

# Package ‘gcspikelite’

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

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**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>

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## Contents

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

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|         |                                           |
|---------|-------------------------------------------|
| targets | <i>Targets file for gcspikelite data.</i> |
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## 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](#)