

# Package ‘mvoutData’

May 8, 2025

**Title** affy and illumina raw data for assessing outlier detector performance

**Description** affy and illumina raw data for assessing outlier detector performance

**Version** 1.45.0

**Author** VJ Carey <stvjc@channing.harvard.edu>

**Depends** R (>= 2.10.0), methods, Biobase (>= 2.5.5), affy (>= 1.23.4), lumi

**Maintainer** VJ Carey <stvjc@channing.harvard.edu>

**License** Artistic-2.0

**biocViews** ExperimentData, MicroarrayData

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

**git\_branch** devel

**git\_last\_commit** 42a93aa

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

**Repository** Bioconductor 3.22

**Date/Publication** 2025-05-08

## Contents

s12c . . . . .	1
<b>Index</b>	<b>3</b>

---

s12c	<i>raw expression data for evaluation of outlier detection algorithms</i>
------	---

---

## Description

raw expression data for evaluation of outlier detection algorithms

## Usage

```
data(s12c); data(ILM1)
```

**Format**

s12c is an AffyBatch created from SpikeIn U133a, with first two samples contaminated. ILM1 is the result of lumi::lumiR applied to the CSV file supplied by Illumina Inc. for the MAQC study (19 arrays).

**Examples**

```
data(s12c)
s12c
```

# Index

\* **datasets**

s12c, [1](#)

ILM1 (s12c), [1](#)

s12c, [1](#)