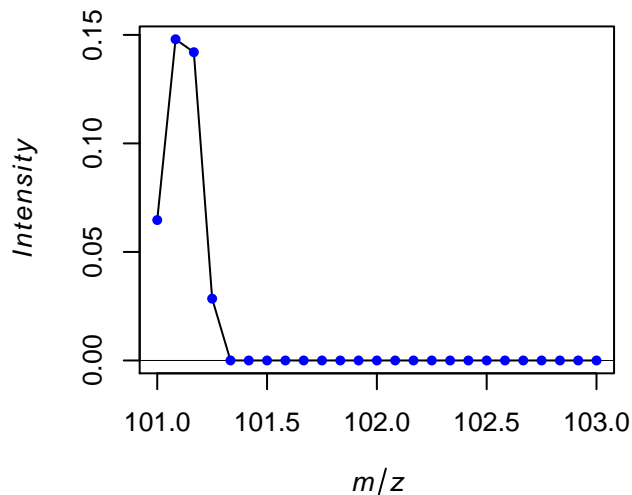
most abundant m/z: 101.5

closest m/z: 101.5

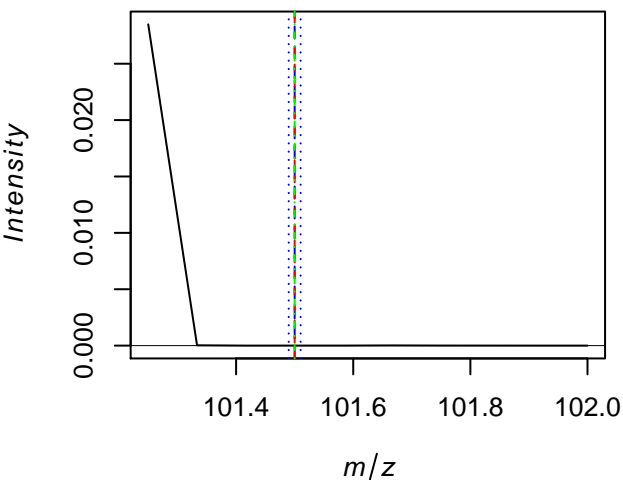
Average spectrum



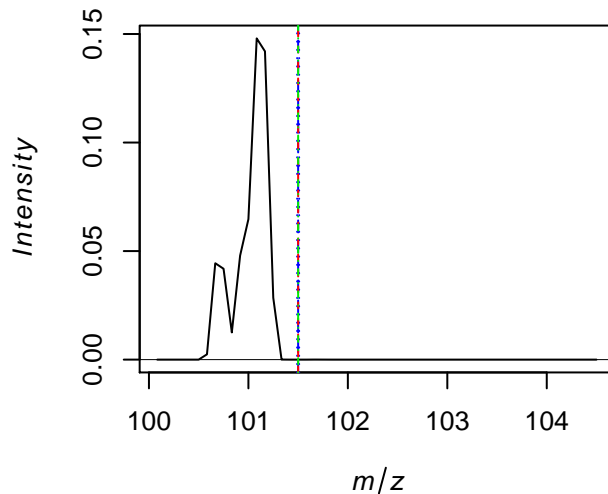
