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Dip-C (Part 1: Chromosome Conformation Capture, for Fixed Nuclei)

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Protocol status: Working

We use this protocol and it's working

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Protocol materials



Sodium dodecyl sulfate solution BioUltra for molecular biology 10% in H₂O **Merck MilliporeSigma (Sigma-Aldrich) Catalog #71736**



Triton X-100, 10% solution **Merck MilliporeSigma (Sigma-Aldrich) Catalog #93443**



NEBuffer 2 (10X) **New England Biolabs Catalog #B7002S**



Mbol (25,000 units/ml) - 2,500 units **New England Biolabs Catalog #R0147M**



T4 DNA Ligase Reaction Buffer - 6.0 ml **New England Biolabs Catalog #B0202S**



BSA, molecular biology grade, 20 mg/ml **New England Biolabs Catalog # B9000S**



T4 DNA Ligase (1 U/μL) **Thermo Fisher Catalog #15224025**



Corning™ Falcon™ Test Tube with 35μm Cell Strainer Snap Cap **Corning Catalog #352235**



Falcon 40 μm Cell Strainer **Corning Catalog #352340**



PBS, pH 7.4 **Thermo Fisher Catalog #10010023**



PBS, pH 7.4 **Thermo Fisher Catalog #10010023**



DAPI (4',6-Diamidino-2-Phenylindole, Dihydrochloride) **Thermo Fisher Catalog #D1306**



96 well LoBind PCR plates Semi-skirted **Eppendorf Catalog #0030129504**



Digestion

40m

- 1 Thaw a tube of fixed nuclei On ice .
- 2 Prepare 0.5% SDS (50 μL per sample; recipe below for 100 μL):
 - 5 μL

Sodium dodecyl sulfate solution BioUltra for molecular biology 10% in H₂O **Merck MilliporeSigma (Sigma-Aldrich) Catalog #71736**
 - (final: 0.5 Mass Percent)
 - 95 μL water
 - Vortex to mix
- 3 Resuspend nuclei in 50 μL 0.5% SDS.

- 4 Incubate at 62 °C for 00:10:00 .

10m

- 5 Add:
 - 145 μL water
 - 25 μL

Triton X-100, 10% solution **Merck MilliporeSigma (Sigma-Aldrich) Catalog #93443**

- 6 Rotate at 37 °C for 00:15:00 .

15m

- 7 Add restriction enzyme and its buffer:
 - 25 μL NEBuffer 2 (10X) **New England Biolabs Catalog #B7002S**
 - 20 μL

MboI (25,000 units/ml) - 2,500 units **New England Biolabs Catalog #R0147M**

- 8 Rotate at 37 °C Overnight .

15m

Ligation


40m




- 9 Centrifuge at 1000 x g, 4°C, 00:05:00 .
- 10 Make Ligation Buffer (2 mL per sample; recipe below for 1 mL):
- 100 μ L
 T4 DNA Ligase Reaction Buffer - 6.0 ml **New England Biolabs Catalog #B0202S**
 - 5 μ L
 BSA, molecular biology grade, 20 mg/ml **New England Biolabs Catalog # B9000S**
 - 865 μ L water
 - Vortex to mix.
- 11 Remove supernatant leaving ~ 50 μ L .
- 12 Resuspend in 1 mL Ligation Buffer.
- 13 Centrifuge at 1000 x g, 4°C, 00:05:00 .
- 14 Remove supernatant leaving ~ 50 μ L .
- 15 Resuspend in 1 mL Ligation Buffer.
- 16 Add 10 μ L T4 DNA Ligase (1 U/ μ L) **Thermo Fisher Catalog #15224025** .
- 17 Pipette to mix.
- 18 Incubate at 16 °C for 04:00:00 , occasionally inverting the tube.

4h



19 Optionally filter with  Falcon 40 µm Cell Strainer **Corning Catalog #352340** or


 Corning™ Falcon™ Test Tube with 35µm Cell Strainer Snap
Cap **Corning Catalog #352235**

to avoid clogging the flow cytometer.

20 Aliquot if needed.


21 Centrifuge at  1000 x g, 4°C, 00:05:00 .



22 Remove supernatant.

23 Store at  -80 °C .

Flow Sorting


40m


24 On the day of flow sorting, thaw a tube of ligated nuclei  On ice .

25 Resuspend in  1 mL  PBS, pH 7.4 **Thermo Fisher Catalog #10010023** .

26 Make 300 µM DAPI:



-  100 µL  PBS, pH 7.4 **Thermo Fisher Catalog #10010023**

-  2.1 µL 14.3 mM (5 mg/mL) DAPI (stock made by dissolving

 DAPI (4',6-Diamidino-2-Phenylindole, Dihydrochloride) **Thermo Fisher Catalog #D1306**

in  2 mL water and stored at  4 °C)

- Vortex to mix.

27 Add  1 µL 300 µM DAPI (final:  0.5 nanomolar (nM)).

28 Pipette to mix.



- 29 Flow sort single nuclei into  96 well LoBind PCR plates Semi-skirted **Eppendorf Catalog #0030129504** either dry or containing lysis buffer (which requires lysis by incubation before storing; see Part 2 for details).
- 30 Proceed directly to Part 2, or store at  -80 °C .