



## ZymoBIOMICS™ RNA Miniprep Kit

Microbiome RNA from any sample

#### Highlights

- ZymoBIOMICS™ innovative lysis system enables efficient and unbiased lysis of microbes including gram positive/negative bacteria, fungi, protozoans, and viruses from any sample including feces, soil, plant, water, biofilms, swabs, saliva, body fluids, etc.
- Rapid and robust, spin-column purification of high-quality RNA (including small/microRNAs) that is inhibitor-free and ready for and microbiome measurements RT/qPCR using Next-Gen sequencing.
- High-sensitivity and increased detection limit of very low abundance organisms.

#### Catalog Numbers: R2001



Scan with your smart-phone camera to view the online protocol/video.





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Revised on: 1/24/2024

### **Product Contents**

| ZymoBIOMICS <sup>™</sup> RNA Miniprep Kit  | <b>R2001</b><br>(50 prep) |
|--|---------------------------|
| ZR BashingBead™ Lysis Tubes (0.1 & 0.5 mm) | 50                        |
| DNA/RNA Shield™                            | 50 ml                     |
| RNA Lysis Buffer                           | 50 ml                     |
| RNA Prep Buffer                            | 25 ml<br>(x2)             |
| RNA Wash Buffer <sup>1</sup> (concentrate) | 24 ml                     |
| ZymoBIOMICS™ DNase/RNase-Free Water        | 30 ml                     |
| ZymoBIOMICS™ HRC Prep Solution             | 30 ml                     |
| DNase I <sup>2</sup> (lyophilized)         | 250 U                     |
| DNA Digestion Buffer                       | 4 ml                      |
| Zymo-Spin <sup>™</sup> III-HRC Filters     | 50                        |
| Zymo-Spin <sup>™</sup> IIICG Columns       | 100                       |
| Collection Tubes                           | 150                       |
| Instruction Manual                         | 1                         |

Storage Temperature - Store all kit components (i.e., buffers, columns) at room temperature.

<sup>1</sup> Add 96 ml 100% ethanol (104 ml 95% ethanol) to the 24 ml RNA Wash Buffer concentrate.

2 Reconstitute lyophilized DNase I with ZymoBIOMICS™ DNase/RNase-Free Water, mix by gentle inversion and store frozen aliquots: #E1009-A (250 U), add 275 µl water

## **Specifications**

- Sample Sources Bacterial, fungal, protozoan, algae, viral, mitochondrial, and host RNA is efficiently isolated from ≤ 250 mg of soil, mammalian feces and plant/seed, ≤ 50-100 mg (wet weight) fungal or bacterial cells¹, biofilms, water, and swabs.
- Sample Homogenization ZymoBIOMICS<sup>™</sup> innovative lysis system ensures complete lysis of the microbial cell walls and accurate microbial analysis, free of bias.
- Sample Preservation DNA/RNA Shield<sup>™</sup> lyses cells, inactivates nucleases and infectious agents, and is ideal for sample storage and transport at ambient temperatures.
- Size Total RNA including small/microRNAs (≥ 17 nt).
- Purity A<sub>260</sub>/A<sub>280</sub> & A<sub>260</sub>/A<sub>230</sub> > 1.8. RNA is ready for Next-Gen Sequencing, RT/qPCR, etc.
- Binding Capacity 100 µg RNA (Zymo-Spin™ IIICG Column).
- Elution Volume ≥ 50 μl ZymoBIOMICS<sup>™</sup> DNase/RNase-Free Water.
- **Equipment Needed** (user provided) Microcentrifuge, Vortex Genie (recommended).
- Recommended Materials (available separately) –

DNA/RNA Shield™ collection devices:

fecal collection tube; R1101

collection tube: R1102

lysis tube (microbe); R1103

lysis tube (microbe) w/ swab; R1104

lysis tube (tissue); R1105

collection tube (1 ml fill) w/ swab; R1106, R1107 collection tube (2 ml fill) w/ swab; R1108, R1109

<sup>1</sup> This equates to approximately 10<sup>9</sup> bacterial cells, 10<sup>8</sup> yeast cells, and 10<sup>7</sup> mammalian cells.

