

# Package ‘fssetup’

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**Title** Package for Producing R Projects and Scripts Using the Defra Farming Stats Template

**Version** 0.1.1

**Description** This package contains the functions needed to create various features in R using the Defra Farming Stats Team templates. Including scripts, projects, readmes and gitignores. It also contains useful functions for linking RStudio and RStudio projects to GitHub.

**License** MIT + file LICENSE

**Encoding** UTF-8

**Imports** cli (>= 3.6.2), credentials (>= 2.0.1), fs (>= 1.6.4), gert (>= 2.1.0), gitcreds (>= 0.1.2), glue (>= 1.7.0), here (>= 1.0.1), magrittr (>= 2.0.3), pkgbuild (>= 1.4.4), quarto (>= 1.4.4), rstudioapi (>= 0.16.0), stringr (>= 1.5.1), usethis (>= 3.0.0)

**Roxygen** list(markdown = TRUE)

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**URL** <https://github.com/jpmoatt/fssetup>

**BugReports** <https://github.com/jpmoatt/fssetup/issues>

**Suggests** testthat (>= 3.0.0)

**Config/testthat/edition** 3

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addin_fs_roxygen	<i>Addin to open new script with roxygen template</i>
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---

## Description

This addin will open a new blank R script which is populated with a standard roxygen header.

## Usage

```
addin_fs_roxygen()
```

## Details

This addin will create a new blank script that is populated with standard roxygen header.

Addins must use the `rstudioapi` package. This addin uses the `documentNew()` function from the `rstudioapi` package to open the new R script.

Note: this opens an normal untitled script. It must be manually saved in the correct directory.

This package also contains another addin to open an R script with a farming stats header, which can be used for creating normal R scripts.

## Value

A script will open in RStudio.

---

addin_fs_script	<i>Addin to open new script with Farming Stats template</i>
-----------------	---

---

## Description

This addin will open a new blank R script which is populated with the Farming Stats header. Unlike the `create_fs_script()` function, this addin will also pre-populate the date.

## Usage

```
addin_fs_script()
```

## Details

This addin will create a new blank script that is populated with the farming stats template header. Unlike the `create_fs_script()` function in this package, this will auto-populate the script with today's date.

Addins must use the `rstudioapi` package. This addin uses the `documentNew()` function from the `rstudioapi` package to open the new R script.

Note: this opens an normal untitled script. It must be manually saved in the correct directory.

This package also contains another addin to open an R script with a roxygen header, which can be used for creating scripts within packages.

**Value**

A script will open in RStudio.

---

create_fs_gitignore	Create a gitignore file based on the Farming Stats template
---------------------	---

---

**Description**

Use this function to create a gitignore for a Farming Stats project.

**Usage**

```
create_fs_gitignore(type = "default", file_path = NULL, custom_txt = NULL)
```

**Arguments**

type	description controlling which gitignore is added. "default" will add the standard template. "custom" will enable the user to provide a custom template via custom_txt.
file_path	optional argument allowing users to specify file path where gitignore should be created. If not entered, will default to current project/working directory.
custom_txt	optional argument allowing users to provide their own gitignore template. Must be provided as a string. Only used if type set to "custom".

**Details**

This function will create a gitignore for a project using a pre-defined Farming Stats template.

Used as default, it will automatically add the data and output folders (as created in the fssetup project template). This is to ensure nor restricted data or unpublished results are accidentally pushed to GitHub.

Alternatively, a custom gitignore can be provided by setting type to "custom" and providing a custom template to custom\_txt as a string.

Note it will replace any existing .gitignore files present in the project already.

**Value**

A gitignore file is added to the project.

---

create\_fs\_readme

*Create a Farming Stats README*


---

### Description

Use this function to create a README for a Farming Stats project.

