

Resource Summary Report

Generated by [NIF](#) on Jul 30, 2026

StoatyDive

RRID:SCR_018796

Type: Tool

Proper Citation

StoatyDive (RRID:SCR_018796)

Resource Information

URL: <https://github.com/BackofenLab/StoatyDive>

Proper Citation: StoatyDive (RRID:SCR_018796)

Description: Software tool to evaluate and classify predicted peak profiles to assess binding specificity of protein to its targets. Can be used for sequencing data such as CLIP-seq or ChIP-Seq, or any other type of peak profile data.

Resource Type: data processing software, software application, data analysis software, software resource

Keywords: Evaluate predicted peak profile, classify predicted peak profile, assess binding specificity, protein-target specificity, sequencing data, CLIP-seq data, CHIP-Seq data, peak profile data, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: StoatyDive

Resource ID: SCR_018796

Alternate IDs: SCR_018800, biotools:StoatyDive

Alternate URLs: <https://bio.tools/StoatyDive>

License: GNU General Public License v3.0

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260731T043035+0000

Ratings and Alerts

No rating or validation information has been found for StoatyDive.

No alerts have been found for StoatyDive.

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](#) .

Heyl F, et al. (2021) StoatyDive: Evaluation and classification of peak profiles for sequencing data. GigaScience, 10(6).