

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## StoatyDive

RRID:SCR\_018796

Type: Tool

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### Proper Citation

StoatyDive (RRID:SCR\_018796)

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### 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 analysis software, data processing software, software application, 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:** 20260905T132842+0000

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### Ratings and Alerts

No rating or validation information has been found for StoatyDive.

No alerts have been found for StoatyDive.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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