### Usage

```
create_fs_readme(
  format = c("markdown", "github", "html"),
  file_path = NULL,
  author = NULL,
  readme_title = NULL
)
```

### Arguments

format	controls the output format of the README. Default is "markdown", but can be "html" or "github".
file_path	string containing file path where README will be saved.
author	string containing authors name. If no string provided author will be set to "add author".
readme_title	string containing README title. If no string provided will be set to "README (edit title)".

### Details

This function will create a README for a project using a pre-defined Farming Stats template. The function will create a Quarto (.qmd) file and do a first render producing a desired output.

The output type can be controlled using the format option (default is markdown). Can also be html or github (gfm).

### Value

Output is a .qmd file containing the desired README template and an initial render of the README in the desired output.

---

create\_fs\_script

*Create a new script with farming stats header.*


---

### Description

This function will create a new script with the farming stats header added. There are multiple options for customization (see details).

## Usage

```
create_fs_script(  
  file_name = NULL,  
  file_path = NULL,  
  author = NULL,  
  email = NULL,  
  date = format(Sys.Date(), "%d/%m/%Y")  
)
```

## Arguments

file_name	string containing desired name for script.
file_path	string containing folder name to save script. This is built on the here function in R, so will follow your root directory. If you want to save in a sub-folder, enter the full folder sequence, e.g. "folder/sub-folder".
author	string containing author's name.
email	string containing author's email.
date	string containing a date. By default, this will be set as today's date.

## Details

This is a simple function to create a new script with the farming stats header template added.

By just calling the function a script will be created with the template entered. All fields of the template will be blank except for the date, which is set to today's date.

There are options for customising and entering some details automatically. The details that can be pre-entered are:

- script name
- location to save
- author
- author's email
- date created

By default these are set to NULL (except for the date). But can be set to any string.

## Value

An R script will be saved in the root directory or in the specified folder.

---

```
create_fs_script_template
```

*Create an default R script template.*

---

## Description

This function can be used to create a default R script template. All scripts opened from this point will have this template applied as a header.

## Usage

```
create_fs_script_template(
  format = c("farming_stats", "custom", "manual_edit", "blank"),
  template = NULL
)
```

## Arguments

format	what format the template will take. There are four options: "farming_stats" (default farming stats template), "custom" will apply a custom template (provided as a string), "manual_edit" will open the template so you can manually edit the template, and "blank" can be used to remove all pre-existing templates. Note: manual edit can also be used to edit existing templates.
template	default is NULL, only used if format = "custom". Used to provide custom template design. Must be provided as a string. Must be provided if using format = "custom" or the function will return an error.

## Details

This function is used to create a new template for R scripts. This will be the default that all subsequent R scripts opened will contain. By doing this it should be easier to follow best practice and properly comment all scripts you create.

The default is stored in the appdata folder on your c drive: "~/AppData/Roaming/RStudio". It will create a "templates" folder where the default will be stored.

The function has various ways it can work which will give you the ability to create whatever header template you wish. It has four ways formatting options:

- farming\_stats
- custom
- manual\_edit
- blank

"farming\_stats" will pre-load the default script with the farming stats template.

"custom" will allow you to provide your own custom template as a string, which will then be added to the default.

"manual" will allow you to manually edit the default template. It will open it in your R studio window and manual edits can be saved. Note, this can also be used to tweak a pre-existing template (e.g. to add your name and email to all scripts).

"blank" will delete the default R script and template. This returns R back to normal, and any script opened subsequently will be blank.

**Value**

New .R file created at "~/AppData/Roaming/RStudio/templates/default.R" containing the script template

---

fs_connect_github	<i>Connect RStudio to your GitHub account</i>
-------------------	---

---

**Description**

This is a simple function which connect RStudio to GitHub, allowing you to work on GitHub repositories. This is an essential part of Reproducible Analytical Pipelines and best practice for coding.

**Usage**

```
fs_connect_github(username, email)
```

**Arguments**

username	string containing GitHub username
email	string containing email address used for gitHub account

**Details**

This is a simple function which will set you GitHub credentials and Personal Access Token (PAT) and connect RStudio to GitHub. This is essential if you want to work in RStudio in projects/repos stored on GitHub.

