

Resource Summary Report

Generated by [NIF](#) on Sep 17, 2026

Rab8[1] red[1] e[4]/TM6B, Sb[1] Tb[1] ca[1]

RRID:BDSC_26173

Type: Organism

Proper Citation

RRID:BDSC_26173

Organism Information

URL: <https://n2t.net/bdsc:26173>

Proper Citation: RRID:BDSC_26173

Description: Drosophila melanogaster with name Rab8[1] red[1] e[4]/TM6B, Sb[1] Tb[1] ca[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Jim Kennison, NIH, National Institute of Child Health and Human Development

Affected Gene: ca, e, Rab8, red, Sb, Tb

Genomic Alteration: Chromosome 3

Catalog Number: 26173

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_26173, BDSC:26173, BL26173

Organism Name: Rab8[1] red[1] e[4]/TM6B, Sb[1] Tb[1] ca[1]

Record Creation Time: 20240911T222433+0000

Record Last Update: 20260912T203449+0000

Ratings and Alerts

No rating or validation information has been found for Rab8[1] red[1] e[4]/TM6B, Sb[1] Tb[1] ca[1].

No alerts have been found for Rab8[1] red[1] e[4]/TM6B, Sb[1] Tb[1] ca[1].

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Chang YC, et al. (2024) Identification of secretory autophagy as a mechanism modulating activity-induced synaptic remodeling. Proceedings of the National Academy of Sciences of the United States of America, 121(16), e2315958121.

Ochi Y, et al. (2022) Stratum is required for both apical and basolateral transport through stable expression of Rab10 and Rab35 in Drosophila photoreceptors. Molecular biology of the cell, 33(10), br17.

Kohrs FE, et al. (2021) Systematic functional analysis of rab GTPases reveals limits of neuronal robustness to environmental challenges in flies. eLife, 10.

Wu Z, et al. (2020) CEP290 is essential for the initiation of ciliary transition zone assembly. PLoS biology, 18(12), e3001034.