

# Package ‘gcspikelite’

July 4, 2024

**Version** 1.43.0

**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** devel

**git\_last\_commit** c52976c

**git\_last\_commit\_date** 2024-04-30

**Repository** Bioconductor 3.20

**Date/Publication** 2024-07-04

## Contents

targets . . . . .	1
<b>Index</b>	<b>3</b>

---

targets	<i>Targets file for gcspikelite data.</i>
---------	---

---

## 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