### **Product Description**

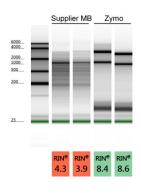
The **ZymoBIOMICS™ RNA Miniprep Kit** is designed for purifying RNA from a wide array of sample inputs (e.g., feces, soil, plant, water, and biofilms) that is ready for microbiome or metagenome analyses.

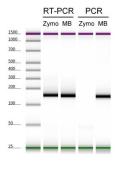
The **ZymoBIOMICS**™ innovative lysis system eliminates bias associated with unequal lysis efficiencies of different organisms (e.g., gram negative/positive bacteria, fungus, protozoans, and algae). The provided **DNA/RNA Shield**™ preserves nucleic acids at ambient temperatures, providing an unbiased molecular snapshot of the sample.

The procedure uses **Zymo-Spin**<sup>™</sup> column technology that results in high-quality total RNA (including small/microRNAs 17-200 nt) that is free of PCR inhibitors (e.g., polyphenols, humic acids and fulvic acids) and is ready for RT-PCR, arrays, sequencing, etc.

#### **High-Quality RNA**







Human stool RNA isolated with the **ZymoBIOMICS**™ **RNA Miniprep Kit** is higher quality (right); compared to Supplier MB procedures (left). Quality assessed by Agilent 2200 TapeStation™.

Human stool RNA was analyzed after RT-PCR and PCR amplification (~150 bp fragment shown) for both Zymo and Supplier MB procedures. Lack of a band in PCR using the ZymoBIOMICS™ RNA Miniprep Kit indicates DNA-free RNA. Quality assessed by Agilent 2200 TapeStation™.

#### **Protocol**

The protocol consists of: (I) Buffer Preparation, (II) Sample Preparation and (III) Total RNA Purification

#### (I) Buffer Preparation

- ✓ Add 96 ml 100% ethanol (104 ml 95% ethanol) to the 24 ml RNA Wash Buffer concentrate.
- ✓ Reconstitute lyophilized DNase I with ZymoBIOMICS™ DNase/RNase-Free Water, mix by gentle inversion and store frozen aliquots:

#E1009-A (250 U), add 275 µl water #E1011-A (1500 U), add 1,500 µl water

#### (II) Sample Preparation

- ✓ Perform all steps at room temperature and centrifugation at 10,000-16,000 x g for 30 seconds, unless specified.
- ✓ The sample input can be scaled up or down, proportionally.
- Add 750 µl DNA/RNA Shield™ to a sample (see table below) in a ZR BashingBead Lysis Tube (0.1 & 0.5 mm) and cap tightly. If a sample is already collected in DNA/RNA Shield™, transfer 750 µl liquid sample into a ZR BashingBead Lysis Tube (0.1 & 0.5 mm) and cap tightly.

**Note:** For samples stored and lysed in **DNA/RNA Shield™ Lysis Tubes**, do not add ZymoBIOMICS™ Lysis Solution and proceed to Step 2.

| Sample Type   | Maximum Input               |
|---|-----------------------------|
| Soil, feces, plant, seed  | ≤ 250 mg                    |
| Cells in DNA/RNA Shield <sup>™</sup> or isotonic buffer/PBS (bacterial 10 <sup>9</sup> , yeast 10 <sup>8</sup> , mammalian 10 <sup>7</sup> )  | ≤ 50-100 mg<br>(wet weight) |
| DNA/RNA Shield <sup>™</sup> collection devices<br>(e.g., cat. #R1101, R1102-R1105) or<br>Biological liquids and swabs collected in DNA/RNA Shield <sup>™</sup><br>(e.g., cat. #R1100, R1106-R1109, R1150) | 750 µl                      |

- 2. For complete lysis of tough-to-lyse samples (microbes, tissue, etc.), perform mechanical homogenization in a **ZR BashingBead Lysis Tube (0.1 & 0.5 mm)** by securing in a high-speed bead beater fitted with a 2 ml tube holder assembly (e.g., MP Bio FastPrep-24, Bertin Precellys, etc.). Process¹ at maximum speed for ≥ 5 minutes.
- 3. Centrifuge and transfer up to 400  $\mu$ I of the supernatant<sup>2</sup> into a nuclease-free tube (not provided).
- 4. Add an equal volume of **RNA Lysis Buffer** to the supernatant<sup>2</sup> (1:1) and mix well. Then proceed to purification (page 6).

