

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

[les](#)

RRID:SCR_001291

Type: Tool

Proper Citation

les (RRID:SCR_001291)

Resource Information

URL: <http://julian-gehring.github.io/les/>

Proper Citation: les (RRID:SCR_001291)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23, 2022. Software package that estimates Loci of Enhanced Significance (LES) in tiling microarray data. These are regions of regulation such as found in differential transcription, ChIP-chip, or DNA modification analysis. The package provides a universal framework suitable for identifying differential effects in tiling microarray data sets, and is independent of the underlying statistics at the level of single probes.

Abbreviations: les

Synonyms: les package: Identifying Differential Effects in Tiling Microarray Data, Loci of Enhanced Significance

Resource Type: software resource

Keywords: loci of enhanced significance, tiling microarray, tiling, microarray, chip-chip, dna modification, probe, dna methylation, differential expression, microarray, transcription, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: les

Resource ID: SCR_001291

Alternate IDs: biotools:les, OMICS_02045

Alternate URLs: <https://bioconductor.org/packages/les/>, <https://bio.tools/les>

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260801T190137+0000

Ratings and Alerts

No rating or validation information has been found for les.

No alerts have been found for les.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.