

GenCatch™ Total RNA Miniprep Kit

User's Guide for **Total RNA Extraction From Cell, Tissue and Bacteria etc.**

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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:

WARNING: strong acids and oxidants (bleach, for example) should not be used together with RX buffer (because this kind of reaction would produce cyanide)!

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 μl β -mercaptoethanol (β -ME) per 1 ml RX Buffer before use.

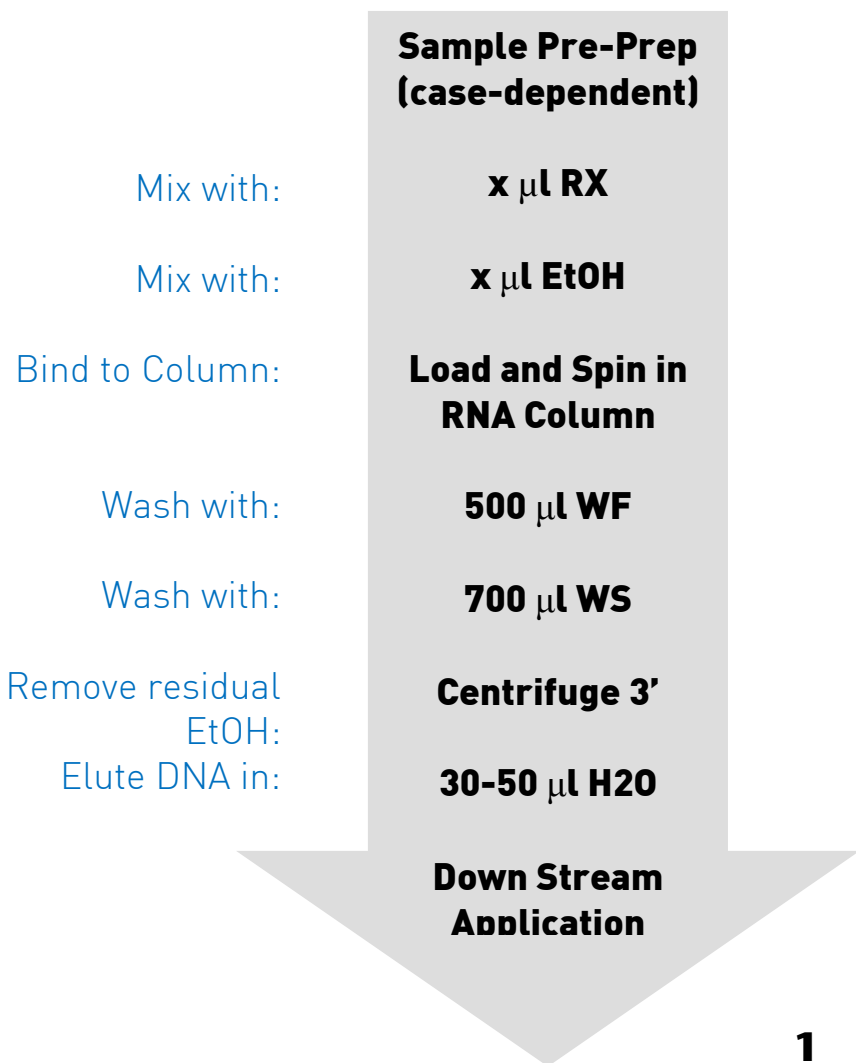




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Overview

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		
	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
Bacteria	Thymus	10 mg	45
	E. coli	1 x 10 ⁹ cells	45
	B. subtilis	1 x 10 ⁹ cells	40

Downstream Applications:

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



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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
	50 preps	250 preps
RX Buffer	36 ml	150 ml
WF Buffer	30 ml	150 ml
WS Buffer	15 ml	45 ml
RNase-free ddH2O	3 ml	25 ml
GenCatch™ Column	50 pieces	250 pieces
Collection Tube	50 pieces	250 pieces
Elution Tube	50 pieces	250 pieces

WARNING, strong acids and oxidants (bleach, for example) should not be used together with RX buffer (because this kind of reaction would produce cyanide)!

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

Storage Conditions:
Store at room temperature

All components are guaranteed for 24 months from the date of purchase, when stored under specified conditions and used as described in this manual. Long term storage of some of the buffer may harden the HDPE plastic bottle. However this will not adversely affect the performance of the kit. GenCatch™ columns should be kept sealed in the zip lock bag provided during storage and away from any heating source.



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Protocol

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

For technical support please reach us at support@epochlifescience.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 plastic ware and containers should be treated properly to make sure RNase-free. Wear gloves when handling RNA.

-Buffers provided in this system contain irritants. Wear appropriate safety apparels such as gloves and lab coat.

-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 Life Science's Shearing Tube.
2. 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 Life Science's Shearing Tube, refer to "Application of Shearing Tube" section on the last page.



3. Centrifuge the lysate for 5 minutes to spin down insoluble materials and use only the supernatant in the following steps.
4. 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.
5. 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.
6. Wash the column once with 0.5 ml WF Buffer by centrifuging for 30-60 seconds. Discard the flow through.
7. 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.
8. Centrifuge the column for another 3 minutes to remove ethanol residue.
9. 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.
10. Stand the column for 1 minute, and centrifuge for 1-2 minutes to elute total RNA.
11. 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 sample by using 20-G needle syringe or Epoch's Shearing Tube (not included, purchase separately Cat. No.: 2610-050).
Add 10 μ l β -mercaptoethanol (β -ME) per 1 ml of RX Buffer. If using Epoch'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 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 300x g to pellet cells.
 - b. Add 180 μ l 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 μ l of RX Buffer (β -ME added) to the lysate and mix by vortexing.
Add 10 μ l β -mercaptoethanol (β -ME) per 1 ml of RX Buffer.
3. Add 450 μ l 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 1×10^9 bacterial cells by centrifuging at $5,000 \times g$ (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 $\mu\text{g}/\text{ml}$ for Gram-negative bacteria; 2 mg/ml for 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 the 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 MgCl_2 , and 50 $\mu\text{g}/\text{ml}$ BSA at 37 $^\circ\text{C}$ 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 $^\circ\text{C}$. 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. The of Shearing Tubes:

Shearing Tube (not included, purchase separately, Cat. No.: 2160-050) 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 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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Trouble Shooting

The following guide addresses some of the most common problems.

For more specific questions please reach us at

support@epochlifescience.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 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^{\circ}\text{C}\$. Improper handling of the sample or storing the sample at \$-20^{\circ}\text{C}\$ will cause RNA degradation.](#)
- d. RNase contamination:
[Use RNase-free solution, pipette tips and centrifuge tubes.](#)

DNA contamination:

Refer to Protocol section V: "Removal of genomic DNA in eluted total RNA by DNase"

A260/A280 ratio of eluted 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 section V: "Removal of genomic DNA in eluted total RNA by DNase"](#)