Animal tissue (fresh and frozen), Bacteria (gram negative, gram positive)

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## **Recommended Starting Material**

To reach optimized results it is better to follow as listed here. The size of recommended material is written in Table.

## Table: Appropriate size of starting material

Sample	Size of Starting Material (min-max)
Liver	1-25 mg
Brain	1-25 mg
Kidney	1-25 mg
Spleen	1-10 mg
Ear	1-25 mg
Muscle	1-25 mg
Skin	1-25 mg
Heart	1-25 mg
Lung	1-25 mg
Mouse tail	1-25 mg
Rat tail	1-25 mg
Bacteria cells	2.5 x 10 <sup>8</sup> to 2 x 10 <sup>9</sup>

## Protocol 1

## Isolation of Genomic DNA (based on silica technology)

Sample type: Animal tissues (fresh and frozen)

## Some tips to know

- All centrifugation steps are carried out at room temperature (15–25°C) in a microcentrifuge.
- If RNase treatment is desired, Prime-RNase A can be ordered separately from ROJETechnologies, Cat NO. EB983013 and EB983014.
- If TL Buffer (TLB) forms precipitate, please warm it to 56°C until the precipitate has fully dissolved. This is due to storage condition and won't influence the efficiency of the buffer.
- If ORB forms precipitate, please warm it to 60°C until the precipitate has fully dissolved. ORB precipitation may occur more often, however this will not influence the efficiency of the buffer.
- Do not forget to add the appropriate amount of Ethanol (96–100%) to BW1 and BW2 buffers (BWB1 and BWB2) as indicated on the bottle, before using for the first time.
- For frozen samples, equilibrate them to room temperature. Avoid repeated freezing and thawing of samples which might lead to a decrease in DNA size.

### Process

1. Select one of the sample preparation methods here (a, b or c) then Place the prepared sample in a 1.5 ml microcentrifuge tube.

- a. Cut the tissue into small pieces. Then, Place the sample in to a clean microcentrifuge tube.
- b. Use Micropestle alternatively homogenizer to grind the tissue in TLB Buffer before addition of RJ-Protease.
- c. Ground the samples under liquid nitrogen (recommended for samples which are difficult to lyse).
- 2. Add 180 µl TLB and then add 25 µl RJ-Protease to the sample. Mix thoroughly by pulse vortexing for 30 s and then incubate at 56°C for 1-2 h until the tissue is completely lysed. Pulse vortex every 10 min during incubation to intersperse the sample, or place it in a thermomixer or shaking water bath.

**Optional:** This protocol is specialized for DNA isolation. For usual and routine applications RNase treatment isn't needed. If RNA-free genomic DNA is required, add 10  $\mu$ l Prime-RNase A (Cat No. EB983013), mix by vortexing, and incubate for 5 min at room temperature before going to step 3.

- 3. Add 200  $\mu I$  ORB to the mixture, mix by pulse vortexing for 10 s and then incubate at 70°C for 10 min.
- 4. Add 200 µl absolute Ethanol, then mix thoroughly by pulse vortexing for 15 s.
- 5. Pipette the mixture from step 4 to a HiPure DR column placed in a 2 ml collection tube (supplied in the kit box). Centrifuge at 13000 rpm for 1 min. Discard flow-through and place the HiPure DR column in the collection tube again.

**Note:** If the lysate did not pass the column, centrifuge at full speed until it passes through the column.

- Add 700 μl BWB1 and centrifuge for 1 min at 13000 rpm, just discard the flowthrough. Place the HiPure DR column in the previous collection tube and go to the next step.
- Add 600 µl BWB2 and centrifuge for 3 min at 14000 rpm. Discard both the flowthrough and the collection tube. Place the HiPure DR column in a new clean 1.5 ml microcentrifuge tube (not provided).

**Note:** To avoid Ethanol carry over, be careful that the column does not come into contact with the flow-through, if it happens discard the flow-through, place the column back in a collection tube and centrifuge for another 1 min at 14000 rpm.