Average spectrum with data points



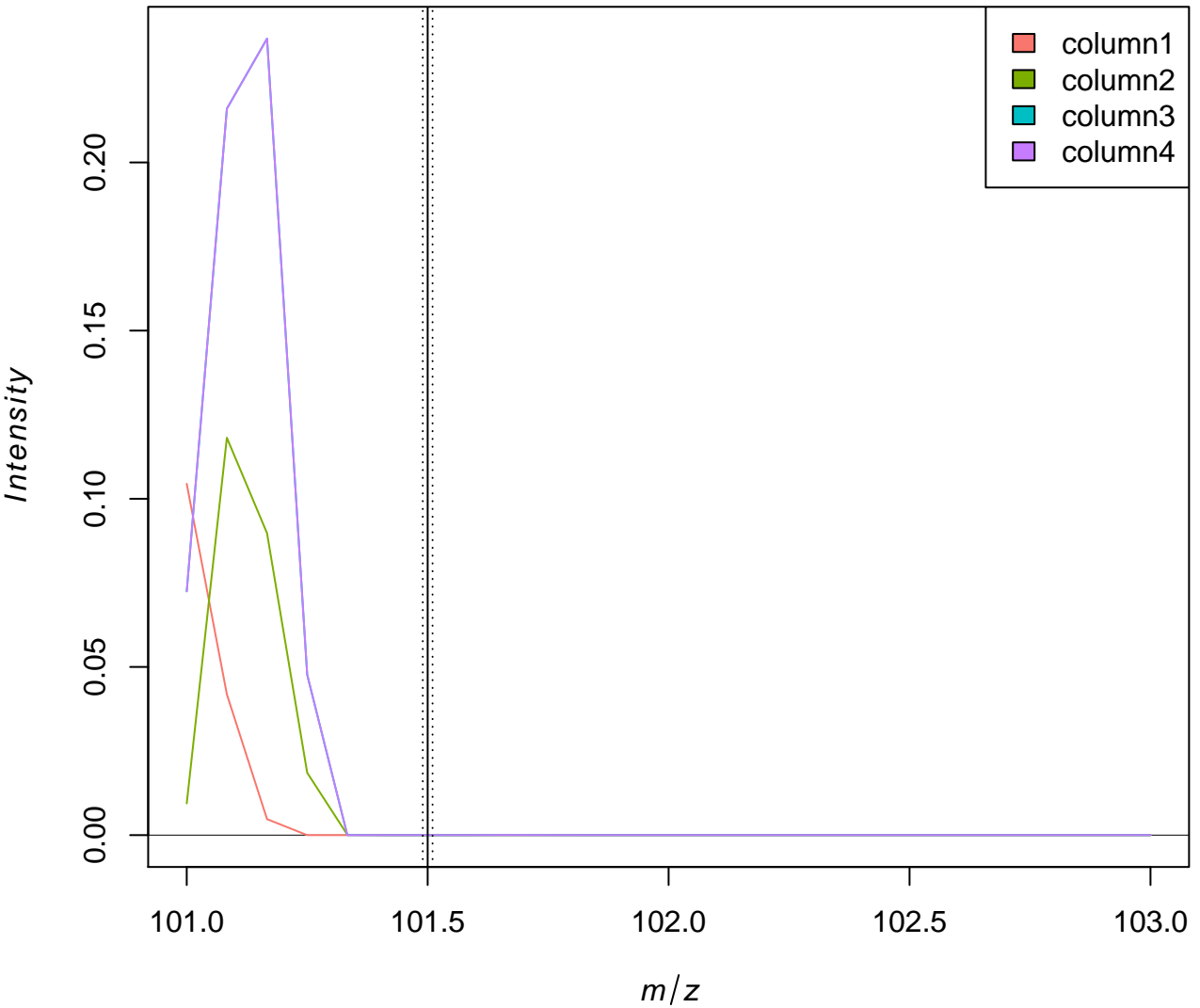
Average spectrum



Average spectrum



Average spectrum per annotation group

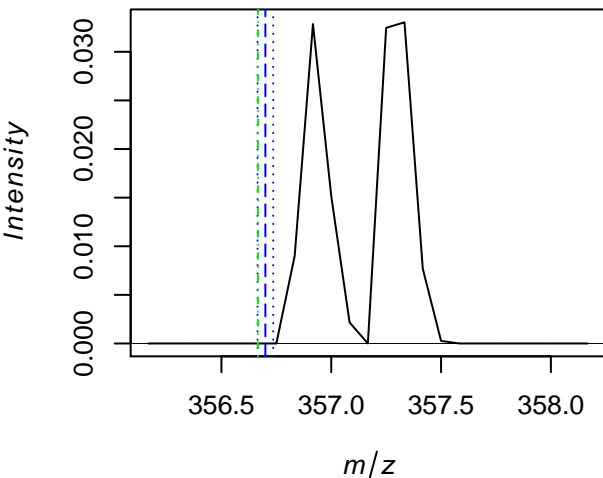


