

# Resource Summary Report

Generated by [NIF](#) on Aug 19, 2026

## NEXT-peak

RRID:SCR\_010862

Type: Tool

### Proper Citation

NEXT-peak (RRID:SCR\_010862)

### Resource Information

**URL:** <http://ww2.odu.edu/~nxkim/nextpeak/>

**Proper Citation:** NEXT-peak (RRID:SCR\_010862)

**Description:** A software program to call peaks from ChIP-seq data for transcription factor binding sites.

**Abbreviations:** NEXT-peak

**Resource Type:** software resource

**Defining Citation:** [PMID:23706083](#)

**Resource Name:** NEXT-peak

**Resource ID:** SCR\_010862

**Alternate IDs:** OMICS\_00450

**Record Creation Time:** 20220129T080301+0000

**Record Last Update:** 20260815T182414+0000

### Ratings and Alerts

No rating or validation information has been found for NEXT-peak.

No alerts have been found for NEXT-peak.

### 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 [NIF](#) .

Tran NT, et al. (2014) A survey of motif finding Web tools for detecting binding site motifs in ChIP-Seq data. Biology direct, 9, 4.