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Nuclei isolation and permeabilisation of fresh frozen human brain samples for 10X Genomics Multiome

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We use this protocol and it's working

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ASAP

Abstract

This protocol details the procedure of nuclei isolation and debris removal from fresh frozen brain tissue in preparation for the 10X Genomics multiome ATAC/GEX assay.

Materials

HOW MANY SAMPLES TO PROCESS? 2

	A	B	C	D	E	F
	2X salt-Tris solution	Sto ck con c		Fin al co nc		V (μ l)
	NaCl	500 0	m M	29 2	m M	122 6,4
	Tris pH7.5	100 0	m M	20	m M	420 ,0
	CaCl	100 0	m M	2	m M	42, 0
	MgCl ₂	100 0	m M	42	m M	882 ,0
	Water					184 29, 6
	Total Volume					210 00, 0

	TST+NP Homogenization buffer	Sto ck con c		Fin al co nc		V (μ l)
	2X salt-Tris solution	2	X	1	X	110 0,0
	Tween-20	10	%	0,0 3	%	6,6
	NP-40	10	%	0,0 3	%	6,6
	BSA	7,5	%	1	%	293 ,3
	KCl	300 0	m M	25	m M	18,3
	Sucrose	200 0	m M	25 0	m M	275 ,0

	DTT	100 0	m M	1	m M	2,2
	cOmplete protease inhibitor	50	X	1	X	44, 0
	Rnase In plus (promega)	40	U/ ul	1	U/ ul	55, 0
	Water					398 ,9
	Total Volume					220 0,0

	Wash buffer 1	Sto ck con c		Fin al co nc		V (μ l)
	2X salt-Tris solution	2	X	1	X	561, 0
	BSA	7,5	%	1	%	149, 6
	KCl	300 0	m M	25	m M	9,4
	Sucrose	200 0	m M	25 0	m M	140, 3
	DTT	100 0	m M	1	m M	1,1
	cOmplete protease inhibitor	50	X	1	X	22, 4
	Rnase In plus (promega)	40	U/ ul	1	U/ ul	28,1
	Water					210, 2
	Total Volume					112 2,0

	GM (Gradient Medium)	Sto ck con c		Fin al co nc		V(μ l)



	CaCl	100 0	m M	1	m M	1,3
	Optiprep	60	%	50	%	108 3,3
	MgCl ₂	100 0	m M	5	m M	6,5
	Tris pH 7.5	100 0	m M	10	m M	13,0
	Sucrose	200 0	m M	75	m M	48, 8
	DTT	100 0	m M	1	m M	1,3
	cOmplete protease inhibitor	50	X	0,5	X	13,0
	Rnase In plus (promega)	40	U/ ul	1	U/ ul	32, 5
	Water					100, 3
	Total Volume					130 0

	ODM (Optiprep Diluent Medium)	Sto ck con c		Fin al co nc		V(μl)
	KCl	300 0	m M	150	m M	62, 0
	MgCl ₂	100 0	m M	30	m M	37,2
	Tris pH 8	100 0	m M	60	m M	74, 4
	Sucrose	200 0	m M	25 0	m M	155, 0
	RNAse in plus (promega)	40	U/ u L	0,5	U/ u L	15,5
	Water					895 ,9



	Total Volume				124 0,0

	29% Cushion	Stock conc	Final conc	V(μl)		
	Optiprep	60	%	29	%	966,7
	ODM					1033,3
	Total Volume					2000

	1x Lysis	Sto ck con c		Fin al co nc		V (μl)
	Tris-HCl (pH 7.4)	100 0	m M	10 0	m M	50, 0
	NaCl	500 0	m M	10 0	m M	10,0
	MgCl ₂	100 0	m M	30	m M	15,0
	Tween-20	10	%	0,1	%	5,0
	NP-40	10	%	0,1	%	5,0
	Digitonin	5	%	0,0 2	%	2,0
	Water					413, 0
	Total Volume					500 ,0

	0.1x Lysis	Sto ck con c		Fin al co nc		V (μl)
	1x lysis	1	x	0,1	x	100, 0



	BSA	7,5	%	1	%	133,3
	DTT	1000	mM	1	mM	1,0
	Rnase Inhibitor (Promega)	40	U/ul	1	U/ul	25,0
	Water					740,7
	Total Volume					1000,0

