

Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

SeWeR - SEquence analysis using WEb Resources

RRID:SCR_004167

Type: Tool

Proper Citation

SeWeR - SEquence analysis using WEb Resources (RRID:SCR_004167)

Resource Information

URL: <http://iubio.bio.indiana.edu/webapps/SeWeR/>

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Description: Sequence analysis using Web Resources (SeWeR) is an integrated, Dynamic HTML (DHTML) interface to commonly used bioinformatics services available on the World Wide Web. It is highly customizable, extendable, platform neutral, completely server-independent and can be hosted as a web page as well as being used as stand-alone software running within a web browser. It doesn't require any server to host itself. The goal of SeWeR is to turn your web-browser into a powerful sequence-analysis tool. It is written entirely in JavaScript1.2. SeWeR can be downloaded and mirrored freely. The whole package is just around 300K. You can even run it from a floppy. SeWeR is not compatible with Netscape 6. SeWeR now generates graphics. Savvy is a plasmid drawing software that generates plasmid map in the revolutionary Scalable Vector Graphics format from W3C.

Synonyms: SEquence analysis using WEb Resources, SeWeR

Resource Type: data or information resource, portal, service resource, topical portal, software resource

Defining Citation: [PMID:11395442](#)

Keywords: nucleic acid, protein, pcr, alignment, sequence, bio.tools

Resource Name: SeWeR - SEquence analysis using WEb Resources

Resource ID: SCR_004167

Alternate IDs: biotools:sewer, nlx_18981

Alternate URLs: <https://bio.tools/sewer>

Record Creation Time: 20220129T080223+0000

Record Last Update: 20260801T042541+0000

Ratings and Alerts

No rating or validation information has been found for SeWeR - SEquence analysis using WEb Resources.

No alerts have been found for SeWeR - SEquence analysis using WEb Resources.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.