

# Package ‘ITALICSData’

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**Version** 2.51.0  
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**Title** ITALICSData  
**Depends** R (>= 2.0.0)  
**Author** Guillem Rigail  
**Maintainer** Guillem Rigail <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>  
**git\_branch** devel  
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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	<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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