

Epoch Life Science, Inc.

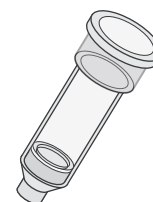
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GenCatch™ Total RNA Miniprep Kit



User's Guide for
Small Scale Total RNA
Extraction from Cell, Tissue and
Bacteria etc.

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1 Quick Start Procedure

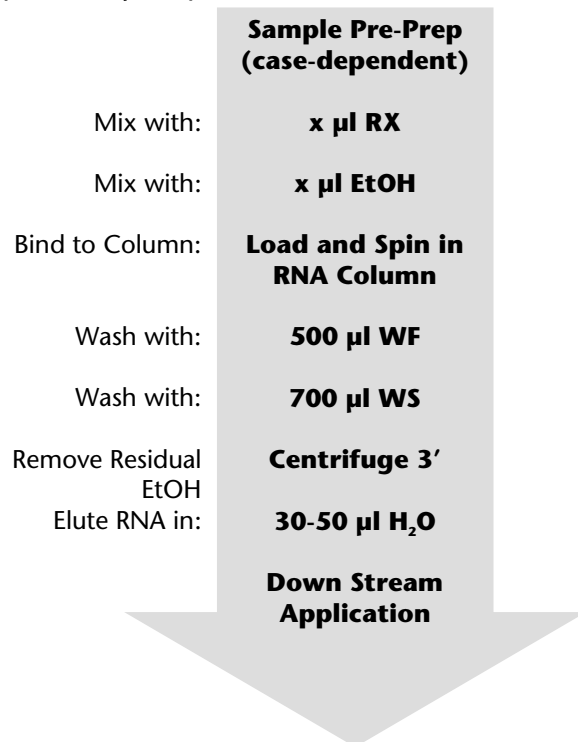
For Experienced Users Only.

The following flow chart is only good as a quick reminder for key steps and buffers to be used. Procedure varies depending on different type of samples. First time users are strongly recommended to read through the detailed instruction protocol in section 4.

Before you start:

Add 60 ml (50 preps) or 180 ml (250 preps) 98-100% ethanol to WS Buffer.

Pipet a required volume of RX Buffer into another tube and add 10 μ l β -mercaptoethanol (β -ME) per 1 ml RX Buffer before use.



5 Troubleshooting Guide

The following guide addresses some of the most common problems. A database of user raised questions and answers are being build at support.epochbiolabs.com.

Little or no RNA eluted:

a. Insufficient disruption or homogenization

Reduce the amount of starting sample and perform more disruption and homogenization.

b. Clogged Total RNA column

Reduce the amount of starting sample and perform more disruption and homogenization. Centrifuge the lysate to remove insoluble materials and use the supernatant only.

c. RNA is degraded

Starting sample should be fresh or frozen in liquid nitrogen and store at -80°C . Improper handling of the sample or storing the sample at -20°C will cause RNA degradation.

d. RNase contamination

Use RNase-free liquid, handling tips and tubes.

DNA contamination:

Refer to Protocol for "Removal of genomic DNA in eluted total RNA by DNase"

A260/A280 ratio of eluted total RNA is low:

a. Use ddH₂O of acidic pH to dilute RNA sample for spectrophotometric analysis

Use 10 mM Tris-HCl of pH 7.5 or TE buffer to dilute RNA sample.

b. DNA is copurified with RNA

Refer to Protocol for "Removal of genomic DNA in eluted total RNA by DNase".

Tube and centrifuge the tube for 1-2 minutes at full speed (10,000 x g or 13,000-14,000 rpm) in a microcentrifuge. When collecting homogenized lysate from the Collection Tube, avoid pipetting any debris and pellet formed at the bottom of the tube.

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Overview

GenCatch™ Total RNA Extraction Miniprep System provides an economical method to purify total RNA from various samples such as cultured cells, tissues, and bacteria. A simple silica-membrane spin-column method can isolate total RNA without need of performing time-consuming phenol/ chloroform extraction and ethanol precipitation. Total RNA longer than 200 nucleotides are isolated, while small RNA such as 5.8S RNA, 5S RNA, and tRNA, which make up 15-20% of the total RNA, are excluded.

