



Research Use Only

DNA/RNA Extraction Kit

User Manual

Manufacturer:

GeneReach Biotechnology Corporation

TEL: 886-4-2463-9869 Email: sales@genereach.com

No. 19, Keyuan 2nd Road, Central Taiwan Science Park, Taichung City 407, Taiwan

Website: www.tacomag.com

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Symbols



Date of manufacturing



Manufacturer



Lot number



Catalogue number



Do not reuse

Kit Components

A. Reagents

taco™ DNA/RNA Extraction Kit		
Cat. No.: atc-d/rna		
Number of reactions: 320		
Reagent Name	Volume	Quantity
Magnetic Bead	18 ml	1 bottle
Lysis Buffer	180 ml	1 bottle
Washing Buffer A ¹	135 ml	2 bottles
Washing Buffer B ²	40 ml	2 bottles
Eluting Buffer	64 ml	1 bottle
User Manual		1 copy

Note: Treat all reagents as potential irritants.

¹ Add 135 ml 95% ethanol to Washing Buffer A before use.

Mark the bottle label after adding ethanol.

² Add 230 ml 95% ethanol to Washing Buffer B before use.

Mark the bottle label after adding ethanol.

B. Consumables**Note: Do not reuse the consumables****(a) For taco™ 32: taco™ Plate & Sleeve**

Product Name	Amount (pcs)	Cat. No.
96-Well Extraction Plate	20	AtcP
Mixing Sleeve	40	
taco™ Sticker	1	

(b) For taco™ 24: taco™ Plate & Comb

Product Name	Amount (pcs)	Cat. No.
48-Well Extraction Plate	40	atcp24
Mixing Comb	40	

Storage

All reagents should be sealed tightly in a cool and dry place at room temperature (16~30°C). The expiration date of the kit and each component are stated on the label of each item. Do not use any reagent of the kit beyond the expiration date. Users should check the expiration date before use, as it could affect the accuracy of the result.

Materials and Equipments Required, but Not Provided

- **taco™** Nucleic Acid Automatic Extraction System: **taco™ 24**, **taco™ 32** or **taco™ mini**
- Step pipette (optional)
- Disposable gloves
- Micro-centrifuge tubes
- Micropipette and filter tips (p1000, p200)
- 95% ethanol
- Phosphate buffer saline (PBS)

Introduction

The **taco™** DNA/RNA Extraction Kit is designed for **taco™** Nucleic Acid Automatic Extraction System. Based on the magnetic separation technology, homogenized sample cells are lysed and nucleic acid is captured by silica coated magnetic beads. Washing Buffer is then applied to remove impurities, and Eluting Buffer to recover nucleic acids from magnetic beads following serial washing steps. This kit can extract viral DNA and RNA from shrimp muscle for research use purpose only. Other sample types must be validated by users.

Intended Use

The **taco™** DNA/RNA Extraction Kit is intended to be used for extracting viral DNA and RNA from various sample types such as shrimp tissue. The **taco™** DNA/RNA Extraction Kit has to be used with the **taco™** Nucleic Acid Automatic Extraction System.

This product is intended to be used by professional users such as well-trained laboratory technicians who are familiar with molecular biology techniques.

Important Notes

- After receiving the kit, please check the kit components for any damage. Contact GeneReach Biotechnology Corporation or your local distributor if reagent bottles are damaged. Do not use damaged kit, as it could affect the accuracy of the result.
- Pipette tips are all for one-time use only. Repeated usage will lead to cross-contamination.
- When working with chemicals, please always wear a suitable lab coat, disposable gloves, and protective goggles.
- Discard gloves if they are contaminated.
- Do not combine components with different batches.
- Avoid microbial contamination of the reagents.
- To minimize the risk of infections from potentially infectious material, we recommend working under a laminar hood until the samples are lysed.
- This kit should only be used by trained personnel.
- Disposal of waste must be compliant with local laws.

Product Limitations

The system performance has been validated by using infected shrimp muscle for viral nucleic acid isolation. The user is responsible for

tacoTM DNA/RNA Extraction Kit

validating the performance of the **taco**TM DNA/RNA Extraction Kit for other particular use.

The kit and plastic parts are not intended for any therapeutic or diagnostic purposes for animals or humans.

