

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on Apr 26, 2025

## ALEXA-Seq

RRID:SCR\_006700

Type: Tool

### Proper Citation

ALEXA-Seq (RRID:SCR\_006700)

### Resource Information

**URL:** [http://www.alexaplatform.org/alex\\_seq/index.htm](http://www.alexaplatform.org/alex_seq/index.htm)

**Proper Citation:** ALEXA-Seq (RRID:SCR\_006700)

**Description:** A method for using massively parallel paired-end transcriptome sequencing for "alternative expression analysis".

**Abbreviations:** ALEXA-Seq

**Resource Type:** software resource

**Funding:**

**Resource Name:** ALEXA-Seq

**Resource ID:** SCR\_006700

**Alternate IDs:** OMICS\_01328

**Record Creation Time:** 20220129T080237+0000

**Record Last Update:** 20250420T014343+0000

### Ratings and Alerts

No rating or validation information has been found for ALEXA-Seq.

No alerts have been found for ALEXA-Seq.

## Data and Source Information

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

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](#).

Shi C, et al. (2020) Alantolactone inhibits cell autophagy and promotes apoptosis via AP2M1 in acute lymphoblastic leukemia. Cancer cell international, 20, 442.

Turpin J, et al. (2016) The ErbB2<sup>Ex16</sup> splice variant is a major oncogenic driver in breast cancer that promotes a pro-metastatic tumor microenvironment. Oncogene, 35(47), 6053.

Alder O, et al. (2014) Hippo signaling influences HNF4A and FOXA2 enhancer switching during hepatocyte differentiation. Cell reports, 9(1), 261.

Pong WW, et al. (2013) F11R is a novel monocyte prognostic biomarker for malignant glioma. PloS one, 8(10), e77571.

Shyr D, et al. (2013) Next generation sequencing in cancer research and clinical application. Biological procedures online, 15(1), 4.

McElwee JL, et al. (2012) Identification of PADI2 as a potential breast cancer biomarker and therapeutic target. BMC cancer, 12, 500.