

Resource Summary Report

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ScienceExchange

RRID:SCR_010620

Type: Tool

Proper Citation

ScienceExchange (RRID:SCR_010620)

Resource Information

URL: <http://scienceexchange.com/>

Proper Citation: ScienceExchange (RRID:SCR_010620)

Description: Access service resource which connects labs needing and offering experimental services. Users can search for academic and government labs and experimental services, request and compare service quotes, and directly communicate with labs to arrange orders.

Abbreviations: Science Exchange

Synonyms: ScienceExchange.com, ScienceExchange

Resource Type: data or information resource, portal, community building portal

Keywords: marketplace, resource, outsource, portal

Resource Name: ScienceExchange

Resource ID: SCR_010620

Alternate IDs: nlx_55676, SciEx_96, SCR_013966

Alternate URLs: <https://www.scienceexchange.com/facilities/science-exchange>

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260812T044217+0000

Ratings and Alerts

No rating or validation information has been found for ScienceExchange.

No alerts have been found for ScienceExchange.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 58 mentions in open access literature.

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

Morales-Molina A, et al. (2024) Deletion of the RGD motif from the penton base in oncolytic adenoviruses enhances antitumor efficacy of combined CAR T??cell therapy. Molecular therapy. Oncology, 32(3), 200863.

Errington TM, et al. (2021) Experiments from unfinished Registered Reports in the Reproducibility Project: Cancer Biology. eLife, 10.

Rodgers P, et al. (2021) What have we learned? eLife, 10.

Kerwin J, et al. (2020) Replication Study: A coding-independent function of gene and pseudogene mRNAs regulates tumour biology. eLife, 9.

Wang H, et al. (2020) Replication Study: Coding-independent regulation of the tumor suppressor PTEN by competing endogenous mRNAs. eLife, 9.

Vervloessem T, et al. (2020) BDA-366, a putative Bcl-2 BH4 domain antagonist, induces apoptosis independently of Bcl-2 in a variety of cancer cell models. Cell death & disease, 11(9), 769.

Sheen MR, et al. (2019) Replication Study: Biomechanical remodeling of the microenvironment by stromal caveolin-1 favors tumor invasion and metastasis. eLife, 8.

Bittremieux M, et al. (2019) Constitutive IP3 signaling underlies the sensitivity of B-cell cancers to the Bcl-2/IP3 receptor disruptor BIRD-2. Cell death and differentiation, 26(3), 531.

Yan X, et al. (2019) Replication Study: The microRNA miR-34a inhibits prostate cancer stem cells and metastasis by directly repressing CD44. eLife, 8.

Essex A, et al. (2019) Replication Study: Wnt activity defines colon cancer stem cells and is regulated by the microenvironment. eLife, 8.

Vanden Heuvel JP, et al. (2018) Replication Study: Systematic identification of genomic markers of drug sensitivity in cancer cells. eLife, 7.

Eaton K, et al. (2018) Replication Study: Intestinal inflammation targets cancer-inducing activity of the microbiota. eLife, 7.

Lewis LM, et al. (2018) Replication Study: Transcriptional amplification in tumor cells with elevated c-Myc. eLife, 7.

Hofseth LJ, et al. (2018) Getting rigorous with scientific rigor. *Carcinogenesis*, 39(1), 21.

Kim J, et al. (2018) Replication study: Melanoma exosomes educate bone marrow progenitor cells toward a pro-metastatic phenotype through MET. *eLife*, 7.

Kandela I, et al. (2017) Replication Study: Discovery and preclinical validation of drug indications using compendia of public gene expression data. *eLife*, 6.

Vervloessem T, et al. (2017) The selective Bcl-2 inhibitor venetoclax, a BH3 mimetic, does not dysregulate intracellular Ca²⁺ signaling. *Biochimica et biophysica acta. Molecular cell research*, 1864(6), 968.

Vervloessem T, et al. (2017) Reciprocal sensitivity of diffuse large B-cell lymphoma cells to Bcl-2 inhibitors BIRD-2 versus venetoclax. *Oncotarget*, 8(67), 111656.

Horrigan SK, et al. (2017) Replication Study: Melanoma genome sequencing reveals frequent PREX2 mutations. *eLife*, 6.

Showalter MR, et al. (2017) Replication Study: The common feature of leukemia-associated IDH1 and IDH2 mutations is a neomorphic enzyme activity converting alpha-ketoglutarate to 2-hydroxyglutarate. *eLife*, 6.