

## HiYield™ Total RNA Maxi Kit (Tissue)

Cat. No.:	YRTM10	YRTM25
<b>Product Name:</b>	HiYield™ Total RNA Maxi Kit (Tissue)	
<b>Reactions:</b>	10	25
<b>Sample:</b>	Up to 200 mg of tissue or paraffin-embedded tissue	
<b>Yield:</b>	Up to 300 ug	
<b>Format:</b>	Spin Column	
<b>Operation:</b>	Centrifuge	
<b>Operation Time:</b>	Within 60 Minutes	

### Introduction

HiYield™ Total RNA Maxi Kit (Tissue) is designed specifically for purifying total RNA from a variety of animal tissue and paraffin-embedded tissue. Provided micropestles can efficiently homogenize tissue samples to shorten the time in the Lysis Step. The entire procedure can be completed within 1 hour without phenol/chloroform extraction or alcohol precipitation. The purified total RNA is ready for use in RT-PCR, Northern Blotting, Primer Extension and cDNA Library Construction.

### Features

Ready-to-use RNA for many downstream applications.

Excellent Recovery of RNA.

Complete removal of contaminants and inhibitors.

### Components

ITEM	YRTM10	YRTM25
RB Buffer	60ml	130ml
W1 Buffer	50ml	130ml
Wash Buffer*	25ml	62.5ml
RNase-Free Water	6ml	30ml
RTM Column	10pcs	25pcs
Lysate Filter Column	10pcs	25pcs

\*Add absolute ethanol to the Wash Buffer prior to initial use (see the bottle label for volume).

### **Applications**

Purified RNA is ready for direct use in RT-PCR, Real-Time RT-PCR, Northern Blotting, Primer Extension, RNase Protection Assays, mRNA Selection, cDNA Synthesis.

### **Quality Control**

The quality of HiYield™ Total RNA Maxi Kit (Tissue) is tested on a lot-to-lot basis. The kits are tested by isolation of total RNA from 100 mg of mouse liver. More than 100 ug of total RNA was quantified with a spectrophotometer and checked by formaldehyde agarose gel analysis. Finally, RT-PCR was used to ensure the quality of total RNA.

### **Caution**

RB Buffer contains chaotropic salt which is a harmful irritant. During operation, always wear a lab coat, disposable gloves, and protective goggles.

### **References**

(1) Vogelstein, B., and Gillespie, D. (1979) Proc. Natl. Acad. Sci. USA 76, 615.

## **Protocol**

### **Things to do before starting**

Add absolute ethanol to the Wash Buffer prior to initial use (see the bottle label for volume).

Additional requirements: absolute ethanol, centrifuge tubes (RNase-free),  $\beta$ -mercaptoethanol, DNase I (2 KU/ml) mixed in a reaction buffer {50 mM Tris-HCl (pH 7.5), 10 mM  $MnCl_2$ , 50  $\mu$ g/ml BSA at 25°C}

### **Optional Steps of DNA Residue Degradation:**

#### **Optional Step 1:**

Add 100  $\mu$ l of DNase I (2 KU/ml) mixed in a reaction buffer {50 mM Tris-HCl (pH 7.5), 10 mM  $MnCl_2$ , 50  $\mu$ g/ml BSA at 25°C} to the center of the RTM Column matrix. Let stand for 10 minutes at room temperature and then proceed to Step 4 Wash.

#### **Optional Step 2:**

Add 2  $\mu$ l of DNase I (2 KU/ml) mixed in a reaction buffer {50 mM Tris-HCl (pH 7.5), 10 mM  $MnCl_2$ , 50  $\mu$ g/ml BSA at 25°C} to the final elution sample. Let stand for 10 minutes at room temperature.

