

# HiPure Soil RNA Kit

### Introduction

The HiPure SoilRNA Kit is suitable for extracting high purity total RNA from Soil or other environmental samples The kit combines the SDS/Phenol and silica gel column purification technology to complete the extraction of high purity total RNA in only 40 minutes. The obtained RNA can be directly used for RT-PCR, Northern blot, Poly-A + purification, nucleic acid protection and in vitro translation experiments.

#### Kit Contents

Product Number	R418301	R418302	R418303
Purification Times	10 preps	50 preps	250 preps
HiPure RNA Mini Columns I	10	50	250
gDNA Filter Column	10	50	250
2ml Collection Tubes	20	100	500
2ml Beads Tubes	10	50	250
Buffer SOL	6 ml	30 ml	150 ml
Buffer SDS	1 ml	4 ml	15 ml
Buffer PHC	6 ml	30 ml	150 ml
Buffer GDP	10 ml	40 ml	150 ml
Buffer RVV1	10 ml	50 ml	200 ml
Buffer RVV2 *	5 ml	20 ml	2 x 50 ml
RNase Free Water	1.8 ml	10 ml	30 ml

## Storage and Stability

Buffer PHC should be stored at 2–8°C upon arrival. However, short-term storage (up to 12 weeks) at room temperature (15–25°C) does not affect their performance. The remaining kit components can be stored dry at room temperature (15–25°C) and are stable for at least 18 months under these conditions. The entire kit can be stored at 2–8°C, but in this case buffers should be redissolved before use. Make sure that all buffers are at room temperature when used.

### Materials and Equipment to be Supplied by User

- absolute ethyl alcohol(96-100%)
- chloroform
- Buffer RW2 was diluted with anhydrous ethanol and stored at room temperature

# [Protocol]

- Transfer 500mg soil, 50-100mg Stool or 200-500mg other environmental samples to 2ml Bead Tubes.
- 2. Add 500µL Buffer SOL, 50µL Buffer SDS and 500µL Buffer PHC to the sample. Lyse sample by vortex at maximum speed for 10 minutes or by Fastpreps 24 ( 6.5 m/s twice for 45s).

For best results, a mixer mill, such as GenoGrinder 2010, Fastprep-24®, or Omni Bead Ruptor should be used.

- 3. Centrifuge for 5 seconds to remove drops of liquid from the lid.
- 4. Add 200µL chloroform and vortex to mix thoroughly. Incubate for 3 minutes.
- 5. Centrifuge at maximum speed ( $\geq$ 13,000 x g) for 10 min at 4°C.
- 6. Transfer the cleared supernatant (~400µL) to a new 2.0 mL microcentrifuge tube.

- 7. Add an equal volume Buffer GDP and mix by inverting the tube 4-6 times.
- 8. Insert a gDNA Filter Column into a 2.0mL Collection Tube (provided).
- Transfer the Mixture from step 7 to a the DNA column placed in a 2 ml collection tube (supplied). Centrifuge for 60 s at ≥12000 x g . Discard the column, and save the flow-through.
- 10. Add 0.5 volume of absolute ethanol to the flow-through, and mix well by pipetting.
- 11. Insert a HiPure RNA Mini Column in a 2ml Collection Tube.
- Add up to 700µl of the sample from Step 10 to the Column. Centrifuge at 12,000 × g for 1 minute at room temperature. Discard the filtrate and reuse collection tube.
- 13. Repeat Step 12 until all of the sample has been transferred to the column.
- Add 700µl Buffer RW1 to the column, centrifuge at 12,000 × g for 1 minute at room temperature. Discard the filtrate and reuse collection tube.
- Add 500µl Buffer RW2 to the column, Centrifuge at 12,000 × g for 1 minute at room temperature. Discard the filtrate and reuse collection tube.
- Add 500µl Buffer RW2 to the column, Centrifuge at 12,000 × g for 1 minute at room temperature. Discard the filtrate and reuse collection tube.
- 17. Centrifuge the empty Column at 12,000  $\times$  g for 2 minute at room temperature to dry the column matrix.
- Transfer the Column to a clean 1.5ml microcentrifuge tube. Add 25~50µl RNase Free Water directly to the center of the column membrane. Let sit at room temperature for 2 minutes.
- 19. Centrifuge at 12,000  $\times$  g for 1 minute at room temperature. Store RNA at -20°C.

If the expected RNA yield is >30  $\mu$ g, repeat step 10 using another 30–50 $\mu$ l RNase-free water, or using the eluate from step 12-13.

### Troubleshooting Guide

#### 1. Clogged HiPure RNA Column

- Too much starting material: In subsequent preparations, reduce the amount of starting material. It is essential to use the correct amount of starting material.
- Inefficient disruption and/or homogenization: Disrupting and homogenizing starting materia as RNeasy Mini Kit pages 18-21.
- 2. RNA does not perform well (e.g. in RT-PCR
- Salt concentration in eluate too high: Modify the wash step by incubating the column for 5 min at room temperature after adding 500ul of Buffer RW2, then centriufge.
- Eluate contains residual ethanol: Ensure that the wash flow-through is drained from the collection tube and that the column is then centrifuged at >12,000 x g for 1 min.
- 3. DNA contamination in downstream experiments
- No DNase treatment: Perform optional on column DNase digestion using RNase-Free DNase Ste at the point individual protocols.