

# Package ‘OhdsiShinyAppBuilder’

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**Type** Package

**Title** Viewing Observational Health Data Sciences and Informatics  
Results via 'shiny' Modules

**Version** 1.0.0

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**Description** Users can build a single 'shiny' app for exploring population characterization, population-level causal effect estimation, and patient-level prediction results generated via the R analyses packages in 'HADES' (see <<https://ohdsi.github.io/Hades/>>). Learn more about 'OhdsiShinyAppBuilder' at <<https://ohdsi.github.io/OhdsiShinyAppBuilder/>>.

**License** Apache License 2.0

**VignetteBuilder** knitr

**Depends** R (>= 3.3.0)

**Imports** checkmate, DatabaseConnector, devtools, dplyr, ParallelLogger  
(>= 2.0.0), ResultModelManager (>= 0.4.0), rlang, shiny,  
shinydashboard, utils

**Suggests** Eunomia, jsonlite, knitr, rmarkdown, testthat, markdown,  
remotes

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addModuleConfig	<i>addModuleConfig</i>
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### Description

Create an R list with the config specification for one or more modules

### Usage

```
addModuleConfig(config, moduleConfig)
```

### Arguments

config	The current config to append the extra module to
moduleConfig	A module config to be added

### Details

User specifies the settings to create a config for a module

**Value**

An R list with the module config settings

---

`createDefaultAboutConfig`  
*createDefaultAboutConfig*

---

**Description**

Create an R list with the about config specification

**Usage**

`createDefaultAboutConfig()`

**Details**

User specifies the settings to create a default config for an about module

**Value**

An R list with the module config settings

---

`createDefaultCharacterizationConfig`  
*createDefaultCharacterizationConfig*

---

**Description**

Create an R list with the characterization config specification

**Usage**

`createDefaultCharacterizationConfig()`

**Details**

User specifies the settings to create a default config for a characterization module

**Value**

An R list with the module config settings

---

`createDefaultCohortDiagnosticsConfig`  
*createDefaultCohortDiagnosticsConfig*

---

**Description**

Create an R list with the characterization config specification

**Usage**

`createDefaultCohortDiagnosticsConfig()`

**Details**

User specifies the settings to create a default config for a characterization module

**Value**

An R list with the module config settings

---

`createDefaultCohortGeneratorConfig`  
*createDefaultCohortGeneratorConfig*

---

**Description**

Create an R list with the cohort generator config specification

**Usage**

`createDefaultCohortGeneratorConfig()`

**Details**

User specifies the settings to create a default config for a cohort generator module

**Value**

An R list with the module config settings

---

`createDefaultCohortMethodConfig`  
*createDefaultCohortMethodConfig*

---

### **Description**

Create an R list with the cohort method config specification (deprecated)

### **Usage**

`createDefaultCohortMethodConfig()`

### **Details**

User specifies the settings to create a default config for a cohort method module

### **Value**

An R list with the module config settings

---

`createDefaultDatasourcesConfig`  
*createDefaultDatasourcesConfig*

---

### **Description**

Create an R list with the datasources config specification

### **Usage**

`createDefaultDatasourcesConfig()`

### **Details**

User specifies the settings to create a default config for a datasources module

### **Value**

An R list with the module config settings

---

```
createDefaultEstimationConfig  
  createDefaultEstimationConfig
```

---

**Description**

Create an R list with the estimation config specification

**Usage**

```
createDefaultEstimationConfig()
```

**Details**

User specifies the settings to create a default config for an estimation module

**Value**

An R list with the module config settings

---

```
createDefaultEvidenceSynthesisConfig  
  createDefaultEvidenceSynthesisConfig
```

---

**Description**

Create an R list with the Evidence Synthesis config specification (deprecated)

**Usage**

```
createDefaultEvidenceSynthesisConfig()
```

**Details**

User specifies the settings to create a default config for a Evidence Synthesis module

