

Package ‘BiocStyle’

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Title Standard styles for vignettes and other Bioconductor documents

Description Provides standard formatting styles for Bioconductor documents. The vignette illustrates use and functionality.

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Author Martin Morgan, Andrzej Oles, Wolfgang Huber

Maintainer Bioconductor Package Maintainer <maintainer@bioconductor.org>

biocViews Software

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| latex | <i>Use Bioconductor styles to format LaTeX vignettes</i> |
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Description

This function inserts code into a document preamble to provide a consistent style to Bioconductor vignettes

Usage

```
latex(..., width=90, short.fignames=FALSE, fig.path)
```

Arguments

| | |
|-----------------------------|--|
| <code>...</code> | Additional arguments, passed to options . |
| <code>width</code> | integer(1), maximum number of columns on a line used in printing. See options . |
| <code>short.fignames</code> | logical, indicates whether <code>\incfig</code> figures should be inserted and referred to using short names equal to corresponding code chunk labels without any additional prefix. |
| <code>fig.path</code> | character, custom prefix to be used for figure file names when used with <code>knitr</code> ; has no effect when compiled with Sweave. For details see opts_chunk . |

Details

Use is described in the ‘Bioconductor LaTeX Style’ vignette.

Value

The return value is a character string directing LaTeX to use the Bioconductor style file.

A side effect is setting any options specified as arguments.

Author(s)

Martin Morgan mtmorgan@fhcrc.org, Andrzej Oles, Wolfgang Huber

Examples

```
## location of the .sty file
system.file(package = "BiocStyle", "sty", "Bioconductor.sty")
```

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