

# Package ‘frmaExampleData’

April 23, 2026

**Title** Frma Example Data

**Version** 1.47.0

**Created** 2010-01-31

**Author** Matthew N. McCall <mmccall@jhsp.h.edu>

**Description** Data files used by the examples in frma and frmaTools packages

**Maintainer** Matthew N. McCall <mmccall@jhsp.h.edu>

**Depends** R (>= 2.10.0)

**License** GPL (>= 2)

**biocViews** Homo\_sapiens\_Data, MicroarrayData

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

**git\_branch** devel

**git\_last\_commit** 0774595

**git\_last\_commit\_date** 2025-10-29

**Repository** Bioconductor 3.23

**Date/Publication** 2026-04-23

## Contents

AffyBatch133a . . . . .	2
AffyBatch133atag . . . . .	2
AffyBatch133plus2 . . . . .	2
AffyBatchExample . . . . .	3

<b>Index</b>	<b>4</b>
--------------	----------

---

AffyBatch133a

*HGU133a Example Data*

---

**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133a)
```

**Examples**

```
data(AffyBatch133a)
```

---

AffyBatch133atag

*HGU133atag Example Data*

---

**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133atag)
```

**Examples**

```
data(AffyBatch133atag)
```

---

AffyBatch133plus2

*HGU133plus2 Example Data*

---

**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133plus2)
```

**Examples**

```
data(AffyBatch133plus2)
```

---

AffyBatchExample      *AffyBatch Example Data*

---

**Description**

An example AffyBatch for use with frma.

**Usage**

```
data(AffyBatchExample)
```

**Examples**

```
data(AffyBatchExample)
```

# Index

## \* datasets

AffyBatch133a, [2](#)

AffyBatch133atag, [2](#)

AffyBatch133plus2, [2](#)

AffyBatchExample, [3](#)

AffyBatch133a, [2](#)

AffyBatch133atag, [2](#)

AffyBatch133plus2, [2](#)

AffyBatchExample, [3](#)