

Resource Summary Report

Generated by [nidm-terms](#) on Aug 9, 2026

Context Likelihood of Relatedness

RRID:SCR_000216

Type: Tool

Proper Citation

Context Likelihood of Relatedness (RRID:SCR_000216)

Resource Information

URL: <https://omictools.com/context-likelihood-of-relatedness-tool>

Proper Citation: Context Likelihood of Relatedness (RRID:SCR_000216)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on July 31, 2025. A software that infers regulatory interactions between transcription factors and their targets using a compendium of gene expression profiles.

Abbreviations: CLR

Resource Type: software resource

Defining Citation: [PMID:17214507](#)

Keywords: transcription factors, gene expression profile, regulatory interactions, likelihood, relatedness

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Context Likelihood of Relatedness

Resource ID: SCR_000216

Alternate IDs: OMICS_01682

Old URLs: <http://gardnerlab.bu.edu/software&tools.html>,

Record Creation Time: 20220129T080200+0000

Record Last Update: 20260808T185716+0000

Ratings and Alerts

No rating or validation information has been found for Context Likelihood of Relatedness.

No alerts have been found for Context Likelihood of Relatedness.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Faith JJ, et al. (2007) Large-scale mapping and validation of Escherichia coli transcriptional regulation from a compendium of expression profiles. PLoS biology, 5(1), e8.