

Resource Summary Report

Generated by [NIF](#) on Sep 2, 2026

Rat Anti-Drosophila vasa Antibody, Unconjugated

RRID:AB_760351

Type: Antibody

Proper Citation

DSHB Cat# anti-vasa, RRID:AB_760351

Antibody Information

URL: http://antibodyregistry.org/AB_760351

Proper Citation: DSHB Cat# anti-vasa, RRID:AB_760351

Target Antigen: Rat Drosophila vasa

Host Organism: rat

Clonality: unknown

Comments: manufacturer recommendations: IgM; IgM, kappa light chain Immunoblotting

Antibody Name: Rat Anti-Drosophila vasa Antibody, Unconjugated

Description: This unknown targets Rat Drosophila vasa

Target Organism: drosophilaarthropod

Antibody ID: AB_760351

Vendor: DSHB

Catalog Number: anti-vasa

Record Creation Time: 20231110T080034+0000

Record Last Update: 20260829T141042+0000

Ratings and Alerts

No rating or validation information has been found for Rat Anti-Drosophila vasa Antibody, Unconjugated.

No alerts have been found for Rat Anti-Drosophila vasa Antibody, Unconjugated.

Data and Source Information

Source: [Antibody Registry](#)

Usage and Citation Metrics

We found 43 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#).

Rice NP, et al. (2026) Rapid disassembly and Piwi-independent reassembly of Drosophila piRNA cluster heterochromatin following acute heat shock. *RNA (New York, N.Y.)*, 32(3), 359.

Ho S, et al. (2026) Aub, Vasa, and Armi concentration in phase-separated nuage is dispensable for piRNA biogenesis and transposon silencing. *RNA (New York, N.Y.)*, 32(3), 341.

Bener MB, et al. (2026) Protocol to evaluate a lineage marking system in the Drosophila testis. *STAR protocols*, 7(2), 104479.

Alizada A, et al. (2025) Transcriptional regulation of the piRNA pathway by Ovo in animal ovarian germ cells. *Genes & development*, 39(3-4), 221.

Easwaran S, et al. (2025) A genome-wide association study implicates the olfactory system in Drosophila melanogaster diapause-associated lifespan extension and fecundity. *eLife*, 13.

Lee SK, et al. (2025) Topoisomerase 3b facilitates piRNA biogenesis to promote transposon silencing and germ cell development. *Cell reports*, 44(4), 115495.

Taira Y, et al. (2025) RNA-binding protein Miso/CG44249 is crucial for minor splicing during oogenesis in Drosophila. *RNA (New York, N.Y.)*, 31(6), 822.

Bener MB, et al. (2025) Detection of dedifferentiated stem cells in the Drosophila testis. *iScience*, 28(10), 113564.

Poursaeid S, et al. (2025) Brinker regulates reciprocal outcomes of BMP signal between stem cells and differentiating cells. *bioRxiv : the preprint server for biology*.

Green NM, et al. (2024) Nuclear actin is a critical regulator of Drosophila female germline stem cell maintenance. *bioRxiv : the preprint server for biology*.

Rojas-R??os P, et al. (2024) piRNAs are regulators of metabolic reprogramming in stem cells. *Nature communications*, 15(1), 8405.

Ridwan SM, et al. (2024) Diffusible fraction of niche BMP ligand safeguards stem-cell differentiation. *Nature communications*, 15(1), 1166.

Jones SD, et al. (2024) Emerin preserves stem cell survival through maintenance of centrosome and nuclear lamina structure. *Development (Cambridge, England)*, 151(22).

Ho S, et al. (2023) Aub, Vasa and Armi localization to phase separated nuage is dispensable for piRNA biogenesis and transposon silencing in *Drosophila*. *bioRxiv* : the preprint server for biology.

Ninova M, et al. (2023) Pervasive SUMOylation of heterochromatin and piRNA pathway proteins. *Cell genomics*, 3(7), 100329.

Ayachit MS, et al. (2023) Atg1 modulates mitochondrial dynamics to promote germline stem cell maintenance in *Drosophila*. *Biochemical and biophysical research communications*, 643, 192.

Sotillos S, et al. (2022) A conserved function of Human DLC3 and *Drosophila* Cv-c in testis development. *eLife*, 11.

Zong Q, et al. (2022) Comparative Ubiquitome Analysis Reveals Deubiquitinating Effects Induced by *Wolbachia* Infection in *Drosophila melanogaster*. *International journal of molecular sciences*, 23(16).

Antel M, et al. (2022) Interchromosomal interaction of homologous Stat92E alleles regulates transcriptional switch during stem-cell differentiation. *Nature communications*, 13(1), 3981.

Fefelova EA, et al. (2022) Impaired function of rDNA transcription initiation machinery leads to derepression of ribosomal genes with insertions of R2 retrotransposon. *Nucleic acids research*, 50(2), 867.