

# Package ‘Hydra’

March 31, 2022

**Type** Package

**Title** Hydrating Package Skeletons into Executable R Study Packages

**Version** 0.3.0

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**Description** An R package and Java library for hydrating package skeletons into executable R study packages based on specifications in JSON format.

**License** Apache License 2.0

**VignetteBuilder** knitr

**URL** <https://ohdsi.github.io/Hydra>, <https://github.com/OHDSI/Hydra>

**BugReports** <https://github.com/OHDSI/Hydra/issues>

**Depends** R (>= 3.5.0),

**Imports** rJava,  
utils

**Suggests** testthat,  
knitr,  
rmarkdown,  
renv,  
DatabaseConnector,  
devtools,  
RJSONIO

**NeedsCompilation** no

**RoxygenNote** 7.1.2

**Language** en-US

**Encoding** UTF-8

## R topics documented:

hydrate . . . . .	2
listSkeletons . . . . .	2
loadSpecifications . . . . .	3
prepareForOfflineStudyPackageExecution . . . . .	3
<b>Index</b>	<b>5</b>

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hydrate	<i>Hydrate a package skeleton according to study specifications</i>
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### Description

Hydrate a package skeleton according to study specifications

### Usage

```
hydrate(
  specifications,
  outputFolder,
  skeletonFileName = NULL,
  packageName = NULL
)
```

### Arguments

specifications	A string containing JSON specifications for hydrating the study. Specifications will be generated by ATLAS.
outputFolder	The folder where the hydrated package will be stored.
skeletonFileName	(Mostly for debugging purposes:) Specify the location of a package skeleton to hydrate. If not specified, the appropriate skeleton will be selected from those inside the Hydra package.
packageName	The name of the R package to create. If provided, will override the packageName variable in the specifications JSON.

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listSkeletons	<i>List skeletons included in Hydra</i>
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### Description

List skeletons included in Hydra

### Usage

```
listSkeletons()
```

### Value

A vector of skeleton names.

### Examples

```
listSkeletons()
```

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loadSpecifications	<i>Load specifications from a JSON file</i>
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**Description**

Load specifications from a JSON file

**Usage**

```
loadSpecifications(fileName)
```

**Arguments**

fileName	Path to the JSON file
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**Value**

A string representation of the JSON.

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prepareForOfflineStudyPackageExecution	<i>Prepare system to run hydrated packages without further internet connection</i>
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**Description**

Prepare system to run hydrated packages without further internet connection

**Usage**

```
prepareForOfflineStudyPackageExecution(
  installRpackages = TRUE,
  installJdbcDrivers = TRUE,
  skeletons = listSkeletons()
)
```

**Arguments**

installRpackages	Install the R packages required by the skeletons?
installJdbcDrivers	Install all JDBC drivers? Requires the DATABASECONNECTOR_JAR_FOLDER environmental variable to be set.
skeletons	A list of skeletons to check, for example 'CohortDiagnosticsStudy_v0.0.1.zip'.

**Details**

Note that when `installJdbcDrivers = TRUE` this will only include the Jar drivers supported by `DatabaseConnector::downloadJdbcDrivers`. Other drivers (like `BigQuery`) will need to be downloaded manually and placed in the folder identified by the `DATABASECONNECTOR_JAR_FOLDER` environmental variable.

Use `list.files(system.file("skeletons", package = "Hydra"), pattern = "*.zip")` for a list of all skeletons

**Value**

This function does not return anything. Instead, it installs all dependencies required to run the hydrated skeletons. (Only those skeletons that use a `renv.lock` file)

# Index

hydrate, [2](#)

listSkeletons, [2](#)

loadSpecifications, [3](#)

prepareForOfflineStudyPackageExecution,  
[3](#)