

Package ‘spatialDmelxsim’

November 18, 2025

Title Spatial allelic expression counts for fly cross embryo

Version 1.17.0

Description Spatial allelic expression counts from Combs & Fraser (2018), compiled into a SummarizedExperiment object. This package contains data of allelic expression counts of spatial slices of a fly embryo, a *Drosophila melanogaster* x *Drosophila simulans* cross. See the CITATION file for the data source, and the associated script for how the object was constructed from publicly available data.

Imports utils

Depends R (>= 4.1), ExperimentHub, SummarizedExperiment

License GPL-3

Encoding UTF-8

Roxygen list(markdown = TRUE)

RoxygenNote 7.1.1

URL <https://github.com/mikelove/spatialDmelxsim>

BugReports <https://github.com/mikelove/spatialDmelxsim/issues>

Suggests knitr, rmarkdown, BiocStyle

VignetteBuilder knitr

biocViews ExperimentHub, SequencingData, RNASeqData, ExpressionData, *Drosophila_melanogaster_Data*, GEO

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

git_branch devel

git_last_commit dcbc1d5

git_last_commit_date 2025-10-29

Repository Bioconductor 3.23

Date/Publication 2025-11-18

Author Michael Love [aut, cre]

Maintainer Michael Love <michaelisaiahlove@gmail.com>

Contents

spatialDmelxsim	2
Index	3

spatialDmelxsim	<i>Spatial allelic expression counts for fly cross embryo</i>
-----------------	---

Description

Allelic expression counts of spatial slices of a fly embryo from Combs & Fraser (2018), a D melanogaster x D simulans reciprocal cross

References

Combs PA, Fraser HB (2018) Spatially varying cis-regulatory divergence in Drosophila embryos elucidates cis-regulatory logic. PLOS Genetics 14(11): e1007631.

Examples

```
suppressPackageStartupMessages(library(SummarizedExperiment))
se <- spatialDmelxsim()
se
```

Index

`spatialDmelxsim`, [2](#)