

# Package ‘CONFESSdata’

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**Title** Example dataset for CONFESS package

**Version** 1.37.0

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**Author** Diana LOW and Efthimios MOTAKIS

**Maintainer** Diana LOW <dlow@imcb.a-star.edu.sg>

**Description** Example text-converted C01 image files for use in the CONFESS Bioconductor package.

**biocViews** HighThroughputImagingData,CellCulture,Homo\_sapiens\_Data,ExpressionData

**NeedsCompilation** no

**Depends** R (>= 3.3)

**License** GPL-2

**LazyData** true

**RoxygenNote** 5.0.1

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

**git\_branch** devel

**git\_last\_commit** c8f8c7e

**git\_last\_commit\_date** 2025-04-15

**Repository** Bioconductor 3.22

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## Description

Text-converted C01 image files generated on the Fluidigm C1 system

**Usage**

```
data("CONFESSdata")
```

**Format**

The format is: List of 6

- \$ BF : chr
- \$ CH1 : chr
- \$ CH2 : chr
- \$ separator : chr
- \$ image.type: chr
- \$ dateIndex : chr

**Value**

data frame describing the text files and other image info

**Source**

This data was generated at RIKEN and can be downloaded at [http://single-cell.clst.riken.jp/bluk\\_download/cell\\_images/](http://single-cell.clst.riken.jp/bluk_download/cell_images/)

**Examples**

```
data(CONFESSdata)
```

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## \* **datasets**

CONFESSdata, [1](#)

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