

Resource Summary Report

Generated by [NIF](#) on Jul 31, 2026

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

Defining Citation: [PMID:20529936](#)

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

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: 20260731T042918+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.