

Resource Summary Report

Generated by SPARC SAWG on Aug 11, 2026

SplitSeek

RRID:SCR_001012

Type: Tool

Proper Citation

SplitSeek (RRID:SCR_001012)

Resource Information

URL: <https://omictools.com/splitseek-tool>

Proper Citation: SplitSeek (RRID:SCR_001012)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 20, 2016. A program for de novo prediction of splice junctions in RNA-seq data.

Resource Type: data analysis software, software application, software resource, sequence analysis software, data processing software

Defining Citation: [PMID:20236510](https://pubmed.ncbi.nlm.nih.gov/20236510/)

Keywords: bioinformatics alignment, sequence analysis software, de novo, prediction, rna seq, rna, splice junction, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SplitSeek

Resource ID: SCR_001012

Alternate IDs: biotools:splitseek, OMICS_01253

Alternate URLs: <https://bio.tools/splitseek>

Old URLs: <http://www.uppmax.uu.se/software/splitseek>

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260811T043119+0000

Ratings and Alerts

No rating or validation information has been found for SplitSeek.

No alerts have been found for SplitSeek.

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 [SPARC SAWG](#) .

Chen Q, et al. (2025) Systems-level immunomonitoring in children with solid tumors to enable precision medicine. Cell, 188(5), 1425.