<sup>1</sup> Processing time will vary based on sample input and bead beater. For low-speed homogenizers (e.g., Vortex Genie), process samples for ≥ 15 minutes. Optimization may be required.

<sup>2</sup> Up to 400 µl sample input can be processed per prep.

#### (III) Total RNA Purification

- ✓ Perform all steps at room temperature and centrifugation at 10,000-16,000 x g for 30 seconds, unless specified.
- Add an equal volume of ethanol (95-100%) to the sample in RNA Lysis Buffer and mix well.

Example: Add 800 µl ethanol to 800 µl mixture (sample mixed in **RNA Lysis Buffer**).

- Transfer the mixture into a Zymo-Spin<sup>™</sup> IIICG Column<sup>1</sup> (green) in a Collection Tube and centrifuge. Discard the flow-through.
- 3. Add 400 µl **RNA Prep Buffer** to the column and centrifuge. Discard the flow-through.
- 4. Add 400 μl **RNA Wash Buffer** to the column and centrifuge. Carefully, transfer the column into a nuclease-free tube (not provided).
- 5. Add 85 μl **ZymoBIOMICS**<sup>™</sup> **DNase/RNase-Free Water** directly to the column matrix, then centrifuge to elute.
- 6. **DNase I**<sup>2</sup> treatment (recommended)
  - (D1) To the eluate, add 10 μl DNA Digestion Buffer and 5 μl DNase I and mix gently by inversion.
  - (D2) Incubate at room temperature (20-30°C) for 15 minutes.
- 7. Add 2 volumes of RNA Lysis Buffer to the sample (2:1) and mix.

Example: Add 200 µl RNA Lysis Buffer to 100 µl mixture (DNase I-treated eluate).

8. Add an equal volume of ethanol (95-100%) (1:1) and mix.

Example: Add 300 µl ethanol to 300 µl mixture (eluate in RNA Lysis Buffer).

- 9. Transfer the mixture into a new **Zymo-Spin**™ **IIICG Column**¹ (green) in a **Collection Tube** and centrifuge. Discard the flow-through.
- 10. Add 400  $\mu$ l **RNA Prep Buffer** to the column and centrifuge. Discard the flow-through.

(continue to purification, page 7)

<sup>1</sup> To process sample volume > 700 µl, **Zymo-Spin**™, columns may be reloaded.

<sup>2</sup> Prior to use, reconstitute the lyophilized **DNase I** (Buffer Preparation, page 4). \* Unit definition – one unit increases the absorbance of a high molecular weight DNA solution at a rate of  $0.001 \, A_{260}$  units/ml of reaction mixture at  $25^{\circ}$ C.

- 11. Add 700 µl **RNA Wash Buffer** and centrifuge. Discard the flow-through.
- 12. Add 400 **RNA Wash Buffer** and centrifuge the column for 1 minute to ensure complete removal of the wash buffer. Carefully transfer the column into a new nuclease-free tube (not provided).
- 13. Add 100 µl **ZymoBIOMICS**™ **DNase/RNase-Free Water** directly to the column matrix and then centrifuge to elute.

Alternatively, for highly concentrated RNA use ≥ 50 µl elution.

- 14. Place a **Zymo-Spin**<sup>™</sup> **III-HRC Filter** in a new **Collection Tube** and add 600 µl **ZymoBIOMICS**<sup>™</sup> **Prep Solution**. Centrifuge at 8,000 x g for 3 minutes.
- 15. Transfer the eluted RNA (step 13) into a prepared **Zymo-Spin**<sup>™</sup> **III-HRC Filter** in a nuclease-free tube (not provided). Then centrifuge at exactly 16,000 x g for 3 minutes.

The filtered RNA can be used immediately or stored frozen.

## **Appendices**

#### Samples stabilized and stored in DNA/RNA Shield™

Recommended: **DNA/RNA Shield**™ effectively lyses cells, inactivates nucleases and infectious agents and is ideal for sample storage/transport at ambient temperatures prior to nucleic acid purification.