 Pipette 50-200 μl RRB directly onto HiPure DR column. Incubate at room temperature for 1-5 min. Centrifuge it at 12000 rpm for 1 min.

**Note:** If the expected DNA yield is more than the yield from pervious step, put the HiPure DR column on a new microtube and add another 50-200  $\mu$ l RRB, incubate for 5 min at room temperature. Then, centrifuge for 1 min at 12000 rpm. However, it is possible to pass the follow-through from step 8 once more to obtain DNA with higher concentrations.

### Protocol 2

## Isolation of Genomic DNA (based on silica technology)

**Sample type:** Bacteria (gram negative)

## Some tips to know

- All centrifugation steps are carried out at room temperature (15–25°C) in a microcentrifuge.
- If RNase treatment is desired, Prime-RNase A can be ordered separately from ROJETechnologies, Cat No. EB983013 and EB983014.
- If TL Buffer (TLB) or OR Buffer (ORB) forms precipitate, please warm it to 56°C until the precipitate has fully dissolved. This is due to storage condition and won't influence the efficiency of the buffer.
- Do not forget to add the appropriate amount of Ethanol (96–100%) to BW1 and BW2 buffers (BWB1 and BWB2) as indicated on the bottle, before using for the first time.
- For frozen samples, equilibrate them to room temperature. Avoid repeated freezing and thawing of samples which might lead to a decrease in DNA size.

### Process

- 1. Calculate the bacteria cell number (refer to appendix 3, in main handbook). Collect the cell by centrifugation at 10000 rpm for 10 min. Discard the supernatant.
- 2. Add 180 µl TLB and then 25 µl RJ-Protease. Mix thoroughly by pulse vortexing for 30 s, then incubate at 56°C for 1-2 h until the lysate becomes clear. Pulse vortex every 10 min during incubation to intersperse the sample, or place it in a thermomixer or shaking water bath.

Animal tissue (fresh and frozen), Bacteria (gram negative, gram positive)

**Optional:** This protocol is specialized for DNA isolation. For usual and routine applications RNase treatment isn't needed. If RNA-free genomic DNA is required, add 10  $\mu$ l Prime-RNase A (Cat No. EB983013), mix by vortexing, and incubate for 5 min at room temperature before going to step 3.

- 3. Add 200 µl ORB, mix by pulse vortexing for 10 s, then incubate at 70°C for 10 min.
- 4. Add 200 µl absolute Ethanol, then mix thoroughly by pulse vortexing for 15 s.
- 5. Pipette the mixture from step 4 to a HiPure DR column placed in a 2 ml collection tube (supplied in the kit box). Centrifuge at 13000 rpm for 1 min. Discard flow-through and place the HiPure DR column in the collection tube again.

**Note:** If the lysate did not pass the column, centrifuge at full speed until it passes through the column.

- Add 700 µl BWB1 and centrifuge for 1 min at 13000 rpm, discard the flow-through.
  Place the HiPure DR column in the previous collection tube and go to the next step.
- Add 600 µl BWB2 and centrifuge for 3 min at 14000 rpm. Discard both the flowthrough and the collection tube. Place the HiPure DR column in a new clean 1.5 ml microcentrifuge tube (not provided).

**Note:** To avoid Ethanol carry over, be careful that the column does not come into contact with the flow-through, if it happens discard the flow-through, place the column back in a collection tube and centrifuge for another 1 min at 14000 rpm.

 Pipette 50-200 μl RRB directly into HiPure DR column. Incubate at room temperature for 1-5 min. Centrifuge it at 12000 rpm for 1 min.

Animal tissue (fresh and frozen), Bacteria (gram negative, gram positive)

**Note:** If the expected DNA yield is more than the yield from pervious step, put the HiPure DR column on a new microtube and add another 50-200 µl RRB, incubate for 5 min at room temperature. Then, centrifuge for 1 min at 12000 rpm. However, it is possible to pass the follow-through from step 8 once more to obtain DNA with higher concentrations.

