

# Package ‘ALLMLL’

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**Title** A subset of arrays from a large acute lymphoblastic leukemia (ALL) study

**Version** 1.46.0

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**Description** This package provides probe-level data for 20 HGU133A and 20 HGU133B arrays which are a subset of arrays from a large ALL study. The data is for the MLL arrays. This data was published in Mary E. Ross, Xi-aodong Zhou, Guangchun Song, Sheila A. Shurtleff, Kevin Girtman, W. Kent Williams, Hsi-Che Liu, Rami Mahfouz, Susana C. Raimondi, Noel Lenny, Anami Patel, and James R. Downing (2003) Classification of pediatric acute lymphoblastic leukemia by gene expression profiling *Blood* 102: 2951-2959

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**Depends** R (>= 2.10), affy (>= 1.23.4)

**License** GPL-2

**biocViews** ExperimentData, CancerData, LeukemiaCancerData, MicroarrayData

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MLL

*AffyBatch* instances *MLL.A* and *MLL.B*

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### Description

These [AffyBatch](#) objects contain a subset of arrays from a large acute lymphoblastic leukemia (ALL) study.

### Usage

```
data(MLL.A)
data(MLL.B)
```

### Format

Each are [AffyBatch](#) containing 20 arrays.

### Source

This package provides probe-level data for 20 HGU133A and 20 HGU133B arrays which are a subset of arrays from a large ALL study. The data is for the MLL arrays. This data was published in:

Mary E. Ross, Xiaodong Zhou, Guangchun Song, Sheila A. Shurtleff, Kevin Girtman, W. Kent Williams, Hsi-Che Liu, Rami Mahfouz, Susana C. Raimondi, Noel Lenny, Anami Patel, and James R. Downing (2003) *Classification of pediatric acute lymphoblastic leukemia by gene expression profiling* Blood 102: 2951-2959

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\* **datasets**

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