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# **User's Manual and Instructions**

### xTractPure™ Cell and Tissue Nucleic Acid Extraction Kit

Catalog Number: Z2212005

### **Storage Conditions**

This kit is shipped at room temperature. Upon receipt, store Proteinase K at -20°C. Store all remaining components of this kit at room temperature.

#### Shelf Life

1 year from the date of receipt under proper storage conditions

#### **Features**

- Non-toxic chemicals
- High DNA recovery
- Short and Scalable Protocol
- Purified DNA is suitable for NGS, PCR, Bisulfite sequencing, etc

#### Description

BioChain's xTractPure™ Cell and Tissue Nucleic Acid Extraction kit allows for fast and efficient genomic DNA extraction from cultured cell and biological tissue samples. The DNA is eluted in EDTA-free buffer, allowing for immediate use in experiments where EDTA is not tolerated. EDTA may then be added separately for ideal long-term storage of the DNA. The magnetic bead-based extraction protocol is also ideally suited for automation using BioChain's AnaPrep 48 instrument. DNA extracted using this kit is suitable for PCR, qPCR, next generation sequencing (NGS), and other applications.

#### **AnaPrep 48 Automation**

The xTractPure™ Cell and Tissue Nucleic Acid Extraction Kit can be used to isolate DNA from up to 48 cell pellets and/or tissue samples in under one hour using the AnaPrep 48 instrument from BioChain (plates and tip combs need to be ordered separately; please see catalog number Z2212099). Enough reagents are included in each kit to process up to 96 cell pellet and/or tissue samples. An alternative protocol below describes the use of the xTractPure™ kit with the AnaPrep 48 to process samples.

#### Contents

This kit contains all necessary reagents for the isolation of genomic DNA from up to 96 cell pellet and/or tissue samples.

#### **Quality Control**

Each component has been tested for purity and efficacy.

#### **Important Notes**

Starting Material (Cell Pellet):

Up to 1 x 10<sup>6</sup> of either fresh or frozen cell pellets may be used

### Starting Material (Tissue):

For high-yield tissues (e.g. spleen), use up to 10 mg of tissue per extraction For lower-yield tissues (e.g. liver), use up to 20 mg of tissue per extraction

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<u>Quantification:</u> Depending on the sample type, this kit will yield up to 20 ug of purified genomic DNA in a single extraction

### **Equipment and Reagents to be Supplied by User**

- 96-well Deep Well plates and 8-strip Tip Combs (if using the AnaPrep 48 only)\*
   \* BioChain Catalog Number: Z2212099
- o Pipettes
- Vortex-Genie 2 or similar vortexing mixer\*
- Magnet stand for molecular applications (e.g. DynaMag<sup>™</sup>-15 or DynaMag<sup>™</sup>-2)
- o 1.5 ml non-stick Eppendorf tubes
- o 15 ml conical tubes
- o 100% EtOH (200 proof)
- o Isopropanol
- o RNase A (optional)

Contact BioChain® Technical Service for additional recommendations for high throughput or automated mixing.

### **Prior to Initial Use**

BFC, FCW1, and FCW2 Buffers are shipped as a concentrate. If precipitate is present in any of these solutions, incubate solution at 37°C for 30 minutes with occasional shaking. Add the following user-supplied reagents to each bottle:

#### **BFC Buffer**

Add 26.4 ml of isopropanol and mix by inverting gently

#### FCW1 Buffer

Add 25.5 ml of 100% ethanol (200 proof) and mix by inverting gently

#### FCW2 Buffer

Add 88 ml of 100% ethanol (200 proof) and mix by inverting gently

Once these alcohols are added, these buffers are stable for one year if stored properly. Be sure to close the bottle tightly for long term storage.