<u>Liquid samples</u>: Mix an equal volume **DNA/RNA Shield**<sup>™</sup> (2X concentrate) and sample (1:1). <u>Solid samples</u>: Submerge sample (not to exceed 10% (v/v or w/v) in **DNA/RNA Shield**<sup>™</sup> (1X).

Mix well/homogenize sample prior to storage. Samples in **DNA/RNA Shield**<sup> $^{\text{TM}}$ </sup> can be stored at ambient temperature  $\geq$  1 month or long term at frozen temperature.

## **Ordering Information**

| Product Description           | Catalog No. | Size      |
|-------------------------------|-------------|-----------|
| ZymoBIOMICS™ RNA Miniprep Kit | R2001       | 50 preps. |

| Individual Kit Components   | Catalog No.                      | Amount               |
|---|----------------------------------|----------------------|
| ZR BashingBead™ Lysis Tubes (0.1 & 0.5 mm)  | S6012-50                         | 50                   |
| DNA/RNA Shield™   | R1100-50<br>R1100-250            | 50 ml<br>250 ml      |
| RNA Lysis Buffer  | R1060-1-50<br>R1060-1-100        | 50 ml<br>100 ml      |
| RNA Prep Buffer   | R1060-2-25<br>R1060-2-100        | 25 ml<br>100 ml      |
| RNA Wash Buffer (concentrate)   | R1003-3-24<br>R1003-3-48         | 24 ml<br>48 ml       |
| ZymoBIOMICS™ DNase/RNase-Free Water   | D4302-5-30<br>D4302-5-50         | 30 ml<br>50 ml       |
| DNase I Set (lyophilized) DNase I (250 U) & DNA Digestion Buffer (4 ml)   | E1010                            | 1 set                |
| OneStep™ PCR Inhibitor Removal Kit  | D6030                            | 50 prep              |
| Zymo-Spin <sup>™</sup> IIICG Columns  | C1006-50-G                       | 50                   |
| Collection Tubes  | C1001-50<br>C1001-500            | 50<br>500            |
| DNA/RNA Shield™ - Fecal Collection Tube   | R1101                            | 10                   |
| DNA/RNA Shield™ Collection Tube DNA/RNA Shield™ Lysis Tube (microbe) DNA/RNA Shield™ Lysis Tube (microbe) w/ swab DNA/RNA Shield™ Lysis Tube (tissue) | R1102<br>R1103<br>R1104<br>R1105 | 50<br>50<br>50<br>50 |
| DNA/RNA Shield™ Collection Tube (1 ml fill) w/ swab   | R1106<br>R1107                   | 10<br>50             |
| DNA/RNA Shield™ Collection Tube (2 ml fill) w/ swab   | R1108<br>R1109                   | 10<br>50             |

## **Complete Your Workflow**

✓ For tough-to-lyse samples, use ZR BashingBead Lysis Tubes:

| ZR BashingBead Lysis Tubes   |                                |
|------------------------------|--------------------------------|
| 2.0 mm beads #S6003-50       | For plant/animal tissue        |
| 0.1 + 0.5 mm beads #S6012-50 | For microbes                   |
| 0.1 + 2.0 mm beads #S6014-50 | For microbes in tissue/insects |

✓ For high-throughput and automatable microbiome DNA and RNA purification from any sample (DNase I Set included):

| ZymoBIOMICS RNA        |   |
|------------------------|---|
| MagBeads #R2137, R2138 | Automatable (Tecan, Hamilton, Kingfisher, etc.) |

✓ For RNA clean-up (purification) from the aqueous phase (e.g., TRIzol, TRI Reagent or similar) or from any enzymatic reaction (e.g., DNase I treated RNA):

| RNA Clean & Concentrator kit |   |
|------------------------------|---|
| Spin-column #R1013-R1014     | DNase I Set included                            |
| MagBeads #R1081, R1082       | Automatable (Tecan, Hamilton, Kingfisher, etc.) |

✓ For NGS:

| Zymo-Seq RiboFree Total RNA Library Prep kit |          |
|--|----------|
| #R3000                                       | 12 preps |
| #R3003                                       | 96 preps |