theor. m/z: 356.7

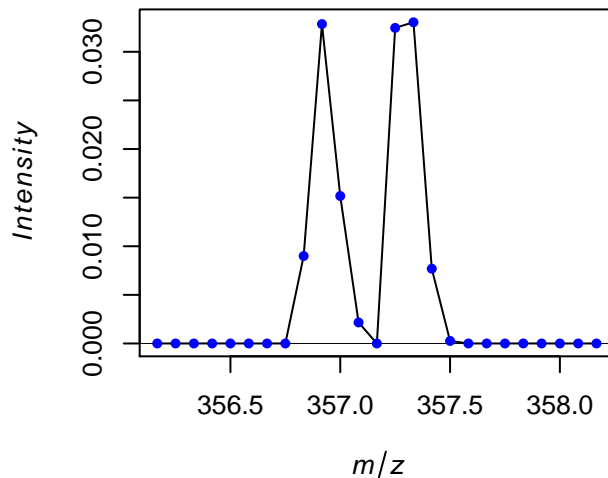
most abundant m/z: NA

closest m/z: 356.6667

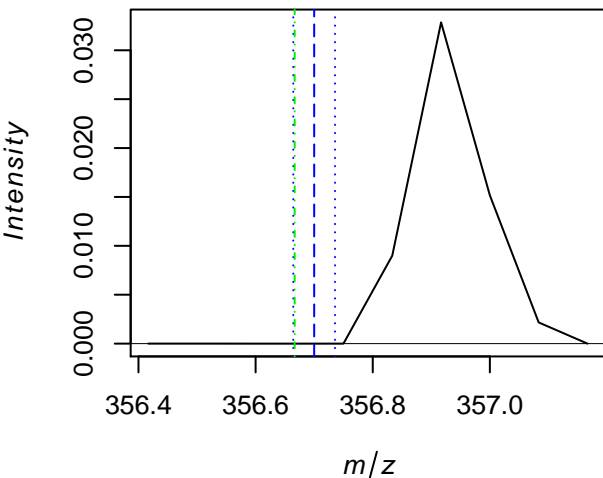
Average spectrum



