

Resource Summary Report

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10x Genomics Xenium Explorer

RRID:SCR_025847

Type: Tool

Proper Citation

10x Genomics Xenium Explorer (RRID:SCR_025847)

Resource Information

URL: <https://www.10xgenomics.com/support/software/xenium-explorer/latest>

Proper Citation: 10x Genomics Xenium Explorer (RRID:SCR_025847)

Description: Desktop application for Windows and MacOS that allows to interactively visualize data generated by the Xenium Analyzer instrument.

Synonyms: Xenium Explorer

Resource Type: software resource

Keywords: 10x Genomics, interactive visualization,

Availability: Restricted

Resource Name: 10x Genomics Xenium Explorer

Resource ID: SCR_025847

License URLs: <https://www.10xgenomics.com/support/software/xenium-explorer/downloads/eula?closeUrl=%2Fsupport%2Fsoftware%2Fxenium-explorer&lastTouchOfferName=Xenium%20Explorer&lastTouchOfferType=Software%20Download&explorer%2Fdownloads>

Record Creation Time: 20241004T053248+0000

Record Last Update: 20260801T191327+0000

Ratings and Alerts

No rating or validation information has been found for 10x Genomics Xenium Explorer.

No alerts have been found for 10x Genomics Xenium Explorer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 12 mentions in open access literature.

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

Johnson KF, et al. (2025) Mapping mesenchymal diversity in the developing human intestine and organoids. bioRxiv : the preprint server for biology.

Vannan A, et al. (2025) Spatial transcriptomics identifies molecular niche dysregulation associated with distal lung remodeling in pulmonary fibrosis. Nature genetics, 57(3), 647.

Hwang A, et al. (2025) Single-cell transcriptomic and chromatin dynamics of the human brain??in PTSD. Nature, 643(8072), 744.

Lin Y, et al. (2025) Defining Keypoints to Align H&E Images and Xenium DAPI-Stained Images Automatically. Cells, 14(13).

Mangoli A, et al. (2025) Disruption of ataxia telangiectasia-mutated kinase enhances radiation therapy efficacy in spatially directed diffuse midline glioma models. The Journal of clinical investigation, 135(12).

Heming M, et al. (2025) Multi-omic identification of perineurial hyperplasia and lipid-associated nerve macrophages in human polyneuropathies. Nature communications, 16(1), 7872.

Huang CH, et al. (2025) Integrative whole slide image and spatial transcriptomics analysis with QuST and QuPath. NPJ precision oncology, 9(1), 70.

Markov NS, et al. (2025) Profibrotic monocyte-derived alveolar macrophages as a biomarker and therapeutic target in systemic sclerosis-associated interstitial lung disease. bioRxiv : the preprint server for biology.

Ebert SN, et al. (2025) Single-Cell Approaches Define the Murine Leptomeninges: Cortical Brain Interface as a Distinct Cellular Neighborhood Composed of Neural and Non-neural Cell Types. eNeuro, 12(8).

Orba Y, et al. (2025) Spatial gene expression analysis reveals pathological niches in Japanese encephalitis virus neuroinvasion. Proceedings of the National Academy of Sciences of the United States of America, 122(43), e2515006122.

Willis A, et al. (2025) Single cell approaches define neural stem cell niches and identify microglial ligands that can enhance precursor-mediated oligodendrogenesis. Cell reports, 44(1), 115194.

Damle SR, et al. (2025) Intratumoral three-cell-type clusters are a conserved feature of endogenous antitumor immunity. Cancer immunology research.