

Resource Summary Report

Generated by T1D on Aug 1, 2026

Webcutter

RRID:SCR_017638

Type: Tool

Proper Citation

Webcutter (RRID:SCR_017638)

Resource Information

URL: <http://heimanlab.com/cut2.html>

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Description: Software tool to find restriction endonucleases. Helps restriction map nucleotide sequences. Tool with customizable interface, platform independent accessibility, interfaces to NCBI's GenBank, DNA sequence database, and NEB's REBase, and restriction enzyme database. In addition to restriction site mapping, Webcutter 2 also performs degenerate digests, including option of finding restriction sites that can be introduced into sequence by silent mutagenesis.

Synonyms: Webcutter 2.0, Webcutter 2

Resource Type: data analysis service, data analysis software, data access protocol, service resource, software application, data processing software, production service resource, software resource, sequence analysis software, analysis service resource, web service

Keywords: Find, restriction, site, endonuclease, map, nucleotide, DNA, sequence, degenarte, digest, silent, mutagenesis

Availability: Free, Freely available

Resource Name: Webcutter

Resource ID: SCR_017638

Record Creation Time: 20220129T080336+0000

Record Last Update: 20260801T042822+0000

Ratings and Alerts

No rating or validation information has been found for Webcutter.

No alerts have been found for Webcutter.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#).

Narancic J, et al. (2024) First Characterization of *Acinetobacter baumannii*-Specific Filamentous Phages. *Viruses*, 16(6).

Foltran BB, et al. (2024) Leucine-rich repeat proteins of *Leptospira interrogans* that interact to host glycosaminoglycans and integrins. *Frontiers in microbiology*, 15, 1497712.

Pertics BZ, et al. (2023) Characterization of a Lytic Bacteriophage and Demonstration of Its Combined Lytic Effect with a K2 Depolymerase on the Hypervirulent *Klebsiella pneumoniae* Strain 52145. *Microorganisms*, 11(3).

Gühmann M, et al. (2021) Preselector.uni-jena.de: optimize your cloning-a resource for identifying restriction enzymes for preselection reactions. *Nucleic acids research*, 49(W1), W541.

Olaya-Abril A, et al. (2020) Role of the Dihydrodipicolinate Synthase DapA1 on Iron Homeostasis During Cyanide Assimilation by the Alkaliphilic Bacterium *Pseudomonas pseudoalcaligenes* CECT5344. *Frontiers in microbiology*, 11, 28.

Gómez GF, et al. (2015) Molecular taxonomy provides new insights into anopheles species of the neotropical arribalzagia series. *PloS one*, 10(3), e0119488.

Somasekharan S, et al. (2013) Functional expression of human NKCC1 from a synthetic cassette-based cDNA: introduction of extracellular epitope tags and removal of cysteines. *PloS one*, 8(12), e82060.

Yun J, et al. (2011) The first report of RPSA polymorphisms, also called 37/67 kDa LRP/LR gene, in sporadic Creutzfeldt-Jakob disease (CJD). *BMC medical genetics*, 12, 108.

Kwambana BA, et al. (2011) Differential effects of frozen storage on the molecular detection of bacterial taxa that inhabit the nasopharynx. *BMC clinical pathology*, 11, 2.

Moreira JL, et al. (2005) Identification to the species level of *Lactobacillus* isolated in probiotic prospecting studies of human, animal or food origin by 16S-23S rRNA restriction profiling. *BMC microbiology*, 5, 15.