

Resource Summary Report

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CLEAR

RRID:SCR_027171

Type: Tool

Proper Citation

CLEAR (RRID:SCR_027171)

Resource Information

URL: <https://github.com/rbundschuh/CLEAR>

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Description: Software workflow that identifies reliably quantifiable transcripts in limiting-cell RNA-seq (lcRNA-seq) data for differentially expressed genes (DEG) analysis. Coverage-based Limiting-cell Experiment Analysis for RNA-seq.

Synonyms: , CLEAR: Coverage-based Limiting-cell Experiment Analysis for RNA-seq, Coverage-based Limiting-cell Experiment Analysis for RNA-seq

Resource Type: software toolkit, source code, software resource

Defining Citation: [PMID:32039730](#)

Keywords: RNA-seq, cell experiment analysis for RNA-seq,

Funding: NCI P30 CA016058;
NCI R50 CA211524;
Ohio State University

Availability: GNU General Public License v3.0

Resource Name: CLEAR

Resource ID: SCR_027171

Record Creation Time: 20250709T061412+0000

Record Last Update: 20260815T183246+0000

Ratings and Alerts

No rating or validation information has been found for CLEAR.

No alerts have been found for CLEAR.

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 [dkNET](#) .

Rogers KA, et al. (2025) Addition of Ianalumab (VAY736) to Ibrutinib in Patients with Chronic Lymphocytic Leukemia on Ibrutinib Therapy: Results from a Phase Ib Study. Clinical cancer research : an official journal of the American Association for Cancer Research, 31(24), 5145.