

# truXTRAC<sup>®</sup> FFPE total NA (tNA) Kit -Magnetic Bead Purification (25)

(PN 520246)

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## **Intended Use**

The truXTRAC FFPE total NA (Nucleic Acid) Kit is intended for research use only. This product is not intended for the diagnosis, prevention, or treatment of any disease.

## Introduction

The truXTRAC FFPE total NA Kit is designed for efficient and sequential extraction of total nucleic acids (RNA and DNA) from Formalin Fixed, Paraffin Embedded (FFPE) tissue samples using Covaris Adaptive Focused Acoustics (AFA).

AFA-energetics<sup>®</sup> enables the active removal of paraffin from FFPE tissue samples in an aqueous buffer, while simultaneously rehydrating the tissue. Compared to traditional passive, chemical-based methods of paraffin removal, this non-contact mechanical process is more efficient as the paraffin is removed and emulsified from the tissue. Uniquely, AFA enables increased yields of nucleic acids and minimizes the degradation of nucleic acids exposed at the FFPE section surface. The truXTRAC protocol results in high yields of high-quality RNA and DNA for sensitive analytical methods such as next-generation sequencing (NGS) or qPCR/RT-qPCR.

This protocol is optimized for FFPE slide sections up to 20 µm in total material thickness (from a 10 mm2 area), scrolls/curls up to 20 µm in total material thickness, or one core up to 1.4 mm in diameter. For samples of larger input sizes, the truXTRAC total NA Plus Kit (<u>PN 520255</u>) using magnetic bead-based purification may be used for extraction and purification of DNA and RNA from FFPE samples.

#### Important Notes on FFPE Samples:

The yield of DNA and RNA from FFPE tissue blocks is highly variable. Factors such as fixation time, size and thickness of the cores, the ratio of tissue to paraffin, the type of tissue, and the age and storage conditions of the FFPE block are the main causes for this variability.

More importantly, however, the quality of DNA and RNA isolated from FFPE samples can also be highly variable. During the fixation process, DNA and RNA are cross-linked to proteins and other nucleic acid molecules to varying degrees. The nucleic acid fragment or strand length isolated from FFPE samples is generally shorter as compared to nucleic acids that are isolated from fresh or frozen tissues [1]. This is particularly evident in older FFPE sample blocks or sample blocks stored at elevated temperatures. Thus, an advanced mechanical deparaffinization process is important to extract the higher quality nucleic acids required for sensitive analytical techniques. Covaris AFA enables non-contact mechanical removal of paraffin from FFPE samples to improve the yield and quality of extracted nucleic acids [2].

#### Note for users:

If you require any assistance with this product, please refer to Troubleshooting (*Appendix B*) in this protocol, check the FAQs found on our website, or contact Covaris Application Support at <u>applicationsupport@covaris.com</u>.



#### **Important Notices**

- Proper Incubation during Proteinase K and De-crosslinking: Incubation times and temperatures during Proteinase K (at 56 °C) and de-crosslinking (at 80 °C) are crucial to the successful extraction and purification of nucleic acids from FFPE. The recommended process of checking and adjusting temperatures on the heat blocks to ensure proper temperatures of lysate in the vessels must be followed (see *Section 3*).
- 2. Separation of RNA and DNA containing Fractions by Centrifugation: A critical step in the truXTRAC FFPE protocol is the separation of RNA and DNA by centrifugation. At this point in the protocol RNA is located in the supernatant and the majority of DNA is still trapped in the tissue pellet. The user must provide equipment that allows centrifugation to protocol specifications (see *Page 5*). Incomplete separation of fractions can result in yield and quality loss.
- 3. **FFPE Sample to Sample Variability:** FFPE tissue samples vary widely due to a variety of reasons. The degree of formaldehydeinduced crosslinking, the tissue type itself (highly connective or granular), as well as the wax to tissue ratio can all impact the yield and quality of the extracted and purified nucleic acids. The truXTRAC FFPE protocol was developed to reduce the impact of these factors. However, the user must follow the recommendations (see *Section 1*) for sample input. In some extreme cases, complete homogenization of the tissue cannot be achieved. After resuspension of the DNA containing pellet and incubation with Proteinase K, chunks of tissue may still be visible. Covaris recommends increasing the Proteinase K incubation time to 90 minutes if the tissue is not completely lysed. The decrosslinking time of RNA and DNA can be increased to 60 and 120 min, respectively.

#### **Kit Contents**

- Tissue Lysis Buffer......6 mL
- Proteinase K (PK Solution) .....1.25 mL
- Magnetic Bead Suspension ......0.5 mL
- Buffer BB3 ......25 mL

- Buffer BE.....7.5 mL
- microTUBE-130 AFA Fiber Screw-Cap FFPE .... 25 count

SDS Information available at: https://www.covaris.com/safety-data-sheets/

#### Storage

Upon kit arrival, store the Proteinase K solution and the Magnetic Bead Suspension at 2 to 8 °C. Store all other kit components at ambient temperature.



