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# Single Nucleus RNAseq Sample Prep from Nodose Ganglia

Sebastian Preissl<sup>1</sup>, Jamie Verheyden<sup>1</sup>, Xin Sun<sup>2</sup><sup>1</sup>University of California, San Diego; <sup>2</sup>University of California-San Diego

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[dx.doi.org/10.17504/protocols.io.v72e9qe](https://dx.doi.org/10.17504/protocols.io.v72e9qe)

SPARC

Tech. support email: [info@neuinfo.org](mailto:info@neuinfo.org)

Jamie Verheyden

How to isolate single nuclei from nodose vagal ganglia neurons in preparation for 10XGenomics sequencing.

DOI

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single nucleus RNAseq

protocol ,

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## MATERIALS

liquid nitrogen Contributed by users

DTT Sigma

Aldrich Catalog #D0632

BSA Sigma

Aldrich Catalog #A7906

RNasin Promega

PBS Contributed by users

Trypan Blue 100 mL Stemcell

Technologies Catalog #7050

cOmplete™, Mini Protease Inhibitor

Cocktail Roche Catalog #11836153001

EDTA Invitrogen - Thermo

Fisher Catalog #AM9261

DRAQ7 BD

Biosciences Catalog #564904

Lysis buffer (LB): -> 550ul per sample

Prepare fresh and put on ice

Reagent	Stock concentration	Final concentration	1 sample	2 samples
TritonX-100	10%	0.2%	11 ul	22 ul
Roche protease inhibitor	25x	1x	22 ul	44 ul
DTT	200mM	1mM	2.75 ul	5.5 ul
RNasin (Promega, N211B)	40 U/ul	0.2U/ul	2.75 ul	5.5 ul
2 % BSA in PBS			512.5 ul	1025 ul

Sort buffer (SB): -> 1ml per sample

Prepare fresh and put on ice

Reagent	Stock concentration	Final concentration	1 sample	5 samples
EDTA (Life technologies)	500mM	1mM	2 ul	10 ul
RNasin (Promega, N211B)	40 U/ul	0.2U/ul	5 ul	25 ul
2 % BSA in PBS			993 ul	4.97 ml

Collection buffer (CB): -> 200ul per sample

**Prepare fresh and put on ice**

Reagent	Stock concentration	Final concentration	1 sample	5 samples
RNAasin (Promega, N211B)	40 U/ul	1U/ul	5 ul	25 ul
5 % BSA in PBS			195 ul	975 ul

Reaction buffer (RB): -> 20ul per sample

**Prepare fresh and put on ice**

Reagent	Stock concentration	Final concentration	1 sample	5 samples	10 samples
5% BSA (Sigma, Molec boil grade, B6917)	5%	1%	4 ul	20 ul	40 ul
RNAasin (Promega, N211B)	40 U/ul	0.2U/ul	0.25 ul	1.25 ul	2.5 ul
PBS			15.75 ul	78.75 ul	157.5 ul

- 1 Nodose ganglia are dissected and flash frozen in liquid nitrogen. Store at -80C until ready to isolate nuclei. Grind up the sample with liquid nitrogen using the 1.5mL centrifuge tube and fitting pestle.
- 2 Add 500 ul Lysis Buffer to pulverized tissue, pipette 10x up and down and place on ice. Incubate 5 min with the overhead shaker in the cold room.
- 3 Spin down 5 min 500g at 4C using soft settings. Remove supernatant.
- 4 Resuspend tissue in 400 ul Sorting Buffer by pipetting 10x up and down.

- 5 Filter with a green 30 um CellTRic (Sysmex) into FACS tube. Wash filter with another 200 ul SB. Gently clap rack against bench to fully transfer the nuclei suspension. Add 6 ul DRAQ7 (Cell signaling) to stain nuclei and place sample on ice.
- 6 For collection of sorted nuclei add 50 ul Collection Buffer  
to a 1.5 ml LoBind tube (collection tube).
- 7 Use 100 um Chip for sorting of nuclei  
Set temperature for sample and collection to 5C  
Identify single nuclei based FSC/SSC signals and DRAQ7 fluorescence in FL-6 (PE-Cy7) channel
- 8 Sort 60,000 (200 ul) nuclei (Gate: "Sorting") with the setting "Purity" into the prepared collection tube. After sorting is finished (10-20 min), mix sample and keep on ice.
- 9 Spin down 15 min 1000xg at 4C (Using swinging bucket rotor). After centrifugation, there should be a tiny blue pellet be visible. Remove supernatant and resuspend in 18 ul RB by pipetting 10x (Recovery of nuclei should be around 50-60%). Leave sample on ice.
- 10  
Check for single cell suspension and count nuclei using hemocytometer. Mix 5 ul sample + 5 ul Trypan blue and count 4 big squares.  
Calculation: (counted number)/4\*20=nuclei/ul. Adjust concentration to 1000 nuclei/ul. Follow 10xGenomics library prep protocol.