

# Resource Summary Report

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## Seek and Blastn

RRID:SCR\_016625

Type: Tool

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### Proper Citation

Seek and Blastn (RRID:SCR\_016625)

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### Resource Information

**URL:** <http://scigendetection.imag.fr/TPD52/>

**Proper Citation:** Seek and Blastn (RRID:SCR\_016625)

**Description:** Web server as semi automated tool that identifies nucleotide sequences in scientific publications, determines whether they are correctly identified. Seeks for nucleotide sequences in pdf files and then call local version of blastn. Text mining and text analysis tool to allow peers to identify errors in published or forthcoming scientific literature. Used to search and extract gene identifiers and nucleotide sequences against human genomic and transcript database.

**Synonyms:** Seek & Blastn, Seek and blastn

**Resource Type:** analysis service resource, data access protocol, data analysis service, production service resource, service resource, software resource, web service

**Keywords:** Nucleotide sequence identification, text mining, text analysis, scientific publication, error identification, ASWG

**Funding:** Universite Grenoble Alpes ;  
France

**Availability:** Free, Freely available

**Resource Name:** Seek and Blastn

**Resource ID:** SCR\_016625

**Alternate IDs:** SCR\_018509

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260829T182808+0000

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### Ratings and Alerts

No rating or validation information has been found for Seek and Blastn.

No alerts have been found for Seek and Blastn.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [NIF](#) .

Weissgerber T, et al. (2021) Automated screening of COVID-19 preprints: can we help authors to improve transparency and reproducibility? Nature medicine, 27(1), 6.