Nucleic Acid Extraction Procedure

A. Use of taco™ Sticker (for taco™ 32)

For your convenience, you may put the **taco™** Sticker on top of reagent bottles and on the rim of 96-Well Extraction Plate to avoid human error.

a. taco™ Sticker

- Plate Sticker:

For **taco™ 32**: Apply the Plate Sticker on the rim of 96-Well Extraction Plate directly.



- Bottle Sticker:

Apply the Sticker on top of each reagent bottle.



b. Abbreviation Definition

LB	Lysis Buffer
M	Magnetic Bead
WA	Washing Buffer A
WAM	Washing Buffer A + Magnetic Bead
WB	Washing Buffer B
E	Eluting Buffer

B. Protocol

Note: The following protocol is for fresh and frozen shrimp samples. For other sample types, please refer to www.tacomag.com for instructions.

- a. Load reagents into 96-Well Extraction Plate/48-Well Extraction Plate according to **Table 1** at the room temperature (16-30°C) for the best performance.

Table 1. Loading Reagent

Step	Reagents
1	Add 200 µl 95% ethanol and 500 µl Lysis Buffer to column #1 (#7)
2	Add 750 µl Washing Buffer A ¹ to column #2 (#8)
3	Add 750 µl Washing Buffer A to column #3 (#9)
4	Add 750 µl Washing Buffer B ² to column #4 (#10)
5	Add 750 µl Washing Buffer B to column #5 (#11)
6	Add 50~200 µl Eluting Buffer to column #6 (#12)
7	Add 50 µl Magnetic Bead ³ to column #2 (#8)

¹ Ensure that 135 ml 95% ethanol has been added to Washing Buffer A before the first time use.

² Ensure that 230 ml 95% ethanol has been added to Washing Buffer B before the first time use.

³ Magnetic Bead must be resuspended before aliquoting.

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- b.** Grind the tissue (40 mg) with 450 µl PBS in a 1.5 ml micro-centrifuge tube with disposable grinder. Centrifuge at 12000 rpm for 5 minutes to spin down the debris (For ethanol preserved sample, please Appendix I).
- c.** Transfer **200 µl of the supernatant** to column #1 (#7) of 96-Well Extraction Plate/48-Well Extraction Plate.
- d.** Open the door of **taco™** to install the loaded plate and Mixing Sleeve/Mixing Comb, ensure they are pushed into position.

Note: Please refer to respective instrument manual for instructions.

- e.** Close the door of **taco™** and press “Start/Operation button to start extraction. The extraction procedure will finished within 1 hour.
- f.** After the extraction procedure, **take out the extraction plate and Mixing Sleeve/Mixing Comb accordingly**, then press “Reset/Operation button to reset the program.

Note: Please refer to respective instrument manual for instructions.

- g.** Transfer the nucleic acids from column #6 and/or #12 to the new micro-centrifuge tubes for further use (See “Purity of

Nucleic Acid”, Appendix II).

Note: Carryover of magnetic beads in eluates will not affect most downstream applications. If the risk of magnetic beads carryover needs to be minimized, transfer the eluates to micro-centrifuge tubes, centrifuge for 1 minute at full speed to pellet down the remaining magnetic beads, and carefully transfer the supernatants to new micro-centrifuge tubes.

h. It is strongly recommended to use freshly extracted nucleic acids for downstream applications such as amplification. Otherwise, the extracted nucleic acids should be kept frozen; for long-term storage -80°C is recommended (See “Storage of Nucleic Acid”, Appendix II).

Note: Do not reuse the Consumables.

Note: Any deviation from the instruction may lead to a low recovery rate of the nucleic acid extract.