## Laboratory Equipment, Chemicals, and Consumables Supplied by User

### **Required Laboratory Equipment and Accessories**

- microTUBE-130 Centrifuge and Heat Block Adapter (Covaris PN 500406)
- Dry block heater for 2 ml tubes or temperature-controlled water bath able to accurately heat between 50 to 90 °C
- Magnetic Stand for 2 mL tubes (e.g., Thermo Fisher Scientific, DynaMag<sup>™</sup>-2 Magnet, PN 12321D)

#### **Required Reagents**

- 100% ethanol, molecular biology grade (e.g., AmericanBio, PN AB00515)
- Nuclease-free water (e.g., Invitrogen, PN AM9930)

#### **Optional Enzymes**

- TURBO<sup>™</sup> DNase (Thermo Fisher Scientific, PN AM2238)
- RNase A, DNase and protease-free (10 mg/mL) (e.g., Thermo Fisher Scientific, PN EN0531)

#### **Required Consumable**

• 2 mL nuclease-free microcentrifuge tubes (e.g., Eppendorf Safe-Lock Tubes, PN 022363352)

#### Covaris Focused-ultrasonicator Accessories and Plate Definitions

The tables below contain the parts and plate definitions necessary to run the protocol. Use the parts and plate definitions specific to your Covaris Focused-ultrasonicator.

Instrument	M220	ME220	S220	E220evolution	E220	LE220/LE220-plus
Holder/Rack Description (PN)	Holder XTU (500414)	Rack 4-place microTUBE Screw- Cap (500522)	Holder microTUBE Screw-Cap (500339)	Rack E220e 4 Place microTUBE Screw Cap (500432)	Rack 24 Place microTUBE Screw- Cap (500308)	Rack 24 Place microTUBE Screw- Cap (500308)
Required Accessories (PN*)	Insert XTU (500489)	ME220 Waveguide 4 Place (500534)	N/A	Intensifier (500141)	Intensifier (500141)	N/A

\*See Appendix C for more information.

Instrument	Plate Definition File Name
ME220	"4 microTUBE-500 Screw-Cap PN 520185.2.rck"
E220evolution	"500484 E220e 4 microTUBE-500 Screw-Cap -9.9mm offset"
E220 / LE220 / LE220-plus / LE220R-plus / LE220RSc	"Name of Instrument*_500452 Rack 24 Place microTUBE-500 Screw-Cap +6mm offset"

\*For example: "LE220plus\_500308 Rack 24 Place microTUBE-130 Screw-Cap +15mm offset"

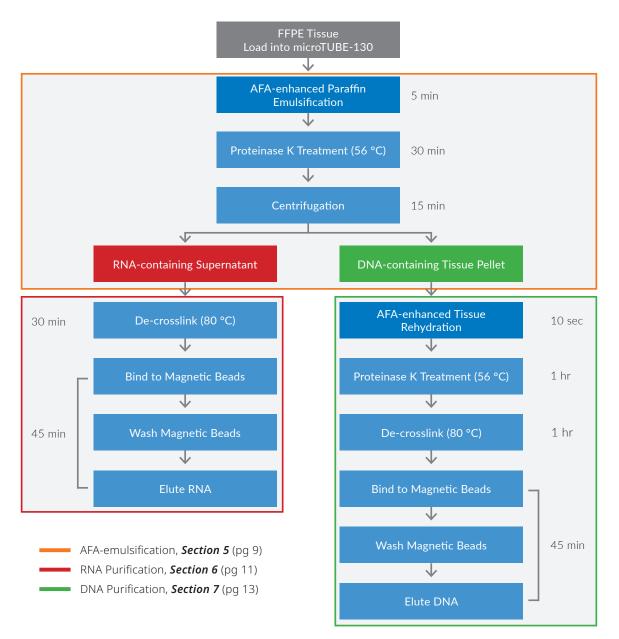


## **FFPE tNA Extraction and Purification Workflow**

Using the Adaptive Focused Acoustics (AFA) process, FFPE tissue samples are prepared in Tissue Lysis Buffer in the presence of Proteinase K, followed by an incubation at 56 °C for a short duration. This results in the release of RNA while minimizing over-digestion of the tissue and loss of genomic DNA.

The RNA-containing supernatant is separated from the DNA-containing tissue by a centrifugation step. RNA is then de-crosslinked and purified using magnetic beads.

Sequentially, DNA is released from the DNA-containing tissue by AFA-enhanced Proteinase K digestion, following a de-crosslinking step. DNA is purified using magnetic beads.



## **1 - FFPE Sample Input Requirements and Guidelines**

The truXTRAC protocol is highly efficient at mechanically removing paraffin, while simultaneously rehydrating the tissue.



