

Resource Summary Report

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DroSpeGe: Drosophila Species Genomes

RRID:SCR_013438

Type: Tool

Proper Citation

DroSpeGe: Drosophila Species Genomes (RRID:SCR_013438)

Resource Information

URL: <http://insects.eugenescience.org/DroSpeGe/>

Proper Citation: DroSpeGe: Drosophila Species Genomes (RRID:SCR_013438)

Description: DroSpeGe provides a preview of newly sequenced Drosophila genomes, with genome maps and BLAST sequence search. Genome maps include D. melanogaster genome homology, homologies to nine eukaryote proteomes, marker gene locations, and Drosophila microsatellites. Current genome assemblies can be BLASTed, with links to genome maps of your BLAST matches. Annotation data are available in Gene Finding Format (GFF) for these whole genome comparisons.

Synonyms: DroSpeGe

Resource Type: database, data or information resource

Keywords: drosophila melanogaster, drosophila microsatellite, eukaryote proteome, d. melanogaster, d. melanogaster genome, drosophila genome, genome map

Resource Name: DroSpeGe: Drosophila Species Genomes

Resource ID: SCR_013438

Alternate IDs: nif-0000-02770

Record Creation Time: 20220129T080316+0000

Record Last Update: 20260803T040616+0000

Ratings and Alerts

No rating or validation information has been found for DroSpeGe: Drosophila Species Genomes.

No alerts have been found for DroSpeGe: Drosophila Species Genomes.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Nie H, et al. (2017) Identification of genes related to high royal jelly production in the honey bee (*Apis mellifera*) using microarray analysis. Genetics and molecular biology, 40(4), 781.