

Resource Summary Report

Generated by SPARC SAWG on Sep 2, 2026

P{ry[+t7.2]=PZ}tara[03881] ry[506]/TM3, ry[RK] Sb[1] Ser[1]

RRID:BDSC_11613
Type: Organism

Proper Citation

RRID:BDSC_11613

Organism Information

URL: <https://n2t.net/bdsc:11613>

Proper Citation: RRID:BDSC_11613

Description: Drosophila melanogaster with name P{ry[+t7.2]=PZ}tara[03881] ry[506]/TM3, ry[RK] Sb[1] Ser[1] from BDSC.

Species: Drosophila melanogaster

Notes: Donor: Berkeley Drosophila Genome Project; Donor's Source: Allan Spradling, Carnegie Institution for Science

Affected Gene: EcoIIacZ, tara, ry, Sb, Ser

Genomic Alteration: Chromosome 3

Catalog Number: 11613

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_11613, BL11613

Organism Name: P{ry[+t7.2]=PZ}tara[03881] ry[506]/TM3, ry[RK] Sb[1] Ser[1]

Record Creation Time: 20240911T222239+0000

Record Last Update: 20260829T185632+0000

Ratings and Alerts

No rating or validation information has been found for P{ry[+t7.2]=PZ}tara[03881] ry[506]/TM3, ry[RK] Sb[1] Ser[1].

No alerts have been found for P{ry[+t7.2]=PZ}tara[03881] ry[506]/TM3, ry[RK] Sb[1] Ser[1].

Data and Source Information

Source: [Integrated Animals](#)

Source Database: Bloomington Drosophila Stock Center (BDSC)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Bose A, et al. (2025) The pioneer transcription factor Zelda controls the exit from regeneration and restoration of patterning in Drosophila. Science advances, 11(23), eads5743.

Bose A, et al. (2024) The pioneer transcription factor Zelda facilitates the exit from regeneration and restoration of patterning in Drosophila. bioRxiv : the preprint server for biology.

Tian Y, et al. (2021) Regulation of growth and cell fate during tissue regeneration by the two SWI/SNF chromatin-remodeling complexes of Drosophila. Genetics, 217(1), 1.