

# Package ‘ITALICSData’

January 7, 2025

**Version** 2.45.0

**Date** 2008-05-01

**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>

**git\_branch** devel

**git\_last\_commit** 910eb87

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

**Repository** Bioconductor 3.21

**Date/Publication** 2025-01-07

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Hind.QuartetEffect      *Hind.QuartetEffect*

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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      *Nsp.QuartetEffect*

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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      *Quartet Info example for the vignette*

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**Description**

An example for the vignette

**Usage**

```
data(quartetInfo)
```

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snpInfo      *snp Info example for the vignette*

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**Description**

An example for the vignette

**Usage**

```
data(snpInfo)
```

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Sty.QuartetEffect	<i>Sty.QuartetEffect</i>
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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	<i>Xba.QuartetEffect</i>
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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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