

Resource Summary Report

Generated by [RRID](#) on Aug 10, 2026

SpliceInfo

RRID:SCR_005100

Type: Tool

Proper Citation

SpliceInfo (RRID:SCR_005100)

Resource Information

URL: <http://spliceinfo.mbc.nctu.edu.tw/>

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Description: A database of mRNA alternative splicing in the human genome. Within it, several modes of mRNA alternative splicing, such as exon skipping, alternative 5"-splicing sites, alternative 3"-splicing sites and mutually exclusive exons are computationally derived and extracted. Finally, for each type of alternative splicing, the flanking intronic sequences are collected and then exploited by motif discovery tools. The tissue-specific information and gene functionalities that correspond to the selected regions are also considered. The database provides a means of investigating alternative splicing and can be used for identifying alternative splicing - related motifs, such as the exonic splicing enhancer (ESE), the exonic splicing silencer (ESS) and other intronic splicing motifs.

Synonyms: SpliceInfo

Resource Type: database, data or information resource

Keywords: exon, human genome, intron, mrna alternative splicing, mrna splicing

Resource Name: SpliceInfo

Resource ID: SCR_005100

Alternate IDs: nif-0000-03488

Record Creation Time: 20220129T080228+0000

Record Last Update: 20260808T190350+0000

Ratings and Alerts

No rating or validation information has been found for SpliceInfo.

No alerts have been found for SpliceInfo.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.