### **Manual Protocol**

### **Sample Preparation (Cell Pellet)**

- 1. Take up to 1 x 10<sup>6</sup> fresh or frozen cells in a 1.5 ml Eppendorf tube and ensure media has been removed as much as possible
- 2. Add 400 µl of CTFC Buffer to the tube containing the cells
- 3. Add 10 µl of Proteinase K
- **4.** Briefly vortex to mix, pipet up and down 10 times, and incubate at 56°C for 10 minutes
- 5. Vortex briefly to mix and incubate at 56°C for an additional 10 minutes
- **6.** Spin tube briefly on a tabletop centrifuge to bring down solution from the lid of the tube
- 7. (Optional) Add 10  $\mu$ l of RNase A (20 mg/ml), pipet up and down briefly to mix, and let incubate at room temperature for 5 minutes
- 8. Proceed to **Binding** steps in protocol

### Sample Preparation (Tissue)

- For high-yield tissues (e.g. spleen), take up to 10 mg of tissue. For lower-yield tissues (e.g. liver), take up to 20 mg of tissue. For maximum yield, mince or grind the tissue into several small pieces
- 2. In a 1.5 ml Eppendorf tube, add 400  $\mu$ l of CTFC Buffer. Then add 10  $\mu$ l of Proteinase K and pipet up and down briefly to mix
- 3. Transfer the fresh or frozen tissue to the above tube
- **4.** Place tube into either a thermal mixer or water bath and incubate at 56°C for 1-4 hours until all of the tissue pieces are dissolved
  - **If using a thermal mixer:** shake continuously at 1,400 rpm during incubation **If using a water bath:** vortex the samples briefly every 10-15 minutes
- 5. Spin tube briefly on a tabletop centrifuge to bring down solution from the lid of the tube
- **6.** (Optional) Add 10 μl of RNase A (20 mg/ml), pipet up and down briefly to mix, and let incubate at room temperature for 5 minutes
- 7. Proceed to **Binding** steps in protocol

### **Binding**

- 1. To the sample tube, add 300 µl of BFC buffer
- 2. Add 10 µl of Magnetic Bead Solution
  - **Important:** Mix beads well prior to adding. There should be no visible sedimentation at the bottom of the solution after mixing. Beads will settle quickly, so be sure to mix the magnetic bead solution after adding to every few samples. Failure to do so may result in inconsistent yields
- 3. Vortex vigorously for 10 minutes at room temperature
- 4. Place tube onto a magnet stand and let sit for 3 minutes
- **5.** While keeping the tube on the magnet stand, remove supernatant. Be careful not to remove magnetic particles
- **6.** Tap magnet stand on bench 5 times and remove remaining supernatant

#### Wash Steps

7. Transfer tube to non-magnetic rack and add 500 µl of FCW1 Buffer
Resuspend beads by vortexing for 15 seconds. Spin tube briefly and transfer back
to magnet stand

\*Centrifuge steps are needed when using vortexing to resuspend beads. Only a brief spin is recommended to remove solution from tube lid. ALWAYS check to make sure the beads are at the

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bottom of the tube before spinning so that they don't become stuck on the side of the tubes and dry out. If beads are attached to the sides, flick the tube until beads are in solution

- 8. Allow beads to attach to magnet stand for 10-30 seconds
- 9. Remove as much supernatant as possible
- 10. Transfer tube to non-magnetic rack and add 500  $\mu$ l of FCW2 Buffer
- Resuspend beads by vortexing for 15 seconds. Spin tube briefly and transfer back to magnet stand
- 12. Allow beads to attach to magnet stand for 10-30 seconds
- 13. Remove as much supernatant as possible
- 14. Transfer tube to non-magnetic rack and add 500 μl of FCW2 Buffer
- 15. Resuspend beads by vortexing for 15 seconds. Spin tube briefly and transfer back to magnet stand
- 16. Allow beads to attach to magnet stand for 10-30 seconds
- 17. Remove as much supernatant as possible
- 18. Tap magnet stand on bench 5 times and remove residual supernatant as much as possible
- 19. Allow beads to dry for 3 minutes

