Installation

artMS

Analytical R Tools
for Mass Spectrometry

Bioconductor
OPEN SOURCE SOFTWARE FOR BIOINFORMATICS

Installation

To install this package, start R (version "4.0") and enter:

```
if (!requireNamespace("BiocManager", quietly = TRUE))
    install.packages("BiocManager")

BiocManager::install("artMS")
```
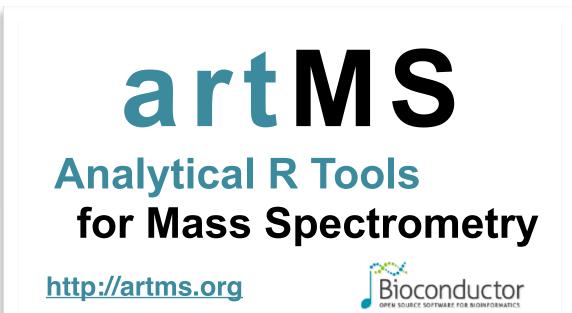
For older versions of R, please refer to the appropriate Bioconductor release.

(Recommended, stable)

```
install.packages("devtools")
library(devtools)
install_github("biodavidjm/artMS")
```

(Development, less stable)

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http://artms.org

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