**CAUTION:** Do NOT exceed the input requirements in the tables below. Overloading will negatively impact the quality and quantity of extractable nucleic acids.

#### **Slide Section Input Requirements:**

Slide Collection Method	Maximum Input per microTUBE-130
Scalpel or razor blade to scrape material from slides	20 μm of total thickness Area: 10 x 10 mm (2 slides at 5 μm thick = 10 μm total thickness)

#### **Curls/Scrolls Input Requirements:**

For best results, minimize the amount of wax present by trimming. We recommend no more than 1-part wax to 1-part tissue.

FFPE Curl/Scroll Thickness	Maximum Scrolls per microTUBE-130
5 µm	3
10 µm	2
15 to 20 μm	1

#### **Core Input Requirements:**

FFPE Core Punch Outer Diameter	Maximum Core Punches per microTUBE-130
≤ 1.4 mm (15 Gauge, outer); Length = 5 mm	1

Core punches may be loaded directly into the microTUBE-130 or transferred into the microTUBE-130 using tweezers or forceps.



## 2 - Preparation of Reagents

NOTE: Follow these instructions before starting the FFPE total NA isolation process.

- 1. Tissue Lysis Buffer: Visually check for a white precipitate that may form during storage before each use. If white precipitate is visible, incubate the Tissue Lysis Buffer at 50 to 60 °C for 5 to 10 minutes before use to dissolve any precipitate.
- 2. 80% Ethanol: Prepare 80% ethanol by mixing 4 parts 100% ethanol with 1 part nuclease free water. One sample requires 1.8 mL of 80% ethanol. To prepare the total amount of 80% ethanol needed, multiply the number of samples to be processed by 2 mL to account for dead volume.

#### **3 - Preparation of Heat Blocks**

- 1. Preheat dry block heaters to 56 °C and 80 °C ± 3 °C. It is critical that these temperatures are accurate to successfully execute the protocol.
- 2. Test the temperature of your heat blocks:
  - a. Place a microcentrifuge tube (1.5 or 2 mL) filled with water into the heat block.
  - b. Immerse a thermometer into the tube.
  - c. Wait until the temperature has reached the plateau.
  - d. Adjust the set-temperature accordingly until the temperature inside the microcentrifuge tube has reached 56 °C or 80 °C ± 3 °C.

CAUTION: The Covaris microTUBE-130 must be used in conjunction with Covaris Centrifuge and Heat Block microTUBE Adapters (PN 500406). It is important to use an accurate heating source for incubation of microTUBE-130 and microcentrifuge tubes during Proteinase K and de-crosslinking incubations. Deviation from the indicated temperatures can adversely impact quality and quantity of purified nucleic acids.



#### 4 - Focused-ultrasonicator Setup

For detailed instructions on how to prepare and use your instrument, please refer to the respective Covaris User Manual. If you do not see a Plate Definition on your system, please contact Covaris Technical Support (<u>techsupport@covaris.com</u>)

NOTE: Refer to Page 5 for Plate Definitions and required Focused-ultrasonicator accessories.

#### 1. Create "Acoustic Paraffin Emulsification" program in SonoLab

Use the settings provided in the table below, specific to your Covaris instrument type, to create a program called "*Acoustic Paraffin Emulsification*" using the Covaris SonoLab method editor. Save the program for later use.

Instrument	M220	ME220	<b>S220</b>	E220evolution	E220	LE220 / LE220-plus
Peak Incident Power (PIP) (Watt)	75	75	175	175	175	450
Duty Factor (%)	20	25	10	10	10	30
Cycles Per Burst (#)	200	1000	200	200	200	200
Treatment time (seconds)	300	300	300	300	300	300
Temperature (°C)	20	20	18	18	18	18
Water Level (run)	Full	Auto	15	10	10	15

#### 2. Create "Acoustic Pellet Resuspension" program in SonoLab

Use the settings provided in the table below, specific to your Covaris instrument type, to create a program called "Acoustic Pellet Resuspension" using the Covaris SonoLab method editor. Save the program for later use.

Instrument	M220	ME220	<b>S220</b>	E220evolution	E220	LE220 / LE220-plus
Peak Incident Power (PIP) (Watt)	75	75	175	175	175	450
Duty Factor (%)	20	25	10	10	10	30
Cycles Per Burst (#)	200	1000	200	200	200	200
Treatment time (seconds)	10	10	10	10	10	10
Temperature (°C)	20	20	18	18	18	18
Water Level (run)	Full	Auto	15	10	10	15



## 5 - Paraffin Emulsification, Tissue Rehydration, and Lysis

- 1. Prepare Tissue Lysis Buffer/Proteinase K Mix by following instructions in *Table 1* below. Mix by inverting 10 times or
  - vortexing for 3 seconds.

**NOTE:** The Tissue Lysis Buffer/Proteinase K Mix should be stored at room temperature and used within 30 minutes after preparation.

