

# Resource Summary Report

Generated by [dkNET](#) on Aug 7, 2026

## INCLUSive

RRID:SCR\_013488

Type: Tool

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### Proper Citation

INCLUSive (RRID:SCR\_013488)

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### Resource Information

**URL:** <http://tomcatbackup.esat.kuleuven.be/inclusive/>

**Proper Citation:** INCLUSive (RRID:SCR\_013488)

**Description:** A suit of algorithms and tools for the analysis of gene expression data and the discovery of cis-regulatory sequence elements.

**Abbreviations:** INCLUSive

**Resource Type:** software resource

**Defining Citation:** [PMID:12824346](#)

**Keywords:** bio.tools

**Availability:** Acknowledgement requested

**Resource Name:** INCLUSive

**Resource ID:** SCR\_013488

**Alternate IDs:** OMICS\_00766, biotools:inclusive

**Alternate URLs:** <https://bio.tools/inclusive>

**Record Creation Time:** 20220129T080316+0000

**Record Last Update:** 20260801T190448+0000

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### Ratings and Alerts

No rating or validation information has been found for INCLUSive.

No alerts have been found for INCLUSive.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Taskiran II, et al. (2024) Cell-type-directed design of synthetic enhancers. *Nature*, 626(7997), 212.

Melendez-Torres GJ, et al. (2022) Effects of a Whole-School Health Intervention on Clustered Adolescent Health Risks: Latent Transition Analysis of Data from the INCLUSIVE Trial. *Prevention science : the official journal of the Society for Prevention Research*, 23(1), 1.

Melendez-Torres GJ, et al. (2022) Locating and testing the healthy context paradox: examples from the INCLUSIVE trial. *BMC medical research methodology*, 22(1), 57.

M??ller JV, et al. (2019) A university's response to people with disabilities in Worcester, Western Cape. *African journal of disability*, 8, 439.

Pilalis E, et al. (2011) Escherichia coli genome-wide promoter analysis: identification of additional AtoC binding target elements. *BMC genomics*, 12(1), 238.

Monsieurs P, et al. (2006) More robust detection of motifs in coexpressed genes by using phylogenetic information. *BMC bioinformatics*, 7, 160.

Vandepoele K, et al. (2006) Identification of novel regulatory modules in dicotyledonous plants using expression data and comparative genomics. *Genome biology*, 7(11), R103.