

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

HRDetect

RRID:SCR_027726

Type: Tool

Proper Citation

HRDetect (RRID:SCR_027726)

Resource Information

URL: <https://github.com/eyzhao/hrdetect-pipeline>

Proper Citation: HRDetect (RRID:SCR_027726)

Description: Software tool for detection and analysis of homologous recombination events.

Resource Type: source code, software application, software resource

Keywords: homologous recombination, detection and analysis of homologous recombination events,

Availability: Free, Available for download, Freely available

Resource Name: HRDetect

Resource ID: SCR_027726

License URLs: <https://github.com/eyzhao/hrdetect-pipeline?tab=License-1-ov-file#readme>

Record Creation Time: 20251209T055452+0000

Record Last Update: 20260815T183309+0000

Ratings and Alerts

No rating or validation information has been found for HRDetect.

No alerts have been found for HRDetect.

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 [RRID](#) .

Culliford R, et al. (2026) Contrasting Features of Papillary and Chromophobe Renal Cell Carcinoma Revealed by Whole-Genome Sequencing. Molecular cancer research : MCR, 24(5), 327.

Yoon KA, et al. (2024) Proteogenomic analysis dissects early-onset breast cancer patients with prognostic relevance. Experimental & molecular medicine, 56(11), 2382.