

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

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

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

Resource Name: ALEXA-Seq

Resource ID: SCR_006700

Alternate IDs: OMICS_01328

Record Creation Time: 20220129T080237+0000

Record Last Update: 20260912T195644+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](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

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??Ex16 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.