<p><b>Step 1 Tissue Dissociation</b></p>	<ul style="list-style-type: none"> <li>★Cut off up to 200 mg of fresh or frozen animal tissue. (If using frozen animal tissue, the sample <b>MUST</b> have been flash frozen in liquid nitrogen and immediately stored at -70°C until use in order to avoid RNA Degradation).</li> <li>★Grind the sample (in liquid nitrogen) to a fine powder. Transfer the powder to a 15 ml centrifuge tube.</li> </ul>
<p><b>Step 2 Lysis</b></p>	<ul style="list-style-type: none"> <li>★Add <b>5 ml of RB Buffer</b> and 50 µl of β-mercaptoethanol to the ground sample and mix by vortex. Incubate at room temperature for 5 minutes.</li> <li>★Place a <b>Lysate Filter Column</b> in a 50 ml centrifuge tube and transfer the sample lysate to the column. Centrifuge for 5 minutes at full speed.</li> <li>★Discard the <b>Lysate Filter Column</b> and Proceed to Step 3 RNA Binding.</li> </ul>
<p><b>Step 3 RNA Binding</b></p>	<ul style="list-style-type: none"> <li>★Add a half volume of absolute ethanol to the clarified filtrate from Step 2 and vortex immediately (ex: add 2.5 ml of absolute ethanol to 5 ml of filtrate).</li> <li>★Place a <b>RTM Column</b> in a 50 ml centrifuge tube.</li> <li>★Transfer the ethanol-added mixture to the <b>RTM Column</b>.</li> <li>★Centrifuge at full speed for 5 minutes and discard the flow-through (<b>if the mixture could not flow through the RTM Column membrane following centrifugation, increase the centrifuge time until the mixture passes completely</b>).</li> </ul> <p style="color: blue;"><b>Optional Step 1 of DNA Residue Degradation</b> (see optional steps of DNA Residue Degradation in page 2)</p>
<p><b>Step 4 Wash</b></p>	<ul style="list-style-type: none"> <li>★Add <b>4 ml of W1 Buffer</b> to the center of the <b>RTM Column</b>.</li> <li>★Centrifuge at full speed for 3 minutes. Discard the flow-through and place the <b>RTM Column</b> back in the 50 ml centrifuge tube.</li> <li>★Add <b>6 ml of Wash Buffer</b> (ethanol added) to the center of the <b>RTM Column</b>. Centrifuge at full speed for 3 minutes.</li> <li>★Discard the flow-through and place the <b>RTM Column</b> back in the 50 ml centrifuge tube.</li> <li>★Add <b>6 ml of Wash Buffer</b> (ethanol added) to the center of the <b>RTM Column</b>. Centrifuge at full speed for 3 minutes.</li> <li>★Discard the flow-through and place the <b>RTM Column</b> back in the 50 ml centrifuge tube. Centrifuge at full speed for 10 minutes to dry the column matrix.</li> </ul>

<b>Step 5 RNA Elution</b>	<ul style="list-style-type: none"> <li>★Place the dried <b>RTM Column</b> in a clean 50 ml centrifuge tube (RNase-free).</li> <li>★Add <b>500 µl of RNase-free water</b> into the center of the column matrix.</li> <li>★Let stand for 5 minutes or until the water has been absorbed by the matrix.</li> <li>★Centrifuge at full speed for 5 minutes to elute the purified RNA.</li> </ul>
	<p><b>Optional Step 2 of DNA Residue Degradation</b> (see optional steps of DNA Residue Degradation in page 2)</p>

### Troubleshooting

Problem	Possible Reasons/Solution
<b>Clogged RTM Column</b>	<ul style="list-style-type: none"> <li>★Inefficient disruption and/or homogenization</li> <li>★Too much starting material</li> <li>★Centrifugation temperature was too low (should be 20-25°C)</li> </ul>
<b>Low RNA Yield</b>	<ul style="list-style-type: none"> <li>★Insufficient disruption and homogenization</li> <li>★Too much starting material</li> <li>★RNA still bound to RTM Column membrane</li> <li>★Ethanol carryover</li> </ul>
<b>RNA Degradation</b>	<ul style="list-style-type: none"> <li>★Harvested sample not immediately stabilized</li> <li>★Inappropriate handling of starting material</li> <li>★RNase contamination</li> </ul>