Reagent	Volume for one sample*	Volume for N samples*
Tissue Lysis Buffer	121 µL	121 µL x N
Proteinase K Solution	11 µL	11 µL x N

 Table 1. Tissue Lysis Buffer/Proteinase K Mix. \*Calculation includes 10% excess in final volume.

- 2. Open the microTUBE-130 Screw-Cap and load the FFPE tissue.
- 3. Add 120 µL Tissue Lysis Buffer/Proteinase K mix to the microTUBE-130. FFPE tissue may also be added directly to microTUBEs containing Lysis buffer. If adding FFPE tissue to microTUBEs containing Lysis buffer, ensure that the FFPE sample is fully immersed in the tube to prevent the sample from getting stuck in the Screw-Cap thread.
- 4. Close the microTUBE-130 tightly with the Screw-Cap and transfer the microTUBE-130 to the appropriate rack or holder/ insert for your Focused-ultrasonicator. Load the rack or holder/insert containing the microTUBE(s) into the Focusedultrasonicator for processing.
- 5. Process the sample using the "Acoustic Paraffin Emulsification" program on the Focused-ultrasonicator.

**NOTE:** It is expected that the solution will turn milky white. See example below.



6. Remove the microTUBE-130 from the Focused-ultrasonicator and load it into the microTUBE-130 Centrifuge and Heat Block adapter.

**NOTE:** When processing in batches, the samples may be kept at ambient temperature for up to two hours prior to PK Solution incubation at 56 °C (*Step 7*).



## 5 - Paraffin Emulsification, Tissue Rehydration, and Lysis (cont.)

7. Incubate sample(s) for 30 minutes at 56 °C. Remove microTUBE-130 together with the microTUBE-130 adapter from the heat block, and let them cool separately at ambient temperature for 3 minutes.

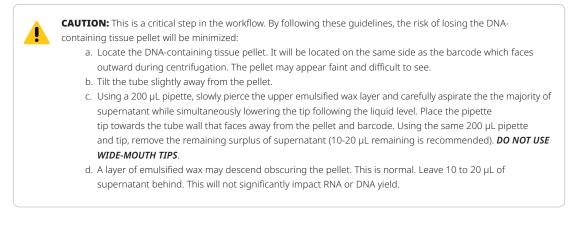
**NOTE:** Do not chill on ice as rapid cooling will cause detergents to precipitate.

8. Place microTUBE-130 in the microTUBE-130 adapter with the bar code on the screw cap sleeve facing outward into a microcentrifuge (fixed angle rotor) and centrifuge at 5,000 x g for 15 minutes.



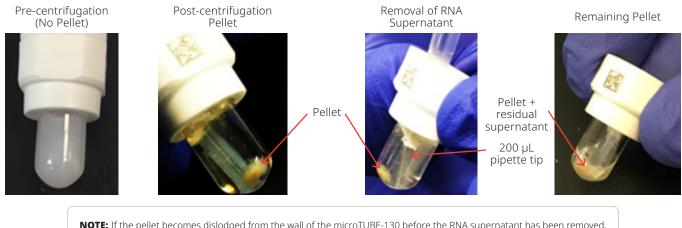
**NOTE:** The centrifuge lid may not close and may need to be left off during centrifugation.

 Remove microTUBE-130(s) from the microcentrifuge, remove Screw-Cap, and carefully transfer 100 µL supernatant into a new 2 mL microcentrifuge tube.





## 5 - Paraffin Emulsification, Tissue Rehydration, and Lysis (cont.)



**NOTE:** If the pellet becomes dislodged from the wall of the microTUBE-130 before the RNA supernatant has been removed, repeat centrifugation (*Step 8*) to re-form pellet. Remove RNA supernatant as described in *Step 9*.

Save the DNA-containing tissue pellet in the microTUBE for subsequent DNA purification as described in *Section-7*.
 Proceed immediately to RNA Purification (*Section-6*).

NOTE: The DNA-containing pellet can be stored on ice or at 2 to 8 °C for up to 1 day. For longer periods, store at -15 to -30 °C.

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## 6 - RNA Purification

- 1. Ensure that dry block heater temperatures have reached 80 °C (*Step 2*) and 56 °C (*Step 22*) as explained in *Section 3*.
- 2. Incubate 2 mL microcentrifuge tube with the RNA-containing supernatant at 80 °C for 20 minutes. Remove tubes from the heat block and cool at room temperature for 3 minutes.
- 3. Prepare BB3/Magnetic Bead Mix according to *Table 2* below.

Reagent	Volume for one sample*	Volume for N samples*
BB3	330 µL	330 µL x N
Magnetic Bead Suspension	8.8 µL	8.8 µL x N

 Table 2. BB3/Magnetic Bead Mix for RNA. \*Calculation includes 10% excess in final volume.

