# Package 'GCSFilesystem'

# October 14, 2021

October 14, 2021
Type Package
Title Mounting a Google Cloud bucket to a local directory
Version 1.2.0
<b>Date</b> 2020-09-16
Description Mounting a Google Cloud bucket to a local directory.  The files in the bucket can be viewed and read as if they are locally stored.  For using the package, you need to install GCSDokan on Windows or gcsfuse on Linux and MacOs.
License GPL (>= 2)
<b>Depends</b> R (>= 4.0.0)
Imports stats
Suggests testthat, knitr, rmarkdown, BiocStyle, GCSConnection
SystemRequirements GCSDokan for Windows, gcsfuse for Linux and macOs
biocViews Infrastructure
RoxygenNote 7.1.1
Encoding UTF-8
VignetteBuilder knitr
git_url https://git.bioconductor.org/packages/GCSFilesystem
git_branch RELEASE_3_13
git_last_commit 7594317
git_last_commit_date 2021-05-19
Date/Publication 2021-10-14
Author Jiefei Wang [aut, cre]
Maintainer Jiefei Wang <szwjf08@gmail.com></szwjf08@gmail.com>
R topics documented:
gcs_list_mountpoints
gcs_mount
gcs_unmount

gcs\_mount

Index 5

```
gcs_list_mountpoints List all GCS mountpoints
```

#### **Description**

List all GCS mountpoints. The function uses 'GCSDokan" on Windows or the command 'df' on Linux to show all mountpoints. Due to the system differences, the function is only able to show the bucket name on Linux and can show the full remote path on Windows.

#### Usage

```
gcs_list_mountpoints()
```

#### Value

a data.frame object with the first column named 'remote' and second named 'mountpoint'

#### **Examples**

```
gcs_list_mountpoints()
```

gcs\_mount

Mount a Google Cloud Storage(GCS) file system

#### Description

The function uses the command-line program 'GCSDokan' on Windows or 'gcsfuse' on Linux system to mount a google cloud bucket path to your local file system.

#### Usage

```
gcs_mount(
  remote,
  mountpoint,
  mode = c("r", "rw"),
  cache_type = c("disk", "memory", "none"),
  cache_arg = NULL,
  billing = NULL,
  refresh = 60,
  implicit_dirs = TRUE,
  key_file = NULL,
  additional_args = NULL
)
```

gcs\_unmount 3

#### Arguments

remote the remote path to a Google Cloud Storage bucket. It can be either the bucket

itself or a directory in the bucket.

mountpoint The mount point where the GCS file system will be mounted to.

mode the permission of the mounted directory. The write permission is only available

on Linux.

cache\_type The location where the file cache will be stored. The cache type 'none' and

'memory' are only available on Windows

cache\_arg The argument of a cache type. If 'cache\_type = "memory"', the argument is the

limit of the memory useage in MB. If 'cache\_type = "disk"', the argument is the

path to a cache directory.

billing The billing project ID.

refresh The refresh rate of the cloud files.

implicit\_dirs Implicitly define directories based on content. This argument is only available

on Linux.

key\_file The service account credentials file

additional\_args

The additional argument that will be passed to the command-line program.

#### Value

no return value

## Examples

```
bucket <- "genomics-public-data"
mountpoint <- paste0(tempdir(),"/GCSFilesystemTest")
## You must have a credentials on Linux and macOs
## To run this code
if(Sys.getenv("GOOGLE_APPLICATION_CREDENTIALS")!=""){
    gcs_mount(bucket, mountpoint)
    gcs_unmount(mountpoint)
}</pre>
```

gcs\_unmount

Unmound a mounted GCS file system

#### **Description**

Unmound a mounted GCS file system

#### Usage

```
gcs_unmount(mountpoint)
```

gcs\_unmount

# Arguments

mountpoint The path to the mounted GCS file system.

### Value

No return value

### **Examples**

```
## Unmount a directory
## No operation will be done
## if the directory does not exist or
## not a mounted directory
gcs_unmount("path_to_your_mounted_directory")
```

# **Index**

```
gcs_list_mountpoints, 2
gcs_mount, 2
gcs_unmount, 3
```