Resource Summary Report

Generated by NIF on May 8, 2025

Columbia University; New York; USA

RRID:SCR_011164
Type: Tool

Proper Citation

Columbia University; New York; USA (RRID:SCR_011164)

Resource Information

URL: http://www.columbia.edu/

Proper Citation: Columbia University; New York; USA (RRID:SCR_011164)

Description: Private Ivy League research university in New York City. Established in 1754 on the grounds of Trinity Church in Manhattan, Columbia is the oldest institution of higher education in New York and the fifth-oldest institution of higher learning in the United States.

Abbreviations: Columbia

Synonyms: Columbia University

Resource Type: university

Funding:

Resource Name: Columbia University; New York; USA

Resource ID: SCR_011164

Alternate IDs: GRID:grid.21729.3f; ISNI: 0000 0004 1936 8729; Crossref Funder ID: 100006474; Wikidata: Q49088;

Alternate URLs: https://ror.org/00hj8s172

Record Creation Time: 20220129T080302+0000

Record Last Update: 20250420T014524+0000

Ratings and Alerts

No rating or validation information has been found for Columbia University; New York; USA.

No alerts have been found for Columbia University; New York; USA.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 14 mentions in open access literature.

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

Nakamura T, et al. (2024) Topologically associating domains define the impact of de novo promoter variants on autism spectrum disorder risk. Cell genomics, 4(2), 100488.

Cardenas R, et al. (2022) The Early Warning and Response System (EWARS-TDR) for dengue outbreaks: can it also be applied to chikungunya and Zika outbreak warning? BMC infectious diseases, 22(1), 235.

Calero-Cuenca FJ, et al. (2021) Ctdnep1 and Eps8L2 regulate dorsal actin cables for nuclear positioning during cell migration. Current biology : CB, 31(7), 1521.

Wang Y, et al. (2021) REVA as A Well-curated Database for Human Expression-modulating Variants. Genomics, proteomics & bioinformatics, 19(4), 590.

Villani V, et al. (2019) Transcutaneous vagus nerve stimulation improves interoceptive accuracy. Neuropsychologia, 134, 107201.

Wang Y, et al. (2019) Detection of epigenetic field defects using a weighted epigenetic distance-based method. Nucleic acids research, 47(1), e6.

He Z, et al. (2018) A semi-supervised approach for predicting cell-type specific functional consequences of non-coding variation using MPRAs. Nature communications, 9(1), 5199.

Riquelme SA, et al. (2017) Cystic Fibrosis Transmembrane Conductance Regulator Attaches Tumor Suppressor PTEN to the Membrane and Promotes Anti Pseudomonas aeruginosa Immunity. Immunity, 47(6), 1169.

Schubach M, et al. (2017) Imbalance-Aware Machine Learning for Predicting Rare and Common Disease-Associated Non-Coding Variants. Scientific reports, 7(1), 2959.

Greene D, et al. (2017) A Fast Association Test for Identifying Pathogenic Variants Involved in Rare Diseases. American journal of human genetics, 101(1), 104.

Fleming SM, et al. (2017) HMeta-d: hierarchical Bayesian estimation of metacognitive

efficiency from confidence ratings. Neuroscience of consciousness, 2017(1), nix007.

Koizumi A, et al. (2016) Is fear perception special? Evidence at the level of decision-making and subjective confidence. Social cognitive and affective neuroscience, 11(11), 1772.

Fleming SM, et al. (2014) Domain-specific impairment in metacognitive accuracy following anterior prefrontal lesions. Brain : a journal of neurology, 137(Pt 10), 2811.

Backenroth D, et al. (2014) CANOES: detecting rare copy number variants from whole exome sequencing data. Nucleic acids research, 42(12), e97.