

# Package ‘RGMQLlib’

October 12, 2020

**Type** Package

**Title** RGMQLlib, java libraries to run GMQL scala API

**Version** 1.8.0

**Author** Simone Pallotta, Marco Masseroli

**Maintainer** Simone Pallotta <simonepallotta@hotmail.com>

**Description** A package that contains scala libraries to call GMQL from R used by RGMQL package. It contains a scalable data management engine written in Scala programming language.

**License** Artistic-2.0

**URL** [http://www.bioinformatics.deib.polimi.it/genomic\\_computing/GMQL/](http://www.bioinformatics.deib.polimi.it/genomic_computing/GMQL/)

**Encoding** UTF-8

**LazyData** true

**RoxygenNote** 6.0.1

**Depends** R(>= 3.4.2)

**VignetteBuilder** knitr

**Suggests** BiocStyle, knitr, rmarkdown

**biocViews** ExperimentData,RepositoryData

**NeedsCompilation** no

**git\_url** <https://git.bioconductor.org/packages/RGMQLlib>

**git\_branch** RELEASE\_3\_11

**git\_last\_commit** bb9aaab

**git\_last\_commit\_date** 2020-04-27

**Date/Publication** 2020-10-12

## R topics documented:

RGMQLlib . . . . .	2
Index	3

---

RGMQLlib

*RGMQL scala API libraries*

---

**Description**

RGMQLlib

**Details**

Is a data package containing compiled libraries in jar format used to invoke GMQL scala API

This library is used in the package RGQML

# Index

RGMQLlib, [2](#)

RGMQLlib-package (RGMQLlib), [2](#)