**NOTE:** Thoroughly vortex the Magnetic Bead Suspension and BB3/Magnetic Bead Mix before using to ensure a homogenous suspension. Beads will settle when left standing.

- 4. Add 308 µL of the BB3/Magnetic Bead Mix to the RNA-containing supernatant and cap the microcentrifuge tube.
- 5. Vortex the microcentrifuge tube for 10 seconds.
- 6. Incubate the microcentrifuge tube at ambient temperature for 10 minutes.
- 7. Place the tube on a magnetic stand and incubate for 5 minutes, ensuring the beads have been pulled to the magnet.

**NOTE:** With some samples, the binding supernatant will appear slightly brown after the 5 minute incubation on the magnet stand due to a small percentage of beads that do not migrate to the magnet. This effect does not reduce the yield significantly.

- 8. With the tube on the magnetic stand, carefully remove and discard the supernatant using a 200 µL pipette. Avoid touching or disturbing the bead pellet.
- 9. Remove the microcentrifuge tube from the magnetic stand and add 600 µL WB4.
- 10. Cap the tube and vortex for 10 seconds. Confirm that all beads are resuspended. If beads are still sticking to the wall continue vortexing until all beads are resuspended.
- 11. Place the tube back on the magnetic stand and incubate for 5 minutes, ensuring the beads have been pulled to the magnet.

**Optional step recommended for FFPE core sample inputs:** Repeat wash steps 8 through 11. The beads may not fully re-suspend in the 2nd wash with WB4 which does not present a problem. If you perform the optional DNase below, this includes a 2nd wash step and is not necessary here.

*Optional DNA removal step:* The truXTRAC FFPE total NA Kit protocol isolates total RNA that may contain trace amounts of genomic DNA. If DNA-free RNA is required for downstream applications such as RNA-seq, an optional DNase treatment may be performed to remove DNA. *Note:* This optional DNase digestion must be performed after *Step 11*.

NOTE: This optional DNase digestion must be performed after Step 11. See Appendix A for step-by-step instructions.

12. With the tube on the magnetic stand, carefully remove and discard the supernatant. Remove as much of the supernatant as possible using a 200 µL pipette. Use a 20 µL pipette to remove any remaining liquid from the bottom of the tube.

**NOTE:** It is critical to remove the wash buffer supernatant completely because it contains residual paraffin. Remaining paraffin residue can result in bead clumping during elution and diminished yield.



## 6 - RNA Purification (cont.)

- 13. Remove the tube from the magnetic stand and add 600  $\mu$ L 80% ethanol.
- 14. Cap the tube and vortex for 10 seconds. Confirm that all beads are resuspended. If beads are still sticking to the wall continue vortexing until all beads are resuspended.
- 15. Place the tube on the magnetic stand and incubate for 2 minutes, ensuring the beads have been pulled to the magnet.
- 16. Remove and discard the supernatant without disturbing the bead pellet.
- 17. Remove the tube from the magnetic stand and add 300  $\mu L$  80% ethanol.
- 18. Cap the tube and vortex for 10 seconds. Confirm that all beads are resuspended. If beads are still sticking to the wall continue vortexing until all beads are resuspended.
- 19. Place the tube on the magnetic stand and incubate for 2 minutes, ensuring the beads have been pulled to the magnet.
- 20. Remove and discard as much of the supernatant as possible. Use a 20 μL pipette to remove any remaining liquid from the bottom of the tube.

**NOTE:** Ensure by visual examination that the ethanol has evaporated before continuing with elution. Residual ethanol can inhibit the elution and impact downstream applications such as PCR.

- 20. Leave the tube open on the magnetic stand and let the beads dry for 6 minutes at room temperature.
- 21. Remove the tube from the magnetic stand and add 50 to 100 µL of RNA Elution Buffer. Resuspend the beads by pipetting up and down 20 times. Ensure that all of the beads are resuspended in the buffer with none still sticking to the wall of the tube.
- 22. Cap the tube and incubate it in the heat block set to 56 °C for 5 minutes.
- 23. Remove the tube from the heat block and place it on the magnetic stand and incubate for 2 minutes, ensuring the beads have been pulled to the magnet.
- 24. Transfer the eluate into a clean elution tube without transferring beads. A small amount of residual paraffin may be visible in the pipette tip. This will not adversely affect downstream processing of the eluted RNA. Keep eluted RNA on ice until further processing. Isolated RNA should be kept at -80 °C for long term storage.

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## 7 - DNA Purification

- Set up the dry-heat blocks as explained in Section-3 and verify the block temperatures to be 56 °C and 80 °C. The heat block set to 56 °C is required for Proteinase K incubation (*Step 7*) and DNA elution after purification via magnetic beads (*Step 33*). The heat block at 80 °C is required for DNA de-crosslinking (*Step 9*). Place the heat block adapters in the heat block set to 56 °C.
- 2. Prepare Tissue Lysis Buffer/Proteinase K Mix in a tube following the instructions in *Table 3* and mix by inverting 10 times or vortexing for 3 seconds.

