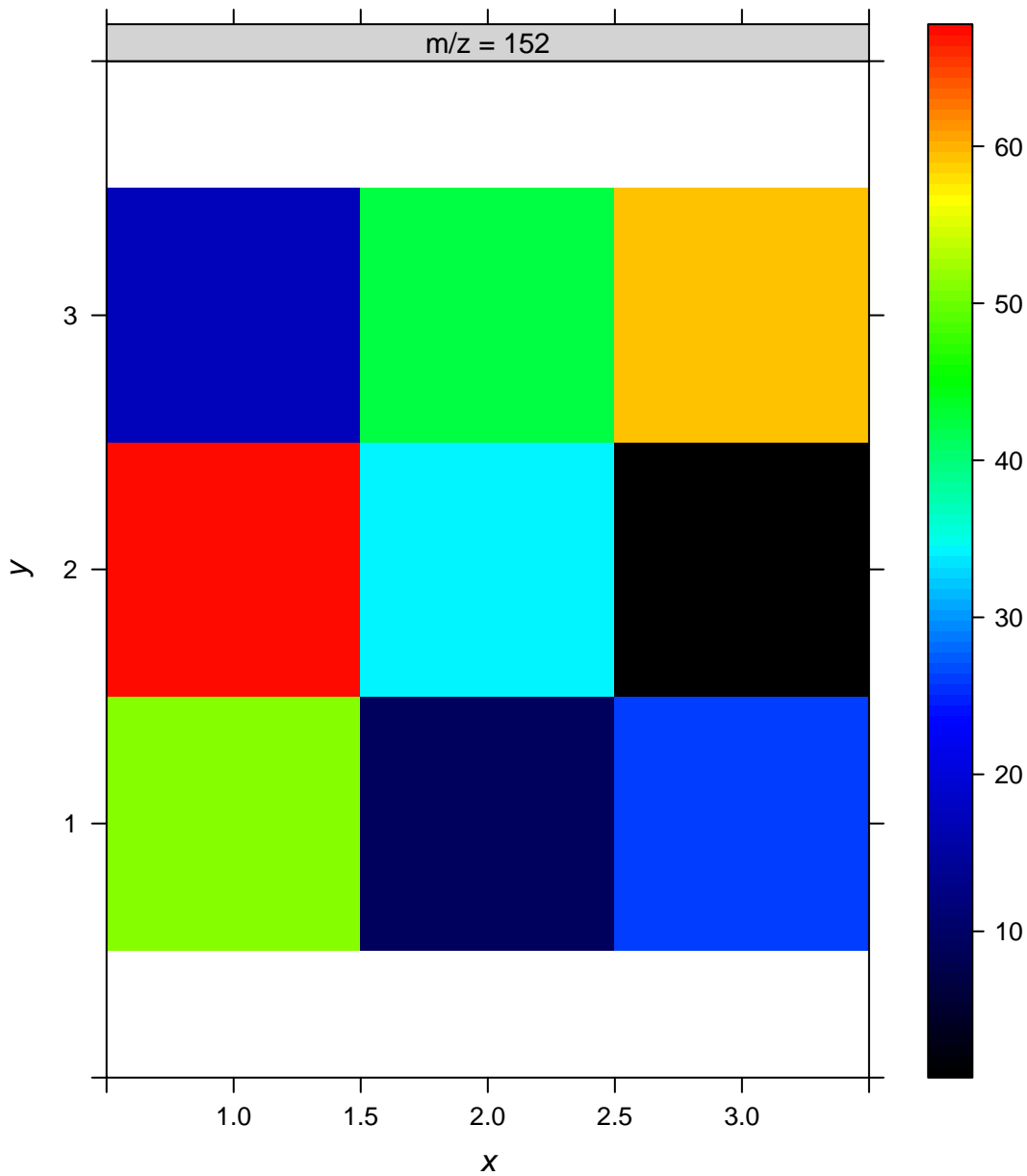


Quality control of MSI data

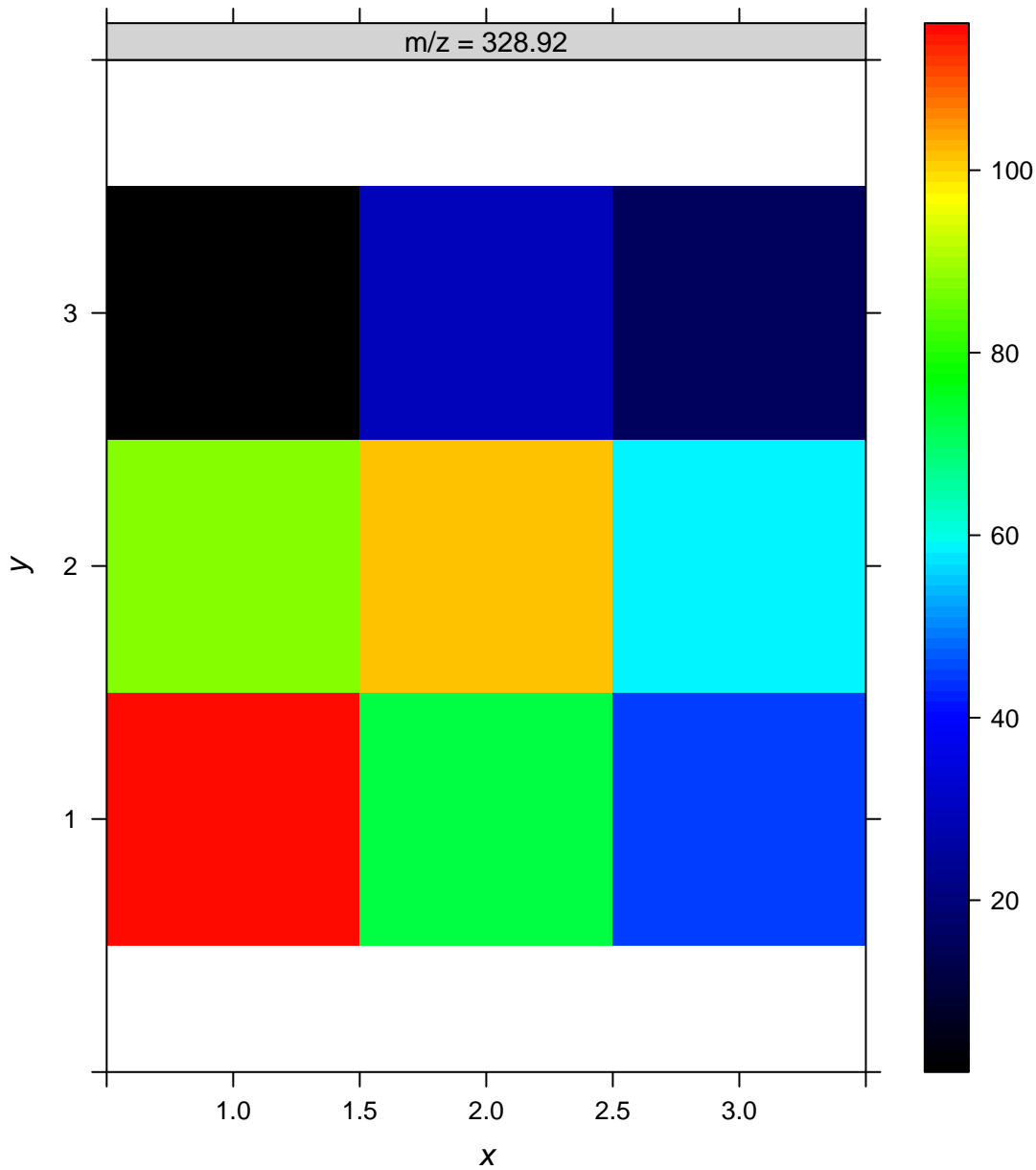
Filename: Testfile_rdata

| properties | values |
|------------------------------------|-----------------|
| Number of mz features | 8399 |
| Range of mz values [Da] | 100.08 – 799.92 |
| Number of pixels | 9 |
| Range of x coordinates | 1 – 3 |
| Range of y coordinates | 1 – 3 |
| Range of intensities | 0 – 318.82 |
| Median of intensities | 0 |
| Intensities > 0 | 30.92 % |
| Number of zero TICs | 0 |
| Preprocessing | |
| Normalization | tic |
| Smoothing | FALSE |
| Baseline reduction | FALSE |
| Peak picking | FALSE |
| Centroided | FALSE |
| # valid peptides in Testfile_rdata | 3/3 |

1) mass1 (152 ± 0.1 Da)



2) mass2 (328.9 ± 0.1 Da)



3) mass3 (185.2 ± 0.1 Da)

