## **Resource Summary Report**

Generated by <u>dkNET</u> on Apr 29, 2025

# **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

Funding:

Resource Name: NEXT-peak

Resource ID: SCR\_010862

Alternate IDs: OMICS\_00450

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

Record Last Update: 20250420T014512+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 <u>dkNET</u>.

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.