

# Package ‘AmpAffyExample’

November 21, 2024

**Version** 1.47.0

**Date** 2009-07-22

**Title** Example of Amplified Data

**Author** Rafael A. Irizarry <rafa@ds.dfci.harvard.edu>

**Maintainer** Robert D Shear <rshear@ds.dfci.harvard.edu>

**URL** <https://bioconductor.org/packages/AmpAffyExample>

**BugReports** <https://github.com/rafalab/AmpAffyExample/issues>

**Depends** R (>= 2.4.0), affy (>= 1.23.4)

**Suggests** hgu133acdf

**Description** Six arrays. Three from amplified RNA, three from the typical procedure.

**License** LGPL (>= 2)

**biocViews** ExperimentData, MicroarrayData

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

**git\_branch** devel

**git\_last\_commit** 57ead8d

**git\_last\_commit\_date** 2024-10-29

**Repository** Bioconductor 3.21

**Date/Publication** 2024-11-21

## Contents

AmpData . . . . .	2
Index	3

---

AmpData

*AffyBatch* instance with six arrays from amplication experiment

---

**Description**

This [AffyBatch](#) represents part of an experiment to determine if one can use amplified RNA with Affymetrix GeneChip arrays.

**Usage**

```
data(AmpData)
```

**Format**

A [AffyBatch](#) containing 6 arrays.

**Source**

Anonymous for now...

# Index

## \* datasets

AmpData, [2](#)

AffyBatch, [2](#)

AmpData, [2](#)