

# Package: NGCHMSupportFiles (via r-universe)

August 19, 2024

**Title** Support Files for Building Next Generation Clustered Heat Maps (NG-CHMs)

**Description** Support files required by the 'NGCHM' package to export Next-Generation Clustered Heat Maps (NG-CHMs) to .ngchm, HTML, and PDF files. These support files are updated mostly independently from the 'NGCHM' package, and are therefore provided separately.

**License** GPL-3

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.1

**SystemRequirements** Java JDK (>= 11)

**URL** <https://github.com/MD-Anderson-Bioinformatics/NGCHMSupportFiles>,  
<https://md-anderson-bioinformatics.r-universe.dev/NGCHMSupportFiles>

**Suggests** knitr, rmarkdown

**Version** 2.24.4

**Repository** <https://marohrdanz.r-universe.dev>

**RemoteUrl** <https://github.com/MD-Anderson-Bioinformatics/NGCHMSupportFiles>

**RemoteRef** HEAD

**RemoteSha** 7425a4ae08a87ae84778ed764eebf8744a52efda

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NGCHMSupportFiles-package

*Support Files for Building Next Generation Clustered Heat Maps (NG-CHMs)*

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## Description

Support Files for 'NGCHM' package

## Details

Files provided by this package:

- ShaidyMapGen.jar - Compiled Java code. Used by `NGCHM::chmExportToFile()`, `NGCHM::chmExportToHTML()`, `NGCHM::chmExportToPDF()` to produce .ngchm, .html, and .pdf files, respectively.
- ngchmWidget-min.js - JavaScript library used to display NG-CHMs in a web browser. Used by `NGCHM::chmExportToHTML()`.

These Support Files are made available to the 'NGCHM' package via environment variables, SHAIIDYMAPGEN and NGCHMWIDGETPATH, which are set when this package loads. These files are required to export a Next-Generation Clustered Heat Map (NG-CHM) to a .ngchm, .html, or .pdf file.

## Installation

This package can be installed from MD Anderson Bioinformatics R-universe repository:

```
install.packages("NGCHMSupportFiles",  
  repos = c("https://md-anderson-bioinformatics.r-universe.dev", "https://cran.r-project.org"))
```

## Author(s)

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**See Also**

Useful links:

- <https://github.com/MD-Anderson-Bioinformatics/NGCHMSupportFiles>
- <https://md-anderson-bioinformatics.r-universe.dev/NGCHMSupportFiles>

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.onAttach	<i>Display message regarding environment variables</i>
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**Description**

Display message to user that environment variables were set for:

- SHAIDYMAPGEN - Path to the included file ShaidyMapGen.jar
- NGCHMWIDGETPATH - Path to the included file ngchmWidget-min.js

**Usage**

```
.onAttach(libname, pkgname)
```

**Arguments**

libname	name of the library
pkgname	name of the package

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.onLoad	<i>Define environment variables</i>
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**Description**

Define environment variables:

- SHAIDYMAPGEN - Path to the included file ShaidyMapGen.jar
- NGCHMWIDGETPATH - Path to the included file ngchmWidget-min.js

**Usage**

```
.onLoad(libname, pkgname)
```

**Arguments**

libname	name of the library
pkgname	name of the package

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