**Value**

An R list with the module config settings

---

`createDefaultHomeConfig`  
*createDefaultHomeConfig*

---

**Description**

Create an R list with the about config specification

**Usage**

`createDefaultHomeConfig()`

**Details**

User specifies the settings to create a default config for an about module

**Value**

An R list with the module config settings

---

`createDefaultPhevaluatorConfig`  
*createDefaultPhevaluatorConfig*

---

**Description**

Create an R list with the phevaluator config specification

**Usage**

`createDefaultPhevaluatorConfig()`

**Details**

User specifies the settings to create a default config for a phevaluator module

**Value**

An R list with the module config settings

---

`createDefaultPredictionConfig`  
*createDefaultPredictionConfig*

---

**Description**

Create an R list with the prediction config specification

**Usage**

`createDefaultPredictionConfig()`

**Details**

User specifies the settings to create a default config for a prediction module

**Value**

An R list with the module config settings

---

`createDefaultReportConfig`  
*createDefaultReportConfig*

---

**Description**

Create an R list with the report config specification

**Usage**

`createDefaultReportConfig()`

**Details**

User specifies the settings to create a default config for a report module

**Value**

An R list with the module config settings



---

```
createDefaultResultDatabaseSettings
      createDefaultResultDatabaseSettings
```

---

### Description

Create an R list with the default table prefixes and tables

### Usage

```
createDefaultResultDatabaseSettings(
  schema = "main",
  vocabularyDatabaseSchema = schema,
  cgTablePrefix = "cg_",
  cgTable = "cohort_definition",
  databaseTable = "DATABASE_META_DATA",
  databaseTablePrefix = "",
  cdTablePrefix = "cd_",
  cTablePrefix = "c_",
  incidenceTablePrefix = "ci_",
  plpTablePrefix = "plp_",
  cmTablePrefix = "cm_",
  sccsTablePrefix = "sccs_",
  esTablePrefix = "es_",
  pvTablePrefix = "pv_"
)
```

### Arguments

schema	The schema containing the OHDSI study results
vocabularyDatabaseSchema	The schema containing the vocabulary database
cgTablePrefix	The table prefix for cohort generator tables
cgTable	The table with the cohort definitions in cohort generator
databaseTable	The table with the database details
databaseTablePrefix	The prefix for the databaseTable
cdTablePrefix	The table prefix for the cohort diagnostic results
cTablePrefix	The table prefix for the characterization results
incidenceTablePrefix	The table prefix for the incidence results
plpTablePrefix	The table prefix for the prediction results
cmTablePrefix	The table prefix for the cohort method results

<code>sccsTablePrefix</code>	The table prefix for the self controlled case series results
<code>esTablePrefix</code>	The table prefix for the evidence synthesis method results
<code>pvTablePrefix</code>	The table prefix for the phevaluator results

**Details**

User specifies the schema name and can override default table prefixes and table names

**Value**

An R list with the module config settings

---

`createDefaultSccsConfig`  
*createDefaultSccsConfig*

---

**Description**

Create an R list with the SCCS config specification (deprecated)

**Usage**

`createDefaultSccsConfig()`

**Details**

User specifies the settings to create a default config for a SCCS module

**Value**

An R list with the module config settings

---

`createModuleConfig`    *createModuleConfig*

---

**Description**

Create an R list with the config specification

**Usage**

```
createModuleConfig(  
  moduleId = "about",  
  tabName = "About",  
  shinyModulePackage = "OhdsiShinyModules",  
  shinyModulePackageVersion = "3.0.0",  
  moduleUiFunction = "aboutViewer",  
  moduleServerFunction = "aboutServer",  
  moduleInfoBoxFile = "aboutHelperFile()",  
  moduleIcon = "info",  
  installSource = "CRAN",  
  gitHubRepo = NULL  
)
```

**Arguments**

moduleId	The shiny id for the tab containing the module UI
tabName	The name of the tab in the shiny app (this will be the side menu button text)
shinyModulePackage	The R package to find the server and UI functions
shinyModulePackageVersion	The minimum version of the shinyModulePackage
moduleUiFunction	The name of the UI function in the R package shinyModulePackage
moduleServerFunction	The name of the server function in the R package shinyModulePackage
moduleInfoBoxFile	The function in the R package shinyModulePackage that contains info text
moduleIcon	The icon to use for the side menu button
installSource	Either CRAN or github specifying where to install shinyModulePackage from
gitHubRepo	If installSource is github then this is the github repo containing the package e.g., 'ohdsi'