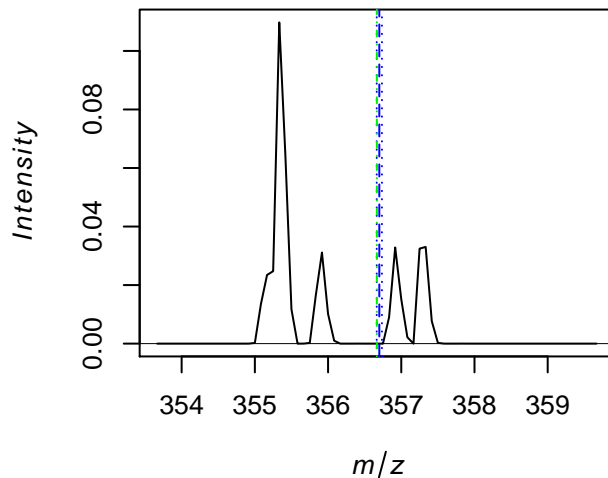
Average spectrum with data points



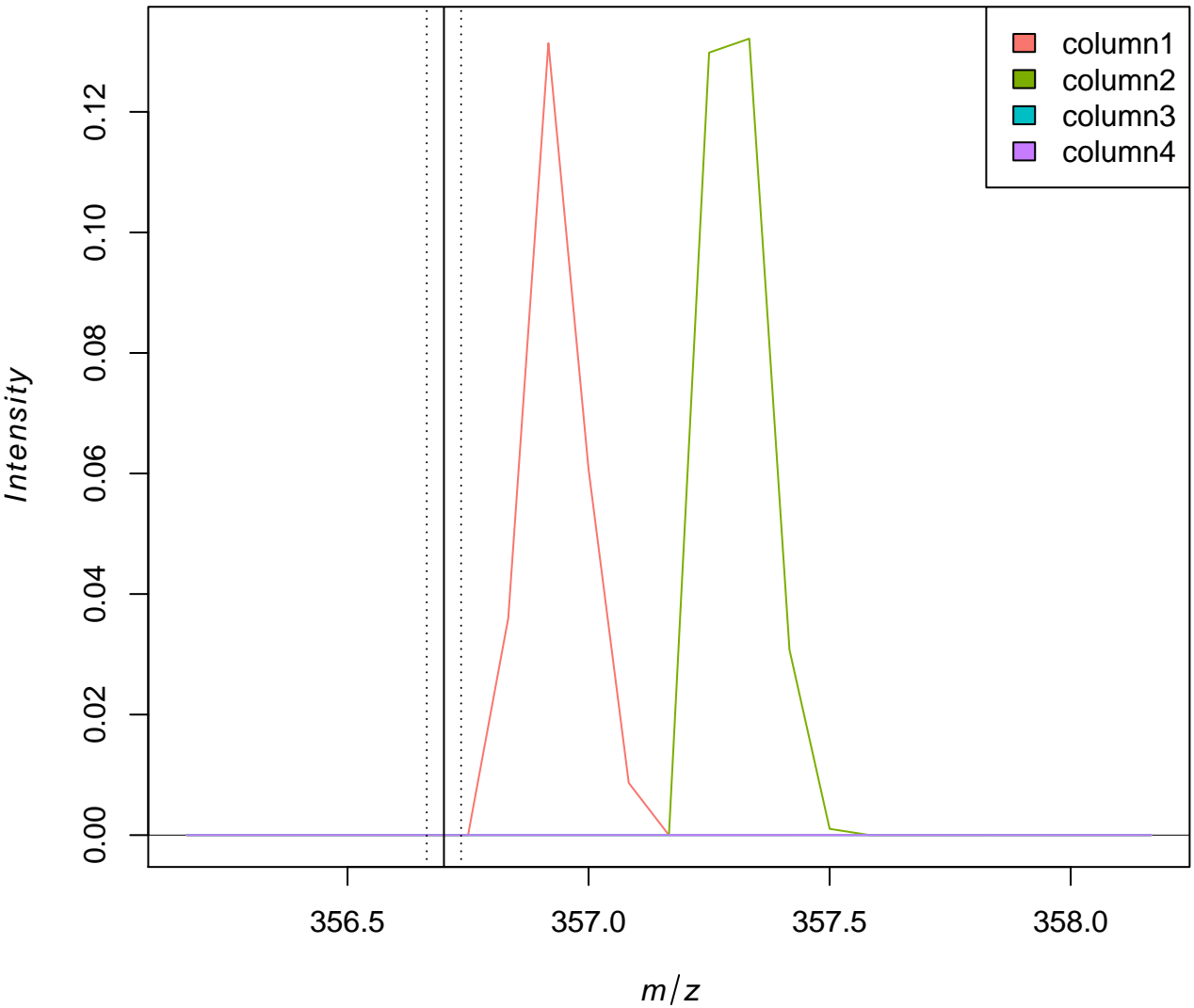
Average spectrum



Average spectrum



Average spectrum per annotation group

