# **Resource Summary Report**

Generated by NIF on Apr 25, 2025

# **Allim**

RRID:SCR\_012114

Type: Tool

### **Proper Citation**

Allim (RRID:SCR\_012114)

#### **Resource Information**

URL: https://code.google.com/p/allim/

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**Description:** A user-friendly software tool to estimate allele-specific gene expression.

Resource Type: software resource

**Defining Citation:** PMID:23615333

Keywords: standalone software, bio.tools

**Funding:** 

Resource Name: Allim

Resource ID: SCR\_012114

Alternate IDs: biotools:allim, OMICS\_05504

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

**Record Creation Time:** 20220129T080308+0000

**Record Last Update:** 20250420T014606+0000

### **Ratings and Alerts**

No rating or validation information has been found for Allim.

No alerts have been found for Allim.

#### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 2 mentions in open access literature.

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

Díaz F, et al. (2023) Transcriptional misexpression in hybrids between species linked by gene flow is associated with patterns of sequence divergence. Genome biology and evolution, 15(5).

Chen J, et al. (2015) Temperature stress mediates decanalization and dominance of gene expression in Drosophila melanogaster. PLoS genetics, 11(2), e1004883.