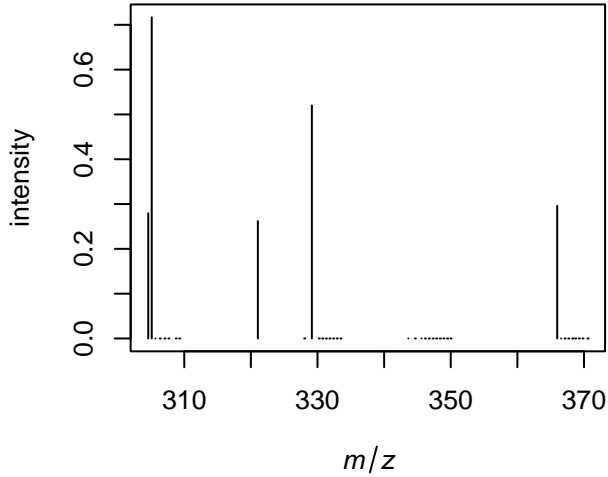


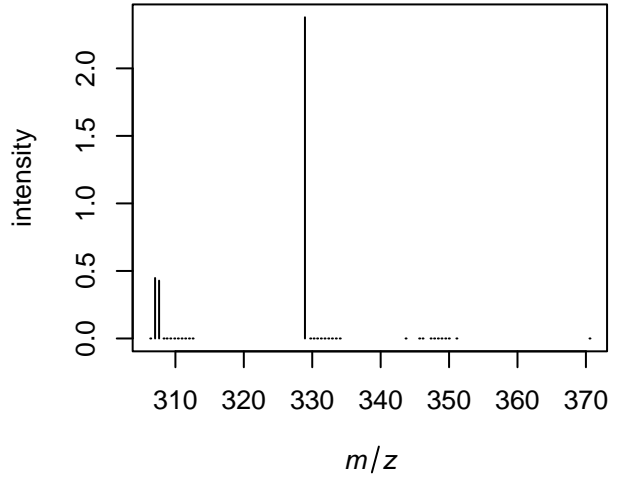
**masspeaks3\_forinput.tabular**

# Input spectra

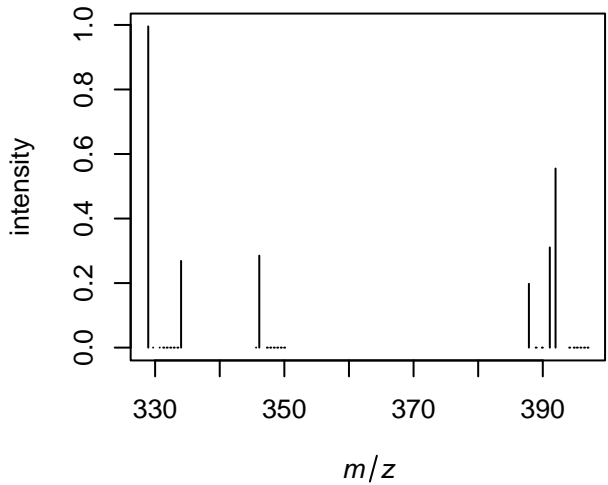
## spectrum xy\_3\_1



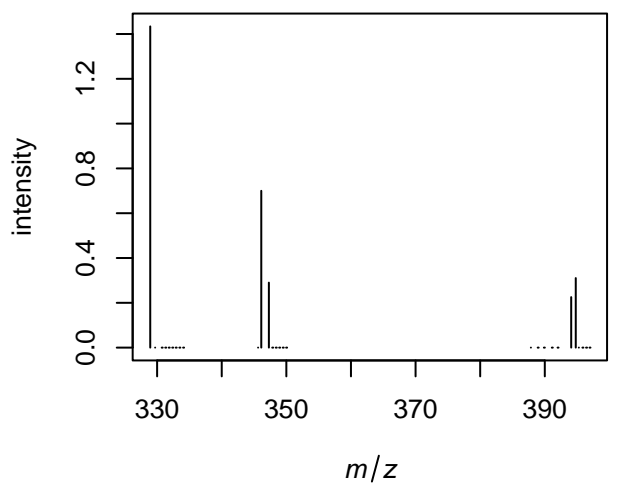
## spectrum xy\_2\_2



## spectrum xy\_2\_3

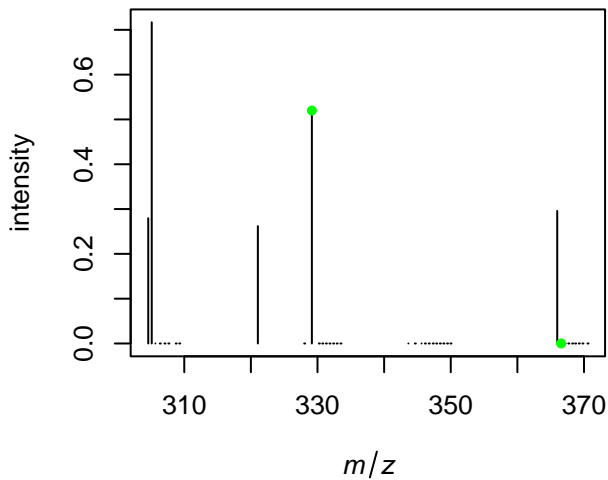


## spectrum xy\_3\_3

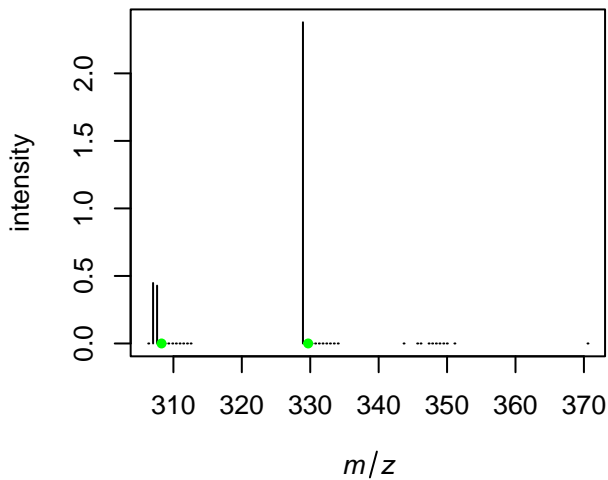


# Monoisotopic peaks in green

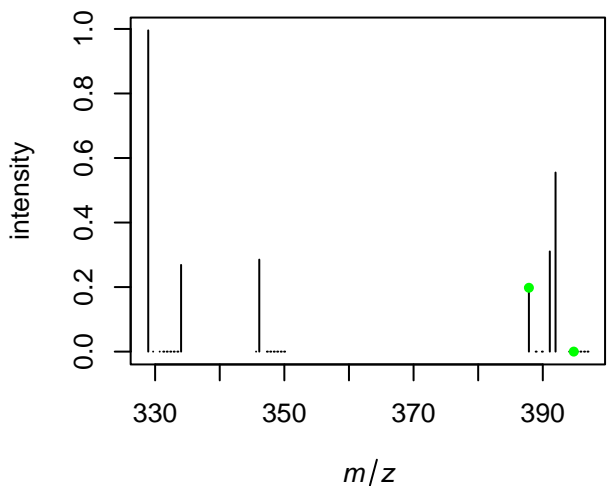
## spectrum xy\_3\_1



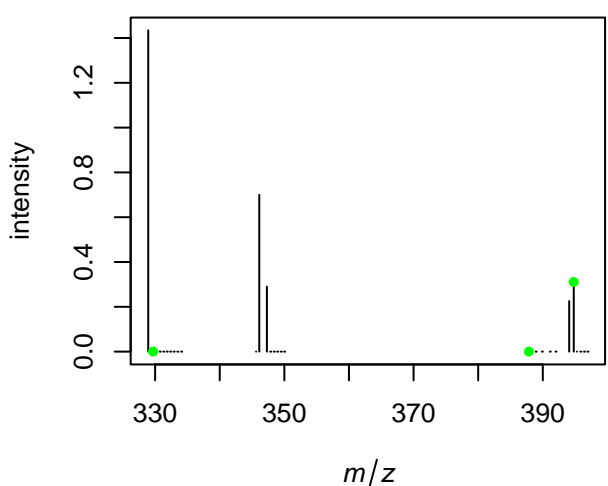