Reagent	Volume for one sample*	Volume for N samples*
Tissue Lysis Buffer	88 µL	88 µL x N
Proteinase K	22 µL	22 µL x N

 Table 3. Tissue Lysis Buffer/Proteinase K Mix for DNA. \*Calculation includes 10% excess in final volume.

**NOTE:** The Tissue Lysis Buffer/PK Solution Mix should be stored at ambient temperature and used within 30 minutes after preparation.

- Open the microTUBE with the DNA-containing tissue pellet (*Section-5 Step 10*) and add 100 μL of the Tissue Lysis Buffer/Proteinase K Mix for DNA. Re-cap the Screw-Cap microTUBE tightly.
- 4. Transfer the microTUBE-130 to the appropriate rack or holder/insert for your Focused-ultrasonicator. Load the rack or holder/insert containing the microTUBE(s) into the Focused-ultrasonicator for processing.
- 5. Process the sample using the "*Acoustic Pellet Resuspension*" program on your Covaris Focused-ultrasonicator.
- 6. Remove the microTUBE-130 from the Focused-ultrasonicator and load the microTUBE into the pre-warmed microTUBE-130 Centrifuge and Heat Block adapter on the heat block set to 56 °C.
- 7. Incubate for 60 minutes at 56 °C.

**NOTE:** The Proteinase K treated sample can be stored at room temperature for up to an additional hour. Do not chill on ice.

**CAUTION:** It is recommended to increase the Proteinase K incubation time to 2 hours or up to overnight for >15 µm scrolls or 1.4 mm core sample inputs.

- 8. Remove microTUBE together with the microTUBE-130 adapter from the heat block and transfer directly to the dry heat block set-up for 80 °C incubation.
- 9. Incubate for 60 minutes at 80 °C.
- 10. Remove the microTUBE-130 from the heat block and let cool for 3 minutes at room temperature. Do not chill on ice.
- 11. Transfer the entire sample into a clean 2 mL microcentrifuge tube.

*Optional RNA removal step:* At this point the sample can be treated with RNase A to remove residual RNA before continuing with DNA purification. Add 5 μL of RNase A solution and incubate for 5 minutes at room temperature, then continue to *Step 12*.



## 7 - DNA Purification (cont.)

12. Prepare BB3/Magnetic Bead Mix according to Table 4.

Reagent	Volume for one sample*	Volume for N samples*
BB3	198 µL	198 µL x N
Magnetic Bead Suspension	8.8 µL	8.8 µL x N

 Table 4. BB3/Magnetic Bead Mix for DNA. \*Calculation includes 10% excess in final volume.

**NOTE:** Thoroughly vortex the Magnetic Bead Suspension and BB3/Magnetic Bead Mix before using to ensure a homogenous suspension. Beads will settle when left standing.

- 13. Add 188 µL of the BB3/Magnetic Bead Mix to the DNA solution in the 2 mL tube.
- 14. Cap the tube and vortex the tube for 10 seconds.
- 15. Incubate the tube on the benchtop at room temperature for 10 minutes to ensure complete binding of the DNA.
- 16. Place the tube on the magnetic stand and incubate for 5 minutes, or until the beads have been pulled to the magnet.

**NOTE:** With some samples, the binding supernatant may appear slightly brown after the 5 minute incubation on the magnet stand due to a small percentage of beads that do not migrate to the magnet. This effect does not reduce the yield significantly.

- With the tube still on the magnetic stand, carefully remove and discard the supernatant using a 200 μL pipette.
   Avoid touching or disturbing the bead pellet.
- 18. Remove tube from the magnetic stand and add 600  $\mu$ L Buffer WB3.
- 19. Cap the tube and vortex for 10 seconds. Confirm that all beads are resuspended. If beads are still sticking to the wall continue vortexing until all beads are resuspended.
- 20. Place the tube on the magnetic stand and incubate for 5 minutes, or until the beads have been pulled to the magnet.
- Remove and discard as much of the supernatant as possible using a 200 μL pipette. Use a 20 μL pipette to remove the remaining liquid from the bottom of the tube.

**NOTE:** It is critical to remove the wash buffer supernatant completely because it contains residual paraffin. Remaining paraffin residue will result in bead clumping during elution and diminished yield.

*Optional step recommended for FFPE cores:* Repeat wash *Steps 18 through 21*. The beads may not fully re-suspend in the 2nd wash with WB3 which does not present a problem.

- 22. Remove tube from the magnetic stand and add 600  $\mu$ L 80% ethanol to the tube.
- 23. Cap the tube and vortex for 10 seconds. Confirm that all beads are resuspended. If beads are still sticking to the wall continue vortexing until all beads are resuspended.
- 24. Place the tube on the magnetic stand and incubate for 2 minutes, or until the beads have been pulled to the magnet.
- 25. Remove and discard the supernatant without disturbing the bead pellet.
- 26. Remove the tube from the magnetic stand and add 300  $\mu L$  80% ethanol.



