

# Package ‘q3ML’

February 16, 2022

**Type** Package

**Title** Open and Parse MRM-MS mzML Files

**Version** 0.1.1

**Description** Parse .mzML files of Multiple Reaction Monitoring (MRM) Mass Spectrometry (MS) data using xml2.

**License** GPL (>= 3)

**Encoding** UTF-8

**LazyData** true

**Depends** magrittr

**Imports** xml2,  
dplyr,  
tidyr,  
purrr,  
tibble,  
stringr,  
base64enc,  
crayon,  
cli

**RoxygenNote** 7.1.1

**Suggests** covr,  
testthat (>= 3.0.0)

**Config/testthat/edition** 3

**Roxygen** list(markdown = TRUE)

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`openFile`*Open .mzML File*

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**Description**

Open and parse a .mzML file containing Selective Reaction Monitoring (SRM) Mass Spectrometry (MS) data using `xml2`

**Usage**

```
openFile(mzml_file)
```

**Arguments**

`mzml_file`      the absolute file path of a valid .mzML file

**Value**

a list of two elements containing peaks and header

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