

# Package ‘ITALICSData’

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**Title** ITALICSData

**Depends** R (>= 2.0.0)

**Author** Guillem Rigai

**Maintainer** Guillem Rigai <italics@curie.fr>

**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      *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)
```

---

snpInfo      *snp Info example for the vignette*

---

**Description**

An example for the vignette

**Usage**

```
data(snpInfo)
```

---

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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