

# Package ‘frmaExampleData’

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**Title** Frma Example Data

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

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**Depends** R (>= 2.10.0)

**License** GPL (>= 2)

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

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

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AffyBatch133a

*HGU133a Example Data*

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**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133a)
```

**Examples**

```
data(AffyBatch133a)
```

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AffyBatch133atag

*HGU133atag Example Data*

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**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133atag)
```

**Examples**

```
data(AffyBatch133atag)
```

---

AffyBatch133plus2

*HGU133plus2 Example Data*

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**Description**

An example AffyBatch for use with frmaTools.

**Usage**

```
data(AffyBatch133plus2)
```

**Examples**

```
data(AffyBatch133plus2)
```

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AffyBatchExample      *AffyBatch Example Data*

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**Description**

An example AffyBatch for use with frma.

**Usage**

```
data(AffyBatchExample)
```

**Examples**

```
data(AffyBatchExample)
```

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