

Package ‘rhdf5filters’

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Type Package
Title HDF5 Compression Filters
Version 1.21.0
Description Provides a collection of additional compression filters for HDF5 datasets. The package is intended to provide seamless integration with rhdf5, however the compiled filters can also be used with external applications.
License BSD_2_clause + file LICENSE
LazyLoad true
VignetteBuilder knitr
Suggests BiocStyle, knitr, rmarkdown, tinytest, rhdf5 (>= 2.47.7)
SystemRequirements GNU make
URL <https://github.com/Huber-group-EMBL/rhdf5filters>
BugReports <https://github.com/Huber-group-EMBL/rhdf5filters/issues>
LinkingTo Rhdf5lib
RoxygenNote 7.3.2
Encoding UTF-8
biocViews Infrastructure, DataImport
git_url <https://git.bioconductor.org/packages/rhdf5filters>
git_branch devel
git_last_commit a4f274e
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Repository Bioconductor 3.22
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available_filters	<i>Return a list of installed filters</i>
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Description

This function can be used to produce a vector of the installed filters.

Usage

```
available_filters()
```

Value

A character vector containing the names of the installed filters.

Examples

```
available_filters()
```

hdf5_plugin_path	<i>Return location of installed filters</i>
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Description

This function can be used to find the location of the installed filters. Setting the environment variable HDF5_PLUGIN_PATH to this value will allow other applications that require the the filters to use the versions distributed with this package.

Usage

```
hdf5_plugin_path()
```

Value

A character string containing the path where the compiled filters are located.

Examples

```
hdf5_plugin_path()
```

*rhdf5filters**rhdf5filters*

Description

This package provides several HDF5 compression filters for use with rhdf5 or other tools using HDF5.

Author(s)

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See Also

Useful links:

- <https://github.com/grimbough/rhdf5filters>
- Report bugs at <https://github.com/grimbough/rhdf5filters>

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