Resource Summary Report

Generated by dkNET on Apr 18, 2025

TBLASTN

RRID:SCR_011822
Type: Tool

Proper Citation

TBLASTN (RRID:SCR_011822)

Resource Information

URL: http://blast.ncbi.nlm.nih.gov/Blast.cgi?PROGRAM=tblastn&PAGE_TYPE=BlastSearch&LINK_LOC=blas

Proper Citation: TBLASTN (RRID:SCR_011822)

Description: Tool to search translated nucleotide databases using a protein query.

Abbreviations: TBLASTN

Synonyms: Translated BLAST: tblastn

Resource Type: analysis service resource, data analysis service, service resource, production service resource

Keywords: protein

Funding:

Resource Name: TBLASTN

Resource ID: SCR_011822

Alternate IDs: OMICS_00999

Record Creation Time: 20220129T080306+0000

Record Last Update: 20250418T055304+0000

Ratings and Alerts

No rating or validation information has been found for TBLASTN.

No alerts have been found for TBLASTN.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5246 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Magyar LB, et al. (2025) Pore-Forming Toxin-Like Proteins in the Anti-Parasitoid Immune Response of Drosophila. Journal of innate immunity, 17(1), 10.

Xiao Q, et al. (2025) Engineered IscB-?RNA system with expanded target range for base editing. Nature chemical biology, 21(1), 100.

Zhang W, et al. (2025) Chromosome-level genome assembly of tetraploid Chinese cherry (Prunus pseudocerasus). Scientific data, 12(1), 136.

Liu J, et al. (2025) Chromosome-level genome assembly of the seasonally polyphenic scorpionfly (Panorpa liui). Scientific data, 12(1), 22.

Dawson RA, et al. (2025) Carbon monoxide-oxidising Pseudomonadota on volcanic deposits. Environmental microbiome, 20(1), 12.

Li L, et al. (2025) A Chromosomal-level genome assembly and annotation of fat greenling (Hexagrammos otakii). Scientific data, 12(1), 78.

Akdeniz Z, et al. (2025) The expanded genome of Hexamita inflata, a free-living diplomonad. Scientific data, 12(1), 192.

Frankenberg SR, et al. (2025) Unearthing the secrets of Australia's most enigmatic and cryptic mammal, the marsupial mole. Science advances, 11(1), eado4140.

Tsukahara S, et al. (2025) Centrophilic retrotransposon integration via CENH3 chromatin in Arabidopsis. Nature, 637(8046), 744.

Yan Y, et al. (2025) Degenerated vision, altered lipid metabolism, and expanded chemoreceptor repertoires enable Lindaspio polybranchiata to thrive in deep-sea cold seeps. BMC biology, 23(1), 13.

Curantz C, et al. (2025) A positive feedback loop between germ cells and gonads induces and maintains sexual reproduction in a cnidarian. Science advances, 11(2), eadq8220.

Inoue S, et al. (2025) A new target of multiple lysine methylation in bacteria. Journal of bacteriology, 207(1), e0032524.

Holthaus KB, et al. (2025) Convergent Evolution Has Led to the Loss of Claw Proteins in Snakes and Worm Lizards. Genome biology and evolution, 17(1).

Lai Y, et al. (2025) Genome assembly of the grassland caterpillar Gynaephora qinghaiensis. Scientific data, 12(1), 158.

Liu C, et al. (2025) A chromosome-scale genome assembly of the pioneer plant Stylosanthes angustifolia: insights into genome evolution and drought adaptation. GigaScience, 14.

Cerqueira de Araujo A, et al. (2025) Genome sequences of four Ixodes species expands understanding of tick evolution. BMC biology, 23(1), 17.

Xu Z, et al. (2025) An orphan viral genome with unclear evolutionary status sheds light on a distinct lineage of flavi-like viruses infecting plants. Virus evolution, 11(1), veaf001.

Zou X, et al. (2025) Chromosome-level genome assembly of the pine wood nematode carrier Arhopalus unicolor. Scientific data, 12(1), 111.

G?siorowski L, et al. (2025) Regeneration in the absence of canonical neoblasts in an early branching flatworm. Nature communications, 16(1), 1232.

Chen Y, et al. (2025) An improved chromosome-level genome assembly and annotation of Hong Kong catfish (Clarias fuscus). Scientific data, 12(1), 193.