- 27. Cap the tube and vortex for 10 seconds. Confirm that all beads are in suspension. If beads are still sticking to the wall continue vortexing until all are suspended.
- 28. Place the tube on the magnetic stand and incubate for 2 minutes, ensuring the beads have been pulled to the magnet.
- Remove and discard as much of the supernatant as possible using a 200 μL pipette. Use a 20 μL pipette to remove the remaining liquid from the bottom of the tube.
- 30. Leave the tube open on the magnetic stand and let the beads dry for 6 minutes at room temperature.

**NOTE:** Ensure by visual examination that the ethanol has evaporated before continuing with elution. Residual ethanol can inhibit the elution and impact downstream applications such as PCR.

- 31. Remove the tube from the magnetic stand and add 50 to 100 µL of Buffer BE (5 mM TrisCl pH 8.5) into the tube.
- 32. Resuspend the beads by pipetting up and down 20 times. Ensure that all of the beads are resuspended in the buffer and none are still sticking to the wall of the tube.
- 33. Cap the tube and incubate the microcentrifuge tube in the heat block set to 56 °C for 5 minutes.
- 34. Remove the tube from the heat block, place it on the magnetic stand, and incubate for 2 minutes.
- 35. Transfer the eluate into a clean elution tube without transferring beads. A small amount of residual paraffin may be visible in the pipette tip. This will not adversely affect downstream processing of the eluted DNA. Isolated DNA should be kept at 2 to 8 °C for short term storage (1 to 2 days) and -20 °C for long term storage.



## **Appendix A: Optional DNase Treatment of Extracted RNA**

The truXTRAC FFPE total NA kit isolates total RNA that may contain small amounts of DNA. An optional DNase treatment protocol is provided if DNA-free RNA is desired.

This procedure is performed after Step 11 in Section-6 (RNA Purification).

#### The protocol below describes removal of DNA specifically using TURBO DNA-free kit (Thermo Fisher Scientific, PN AM1907).

1. Prepare a 1 X TURBO DNase master mix:

Reagent	Volume for one sample*	Volume for N samples*
RNase-free H2O	93.5 μL	93.5 μL x N
10X TURBO DNase buffer	11 µL	11 μL x N
TURBO DNase	5.5 µL	5.5 μL x N

Table 5. DNase master mix. \*Calculation includes 10% excess in final volume.

- 2. Add 100 µL of DNase master mix to each bead pellet.
- 3. Resuspend the beads by pipetting up and down 20 times.
- 4. Incubate at room temperature for 30 minutes.
- 5. Add 300  $\mu$ L of BB3 and vortex for 5 seconds.
- 6. Incubate at room temperature for 10 minutes.
- 7. Place the tube on a magnetic stand and incubate for 5 minutes until the beads have been pulled to the magnet.
- 8. Carefully remove and discard the supernatant using a 200 µL pipette. Avoid disturbing the bead pellet.
- 9. Remove the tube from the magnetic stand and add 600 µl WB4 to the tube.
- 10. Cap it and vortex thoroughly until all beads are resuspended.
- 11. Place the tube back on the magnetic stand and incubate for 5 minutes until the beads have been pulled to the magnet.
- 12. Remove as much of the supernatant as possible using a 200 μL pipette. Use a 20 μL pipette to remove the remaining liquid from the bottom of the tube.
- 13. Proceed with Step 12 in Section-6 (RNA Purification).

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# Appendix B: Troubleshooting Guide

