Resource Summary Report

Generated by NIF on May 29, 2025

SciBite

RRID:SCR_003952
Type: Tool

Proper Citation

SciBite (RRID:SCR_003952)

Resource Information

URL: http://scibite.com/

Proper Citation: SciBite (RRID:SCR_003952)

Description: An informatics service focusing on competitor intelligence and text-mining within the life sciences. It mines a diverse array of scientific content to look for mentions of targets, indications, companies, drugs and technologies of interest to those involved in human health and drug discovery. A free newsletter service is provided to allow everyone to explore the therapeutic landscape and stay informed and a range of professional products and services are offered for those who need more in-depth analysis of the therapeutic landscape. Its text-mining engine can be configured to a variety of different use cases and performs with very high throughput. SciBite has a number of big-pharma and scientific publishers as current customers. SciBITES is free on the website but for extended functionality, including a data mining API and access to internal or commercial documents, you may wish to consider an installation of the SciBITES server software within your organization.

Abbreviations: SciBite

Synonyms: SciBite Limited

Resource Type: commercial organization

Keywords: text-mining, biomedical, drug discovery, pharmaceutical, data mining, biotechnology, drug, newsletter, semantics, ontology, termite, text-mining software

Funding:

Availability: Free, Subscription, API access, License

Resource Name: SciBite

Resource ID: SCR_003952

Alternate IDs: nlx_158349

Record Creation Time: 20220129T080221+0000

Record Last Update: 20250525T030806+0000

Ratings and Alerts

No rating or validation information has been found for SciBite.

No alerts have been found for SciBite.

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 NIF.

Ferrero E, et al. (2017) In silico prediction of novel therapeutic targets using gene-disease association data. Journal of translational medicine, 15(1), 182.