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Sequencing open chromatin of single cell nuclei: snATAC-seq V.2

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Protocol status: Working
We use this protocol and it's working

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Keywords: open chromatin of single cell nuclei, chromatin of single cell nuclei, sequencing open chromatin, single cell nuclei, center for epigenomic, chromatin, epigenomic, coordination with the brain initiative cell census network, brain initiative cell census network, nuclei, seq method to the analysis, seq, seq protocol, mouse bain atlas, based atac, seq method

Abstract

Protocol for applying the TN5-based ATAC-seq method to the analysis of the chromatin of single cell nuclei isolated from brain tissue, as performed within the Center for Epigenomics of the Mouse Bain Atlas (CEMBA) and in coordination with the BRAIN Initiative Cell Census Network (BICCN).

Attachments

Ren_Lab_snATACseq_CE...
116KB

Troubleshooting