

Resource Summary Report

Generated by T1D on Aug 2, 2026

IntegromeDB

RRID:SCR_004620

Type: Tool

Proper Citation

IntegromeDB (RRID:SCR_004620)

Resource Information

URL: <http://integromedb.org/>

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Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented May 26, 2016. Search engine that integrates over 100 curated and publicly contributed data sources and provides integrated views on the genomic, proteomic, transcriptomic, genetic and functional information currently available. Information featured in the database includes gene function, orthologies, gene expression, pathways and protein-protein interactions, mutations and SNPs, disease relationships, related drugs and compounds.

Synonyms: Integrome DB

Resource Type: data or information resource, database

Defining Citation: [PMID:22260095](#), [PMID:20427517](#)

Keywords: catalog, search engine, gene, protein, gene regulation, gene expression, protein-protein interaction, pathway, metagenomics, mutation, disease, transcriptional regulation, genomics, transcriptomics, genetics, function, interaction, ortholog

Funding: NIH ;
NIGMS R01 GM084881

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: IntegromeDB

Resource ID: SCR_004620

Alternate IDs: nlx_63198

Record Creation Time: 20220129T080225+0000

Record Last Update: 20260801T190922+0000

Ratings and Alerts

No rating or validation information has been found for IntegromeDB.

No alerts have been found for IntegromeDB.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Esch M, et al. (2015) LAILAPS: the plant science search engine. Plant & cell physiology, 56(1), e8.

Kozhenkov S, et al. (2011) BiologicalNetworks--tools enabling the integration of multi-scale data for the host-pathogen studies. BMC systems biology, 5, 7.

Kozhenkov S, et al. (2010) BiologicalNetworks 2.0--an integrative view of genome biology data. BMC bioinformatics, 11, 610.