Issue	Cause	Solution	Comments / Suggestions
Low DNA yield	First proteinase K incubation too long.	Optimize the 1st proteinase K digestion step for your tissue samples.	During the 1st incubation step with Proteinase K at 56C, the RNA is released, and most of the DNA stays in the remaining tissue. If the PK digestion step is too long, the tissue will be over digested resulting in the release of the DNA into the solution.
	Parts of entire tissue pellet lost during supernatant removal.	Repeat using narrow mouth 200 µL pipette tip to take off RNA-containing supernatant.	Follow guidelines in the protocol closely. Make sure laboratory personnel are trained in this procedure.
	Loss of magnetic beads during purification steps.	Remove supernatant of the bind and wash steps slowly and carefully. If beads appear in the pipette tip, eject the liquid back into the tube, wait for 1 minute, and try aspirating the supernatant again.	The viscosity of buffers BB3, WB3 and WB4, as well as the presence of the paraffin emulsion can make supernatant removal difficult.
	Low tissue to wax ratio in FFPE section.	Trim off any excess paraffin before sectioning a FFPE tissue block. Repeat the procedure using additional sections until desired yield is achieved.	In your initial use of the truXTRAC FFPE total NA kit, use FFPE blocks that have been well characterized for yield and quality.
	Insufficient tissue input.	Select FFPE section with higher tissue to wax ratio or add additional section.	See sample input guidelines in Section-1.
Low RNA yield	Loss of magnetic beads during purification steps.	Remove supernatant of the bind and wash steps slowly and carefully. If beads appear in the pipette tip, eject the liquid back into the tube, wait for 1 minute, and try aspirating the supernatant again.	The viscosity of buffers BB3, WB3 and WB4, as well as the presence of the paraffin emulsion can make supernatant removal difficult.
	Low tissue to wax ratio in FFPE section.	Trim off any excess paraffin before sectioning a FFPE tissue block. Repeat the procedure using additional sections until desired yield is achieved.	In your initial use of the truXTRAC FFPE total NA kit use FFPE blocks that have been well characterized for yield and quality.
	Insufficient tissue input.	Select FFPE section with higher tissue to wax ratio or add additional section.	See sample input guidelines in Section-1.
Beads clumpy during elution and DNA and/ or RNA yield low	Residual paraffin in elution.	For samples with a high paraffin content, a second wash buffer wash may be required to completely remove the paraffin.	If the paraffin emulsion was not completely removed in the wash steps, residual wax can be carried through to the elution step.
Eluates are cloudy.	Residual paraffin in elution.	Spin the eluate for 30 seconds at 10,000 rcf. The residual wax will form a layer on top of the liquid and the aqueous solution can be transferred to a new tube.	If the paraffin emulsion was not completely removed in the wash steps, residual wax can be carried through to the elution step.



## Appendix C: Removing or Installing the Intensifier from an E-Series Instrument (PN 500141)

The 500141 Intensifier is a small, inverted stainless-steel cone centered over the E-Series transducer by four stainless wires. The wires are held in place by a black plastic ring pressed into the transducer well.

If an AFA protocol requires "no Intensifier", please remove the Intensifier, using the following steps:

- 1. Empty the water bath. Start the instrument and start the SonoLab software.
- 2. Wait for the homing sequence to complete (the transducer will be lowered with the rack holder at the home position, allowing easy access to the Intensifier).
- 3. Grasp opposite sides of plastic ring and gently pull the entire assembly out of the transducer well. Do not pull on the steel cone or the wires. The ring is a friction fit in the well no hardware is used to hold it in place.



The 500141 Intensifier (left) shown installed in the E-Series transducer well and (right) removed. Note the "UP" marking at the center of the Intensifier.

If a protocol requires the Intensifier to be present, simply reverse this process:

- 4. Align the black plastic ring with the perimeter of the transducer well. Note that the flat side of the center cone (marked UP) should be facing up (away from the transducer).
- 5. Gently press each section of the ring into the well until the ring is seated uniformly in contact with the transducer, with approximately 2 mm of the ring evenly exposed above the transducer assembly. Do not press on the cone or wires. The rotation of the ring relative to the transducer assembly is not important.
- 6. Refill the tank. Degas and chill the water before proceeding.



## **Tips for Determining Quality and Quantity of the Purified RNA**

- To determine DNA and RNA yields, a fluorometric assay such as Qubit<sup>™</sup> (Life Technologies) should be used.
- In addition, spectrophotometric analysis determining the A260/280 and A260/230 ratios will determine if protein or peptide/salt contamination is present in the sample.
- qPCR can be used to assess the amplifiability of isolated DNA as well as the presence of inhibitors. Note that DNA from FFPE tissue itself can act as inhibitor at high input concentrations due to the extensive damage (nicks, depurination, etc.) that occurs. Therefore, a dilution series over at least 5 orders of magnitude starting with undiluted material of the extracted DNA should always be done when assessing quality by qPCR. An example is shown in Dietrich et al. Figure 1 [3].

## **Additional Notes**

- 1. See following link: <u>https://www.covaris.com/protocols/</u> for updates to this document.
- 2. The treatment settings listed in this document are recommended guidelines. Actual results may vary depending on the tissue type, mass, and previous handling of FFPE samples.
- 3. Covered by US Patent 9,080,167
- 4. Other patents pending

#### References

- 1. Carrick et al. (2015) Robustness of Next Generation Sequencing on Older Formalin-Fixed Paraffin-Embedded Tissue. PLoS ONE 10(7): e0127353.
- 2. Landolt et al. (2016) RNA extraction for RNA sequencing of archival renal tissues. Scand J Clin Lab Invest 76(5):426-434.
- 3. Dietrich et al. (2013) Improved PCR Performance Using Template DNA from Formalin-Fixed and Paraffin-Embedded Tissues by Overcoming PCR Inhibition. PLOS one 8(10): e77771

## **Technical Assistance**

**Technical Support** – Ongoing assistance with the operation or application of the equipment and/or troubleshooting is provided via:

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- · E-mail instrumentation queries to techsupport@covaris.com or application queries to applicationsupport@covaris.com



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