

Package ‘zebrafishRNASeq’

October 10, 2024

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

Version 1.25.1

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

biocViews ExperimentData, Danio_rerio_Data, RNASeqData

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License GPL

VignetteBuilder knitr

Suggests BiocStyle, knitr

Depends R (>= 2.10)

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

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* datasets

zfGenes, [1](#)

zebrafishRNASeq (zfGenes), [1](#)

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