

Resource Summary Report

Generated by SPARC SAWG on Aug 30, 2026

P{ry[+t7.2]=PZ}CrebA[03576] ry[506]/TM3,
P{ry[+t7.2]=ftz-lacC}SC1, ry[RK] Sb[1] Ser[1]

RRID:BDSC_10183

Type: Organism

Proper Citation

RRID:BDSC_10183

Organism Information

URL: <https://n2t.net/bdsc:10183>

Proper Citation: RRID:BDSC_10183

Description: Drosophila melanogaster with name P{ry[+t7.2]=PZ}CrebA[03576] ry[506]/TM3, P{ry[+t7.2]=ftz-lacC}SC1, ry[RK] Sb[1] Ser[1] from BDSC.

Species: Drosophila melanogaster

Notes: May be segregating w[1118]. Donor: Berkeley Drosophila Genome Project; Donor's Source: Allan Spradling, Carnegie Institution for Science

Affected Gene: EcoIIacZ, ftz, CrebA, ry, Sb, Ser

Genomic Alteration: Chromosome 3

Catalog Number: 10183

Database: Bloomington Drosophila Stock Center (BDSC)

Database Abbreviation: BDSC

Availability: available

Alternate IDs: BDSC_10183, BL10183

Organism Name: P{ry[+t7.2]=PZ}CrebA[03576] ry[506]/TM3, P{ry[+t7.2]=ftz-lacC}SC1, ry[RK] Sb[1] Ser[1]

Record Creation Time: 20240911T222230+0000

Record Last Update: 20260829T185620+0000

Ratings and Alerts

No rating or validation information has been found for P{ry[+t7.2]=PZ}CrebA[03576] ry[506]/TM3, P{ry[+t7.2]=ftz-lacC}SC1, ry[RK] Sb[1] Ser[1].

No alerts have been found for P{ry[+t7.2]=PZ}CrebA[03576] ry[506]/TM3, P{ry[+t7.2]=ftz-lacC}SC1, 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 4 mentions in open access literature.

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

Li J, et al. (2026) Profiling the host defense responses against *Candida auris* in a reliable *Drosophila melanogaster* infection model. *mLife*, 5(1), 51.

Wang X, et al. (2026) An insulin receptor activity surge in follicle cells drives vitellogenesis by upregulating CrebA. *EMBO reports*, 27(3), 748.

Jackson DJ, et al. (2025) CrebA regulation of secretory capacity: Genome-wide transcription profiling coupled with in vivo DNA binding studies. *bioRxiv : the preprint server for biology*.

Jackson DJ, et al. (2025) CrebA regulation of secretory capacity: genome-wide transcription profiling coupled with in vivo DNA binding studies. *Genetics*, 231(4).