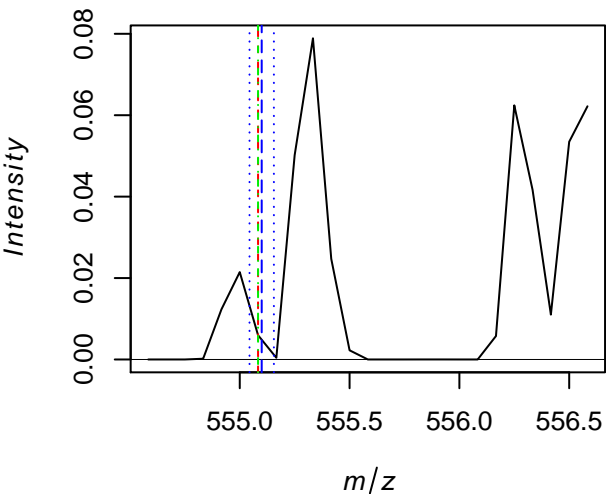


theor. m/z: 555.1

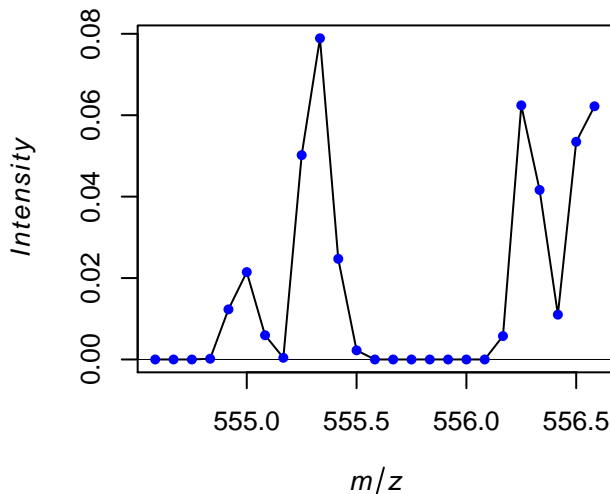
most abundant m/z: 555.0834

closest m/z: 555.0834

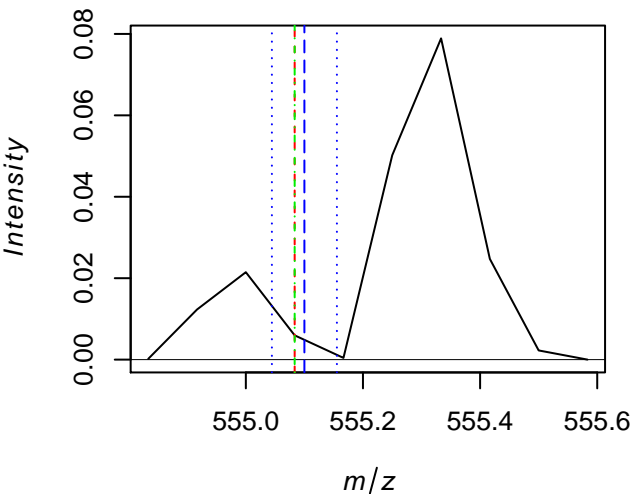
