

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

NucleoFinder

RRID:SCR_000368

Type: Tool

Proper Citation

NucleoFinder (RRID:SCR_000368)

Resource Information

URL: <https://sites.google.com/site/beckerjeremie/home/nucleofinder>

Proper Citation: NucleoFinder (RRID:SCR_000368)

Description: A software for a statistical approach for the detection of nucleosome positions in a cell population. The software identifies important features of nucleosome organization such as the spacing downstream of active promoters and the enrichment and depletion of GC/AT dinucleotides of in vitro nucleosomes.

Abbreviations: NucleoFinder

Resource Type: software resource

Defining Citation: [PMID:23297036](#)

Keywords: nucleosome, position, promoter, analysis, downstream, nucleotide, bio.tools

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: NucleoFinder

Resource ID: SCR_000368

Alternate IDs: biotools:nucleofinder, OMICS_00510

Alternate URLs: <https://omictools.com/nucleofinder-tool>, <https://bio.tools/nucleofinder>

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260808T185720+0000

Ratings and Alerts

No rating or validation information has been found for NucleoFinder.

No alerts have been found for NucleoFinder.

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 [SPARC SAWG](#) .

Becker J, et al. (2013) NucleoFinder: a statistical approach for the detection of nucleosome positions. Bioinformatics (Oxford, England), 29(6), 711.