

# Package ‘ITALICSData’

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**Title** ITALICSData

**Depends** R (>= 2.0.0)

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**Description** Data needed to use the ITALICS package

**License** GPL

**URL** <http://bioinfo.curie.fr>

**biocViews** ExperimentData, HapMap

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

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Hind.QuartetEffect	<i>Hind.QuartetEffect</i>
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**Description**

It contains computed quartet effect for Hind array. Those were computed on some normal samples of the hapmap dataset.

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Nsp.QuartetEffect	<i>Nsp.QuartetEffect</i>
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**Description**

It contains computed quartet effect for Nsp array. Those were computed on some normal samples of the hapmap dataset.

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quartet	<i>Quartet Info example for the vignette</i>
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**Description**

An example for the vignette

**Usage**

```
data(quartetInfo)
```

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snpInfo	<i>snp Info example for the vignette</i>
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**Description**

An example for the vignette

**Usage**

```
data(snpInfo)
```

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Sty.QuartetEffect      *Sty.QuartetEffect*

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**Description**

It contains computed quartet effect for Sty array. Those were computed on some normal samples of the hapmap dataset.

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Xba.QuartetEffect      *Xba.QuartetEffect*

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**Description**

It contains computed quartet effect for Xba array. Those were computed on some normal samples of the hapmap dataset.

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