



Formalin-fixed and non-paraffin-embedded tissue DNA extraction

This method is applicable for scientists that need to extract DNA from formalin-fixed tissue.

Purpose

Due to fragmentation of the DNA and cross-linked protein-DNA complexes during the tissue formalin fixation process, getting any amount of DNA from formalin-fixed and non-paraffin-embedded tissue can be a big challenge. Here we demonstrated a DNA extraction protocol from formalin-fixed and nonparaffin-embedded brain tissue using FormaPure DNA XL kit. In this protocol, protein-DNA cross-links are reversed using heat.

Materials Used

Material	Part Number	Supplier
FormaPure XL DNA	C35996	Beckman Coulter
Microcentrifuge tubes 1.5 mL	357448	Beckman Coulter
Tube Magnet (1.5, 1.7, and 2ml)	A29182	Beckman Coulter
SPRIPlate 96R Ring Super Magnet Plate	A32782	Beckman Coulter
1.2 mL 96-well Plate	AB1127, or equivalent	Thermo Fisher
Ethanol	AB-00138	American Bioanalytical
DNase I (RNase-free)	AM2222 or AM2224	Thermo Fisher
Nuclease-free water (Molecular Grade)	AM9932	Thermo Fisher
Microcentrifuge	NA	NA
Tissuelyser	NA	NA
Hand homogenizer	NA	NA

Protocol

1. Homogenization/Lysis

- a. Transfer the fixed tissue from the original container into a 1.5mL tube. Wash the fixed tissue with 1 mL 1x PSB for 30sec. Repeat for a total of 2 washes
- b. Hydrate the fixed tissue with gradient of Ethanol (from 100%, 95%, 90%, 80%, 70%, to 50%) and finally in H₂O
- c. Transfer 30 mg fixed tissue to a 1.5 mL centrifuge tube or 96-well plate
- d. Add 200 μL LBA and 30 μL Proteinase K to the tissue
- e. Homogenize the tissue use either tissuelyser (steel ball). Adjust the speed and time to make sure of proper tissue homogenization. After homogenization, remove the steel ball. Hand homogenizer is also recommended if low throughput is desired.
- f. Centrifuge the sample at 12000 rpm for 1 min
- g. Transfer the supernatant to a new well or tube. Leave the non-homogenized tissue at the bottom of the tube or plate
- h. Incubate the sample at 60°C overnight

2. Decrosslinking

- a. Incubate the sample for 60 min at 80°C
- b. Remove the sample from the heat source

3. RNase A Treatment

- a. Add $5 \mu L$ of RNase A to the sample
- b. Mix by pipetting up and down 10 times, or until thoroughly mixed
- c. Incubate for 5 min at room temperature

4. Bind

- a. Add $300 \mu L$ of BBA to the sample
- b. Mix by pipetting up and down 10 times, or until thoroughly mixed.
- c. Incubate for 5 min at room temperature
- d. Place the sample on a **magnet** for **10 min** (or until supernatant is clear)
- e. Remove and discard the supernatant without disrupting the beads
- f. Remove the sample from the magnet

5. Wash

- a. Add 400 µL of Wash buffer WBA to the sample
- b. Mix by pipetting up and down 10 times, or until thoroughly mixed
- c. Place the sample on a **magnet** for **10 minutes** (or the supernatant is clear)
- d. Remove and discard the supernatant without disrupting the beads
- e. Remove the sample from the magnet

6. Ethanol Wash

- a. Add 750 μ L of freshly prepared 80% ethanol to each sample
- b. Mix by pipetting up and down 10 times, or until thoroughly mixed
- c. Place the sample on a magnet for 10 minutes (or the supernatant is clear)
- d. Remove and discard the supernatant without disrupting the beads
- e. Remove the sample from the magnet
- f. Air dry the samples on the magnet for 10 min

7. Elute

- a. Add 40 μL of nuclease free water to the sample
- b. Mix by pipetting up and down 10 times, or until thoroughly mixed
- c. Incubate for 2 minutes at 60°C
- d. Place the sample on a magnet for **5 minutes** (or until supernatant is clear)
- e. Remove and **save** the supernatant without disrupting the beads

Results

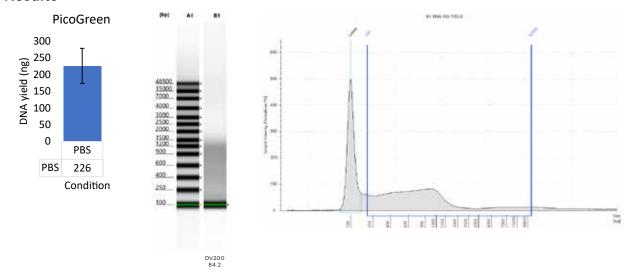


Figure. Formalin-fixed and non-paraffin-embedded brain tissue (customer sample) derived DNA yield and integrity using FormaPure DNA XL. Left: DNA yield was measured by Quant-iT™ PicoGreen® dsDNA Assay. Middle: DNA integrity and DV200 were measured by gDNA TapeStation. Extraction experiments were done in triplicate.

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