

Resource Summary Report

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Codex

RRID:SCR_027783

Type: Tool

Proper Citation

Codex (RRID:SCR_027783)

Resource Information

URL: <https://codex.flywire.ai>

Proper Citation: Codex (RRID:SCR_027783)

Description: Web based platform for exploring full brain connectomic datasets. Used for exploring and analyzing large connectomics datasets.

Synonyms: Codex - Connectome Data Explorer

Resource Type: data or information resource, web application, software resource

Keywords: exploring and analyzing large connectomics datasets, full brain, connectomic datasets,

Availability: Restricted

Resource Name: Codex

Resource ID: SCR_027783

Record Creation Time: 20251213T054634+0000

Record Last Update: 20260729T044717+0000

Ratings and Alerts

No rating or validation information has been found for Codex.

No alerts have been found for Codex.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 18 mentions in open access literature.

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

Trigila AP, et al. (2025) Comparative genomics sheds light on mammalian and avian gene regulation and phenotypic evolution. *Nature communications*, 16(1), 9111.

Wang T, et al. (2025) scProAtlas: an atlas of multiplexed single-cell spatial proteomics imaging in human tissues. *Nucleic acids research*, 53(D1), D582.

Trehan R, et al. (2025) SPP1 + macrophages cause exhaustion of tumor-specific T cells in liver metastases. *Nature communications*, 16(1), 4242.

Barlow GL, et al. (2025) The extra-islet pancreas supports autoimmunity in human type 1 diabetes. *eLife*, 13.

Giakoumas DS, et al. (2025) Connectomic mapping of pharyngeal and gut sensory circuits in adult *Drosophila*. *bioRxiv : the preprint server for biology*.

Roider T, et al. (2024) Multimodal and spatially resolved profiling identifies distinct patterns of T cell infiltration in nodal B cell lymphoma entities. *Nature cell biology*, 26(3), 478.

Zahraeifard S, et al. (2024) Loss of tumor suppressors promotes inflammatory tumor microenvironment and enhances LAG3+T cell mediated immune suppression. *Nature communications*, 15(1), 5873.

Meschi E, et al. (2024) Compensatory enhancement of input maintains aversive dopaminergic reinforcement in hungry *Drosophila*. *Neuron*, 112(14), 2315.

Wu R, et al. (2024) Gasdermin C promotes Stemness and Immune Evasion in Pancreatic Cancer via Pyroptosis-Independent Mechanism. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 11(42), e2308990.

Hickey JW, et al. (2023) Organization of the human intestine at single-cell resolution. *Nature*, 619(7970), 572.

Sexton T, et al. (2023) Competition between transcription and loop extrusion modulates promoter and enhancer dynamics. *Research square*.

Antaki D, et al. (2022) A phenotypic spectrum of autism is attributable to the combined effects of rare variants, polygenic risk and sex. *Nature genetics*, 54(9), 1284.

Mantsoki A, et al. (2021) Identification and Characterisation of Putative Enhancer Elements in Mouse Embryonic Stem Cells. *Bioinformatics and biology insights*, 15, 1177932220974623.

Zhang L, et al. (2021) Sex-specific DNA methylation differences in Alzheimer's disease pathology. *Acta neuropathologica communications*, 9(1), 77.

Rendeiro AF, et al. (2020) Chromatin mapping and single-cell immune profiling define the temporal dynamics of ibrutinib response in CLL. *Nature communications*, 11(1), 577.

Moreno-Cabrera JM, et al. (2020) Evaluation of CNV detection tools for NGS panel data in genetic diagnostics. *European journal of human genetics : EJHG*, 28(12), 1645.

Schürch CM, et al. (2020) Coordinated Cellular Neighborhoods Orchestrate Antitumoral Immunity at the Colorectal Cancer Invasive Front. *Cell*, 182(5), 1341.

Schick S, et al. (2019) Systematic characterization of BAF mutations provides insights into intracomplex synthetic lethality in human cancers. *Nature genetics*, 51(9), 1399.