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DNA Extraction from FANS sorted nuclei

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1 Works for me

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ABSTRACT

Here we describe an optimised protocol for the extraction of genomic DNA from frozen nuclei samples collected from different neural cells using Fluorescence-assisted Nuclei Sorting (FANS) of post mortem human brain. The same protocol is also suitable for DNA extraction from cells allowing optimal recovery in terms of DNA purity and yield.

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MATERIALS TEXT

Reagents required

Slagboom buffer (50mL):

10x STE buffer (5mL) 5% SDS (5mL) Water (40 mL)

- RNAase A (stock concentration: 10mg/ml)
- Proteinase K (stock concentration: 20 mg/mL)
- Majiik mix (1:1 ratio yeast Reagent 3 (Autogen Bioclear, Caine, Wiltshire, UK) + 100% ethanol)
- 100% Isopropanol
- 80% Ethanol
- TE or water

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NOTE- Slagboom buffer can be made in advance while 80% Ethanol should be prepared fresh the same day.

ABSTRACT

Here we describe an optimised protocol for the extraction of genomic DNA from frozen nuclei samples collected from different neural cells using Fluorescence-assisted Nuclei Sorting (FANS) of post mortem human brain. The same protocol is also suitable for DNA extraction from cells allowing optimal recovery in terms of DNA purity and yield.

DNA extraction from FANS purified nuclei

Defrost frozen FANS sorted nuclei contained in a 1.5ml eppendorf tube § On ice



NOTE 1-2 nuclei aliquots (~200,000 nuclei/tube) per each nuclei population should yield between 300-500ng DNA).

- 2 To each sample add either 250μ L / 500μ L / 1mL of Slagboom buffer (SB) depending on size of cell pellet. Normally 250μ L is sufficient for 200,000 nuclei aliquot or $1x10^6$ cells
- 3 Add 500µL of SB to each nuclei sample (if nuclei pellet)*
 - <u></u>
- * If nuclei are stored in running buffer: calculate the volume of 10x STE and 5% SDS to add to each sample to be consistent with the composition of SB (1x STE; 0.5% SDS).
- e.g. If you have collected a 400 μ L sample you would add 50 μ L 10x STE and 50 μ L 5% SDS to make the sample up to 500 μ L
- 4 Add 1µL of RNase A per 500uL buffer
- 5 Incubate at § 37 °C for © 00:45:00 (using a heat block)
- Add proteinase K to a final concentration 2mg/mL (e.g $5\mu L$ for every $500\mu L$ of SB)
- 7 Mix by inverting 10 times (do not vortex or pipette mix)

- Incubate at § 60 °C for © 01:00:00 (water bath) 9 Move samples from the water bath and leave at § Room temperature for ⊚ 00:05:00 10 Briefly spin down tubes (pulse spin) For every 500 µL of SB used, add 100 µL of Majiik Mix (e.g. 200 µL for 1 mL of SB) 11 Mix by vigorous inversions (DO NOT vortex) 12 13 Centrifuge at **317000 x g** for **00:10:00** at **8 Room temperature** Carefully recover supernatant and transfer it to a new labelled 1.5ml tube (leaving ~50µL at the bottom of the tube) 14 Repeat step above by adding another 100µL of Majiik Mix to the new tube 15 Mix by vigorous inversions (DO NOT vortex) 16 17 Centrifuge at **®17000 x g** for **⊘00:10:00 § Room temperature** Recover supernatant and transfer to a new labelled 1.5ml tube leaving ~50ul at the bottom of the old tube. 18 NOTE - If supernatant exceeds 1mL, split into 2 tubes
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Add an equal volume of 100% Isopropanol to each tube (e.g. 1mL supernatant + 1 mL 100% Isopropanol)

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After adding the Isopropanol, add 0.7-1µL Glycogen Blue per tube (this step is optional – but advised as it helps to 20 visualize DNA pellet at the end) Mix by inversion (10 times) 21 22 Centrifuge at **⊗17000 x g** for **⊘00:15:00** at **& Room temperature** (hinges of tubes facing upwards) Carefully remove supernatant and discard 23 NOTE - Care must be taken when pouring supernatant since the DNA pellet only weakly adheres to the side of the tube Proceed by adding 500µL of 80% ethanol to each tube 24 25 Mix gently by pipetting 26 Centrifuge at **③17000 x g** for **⊙00:05:00** at **§ Room temperature** 27 Carefully discard supernatant 28 Pulse centrifuge to collect remaining liquid at the bottom of the tube 29 Recover and discard residual liquid from the bottom of the tube making sure not to disturb the DNA pellet Leave DNA pellets to air dry for 5-15 minutes (leave lids open) 30 OPTIONAL - Leave to dry for additional © 00:10:00 at § 37 °C but be careful not to over-dry pellets

- Resuspend pellet in 20uL of ddH_2O or 1xTE ($15\mu L$ per tube if expected very low yield). Avoid pipette mixing, only gently flicking
- $32\,$ $\,$ Pulse spin tubes to help pellet resuspend/ dislodge from the tube wall
- 33 Leave tubes at § 4 °C overnight to fully resuspend before quantifying DNA samples (Nanodrop or Qubit measurements).