Binding Capacity and Typical Yield:

The binding capacity of the **GenCatch™** total RNA mini column is 100 µg of total RNA, in order to avoid exceeding the binding capacity, use the sample preparation guide listed below:

Sample	Recommended amount of sample	Yield (µg)	
Animal cells	NIH-3T3	1 x 10 ⁶ cells	12
	HeLa	1 x 10 ⁶ cells	15
	COS-7	1 x 10 ⁶ cells	30
	LMH	1 x 10 ⁶ cells	12

Animal tissues	Mouse/rat tissues	Yield (µg)	
	Embryo	10 mg	30
	Heart	10 mg	10
	Brain	10 mg	10
	Kidney	10 mg	35
	Liver	10 mg	45
	Spleen	10 mg	35
	Lung	10 mg	10
	Thymus	10 mg	45

Bacteria	Yield (µg)	
E. coli	1 x 10 ⁹ cells	65
B. subtilis	1 x 10 ⁹ cells	40

Downstream Applications:

- Northern-blotting
- RT-PCR
- PolyA+ RNA selection
- cDNA synthesis
- Primer Extension
- *In vitro* translation

Gram-positive bacteria, and incubate at room temperature for 10 minutes.

4. Add 350 µl RX Buffer (β-ME added) to the sample and mix by vortexing.
Add 10 µl β-mercaptoethanol (β-ME) per 1 ml of Buffer RX.
5. Centrifuge lysate for 5 minutes to spin down insoluble materials and use only the supernatant in the following steps.
6. Add 250 µl of 98 % ethanol to the sample and mix by vortexing.
7. Follow the Animal Tissue Protocol starting from Step 4.

V. Removal of genomic DNA in eluted total RNA by DNase

1. Incubate total RNA with RNase-free DNase I (1 unit per µg of total RNA) in 50 mM Tris-HCl (pH 7.5), 10 mM MnCl₂, and 50 µg/ml BSA at 37°C for 15-30 minutes.
2. Remove DNase I by adding an equal volume of phenol:chloroform (1:1) and mix well. Centrifuge for 5 minutes. Transfer the upper aqueous layer to a new eppendorf tube.
3. Add 1/10 volume of 3 M sodium acetate (pH 5.2) and 1 volume of ice-cold isopropanol to the solution and mix well. Chill on ice for 30 minutes.
4. Centrifuge for 10 minutes at 4°C. Discard the supernatant. Wash the pellet twice with 1 ml of 70 % ethanol and recentrifuge.
5. Remove all supernatant. Air dry the RNA pellet. Redissolve RNA in RNase-free ddH₂O.

VI: Application of Shearing Tube

Shearing Tube is designed for simple and fast homogenization of tissue and cell lysate. The lysate is loaded into a Shearing Tube sitting in a 2-ml Collection

sample by using 20-G needle syringe or Epoch Biolabs's Shearing Tube.

Add 10 µl β-mercaptoethanol (β-ME) per 1 ml of RX Buffer. If using Epoch Biolabs's Shearing Tube, apply the disrupted lysate to a Shearing Tube and centrifuge for 1 minute to shear genomic DNA.

3. Follow the Animal Tissue Protocol starting from Step 2.

III. Animal Cell Cytoplasm Protocol:

1. Prepare cytoplasm lysate.

Prepare cell lysis buffer: (provide by user) 20 mM Tris-HCl pH 8.0, 1 mM MgCl₂, 0.5% NP-40. Keep at 4°C.

Only fresh cells are used for preparing cytoplasm lysate.

a. Harvest 5 x 10⁶ - 1 x 10⁷ cells and centrifuge at 300 x g to pellet cells.

b. Add 180 µl of cell lysis buffer to the cell pellet, resuspend and lysis cells by gentle pipetting. Incubate the lysate on ice for 5 minutes.

c. Centrifuge the lysate at 300 x g at 4°C for 3 minutes, transfer the supernatant to a new tube, and use the supernatant (lysate) in the following steps.

2. Add 600 µl of RX Buffer (β-ME added) to the lysate and mix by vortexing.

Add 10 µl β-mercaptoethanol (β-ME) per 1 ml of RX Buffer.

3. Add 450 µl of 98% ethanol to the sample and mix by vortexing.

4. Follow the Animal Tissue Protocol starting from Step 4.

IV. Bacteria Protocol:

1. Pellet up to 1 x 10⁹ bacterial cells by centrifuging at 5,000 x g (7,500 rpm) for 5 minutes. Remove all the supernatant.

