

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

Hollywood

RRID:SCR_002920

Type: Tool

Proper Citation

Hollywood (RRID:SCR_002920)

Resource Information

URL: <http://hollywood.mit.edu/hollywood/>

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Description: Database built upon genomic annotation of splicing patterns of known genes derived from spliced alignment of complementary DNAs (cDNAs) and expressed sequence tags for human and mouse. It links features such as splice site sequence and strength, exonic splicing enhancers and silencers, conserved and non-conserved patterns of splicing, and cDNA library information for inferred alternative exons. A query tool allows searches for sets of exons with specific splicing characteristics or splicing regulatory element composition, or gives a graphical or sequence-level summary of splicing patterns for a specific gene. A graphical representation of gene splicing patterns is provided, and these patterns can alternatively be layered onto existing information in the UCSC Genome Browser.

Abbreviations: Hollywood

Synonyms: Hollywood.mit.edu, Hollywood mRNA Alternative Splicing, Hollywood.mit.edu - Alternatively spliced mRNA

Resource Type: data or information resource, database

Defining Citation: [PMID:16381932](#)

Keywords: alternative splicing, exon, mrna, splicing, annotation, regulatory element, ortholog, splicing pattern, cdna

Availability: Account required

Resource Name: Hollywood

Resource ID: SCR_002920

Alternate IDs: OMICS_01889, nif-0000-02972

Record Creation Time: 20220129T080216+0000

Record Last Update: 20260919T195637+0000

Ratings and Alerts

No rating or validation information has been found for Hollywood.

No alerts have been found for Hollywood.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.