

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

[polyaPeak](#)

RRID:SCR_007687

Type: Tool

Proper Citation

polyaPeak (RRID:SCR_007687)

Resource Information

URL: <http://web1.sph.emory.edu/users/hwu30/polyaPeak.html>

Proper Citation: polyaPeak (RRID:SCR_007687)

Description: An R package for ranking ChIP-seq peaks with shape information.

Abbreviations: polyaPeak

Resource Type: software resource

Resource Name: polyaPeak

Resource ID: SCR_007687

Alternate IDs: OMICS_00456

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260725T190648+0000

Ratings and Alerts

No rating or validation information has been found for polyaPeak.

No alerts have been found for polyaPeak.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Scialdone A, et al. (2016) Resolving early mesoderm diversification through single-cell expression profiling. Nature, 535(7611), 289.

Wu H, et al. (2014) PolyPeak: detecting transcription factor binding sites from ChIP-seq using peak shape information. PloS one, 9(3), e89694.