

Resource Summary Report

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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: 20260919T195116+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 [nidm-terms](#) .

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.