

# Package ‘gcspikelite’

February 22, 2024

**Version** 1.40.0

**Date** 2010/10/16

**Title** Spike-in data for GC/MS data and methods within flagme

**Author** Mark Robinson <mrobinson@wehi.edu.au>

**Maintainer** Mark Robinson <mrobinson@wehi.edu.au>

**Depends** R (>= 2.5.0)

**Description** Spike-in data for GC/MS data and methods within flagme

**License** LGPL

**biocViews** MassSpectrometryData

**git\_url** <https://git.bioconductor.org/packages/gcspikelite>

**git\_branch** RELEASE\_3\_18

**git\_last\_commit** 9a2fd3b

**git\_last\_commit\_date** 2023-10-24

**Repository** Bioconductor 3.18

**Date/Publication** 2024-02-22

## R topics documented:

targets . . . . . 1

**Index** 3

---

targets *Targets file for gcspikelite data.*

---

## Description

List of filenames and experimental group

**Usage**

```
data(targets)
```

**Format**

Data frame with columns:

Filename a character vector

Group a character vector

# Index

\* **datasets**  
  targets, [1](#)

targets, [1](#)