	Wash Buffer 2	Stock conc		Final conc		V (μl)
	Tris-HCl (pH 7.4)	1000	mM	10	mM	22,0
	NaCl	5000	mM	10	mM	4,4
	MgCl ₂	1000	mM	3	mM	6,6
	BSA	7,5	%	1	%	293,3
	Tween-20	10	%	0,01	%	2,2
	DTT	1000	mM	1	mM	2,2
	Rnase Inhibitor (SIGMA)	40	U/ul	1	U/ul	55,0
	Water					1814,3
	Total Volume					2200,0

	1X Diluted Nuclei buffer	Stock conc		Final conc	V(μl)	



	Nuclei buffer	20	X	1	X	7,5
	DTT	100	mM	1	mM	1,5
	Rnase Inhibitor (SIGMA)	40	U/ul	1	U/ul	3,75
	Water					137,25
	Total Volume					150

Preparation

- 1 Prepare the 2X salt-Tris solution and aliquot to 1 ml tube and freeze at -20°C
- 2 Prepare the BSA in PBS and filter through 0.2um filter.
- 3 Prepare the Tween-20 Homogenization lysis buffer fresh for each preparations
- 4 Filter sucrose 2M solution through 0.2um/0.45um filter
- 5 Prepare cOmplete protease inhibitor (50X) by dissolving a tablet in 1 mL of water.
- 6 Filter protease inhibitor solution through 0.2um filter
- 7 Place the homogenizer at -80°C at the start and leave 10 min before homogenization on ice.

Homogenization (Perform all steps on ice)

- 8 Cut brain piece on dry ice (if a large section), then transfer immediately in homogenizer containing 250 uL of TST+NP Homogenization Lysis Buffer. Immediately add 500uL TST+NP Homogenization Lysis Buffer
- 9 Allow the tissue to thaw for 2 mins in the homogenization buffer.
- 10 Homogenize tissue with pestle A (10X), wait 1min, and with pestle B (10X).
- 11 Put the homogenate through 70µm cell strainer and place the falcon on ice. Incubate on ice for 5 mins. Rinse dounce and filter with 250uL TST+NP HB.
- 12 During the incubation (in about 4 mins), use LUNA-fl counter to assess the viability of the nuclei.



- 13 Transfer the contents from the 50 mL falcon to a 2 mL protein lo-bind tube.
- 14 Centrifuge @ 500xg for 5 mins. Discard the supernatant.
- 15 Resuspend the pellet in 200 uL of Wash Buffer 1, and transfer to a 2mL DNA lobind tube.
- 16 Add additional 320 uL Wash Buffer 1 to a final volume of 520 uL.
- 17 Add 520uL of Gradient medium to sample ($V_f = 1040$ uL).

Isolation by centrifugation

- 18 Layer 770 uL of 29% Cushion in the ultracentrifuge tube.
- 19 Layer the sample on top of cushion using a P1000, without disturbing the cushion.
- 20 Optional: Check tube weight, adjust weight to counter differences.
- 21 Centrifuge at least 3,000 rcf in a swinging bucket, at 4°C for 20 minutes with brake off.
- 22 Remove supernatant, remove the lower supernatant with P200, leaving about 50-100 uL.

Resuspension and permeabilisation

- 23 Gently resuspend nuclei in ultracentrifuge tube and transfer to 1.5 mL DNA lobind tube.



- 24 Rinse ultracentrifuge tube with Wash buffer 2 and transfer also to 1.5 mL DNA lobind tube.
- 25 Centrifuge @ 350-450xg for 5 mins. Discard the supernatant.
- 26 Resuspend in 0.1x lysis buffer (200 uL) and gently pipetmix 5x
- 27 Incubate on ice for 2 min
- 28 Add Wash buffer 2 (1mL) & gently pipetmix 5x
- 29 Centrifuge @ 350-450xg for 5 mins. Discard the supernatant.
- 30 Resuspend in 1X Diluted Nuclei buffer. Count