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Rapid Nuclei Isolation from Human Brain

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Human Cell Atlas Method Development Community

Neurodegeneration Method Development Community

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ABSTRACT

This protocol outlines our preparation of single-cell suspension from surgically acquired fresh human adult brain tissue.

This method is adapted from Habib et al¹.

1 Habib, N. *et al.* Massively parallel single-nucleus RNA-seq with DroNc-seq. *Nat Methods* **14**, 955-958, doi:10.1038/nmeth.4407 (2017).

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

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- 1 Prepare Nuclei Suspension Buffer (NSB):
For drop-seq: 1x PBS, 0.01% BSA (ultra pure), 0.2U/ul RNase inhibitor
For 10X Genomics: 1X PBS, 1%BSA (ultra pure), 0.2U/ul RNase inhibitor
- 2 Dounce homogenize tissue samples (smaller than 0.5 cm) or cell pellets in 2 ml of ice-cold Nuclei EZ lysis buffer (Sigma, #EZ PREP NUC-101). Dounce tissue 20-25 times with pestle A, followed by 20-25 times with pestle B while on ice.
- 3 Move sample to a 15 ml conical tube, add 2 ml of ice-cold Nuclei EZ lysis buffer and incubate on ice for 5 min.
- 4 Collect nuclei by centrifugation at 500 x g for 5 minutes at 4°C. Discard supernatant and carefully resuspend nuclei in 4 ml of ice-cold

Nuclei EZ lysis buffer. Incubate on ice for 5 minutes.

- 5 Repeat the centrifugation at 500 x g for 5 minutes at 4°C. Resuspend isolated nuclei in 1 ml of NSB, and filter through a Flowmi Cell Strainer (Bel-Art, H13680-0040).
- 6 Stain 10µl of nuclei suspension with DAPI (1:5000) or 0.4% Trypan Blue to count.
- 7 Adjust the volume with NSB to 1000 nuclei/µl (for 10X Genomics) and 300 nuclei/µl (for drop-seq).