Resource Summary Report

Generated by dkNET on Apr 25, 2025

EPIGEN

RRID:SCR_000093

Type: Tool

Proper Citation

EPIGEN (RRID:SCR_000093)

Resource Information

URL: http://www.epilepsygenetics.eu/

Proper Citation: EPIGEN (RRID:SCR_000093)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on August 16,2023. Group of clinical care and epilepsy research centers who are committed to improving the lives of people with epilepsy through an understanding of the genetics of epilepsy. The consoritum was in an effort to speed discovery to epilepsy genetics by pooling the resources of several research centres.

Abbreviations: EPIGEN

Synonyms: EPIGEN: An international consortium dedicated to tackling epilepsy through

genetics, EPIGEN Consortium

Resource Type: data or information resource, organization portal, portal, consortium

Keywords: epilepsy, genetics, gene, mri, genetic variation, clinical

Related Condition: Epilepsy

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: EPIGEN

Resource ID: SCR_000093

Alternate IDs: nlx_143740

Record Creation Time: 20220129T080159+0000

Record Last Update: 20250425T055124+0000

Ratings and Alerts

No rating or validation information has been found for EPIGEN.

No alerts have been found for EPIGEN.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 24 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Ahrens JM, et al. (2017) Differential activity of 2-methylene-19-nor vitamin D analogs on growth factor gene expression in rhino mouse skin and comparison to all-trans retinoic acid. PloS one, 12(11), e0188887.

França GVA, et al. (2017) Genomic ancestry and education level independently influence abdominal fat distributions in a Brazilian admixed population. PloS one, 12(6), e0179085.

Zhang H, et al. (2017) Targeting naturally occurring epitope variants of hepatitis C virus with high-affinity T-cell receptors. The Journal of general virology, 98(3), 374.

Lin A, et al. (2016) The LINK-A IncRNA activates normoxic HIF1? signalling in triple-negative breast cancer. Nature cell biology, 18(2), 213.

Sas-Chen A, et al. (2016) LIMT is a novel metastasis inhibiting lncRNA suppressed by EGF and downregulated in aggressive breast cancer. EMBO molecular medicine, 8(9), 1052.

Haukaas TH, et al. (2016) Metabolic clusters of breast cancer in relation to gene- and protein expression subtypes. Cancer & metabolism, 4, 12.

Noberini R, et al. (2016) Pathology Tissue-quantitative Mass Spectrometry Analysis to Profile Histone Post-translational Modification Patterns in Patient Samples. Molecular & cellular proteomics: MCP, 15(3), 866.

Fagnocchi L, et al. (2016) A Myc-driven self-reinforcing regulatory network maintains mouse

embryonic stem cell identity. Nature communications, 7, 11903.

Gangisetty O, et al. (2015) Fetal Alcohol Exposure Reduces Dopamine Receptor D2 and Increases Pituitary Weight and Prolactin Production via Epigenetic Mechanisms. PloS one, 10(10), e0140699.

Lima-Costa MF, et al. (2015) Genomic ancestry and ethnoracial self-classification based on 5,871 community-dwelling Brazilians (The Epigen Initiative). Scientific reports, 5, 9812.

Kappler CS, et al. (2015) Oncogenic signaling in amphiregulin and EGFR-expressing PTEN-null human breast cancer. Molecular oncology, 9(2), 527.

Lei J, et al. (2015) Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with prognosis of estrogen receptor-negative breast cancer after chemotherapy. Breast cancer research: BCR, 17(1), 18.

Meena JK, et al. (2015) Telomerase abrogates aneuploidy-induced telomere replication stress, senescence and cell depletion. The EMBO journal, 34(10), 1371.

Liu Y, et al. (2014) Functional variants in DPYSL2 sequence increase risk of schizophrenia and suggest a link to mTOR signaling. G3 (Bethesda, Md.), 5(1), 61.

Beznoussenko GV, et al. (2014) Transport of soluble proteins through the Golgi occurs by diffusion via continuities across cisternae. eLife, 3.

Gangisetty O, et al. (2014) Fetal alcohol exposure alters proopiomelanocortin gene expression and hypothalamic-pituitary-adrenal axis function via increasing MeCP2 expression in the hypothalamus. PloS one, 9(11), e113228.

He W, et al. (2013) Molecular basis of live-attenuated influenza virus. PloS one, 8(3), e60413.

O Neill V, et al. (2013) Vitamin D Receptor Gene Expression and Function in a South African Population: Ethnicity, Vitamin D and Fokl. PloS one, 8(6), e67663.

Boruah BM, et al. (2013) Single domain antibody multimers confer protection against rabies infection. PloS one, 8(8), e71383.

Mozzetta C, et al. (2013) Fibroadipogenic progenitors mediate the ability of HDAC inhibitors to promote regeneration in dystrophic muscles of young, but not old Mdx mice. EMBO molecular medicine, 5(4), 626.