

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

InsPecT

RRID:SCR_024038

Type: Tool

Proper Citation

InsPecT (RRID:SCR_024038)

Resource Information

URL: <http://proteomics.ucsd.edu/Software/Inspect/>

Proper Citation: InsPecT (RRID:SCR_024038)

Description: Software tool to addresses several algorithmic problems in order to identify modified proteins. Software MS/MS database search tool specifically designed to address two crucial needs of the proteomics community: post-translational modification identification and search speed.

Synonyms: Inspect, inspect

Resource Type: software application, software resource

Defining Citation: [PMID:16013882](#)

Keywords: addresses algorithmic problems, identify modified proteins, post-translational modification identification,

Availability: Free, Available for download, Freely available,

Resource Name: InsPecT

Resource ID: SCR_024038

Alternate IDs: OMICS_02397

Old URLs: <https://sources.debian.org/src/inspect/>

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260905T133316+0000

Ratings and Alerts

No rating or validation information has been found for InsPecT.

No alerts have been found for InsPecT.

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 [PRECISE-TBI](#) .

Tan JY, et al. (2020) Splicing of enhancer-associated lincRNAs contributes to enhancer activity. Life science alliance, 3(4).