

# Package ‘rawReader’

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**Type** Package

**Title** Read Thermo .RAW files in R

**Version** 0.1

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**Description** A series of C++ functions are used to access the Thermo MSFilleReader.dll. Each C++ function is designed for single file and single scan input; corresponding R functions and system calls are used as multi scan and multi file wrappers to the C++ functions, via the pre-compiled binary executables.

**License** GPL (>= 3)

**URL** <https://github.com/wilsontom/rawReader>

**BugReports** <https://github.com/wilsontom/rawReader/issues>

**LazyData** TRUE

**RoxygenNote** 5.0.1

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`getBValues`*Get B Value*

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**Description**

Get the FT-MS conversion parameter B for each scan in a given range

**Usage**

```
getBValues(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the B value for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getBValues(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

---

`getCValues`*Get C Value*

---

**Description**

Get the FT-MS conversion parameter C for each scan in a given range

**Usage**

```
getCValues(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the C value for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getCValues(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

---

*getITtime**Get Ion Injection Time*

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**Description**

Get the ion injection time (IT) for each scan in a given range

**Usage**

```
getITtime(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the Ion Injection Time for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getITtime(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

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getMzInt	<i>Get Mass and Intensity</i>
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**Description**

Get the profile data (m/z and intensity) across a given scan range

**Usage**

```
getMzInt(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a list of matrices for m/z and intensity.

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getMzInt(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

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getMzIntNoise	<i>Get Mass, Intensity and Noise</i>
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**Description**

Get the profile data (m/z, intensity and noise) across a given scan range

**Usage**

```
getMzIntNoise(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a list of matrices for m/z, intensity and noise

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getMzIntNoise(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

---

getResComp

*Get Resolution Compensation*

---

**Description**

Get the Resolution mass compensation (ppm) for each scan in a given range

**Usage**

```
getResComp(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the Resolution compensation for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getResComp(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

---

`getRFComp`*Get RF Compensation*

---

**Description**

Get the RF mass compensation (ppm) for each scan in a given range

**Usage**

```
getRFComp(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the RF compensation for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getRFComp(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

---

`getSpaceComp`*Get Space Charge Compensation*

---

**Description**

Get the space charge mass compensation (ppm) for each scan in a given range

**Usage**

```
getSpaceComp(filename, scans = c())
```

**Arguments**

filename	a .raw file
scans	a numeric vector of scan numbers to extract

**Value**

a numeric vector of the space charge compensation for each scan

**Author(s)**

Tom Wilson <tpw2@aber.ac.uk>

**Examples**

```
## Not run:  
getSpaceComp(QC.raw, scans = c(2:24))  
  
## End(Not run)
```

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