

# Resource Summary Report

Generated by [dkNET](#) on Aug 26, 2026

## Yeast Intron Database

RRID:SCR\_007144

Type: Tool

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### Proper Citation

Yeast Intron Database (RRID:SCR\_007144)

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### Resource Information

**URL:** [http://compbio.soe.ucsc.edu/yeast\\_introns.html](http://compbio.soe.ucsc.edu/yeast_introns.html)

**Proper Citation:** Yeast Intron Database (RRID:SCR\_007144)

**Description:** Database of information about the spliceosomal introns of the yeast *Saccharomyces cerevisiae*. Listed are known spliceosomal introns in the yeast genome and the splice sites actually used are documented. Through the use of microarrays designed to monitor splicing, they are beginning to identify and analyze splice site context in terms of the nature and activities of the trans-acting factors that mediate splice site recognition. In version 3.0, expression data that relates to the efficiency of splicing relative to other processes in strains of yeast lacking nonessential splicing factors is included. These data are displayed on each intron page for browsing and can be downloaded for other types of analysis.

**Abbreviations:** Yeast Intron Database

**Synonyms:** Ares lab Yeast Intron Database

**Resource Type:** data or information resource, database

**Defining Citation:** [PMID:11988574](#)

**Keywords:** intron, spliceosomal, splicing, genome, intron splice signal, sequence, splice site

**Funding:** W. M. Keck Foundation ;  
Packard Foundation ;  
NIH

**Availability:** The community can contribute to this resource

**Resource Name:** Yeast Intron Database

**Resource ID:** SCR\_007144

**Alternate IDs:** nif-0000-03649, OMICS\_01890

**Alternate URLs:** [http://www.cse.ucsc.edu/research/compbio/yeast\\_introns.html](http://www.cse.ucsc.edu/research/compbio/yeast_introns.html)

**Record Creation Time:** 20220129T080240+0000

**Record Last Update:** 20260821T193818+0000

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## Ratings and Alerts

No rating or validation information has been found for Yeast Intron Database.

No alerts have been found for Yeast Intron Database.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Barrass JD, et al. (2015) Transcriptome-wide RNA processing kinetics revealed using extremely short 4tU labeling. Genome biology, 16, 282.

Ladd AN, et al. (2002) Finding signals that regulate alternative splicing in the post-genomic era. Genome biology, 3(11), reviews0008.