

Package ‘zebrafishRNASeq’

October 5, 2024

Title Zebrafish RNA-Seq Experimental Data from Ferreira et al. (2014)

Version 1.25.1

Author Davide Risso

Description Gene-level read counts from RNA-Seq for gallein-treated and control zebrafish.

biocViews ExperimentData, Danio_rerio_Data, RNASeqData

Maintainer Davide Risso <risso.davide@gmail.com>

License GPL

VignetteBuilder knitr

Suggests BiocStyle, knitr

Depends R (>= 2.10)

git_url <https://git.bioconductor.org/packages/zebrafishRNASeq>

git_branch devel

git_last_commit ad13a44

git_last_commit_date 2024-06-07

Repository Bioconductor 3.20

Date/Publication 2024-10-05

Contents

zfGenes	1
Index	3

zfGenes	<i>Zebrafish RNA-Seq Experimental Data from Ferreira et al. (2014)</i>
---------	------------------------------------------------------------------------

Description

Gene-level read counts from the RNA-Seq experiment of Ferreira et al. (2014). The data correspond to RNA libraries for three pairs of gallein-treated and control embryonic zebrafish cell pools. Please see the vignette for details on pre-processing.

Usage

```
data(zfGenes)
```

Source

A citation for the dataset can be generated using `citation("zebrafishRNASeq")`.

Index

* datasets

zfGenes, [1](#)

zebrafishRNASeq (zfGenes), [1](#)

zfGenes, [1](#)