

# Package ‘zebrafishRNASeq’

April 12, 2018

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

**Version** 0.112.0

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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)

**NeedsCompilation** no

## R topics documented:

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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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## \*Topic **datasets**

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