Resource Summary Report

Generated by NIF on May 24, 2025

SPEX2

RRID:SCR_014923

Type: Tool

Proper Citation

SPEX2 (RRID:SCR_014923)

Resource Information

URL: http://www.sailing.cs.cmu.edu/main/?page_id=511

Proper Citation: SPEX2 (RRID:SCR_014923)

Description: Automatic software program for profiling spatial gene expression patterns from Fly embryo ISH images. It utilizes image-based genome-scale profiling of whole-body mRNA patterns.

Resource Type: sequence analysis software, software application, data analysis software, data processing software, software resource

Defining Citation: PMID:20529936

Keywords: software, spatial gene expression, fly, embryo, extraction, mrna, bio.tools

Funding:

Resource Name: SPEX2

Resource ID: SCR_014923

Alternate IDs: biotools:spex2

Alternate URLs: https://bio.tools/spex2

Record Creation Time: 20220129T080323+0000

Record Last Update: 20250524T060555+0000

Ratings and Alerts

No rating or validation information has been found for SPEX2.

No alerts have been found for SPEX2.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Wang K, et al. (2021) SUV39H2/KMT1B Inhibits the cardiomyocyte senescence phenotype by down-regulating BTG2/PC3. Aging, 13(18), 22444.