### **Elution Step**

- 20. Transfer tube to non-magnetic rack and add 50  $\mu$ l of FCE Buffer Important:: A minimum of 50  $\mu$ l of FCE Buffer is recommended to elute DNA to ensure optimal yields
- 21. Vortex vigorously for 10-15 seconds to loosen bead clumps
- 22. Incubate at 50°C for 10 minutes
- 23. Vortex vigorously for 10-15 seconds to loosen any remaining bead clumps
- 24. Spin tube briefly
- 25. Place tube on magnetic rack for 3 minutes
- 26. Transfer eluate\* into a new 1.5 ml Eppendorf tube

<sup>\*</sup> **Note:** To maximize the utilization of the final product in downstream applications, the Elution Buffer does not contain EDTA. If long-term storage of the DNA is desired, we recommend adding EDTA to a final volume of 1 mM (e.g. add 1 ul of 100 mM EDTA pH 8.0 stock solution for every 100 ul of elution volume).



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### **AnaPrep 48 Protocol**

The xTractPure™ Cell and Tissue Nucleic Acid Extraction Kit can be used to isolate DNA from up to 96 cell pellet and/or tissue samples using the AnaPrep 48 instrument with 96-well Deep Well Plates and 8-strip Tip Combs. Each Deep Well Plate can be used to extract DNA from up to 16 samples, allowing for higher throughput processing with minimal usage of consumable plastics. The instrument can be loaded with 3 Deep Well plates, thus allowing for simultaneous processing of up to 48 samples. This guide describes the use of the kit with the AnaPrep 48 to process cell and tissue samples.

### **Additional Materials Required**

RNase A (optional)

1 package of Deep Well Plates and Tip Combs (Catalog No. Z2212099)

#### **Prior to Initial Use**

BFC, FCW1, and FCW2 Buffers are shipped as a concentrate. If precipitate is present in any of the solutions, incubate solution at 37°C for 30 minutes with occasional shaking. Add the following user-supplied reagents to each bottle:

BFC Buffer

Add 26.4 ml of isopropanol and mix by inverting gently

FCW1 Buffer

Add 25.5 ml of 100% ethanol (200 proof) and mix by inverting gently

FCW2 Buffer

Add 88 ml of 100% ethanol (200 proof) and mix by inverting gently

### **Sample Preparation (Cell Pellet)**

- 1. Take up to  $1 \times 10^6$  fresh or frozen cells in a 1.5 ml Eppendorf tube and ensure media has been removed as much as possible
- 2. Add 400 µl of CTFC Buffer to the tube containing the cells
- 3. Add 10 µl of Proteinase K
- 4. Briefly vortex to mix, pipet up and down 10 times, and incubate at 56°C for 10 minutes
- 5. Vortex briefly to mix and incubate at 56°C for an additional 10 minutes
  - During this incubation period, set up 96-well Deep Well plates with appropriate reagents as shown in Plate Set Up section
- 6. Spin tube briefly on a tabletop centrifuge to bring down solution from the lid of the tube
- 7. (Optional) Add 10  $\mu$ l of RNase A (20 mg/ml), pipet up and down briefly to mix, and let incubate at room temperature for 5 minutes
- 8. Spin tube briefly to pellet any residual cell debris. Transfer resulting supernatant ( $\sim$ 400  $\mu$ l) to the **Binding** position of the 96-well Deep Well plate

### Sample Preparation (Tissue)

- 1. For high-yield tissues (e.g. spleen), take up to 10 mg of tissue. For lower-yield tissues (e.g. liver), take up to 20 mg of tissue. For maximum yield, mince or grind the tissue into several small pieces
- 2. In a 1.5 ml Eppendorf tube, add 400 μl of CTFC Buffer. Then add 10 μl of Proteinase K and pipet up and down briefly to mix
- 3. Transfer the fresh or frozen tissue to the above tube
- **4.** Place tube into either a thermal mixer or water bath and incubate at 56°C for 1-4 hours until all of the tissue pieces are dissolved

**If using a thermal mixer:** shake continuously at 1,400 rpm during incubation **If using a water bath:** vortex the samples briefly every 10-15 minutes