2. Resuspend cells in 100 µl of TE buffer by vortexing.

3. Add lysozyme (provide by user) to a final concentration of 500 µg/ml for Gram-negative bacteria; 2 mg/ml for

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Product Contents

GenCatch™ Total RNA Extraction Miniprep Kit contains sufficient reagents for 50 (Cat. No. 1660050) and 250 (Cat. No. 1660250) total RNA extraction applications respectively.

Catalog Number	1660050	1660250
RX Buffer	36 ml	150 ml
WF Buffer	30 ml	150 ml
WS Buffer	15 ml	45 ml
RNase-free ddH ₂ O	3 ml	25 ml
GenCatch™ Column	50 pieces	250 pieces
Collection Tube	50 pieces	250 pieces
Elution Tube	50 pieces	250 pieces
Protocol	1	1

Storage Conditions:

Store at room temperature

All components are guaranteed for 48 months from the date of purchase, when stored under specified conditions and used as described in this manual.

4 Protocol

First time users are strongly recommended to read through this detailed protocol instruction.

For technical support and user raised common questions and answers please visit: support.epochbiolabs.com

Before you start:

Add 60 ml (50 preps) or 180 ml (250 preps) 98-100% ethanol to Buffer WS before use (see instructions on bottle label).

Please read the following notes before starting the procedures.

- Pipet a required volume of RX Buffer into another tube and add 10 μ l β -mercaptoethanol (β -ME) per 1 ml RX Buffer before use.
- Complete disruption and homogenization of sample is essential for total RNA extraction.
- All plasticware and containers should be treated properly to make sure RNase-free. Gloves should be worn when handling RNA.
- Buffers provided in this system contain irritants. Appropriate safety apparels such as gloves and lab coat should be worn.
- All centrifugation steps except cell pelleting are done at full speed (10,000 x g or 13,000-14,000 rpm) in a microcentrifuge.
- Some genomic DNA (and plasmid DNA, if any) will also be copurified with RNA. DNase treatment is therefore required when DNA-free RNA is desired. DNase can then be removed by phenol/chloroform extraction (refer to Protocol for "Removal of genomic DNA in eluted total RNA by DNase").

I. Animal Tissue Protocol:

1. Add 350 μ l RX Buffer (β -ME added) to 10 mg of liquid-nitrogen-frozen or fresh tissue. Disrupt and homogenize the sample by grinding and shearing using 20-G needle syringe or Epoch Biolabs' Shearing Tube.
Add 10 μ l β -mercaptoethanol (β -ME) per 1 ml of Buffer RX. If using 20 mg of tissue add 700 μ l of Buffer RX. If using Epoch Biolabs's Shearing Tube, refer to "Application of Shearing Tube" section on the last page.
2. Centrifuge the lysate for 5 minutes to spin down insoluble materials and use only the supernatant in the following steps.

3. Determine the final volume of the supernatant. Add an equal volume of 70% ethanol to the clear lysate and mix by vortexing.
If lysate is lost during the preparation, reduce the volume of ethanol accordingly. Do not centrifuge the ethanol added lysate.
4. Place a Total RNA Mini Column onto a Collection Tube. Add 700 μ l of the ethanol-added sample (including any precipitate) into the column. Centrifuge for 30-60 seconds. Discard the flow-through.
Repeat this step for the rest of the sample. If some sample still retains in the column, repeat centrifugation until all sample pass the column.
5. Wash the column once with 0.5 ml WF Buffer by centrifuging for 30-60 seconds. Discard the flow-through.
6. Wash the column once with 0.7 ml WS Buffer by centrifuging for 30-60 seconds. Discard the flow-through.
Add 60 ml (50 preps) or 180 ml (250 preps) of 98-100% ethanol into WS Buffer bottle when first open.
7. Centrifuge the column for another 3 minutes to remove ethanol residue.
8. Place the column onto a 1.5-ml RNase-free Elution Tube. Add 30-50 μ l RNase-free ddH₂O (provided) onto the center of the membrane.
For effective elution, make sure that the elution solution is dispensed onto the center of the membrane.
9. Stand the column for 1 minute, and centrifuge for 1-2 minutes to elute total RNA.
10. Store RNA at -70 °C.

II. Animal Cells Protocol:

1. Pellet 1 to 5 x 10⁶ cells by centrifuging at 300 x g for 5 minutes. Remove all the supernatant.
2. Disrupt cells by adding 350 μ l RX Buffer (β -ME added) to the cell pellet and vortex the sample. Homogenize the