It uses `system()` to run the necessary code in the terminal to set your credentials, and `gitcreds_set()` and `set_github_pat()` from the `gitcreds` and `credentials` packages to connect your RStudio to GitHub.

`gitcreds_set()` is an interactive function and will prompt users for input. To replace existing credentials/PAT choose option 2. You will then be prompted for you PAT. PAT should be changed every 30 days to ensure security.

An additional feature I have added, not mentioned in the Defra instructions is to add the `set_github_pat()` function call to your .Rprofile. This will ensure your PAT is set for every R session, meaning you wont need to provide your PAT when running functions such as `install_github()` from the dev-tools package.

Note: For this function to work you must:

- have git installed on your local machine
- have a GitHub account
- have created a Personal Access Token (PAT) on GitHub.

Guidance on how to create a PAT can be found here: [ADD LINK](#).

**Value**

GitHub credentials and PAT set

---

fs_proj	<i>Function for RStudio project template</i>
---------	--

---

### Description

This is a function that is called in the "New project" viewer pane when the user chooses a Farming Stats Project template. It should not be used away from the RStudio "New project" viewer.

I have not included additional information on how to use this function, as it is not intended to be used outside the template call.

To subsequently link this to a github repo, the best plan is to use `fs_use_github()`.

### Usage

```
fs_proj(path, ...)
```

---

fs_use_github	<i>Create a GitHub repository from local project.</i>
---------------	---

---

### Description

This simple function will turn your local R project into a git repo, then create a repo on GitHub and perform the initial set up and commit.

Note: by default creates a private repo to ensure security. Can be change if the repo needs to be public. Can also be done in GitHub at a later date.

### Usage

```
fs_use_github(message = "Initial commit", private = TRUE)
```

### Arguments

message	initial commit message. Default is "Initial commit".
private	if TRUE creates private repo.

### Details

This function will take an existing R project on you local machine, turn it into a git repo and create a GitHub repository. It will do the following:

- add a gitignore
- initialise the git repo
- stage any uncommitted files
- commit the files
- create a GitHub repo
- restart RStudio to activate the git pane in R



The gitignore is added using the `create_fs_gitignore()` function within this package.

The repo initialisation, staging and committing is all done using the gert package.

Creating the GitHub repo uses the `use_github()` function from the usethis package.

For this function to work you must:

- have git installed on your machine
- have a GitHub account
- have your GitHub credentials entered into RStudio
- have your Personal Access Token (PAT) entered in RStudio

Note: occasionally the function seems to fail to set the master branch and users are unable to push changes. If this happens try running `git push -u origin master` in the terminal, this should set your current branch as the master. We're not sure why this happens, but it is advisable to use `fs_connect_github()` to set your credentials properly before trying this function.

## Value

R project is turned into a git repo and an associated github repo is created in the users github account.

---

set_databricks_pat	<i>Set databricks PAT</i>
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## Description

Simple function to set databricks PAT. Done via function to avoid committing to GitHub accidentally.

## Usage

```
set_databricks_pat()
```

## Value

databricks PAT written to environment

## Author(s)

Josh Moatt

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`uc_volume_pull`*Pull data from the DASH Unity Catalog into an RStudio cluster*

---

**Description**

Currently the DASH Unity Catalog (UC) and RStudio server cannot directly interact. So you must interact with the databricks API and copy data from DASH UC to the RStudio server. This is a simple function to read data from UC into RStudio cluster on databricks.

Note, you must have set up a databricks PAT in the databricks settings before doing this.

**Usage**

```
uc_volume_pull(workspace, volume, token, out_file)
```

**Arguments**

<code>workspace</code>	databricks workspace (string)
<code>volume</code>	file path to data folder (string)
<code>token</code>	databricks PAT (string)
<code>out_file</code>	filepath on RStudio Cluster that data should be stored.

**Value**

data saved in Rstudio cluster.

**Author(s)**

Josh Moatt

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