

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 1, 2026

## SEBIDA: Sex Bias Database

RRID:SCR\_007918

Type: Tool

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### Proper Citation

SEBIDA: Sex Bias Database (RRID:SCR\_007918)

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### Resource Information

**URL:** <http://www.sebida.de/>

**Proper Citation:** SEBIDA: Sex Bias Database (RRID:SCR\_007918)

**Description:** A database for the functional and evolutionary analysis of sex-biased genes. Sebida integrates data from multiple microarray studies comparing male versus female gene expression in *D. melanogaster*, *D. simulans*, and *A. gambiae*. In addition to the ratio of male to female (or testes to ovaries) expression for each gene, Sebida provides information useful for evolutionary studies, including measures of recombination, codon bias, and interspecific divergence.

**Abbreviations:** SEBIDA

**Synonyms:** Sex Bias Database

**Resource Type:** data or information resource, database

**Resource Name:** SEBIDA: Sex Bias Database

**Resource ID:** SCR\_007918

**Record Creation Time:** 20220129T080244+0000

**Record Last Update:** 20260801T190952+0000

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### Ratings and Alerts

No rating or validation information has been found for SEBIDA: Sex Bias Database.

No alerts have been found for SEBIDA: Sex Bias Database.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

Hu TT, et al. (2013) A second-generation assembly of the *Drosophila simulans* genome provides new insights into patterns of lineage-specific divergence. *Genome research*, 23(1), 89.

Grath S, et al. (2012) Rate of amino acid substitution is influenced by the degree and conservation of male-biased transcription over 50 myr of *Drosophila* evolution. *Genome biology and evolution*, 4(3), 346.