## **Troubleshooting Guide**

| Incomplete lysis and/or high-mass input:  - If precipitation occurs (upon adding ethanol to the lysate) or if the lysate is extremely viscous, increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization until lysate is transparent (see image).  - Ethanol and/or salt contamination. After centrifugation steps, carefully remove the column from the collection tube to prevent buffer carryover. Alternatively, blot emptied collection tube with a tissue or towel.  - Make sure lysate and wash buffers have passed completely through the matrix of the column. This may require centrifuging at a higher speed and/or longer time.  Incomplete lysis and/or cellular debris:  - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  - Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  - To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page.4), then clean-up the treated sample.  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be stored frozen for later processing. | Problem                            | Possible Causes and Suggested Solutions   |
|---|------------------------------------|---|
| - If precipitation occurs (upon adding ethanol to the lysate) or if the lysate is extremely viscous, increase the volume of DNA/RNA shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization until lysate is transparent (see image).  - Ethanol and/or salt contamination. After centrifugation steps, carefully remove the column from the collection tube to prevent buffer carryover. Alternatively, blot emptied collection tube with a tissue or towel.  - Make sure lysate and wash buffers have passed completely through the matrix of the column. This may require centrifuging at a higher speed and/or longer time.  Incomplete lysis and/or cellular debris:  - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  DNA contamination  To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page 4), then clean-up the treated sample.  RNA degradation  To prevent RNA degradation:  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be                  |                                    | Incomplete lysis and/or high-mass input:  |
| - Ethanol and/or salt contamination. After centrifugation steps, carefully remove the column from the collection tube to prevent buffer carryover. Alternatively, blot emptied collection tube with a tissue or towel.  - Make sure lysate and wash buffers have passed completely through the matrix of the column. This may require centrifuging at a higher speed and/or longer time.  Incomplete lysis and/or cellular debris:  - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  DNA contamination  To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page 4), then clean-up the treated sample.  RNA degradation  To prevent RNA degradation:  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be   | iysate                             | is extremely viscous, increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization until |
| - Ethanol and/or salt contamination. After centrifugation steps, carefully remove the column from the collection tube to prevent buffer carryover. Alternatively, blot emptied collection tube with a tissue or towel.  - Make sure lysate and wash buffers have passed completely through the matrix of the column. This may require centrifuging at a higher speed and/or longer time.  Incomplete lysis and/or cellular debris:  - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  DNA contamination  To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page 4), then clean-up the treated sample.  RNA degradation  To prevent RNA degradation:  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be   |                                    | Sample handling:  |
| matrix of the column. This may require centrifuging at a higher speed and/or longer time.  Incomplete lysis and/or cellular debris:  - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  DNA contamination  To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page 4), then clean-up the treated sample.  RNA degradation  To prevent RNA degradation:  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be  | (A260/A230 IIIII, A260/A280 IIIII) | remove the column from the collection tube to prevent buffer carryover.   |
| - Increase the volume of DNA/RNA Shield and/or RNA Lysis Buffer to ensure complete lysis and homogenization. Be sure to centrifuge any cellular debris and then process the cleared lysate.  Low yield  Sample input:  - Too much input or incomplete lysis/homogenization can cause cellular debris to clog or overload the column and result in compromised RNA recovery. Use less input material and/or increase DNA/RNA Shield and/or RNA Lysis Buffer.  DNA contamination  To remove DNA:  - Perform in-column DNase I treatment or perform DNase I treatment post-purification (R1013, page 4), then clean-up the treated sample.  RNA degradation  To prevent RNA degradation:  - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be   |                                    | matrix of the column. This may require centrifuging at a higher speed   |
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| - Immediately collect and lyse fresh sample into DNA/RNA Shield to ensure nucleic acid stability. Homogenized samples in DNA/RNA Shield can be  |                                    |   |
| nucleic acid stability. Homogenized samples in DNA/RNA Shield can be  | RNA degradation                    | To prevent RNA degradation:   |
|   |                                    | nucleic acid stability. Homogenized samples in DNA/RNA Shield can be  |

For technical assistance, please contact 1-888-882-9682 or email tech@zymoresearch.com

| Notes |  |
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