### Protocol 3

## Isolation of Genomic DNA (based on silica technology)

**Sample type:** Bacteria (gram positive)

## Some tips to know

- All centrifugation steps are carried out at room temperature (15–25°C) in a microcentrifuge.
- For gram positive bacteria like *B. subtilis*, Lysozyme (Cat No. EB983017) should be ordered separately.
- Prepare the lysis buffer as follows:

20 mM Tris.Cl, pH 8.0

2 mM sodium EDTA

1.2% Triton® X-100

add lysozyme to 20 mg/ml (immediately before use).

- Preheat a heat block or water bath to 37 °C.
- If RNase treatment is desired, Prime-RNase A can be ordered separately from ROJETechnologies, Cat No. EB983013.
- If TL Buffer (TLB) or OR Buffer (ORB) forms precipitate, please warm them to 56°C until the precipitate has fully dissolved. This is due to storage condition and won't influence the efficiency of the buffer.
- Do not forget to add the appropriate amount of Ethanol (96–100%) to BW1 and BW2 buffers (BWB1 and BWB2) as indicated on the bottle, before using for the first time.

• For frozen samples, equilibrate them to room temperature. Avoid repeated freezing and thawing of samples which might lead to a decrease in DNA size.

#### Process

- 1. Calculate the bacteria cell number (refer to appendix 3, in main handbook). Collect the cell by centrifugation at 10000 rpm for 10 min. Discard the supernatant.
- 2. Resuspend the pellet, add 180  $\mu l$  enzymatic lysis buffer. Incubate 30-60 min at 37  $^{\circ}\text{C}.$
- 3. Add 200 µl ORB and then 25 µl RJ-Protease. Mix thoroughly by pulse vortexing for 30 s and then incubate at 56 °C for 30-60 min until the sample is completely lysed. Pulse vortex every 10 min during incubation to intersperse the sample, or place it in a thermomixer or shaking water bath.

**Optional:** This protocol is specialized for DNA isolation. For usual and routine applications RNase treatment isn't needed. If RNA-free genomic DNA is required, add 10  $\mu$ l Prime-RNase A (Cat No. EB983013), mix by vortexing, and incubate for 5 min at room temperature before going to step 4.

- 4. Centrifuge at 15000 rpm for 2 min. Pour supernatant to clean tube.
- 5. Add 200  $\mu$ l absolute Ethanol, then mix thoroughly by pulse vortexing for 15 s.
- 6. Pipette the mixture from step 5 to a HiPure DR column placed in a 2 ml collection tube (supplied in the kit box). Centrifuge at 13000 rpm for 1 min. Discard flow-through and place the HiPure DR column in the collection tube again.

**Note:** If the lysate did not pass the column, centrifuge at full speed until it passes through the column.

- Add 700 µl BWB1 and centrifuge for 1 min at 13000 rpm, just discard the flowthrough. Place the HiPure DR column in the previous collection tube and go to the next step.
- Add 600 µl BWB2 and centrifuge for 3 min at 14000 rpm. Discard both the flowthrough and the collection tube. Place the HiPure DR column in a new clean 1.5 ml microcentrifuge tube (not provided).

**Note:** To avoid Ethanol carry over, be careful that the column does not come into contact with the flow-through, if it happens discard the flow-through, place the column back in a collection tube and centrifuge for another 1 min at 14000 rpm.

9. Pipette 50-200 μl RRB directly into HiPure DR column. Incubate at room temperature for 1-5 min. Centrifuge it at 12000 rpm for 1 min.

**Note:** If the expected DNA yield is more than the yield from pervious step, put the HiPure DR column on a new microtube and add another 50-200  $\mu$ l RRB, incubate for 5 min at room temperature. Then, centrifuge for 1 min at 12000 rpm. However, it is possible to pass the follow-through from step 9 once more to obtain DNA with higher concentrations.