Package 'curatedAdipoRNA'

April 3, 2025

```
Type Package
Title A Curated RNA-Seq Dataset of MDI-induced Differentiated
     Adipocytes (3T3-L1)
Version 1.23.0
Year 2019
Description A curated dataset of RNA-Seq samples. The samples are MDI-induced
     pre-phagocytes (3T3-L1) at different time points/stage of differentiation.
     The package document the data collection, pre-processing and processing. In
     addition to the documentation, the package contains the scripts that was used
     to generated the data.
License GPL-3
URL https://github.com/MahShaaban/curatedAdipoRNA
BugReports https://github.com/MahShaaban/curatedAdipoRNA/issues
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RoxygenNote 6.1.1
LazyData TRUE
Depends R (>= 3.6), SummarizedExperiment
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Author Mahmoud Ahmed [aut, cre] (ORCID:
     <https://orcid.org/0000-0002-4377-6541>)
Maintainer Mahmoud Ahmed <mahmoud.s.fahmy@students.kasralainy.edu.eg>
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2 adipo_counts

Contents

	adipo_counts curatedAdipoI																					
Index																						4
adipo	_counts	Gei	ne c	сои	nts	in	dij	ffer	en	tia	tir	ıg a	adi	ро	cyi	tes						

Description

Gene counts in differentiating adipocytes

Usage

adipo_counts

Format

A RangedSummarizedExperiment object contains:

```
assay The gene counts matrix.
```

colData The phenotype data and quality control data of the samples.

rowRanges The feature data at gene level.

metadata The study level metadata which contains one object called studies. This is a data. frame of bibliography information of the studies from which the samples were collected.

Examples

```
# load the data object
data('adipo_counts')
# print the object
adipo_counts
```

curatedAdipoRNA 3

curatedAdipoRNA	curatedAdipoRNA package	

Description

A Curated RNA-Seq Dataset of MDI-induced Differentiated Adipocytes (3T3-L1)

Details

A curated dataset of RNA-Seq samples. The samples are MDI-induced pre-phagocytes (3T3-L1) at different time points/stage of differentiation. The package document the data collection, pre-processing and processing. In addition to the documentation, the package contains the scripts that was used to generated the data. The datasets and the pipeline used to process it are documented in adipo_counts and the package vignette.

Index