Average spectrum



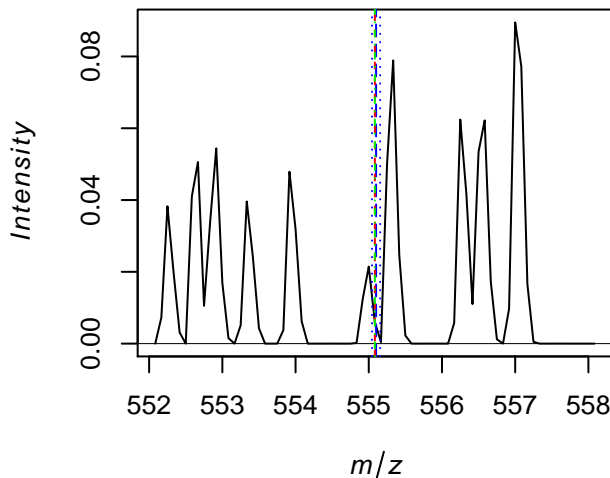
Average spectrum with data points



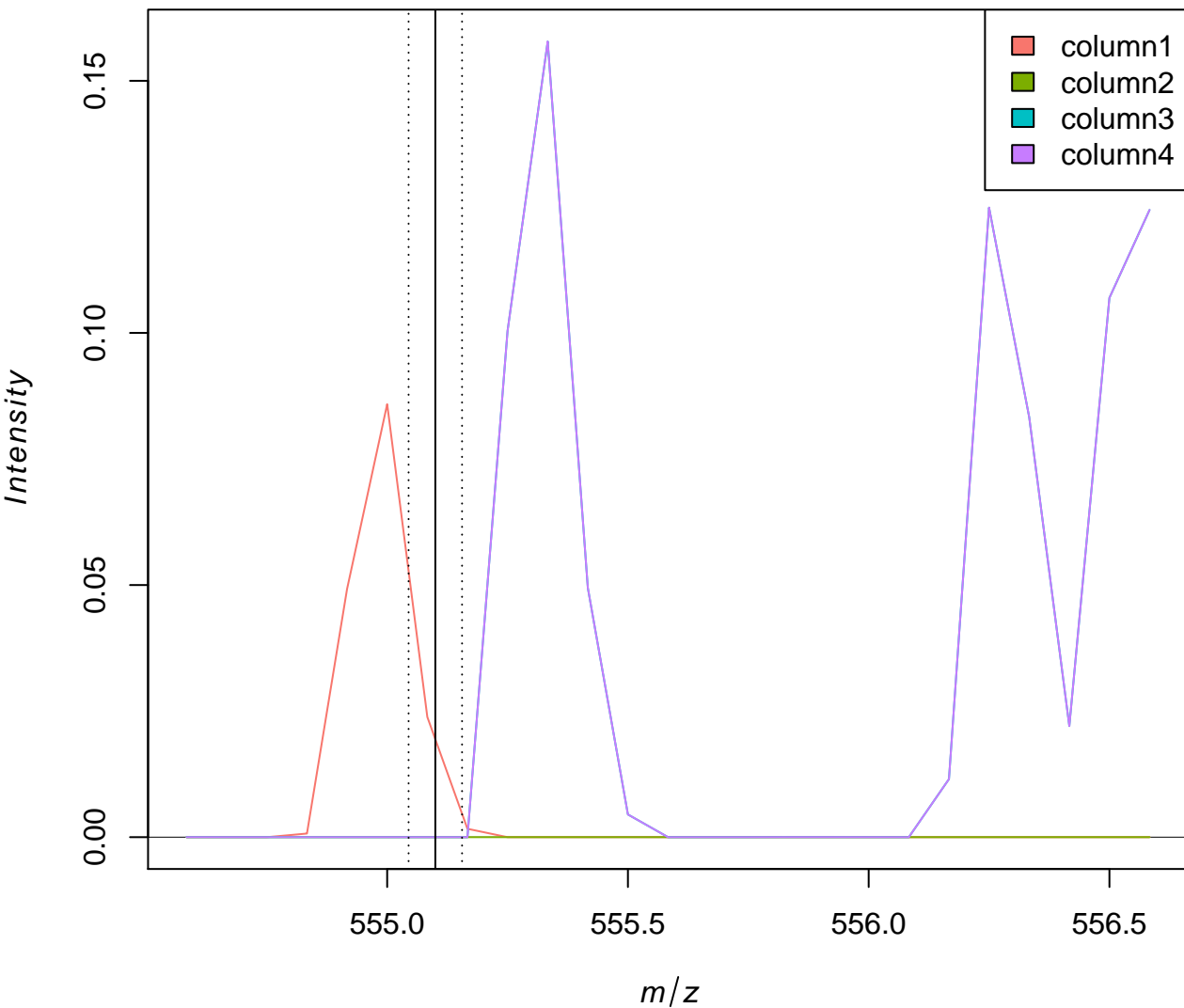
Average spectrum



Average spectrum