Troubleshooting

Comments and suggestions

Low DNA/RNA yield

- | | |
|--|--|
| (a) Magnetic bead was not resuspended completely | Before starting the procedure, ensure that magnetic bead is fully resuspended. Vortex for at least 5 seconds before first use, and perform mild agitation before subsequent uses. |
| (b) Washing Buffer A and B did not contain ethanol | Ensure the correct volume of ethanol is added to Washing Buffer A and B; tightly seal the reagent bottles to prevent ethanol from evaporating. Repeat the extraction procedure with proper reagent is necessary when the ethanol was not added to Wash Buffer A and Wash Buffer B before use. (For the proper procedure of extraction, please see “Protocol”). |

Comments and suggestions

- | | |
|---|---|
| (c) Reagents were loaded in wrong order | Restart the loading procedure with a new extraction plate. Ensure that all reagents were loaded on the well in the correct order. Repeat the extraction procedure with new samples. |
| (d) Poor sample quality | Using fresh sample for extraction is recommended. Poor sample quality may influence nucleic acid quality. |
| (e) Incorrect sample volume | The kit performance would be affected if user did not use the right volume of sample. User should optimize the sample quantity when dealing with different sample types. |
| (f) Mixing Sleeve/Mixing Comb was not installed | Contact your local distributor or GeneReach Biotechnology Corporation for assistance. |
| (g) Inappropriate operation environment | Operation temperature could affect the recovery rate. Please ensure the operation environment is under room temperature (16-30°C). |

Comments and suggestions

- | | |
|---|---|
| (h) Use non-recommended extraction instrument | Using non-recommended instrument may influence the performance of taco™ DNA/RNA Extraction Kit. We strongly recommend user to apply DNA/RNA Extraction Kit on taco™ extraction systems. |
|---|---|

Poor DNA/RNA performance in downstream applications

Note: A spectrophotometer is required for following check-up.

- | | |
|--|---|
| (a) Insufficient DNA/RNA is used in downstream application | Quantify the extracted DNA/RNA by spectrophotometer of the absorbance at 260 nm (See “Quantification of Nucleic Acid”, Appendix II). |
| (b) Excess DNA/RNA used in downstream application | Excess DNA/RNA can inhibit some enzymatic reactions. Quantify the extracted DNA/RNA by spectrophotometer of the absorbance at 260 nm (See “Quantification of Nucleic Acid”, Appendix II). |

Comments and suggestions

Low A_{260}/A_{280} ratio

- | | |
|---|--|
| (a) Absorbance reading at 320 nm was not subtracted from the absorbance readings at 260 nm and 280 nm | To correct for the presence of magnetic beads particles in the eluted solution, an absorbance reading at 320 nm should be taken and subtracted from the absorbance readings obtained at 260 nm and 280 nm. |
|---|--|

Appendix I

Sample Preparation

- Animal Tissue: ethanol preserved shrimp muscle
 - i. Grind the tissue (20 mg) with 500 µl Lysis Buffer in 1.5 ml micro-centrifuge tube with disposable grinder.
 - ii. Centrifuge at 12000 rpm for 5 minutes to spin down the debris.
 - iii. Transfer 400 µl of the supernatant and 200 µl 95% ethanol to column #1 (#7) of 96-Well Extraction Plate.
 - iv. Follow **Table 1** from **Step 2 to 7** for loading reagents.

Note: The above sample preparation method is recommended for general muscle tissues which contain high volume of protein; other sample types need to be validated by users.

Note: For other sample types, please refer to www.tacomag.com.

Appendix II

A. Storage of Nucleic Acid

For long-term storage, extracted nucleic acids should be stored at -80°C.

B. Quantification of Nucleic Acid

Note: A spectrophotometer is required for following check-up.

The concentration of nucleic acids should be determined by measuring the absorbance at 260 nm in a spectrophotometer.

Use Eluting Buffer as the blank to calibrate the spectrophotometer. If the purified nucleic acids need to be diluted before the quantification, the Eluting Buffer also has to be diluted before use. Also, the same dilution factor needs to be applied for calculation.

Collect the absorbance reading of purified nucleic acid at 260 nm and 280 nm. The ratio between the absorbance values at 260 nm and 280 nm gives an estimation of nucleic acid purity (See “Purity of Nucleic Acid”).

Carryover of magnetic beads may affect the A_{260} reading, but should not affect the performance of nucleic acid in downstream applications.

C. Purity of Nucleic Acid

Purity is determined by calculating the ratio of absorbance at 260 nm to absorbance at 280 nm with a background correction at 320 nm, i.e., $(A_{260}-A_{320}) / (A_{280}-A_{320})$. A subtracted absorbance reading at 320

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nm is to correct the presence of magnetic beads particles in the eluted solution. An A_{260} / A_{280} ratio of 1.6~2.0 is indicative of highly purified nucleic acid.

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