Resource Summary Report

Generated by NIF on May 24, 2025

Centrillion Biosciences Inc.

RRID:SCR_012358
Type: Tool

Proper Citation

Centrillion Biosciences Inc. (RRID:SCR_012358)

Resource Information

URL: http://www.scienceexchange.com/facilities/centrillion-biosciences-inc

Proper Citation: Centrillion Biosciences Inc. (RRID:SCR_012358)

Description: Centrillion offers a portfolio of genomic services to academic, clinical and industrial researchers. Core provides experimental design consultation, data production services, and bioinformatics analyses for a wide variety of genomic applications. Core offers access to next-gen sequencing, genotyping and bioinformatics analysis.

Abbreviations: Centrillion

Synonyms: Centrillion Biosciences

Resource Type: core facility, service resource, access service resource

Keywords: genomic, consultation, consulting, data production, bioinformatics analyses, sequencing

Funding:

Availability: Available to external user

Resource Name: Centrillion Biosciences Inc.

Resource ID: SCR_012358

Alternate IDs: SciEx_12068

Alternate URLs: http://www.centrilliontech.com/

Record Creation Time: 20220129T080309+0000

Record Last Update: 20250524T060435+0000

Ratings and Alerts

No rating or validation information has been found for Centrillion Biosciences Inc..

No alerts have been found for Centrillion Biosciences Inc..

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 NIF.

Hill J, et al. (2019) Unprecedented reorganization of holocentric chromosomes provides insights into the enigma of lepidopteran chromosome evolution. Science advances, 5(6), eaau3648.

Nallu S, et al. (2018) The molecular genetic basis of herbivory between butterflies and their host plants. Nature ecology & evolution, 2(9), 1418.

Bakr S, et al. (2018) A radiogenomic dataset of non-small cell lung cancer. Scientific data, 5, 180202.

Pasqualucci L, et al. (2014) Genetics of follicular lymphoma transformation. Cell reports, 6(1), 130.

Haraksingh RR, et al. (2014) Exome sequencing and genome-wide copy number variant mapping reveal novel associations with sensorineural hereditary hearing loss. BMC genomics, 15(1), 1155.

Ehmer U, et al. (2014) Organ size control is dominant over Rb family inactivation to restrict proliferation in vivo. Cell reports, 8(2), 371.