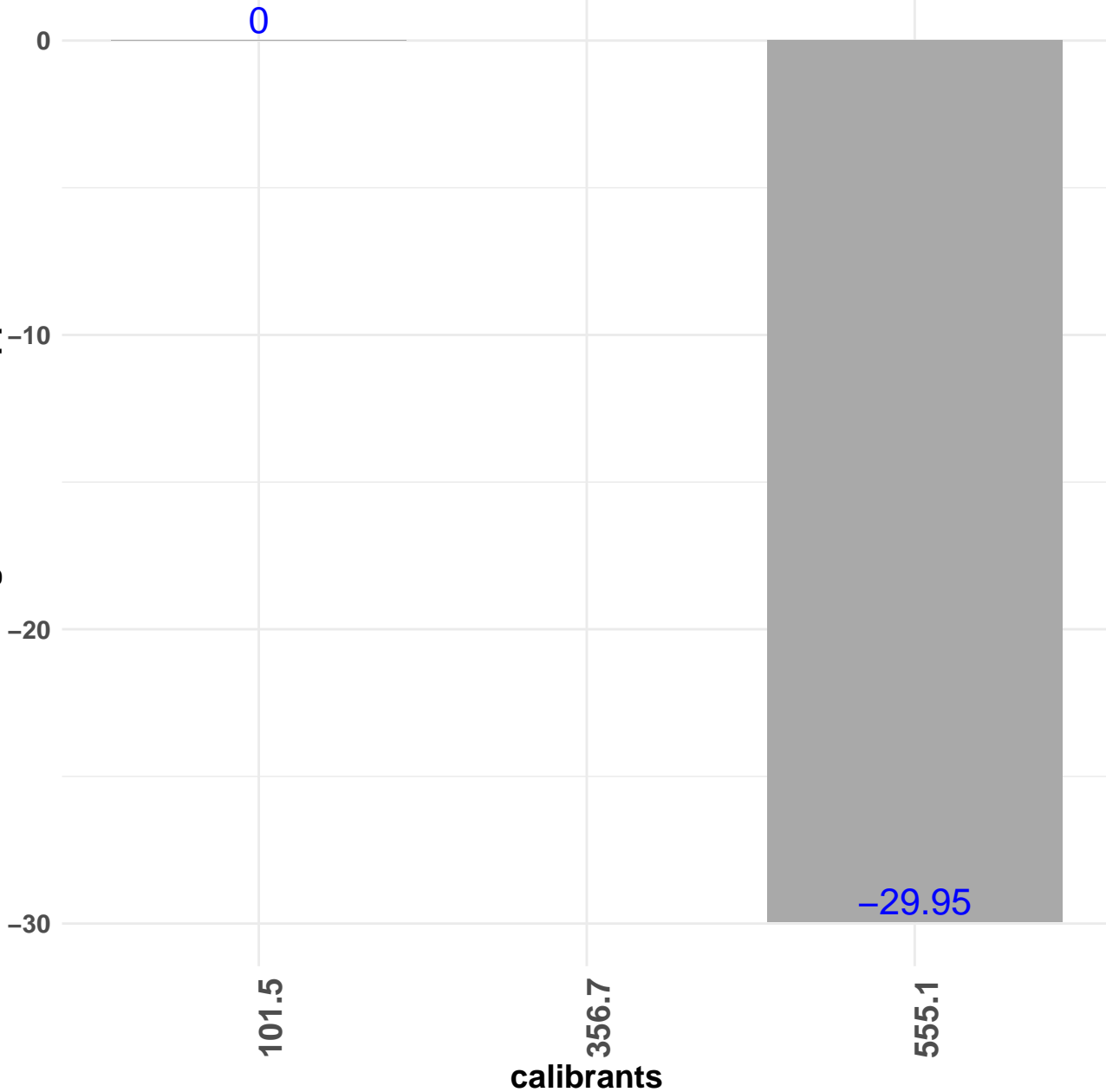


Average spectrum per annotation group

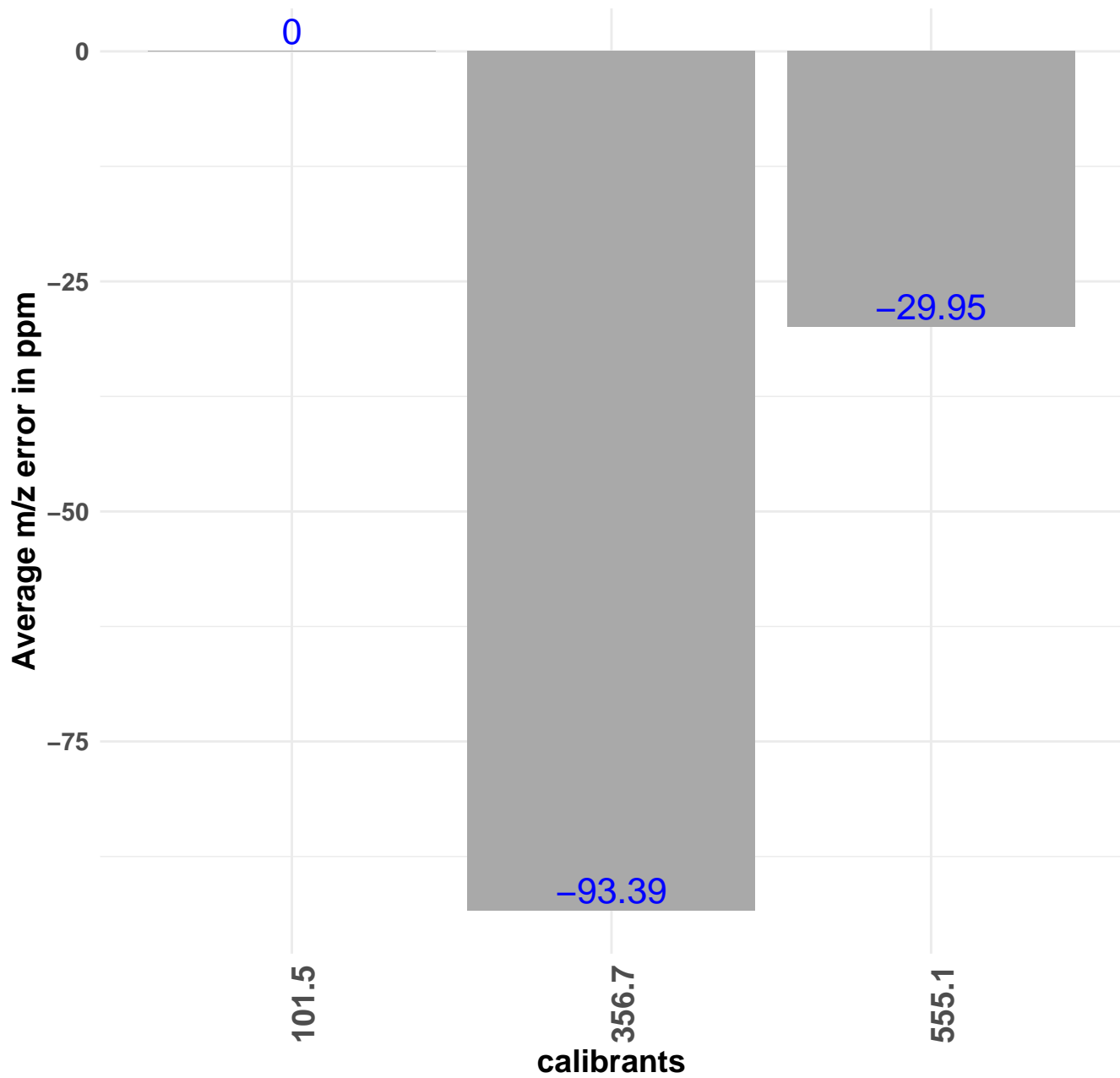


Average m/z error (max. average intensity vs. theor. calibrant m/z)

Average m/z error in ppm



Average m/z error (closest measured m/z vs. theor. calibrant m/z)



**Difference m/z with max. average intensity vs. theor. m/z
(per spectrum)**