## spectrum xy\_2\_2



## spectrum xy\_2\_3

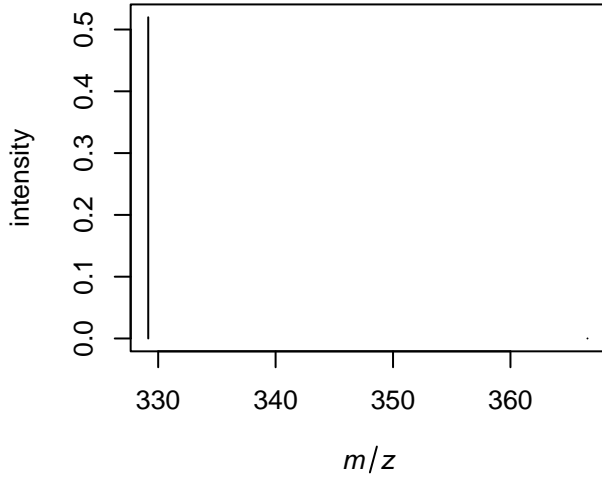


## spectrum xy\_3\_3

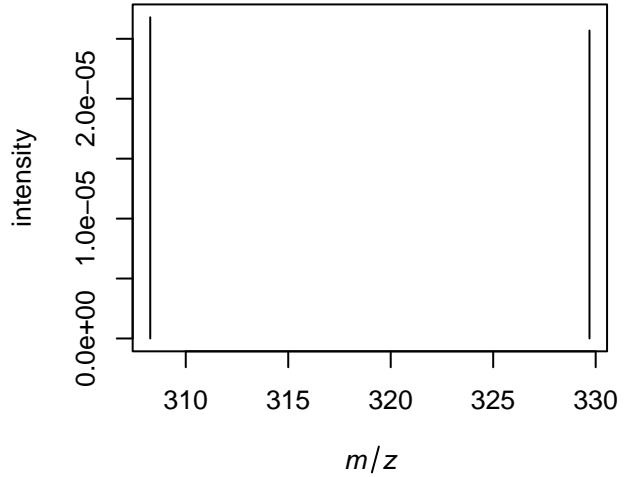


# Monoisotopic peaks

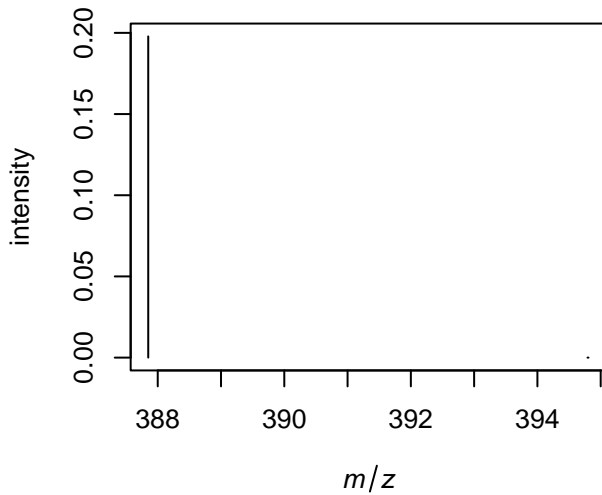
## spectrum xy\_3\_1



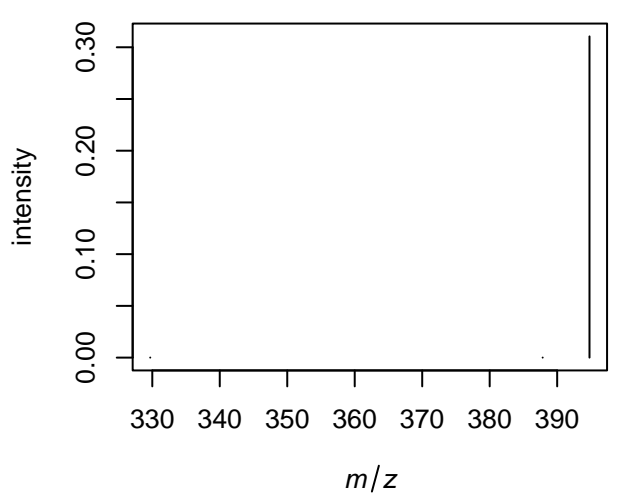
## spectrum xy\_2\_2



## spectrum xy\_2\_3



## spectrum xy\_3\_3



|                     | <b>min m/z</b> | <b>max mz</b> | <b># features</b> | <b>median<br/># peaks (int.&gt;0)</b> | <b>median<br/>intensity</b> | <b># spectra</b> |
|---------------------|----------------|---------------|-------------------|---------------------------------------|-----------------------------|------------------|
| <i>inputdata</i>    | 304.5953       | 396.9167      | 60                | 33                                    | 0                           | 9                |
| <i>monoisotopes</i> | 308.2667       | 394.7917      | 9                 | 2.11                                  | 0.2                         | 9                |