- A few minutes before the incubation period is complete, set up 96-well Deep Well plates with appropriate reagents as shown in Plate Set Up
- 5. Spin tube briefly on a tabletop centrifuge to bring down solution from the lid of the tube
- **6.** (Optional) Add 10  $\mu$ l of RNase A (20 mg/ml), pipet up and down briefly to mix, and let incubate at room temperature for 5 minutes
- 7. Spin tube briefly to pellet any residual tissue debris. Transfer resulting supernatant ( $\sim$ 400  $\mu$ l) to the **Binding** position of the 96-well deep-well plate

### Plate Set up

Set up 96-well Deep Well Plates by adding appropriate reagents according to table below.

Each plate can accommodate 16 samples. If processing 8 or less samples on a plate, ensure that the reagents are added only to columns 1-6 or 7-12 on the plate. In other words, if FCW1 is added to column 2, then FCW2 buffers should be added to columns 3 and 4, not 9 and 10.

Step	Column Position on Plate	Reagent	Volume per well		
		Cell or Tissue Sample	400 μΙ		
Binding	1, 7	BFC Buffer	300 μΙ		
		Magnetic Bead Solution	10 μΙ		
Wash 1	2, 8	FCW1 Buffer	500 μΙ		
Wash 2	3, 9	FCW2 Buffer	500 μl		
Wash 3	4, 10	FCW2 Buffer	500 μΙ		
Elution	6, 12	FCE Buffer	50 μl*		

<sup>\*</sup> **Important**: A minimum of 50  $\mu$ l must be present within the well of the plate in order for the elution step to take place.

### Instrument Set up

- Place 96-well Deep Well plate(s) into the instrument, ensuring that the heat strips are aligned with columns 1, 6, 7, and 12 of the plate
- Load 8-strip tip comb(s) into holder inside the instrument
- Touch the "CellTissueDNA" button and then touch Continue to begin the run
- After the run is complete, carefully remove plate(s) from the instrument
- Transfer the eluates (columns 6 and 12) from the plate to fresh tubes
- Isolated DNA is ready for immediate use

<sup>\*</sup> **Note:** To maximize the utilization of the final product in downstream applications, the Elution Buffer does not contain EDTA. If long-term storage of the DNA is desired, we recommend adding EDTA to a final volume of 1 mM (e.g. add 1 ul of 100 mM EDTA pH 8.0 stock solution for every 100 ul of elution volume).



## **Kit Components**

Item	Cat#	Amount	Storage		
1. CTFC Buffer	Z2212005-1	40 ml	Room Temp		
2. BFC Buffer	Z2212005-2	6 ml	Room Temp		
3. Proteinase K	Z2212005-3	1 ml	-20°C		
4. Magnetic Bead Solution	Z2212005-4	1 ml	Room Temp		
5. FCW1 Buffer	Z2212005-5	25 ml	Room Temp		
6. FCW2 Buffer	Z2212005-6	20 ml	Room Temp		
7. FCE Buffer	Z2212005-7	15 ml	Room Temp		

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## Appendix: Cell and Tissue Nucleic Acid Program Settings for AnaPrep 48

The following shows the AnaPrep 48 program settings for the xTractPure™ Cell and Tissue Nucleic Acid Extraction Kit.

Information of this step			Temperature Control		Mix		Waiting time		Magnetic binding			
					Temp			Time		Time	Precipitate	Repeat
Step	Well	Volume	Mode	Setup	(°C)	Wait	Speed	(min)	Mode	(min)	(sec)	(times)
Bind	1	710	IV	Stop	0	N	Fast	10	0	0	20	1
Wash 1	2	500	IV	Stop	0	N	Fast	2	0	0	20	1
Wash 2	3	500	IV	Stop	0	N	Fast	2	0	0	20	1
Wash 3	4	500	IV	Stop	0	N	Fast	2	0	0	20	1
Elute	6	50	IV	Elute	50	N	Fast	10	0	3	30	3
End	1	50	End	Stop	0	N	Stop	0	0	0	1	1