

Package ‘zebrafishRNASeq’

October 12, 2023

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

Version 1.20.0

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 RELEASE_3_17

git_last_commit c46831b

git_last_commit_date 2023-04-25

Date/Publication 2023-10-12

R topics documented:

zfGenes 1

Index 3

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

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](#)