**Details**

User specifies the settings to create a config for a module

**Value**

An R list with the module config settings

---

<code>createShinyApp</code>	<i>createShinyApp</i>
-----------------------------	-----------------------

---

## Description

Create a shiny app for a shiny server

## Usage

```
createShinyApp(
  config,
  connection,
  resultDatabaseSettings = createDefaultResultDatabaseSettings(),
  connectionDetails = NULL,
  usePooledConnection = TRUE,
  studyDescription = NULL,
  title = "OHDSI Analysis Viewer",
  protocolLink = "http://ohdsi.org",
  themePackage = "OhdsiShinyAppBuilder",
  reportSummaryDetails = NULL
)
```

## Arguments

<code>config</code>	The json with the app config
<code>connection</code>	A connection to the results
<code>resultDatabaseSettings</code>	A list with the result schema and table prefixes
<code>connectionDetails</code>	A <code>DatabaseConnector::connectionDetails</code> connection to the results database
<code>usePooledConnection</code>	Use a pooled database connection or not - set to true for multi-user environments (default)
<code>studyDescription</code>	A human-readable character string describing the study/analysis
<code>title</code>	The title for the app. Defaults to: OHDSI Analysis Viewer
<code>protocolLink</code>	A link to a site containing the study protocol
<code>themePackage</code>	A package containing custom theme elements
<code>reportSummaryDetails</code>	NULL or a <code>data.frame</code> with the columns <code>reportName</code> and <code>reportLocation</code>

## Details

User specifies the json config and connection

**Value**

Shiny app instance

---

`initializeModuleConfig`  
*initializeModuleConfig*

---

**Description**

Creates an empty config

**Usage**

```
initializeModuleConfig()
```

**Details**

An empty config that can be used to add shiny module configs to

**Value**

An empty list

---

`loadConfig`            *loadConfig*

---

**Description**

Load the config specification

**Usage**

```
loadConfig(configLocation)
```

**Arguments**

`configLocation` The location of the config json file

**Details**

User specifies the config file and this function returns the R list with the config

**Value**

An R list with the config settings

---

OhdsiShinyAppBuilder    *OhdsiShinyAppBuilder*

---

**Description**

A package for creating viewers for the OHDSI shiny apps

**Author(s)**

**Maintainer:** Jenna Reps <jreps@its.jnj.com>

Authors:

- Nathan Hall
- Josh Ide
- Jamie Gibert

---

saveConfig                    *saveConfig*

---

**Description**

Save the R list with the config specification as a json

**Usage**

```
saveConfig(configList, configLocation)
```

**Arguments**

`configList`        An R list with the config settings  
`configLocation`    The location to save the config json file

**Details**

User saves the R list with the config as a json file

**Value**

An R list with the config settings

---

 viewShiny

*viewShiny*


---

**Description**

Open the shiny app

**Usage**

```
viewShiny(
  config,
  connection,
  resultDatabaseSettings = createDefaultResultDatabaseSettings(),
  connectionDetails = NULL,
  usePooledConnection = TRUE,
  studyDescription = NULL,
  title = "OHDSI Analysis Viewer",
  protocolLink = "http://ohdsi.org",
  themePackage = "OhdsiShinyAppBuilder"
)
```

**Arguments**

<code>config</code>	The json with the app config
<code>connection</code>	A connection to the results
<code>resultDatabaseSettings</code>	A list with the result schema and table prefixes
<code>connectionDetails</code>	A DatabaseConnector::connectionDetails connection to the results database
<code>usePooledConnection</code>	Use a pooled database connection or not - set to true for multi-user environments (default)
<code>studyDescription</code>	A human-readable character string describing the study/analysis
<code>title</code>	The title for the app. Defaults to: OHDSI Analysis Viewer
<code>protocolLink</code>	A link to a site containing the study protocol
<code>themePackage</code>	A package containing custom theme elements

**Details**

User specifies the json config and connection

**Value**

The shiny app will open

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