

#### Manual

# MagnifiQ<sup>™</sup> 1 Total RNA Plus instant kit

Kit for automated, magnetic isolation of total RNA in the strip format. Contains ready-to-use, reagent-filled strips and all necessary consumables.

catalog#	size	compatible devices *		
614A-1V-32	32 isolations	Auto-Pure Mini		
614A-1V-160	160 isolations	Auto-Pure Mini		

#### \* Compatible devices

The kit has been tested with specific Allsheng brand isolation devices. This does not preclude it from working with other devices. If your device is not listed, please contact us at info@aabiot.com.

For research use only.

#### Guarantee

A&A Biotechnology provides a guarantee on this product.

The company does not guarantee the correct performance of this kit in the event of:

- not adhering to the supplied protocol
- use of not recommended equipment or materials
- use of other reagents than recommended or which are not a component of the product
- use of expired or improperly stored product or its components.



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## Advantages

- Requires only a few minutes of manual work while adding the samples. The rest of the procedure is carried out by automatic extraction device.
- It does not require the initial preparation of buffers. Just add samples to the plate and after about 30 min you get extracted material.
- Enables RNA isolation from single sample.

### Sample type

## Specification

	sample size		
<u>Bacteria G+ (cultures)</u>	up to $2 \times 10^8$	protocol time	~ 30 min.
Bacteria G- (cultures)	up to $2 \times 10^8$	elution volume	100 µl 1
Cell cultures	up to 2 x 10 <sup>6</sup>	elution solution	Tris buffer
Fresh blood (not frozen)	up 1 ml	binding capacity	30 µg RNA
<u>Solid tissue</u>	20 - 50 mg	downstream applications	reverse transcription RT-qPCR, transcriptome seque

<sup>1</sup> The elution volume prepared on the strip is 100 µl. To obtain a smaller elution volume, subtract the appropriate amount of elution solution from the well 6 on the XS-R strip. Attention! Do not reduce the elution volume below 50 µl.

## Description

MagnifiQ<sup>™</sup> 1 Total RNA Plus instant kit is designed for total RNA isolation from various types of biological materials. The isolated material is perfect for further analyzes and tests by RT-PCR methods and for sequencing.

The MagnifiQ<sup>™</sup> product series is based on the automated isolation of nucleic acids with use of magnetic beads. This method significantly shortens working time and reduces risk of mistake in comparison to manual methods.

## Contents

	620A-1V-32 620A-1V-160				
component	quantity	cat #	quantity	cat#	storage
XS-R - extraction strip	4 x 8 pcs	K-S1V22XR	20 x 8 pcs	K-S1V22XR	15-25 °C
Fenozol Plus	20 ml	K-FENP-20	90 ml	K-FENP-90	4-8 °C
ultrapure water	8 ml	K-WUP-8	40 ml	K-WUP-40	-20-25 °C
tip comb 8	8 x 2 pcs	K-C8U-2	40 x 2 pcs	K-C8U-2	15-25 ℃

## Additional equipment and reagents

#### Necessary

- 1.5 ml Eppendorf tubes (sample lysis)
- pipette
- pipette tips
- vortex
- thermoblock
- centrifuge

## **Material preparation**

#### Bacteria G+ (cultures)

Additional reagents you will need:

• BacBreaker bacteria lysis enzyme mix (20 µl per sample), cat # BACB-15A

Option:

• Lysostaphin (20 μl per sample), cat # 1007-3; For Staphylococcus spp. we recommend using lysostaphin.

Transfer the sample of the bacterial culture to the 1.5 ml tube (not included). Centrifuge for <b>3 min</b> at <b>10 000 RPM</b> . Discard the supernatant.
Suspend the bacterial pellet in <b>50 µl</b> of <b>ultrapure water</b> .
Add <b>20 µl BacBreaker</b> enzyme mix.
Attention. For lysis of Staphylococcus bacteria, add 20 µl of lysostaphin.
Vortex the sample for <b>10 s</b> and incubate for <b>10 min</b> at <b>37 °C</b> .
Add <b>500 μl</b> of <b>Fenozol Plus.</b>
Vortex the sample for <b>10 s</b> and incubate for <b>5 min</b> at <b>50 °C</b> .
Add <b>170 µl</b> of <b>ultrapure water</b> . Vortex the sample for <b>10 s.</b>
Note. Adding water results in DNA precipitation.
Centrifuge the sample for <b>10 min</b> at <b>12 000 RPM</b> .
Attention. In the isolation protocol, use the supernatant as the sample. Take the filtrate starting from the top so that the precipitate is not carried over to the next purification step. The step is important for RNA purity.
Follow point 1. <u>of the protocol</u> .

#### Bacteria G- (cultures)

1.	Transfer the sample of bacterial culture to 1.5 ml Eppendorf tube (not included). Centrifuge for <b>3 min</b> at <b>10 000 RPM</b> . Discard the supernatant.
2.	Add 500 μl of Fenozol Plus.
3.	Vortex the sample for <b>10 s</b> and incubate for <b>5 min</b> at <b>50 °C</b> .
4.	Add <b>170 μl</b> of <b>ultrapure water.</b> Vortex the sample for <b>10 s.</b> <b>Note.</b> Adding water results in DNA precipitation.
5.	Centrifuge the sample for <b>10 min</b> at <b>12 000 RPM</b> .
6.	Attention. In the isolation protocol, use the supernatant as the sample. Take the filtrate starting from the top so that the precipitate is not carried over to the next purification step. The step is important for RNA purity.
	Follow point 1. <u>of the protocol</u> .

#### **Cell cultures**

1.	Transfer the sample of the cell culture to 1.5 ml Eppendorf tube (not included). Centrifuge for <b>3 min</b> at <b>10 000 RPM</b> . Discard the supernatant.
2.	Add <b>500 μl</b> of <b>Fenozol Plus.</b>
3.	Vortex the sample for $10 s$ and incubate for $5$ min at $50$ °C.
4.	Add <b>170 µl</b> of <b>ultrapure water</b> . Vortex the sample for <b>10 s.</b>
	Note. Adding water results in DNA precipitation.
5.	Centrifuge the sample for <b>10 min</b> at <b>12 000 RPM</b> .
6.	Attention. In the isolation protocol, use the supernatant as the sample. Take the filtrate starting from the top so that the precipitate is not carried over to the next purification step. The step is important for RNA purity.
	Follow point 1. <u>of the protocol</u> .

### Fresh blood (not frozen)

#### Additional reagents you will need:

• **RBCL** (max. 5 ml per sample), <u>cat # 213-100</u>

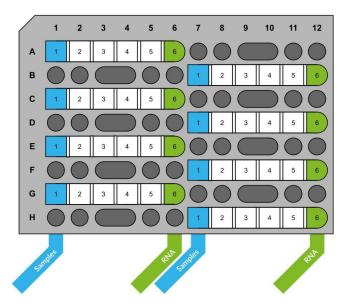
1.	Add <b>the appropriate amount</b> of <b>RBCL</b> to <b>1 ml</b> of <b>blood sample</b> . <b>Attention</b> . We recommend using 5 volumes of RBCL to 1 volume of blood sample.
2.	Mix and incubate on ice for <b>15 min</b> . <b>Note.</b> Note the changing appearance of the sample during the incubation. The initially opaque solution should turn to a completely transparent ruby-red at the incubation end.
3.	Centrifuge for <b>10 min</b> at <b>3000 x g.</b> Carefully discard supernatants.
4.	Add <b>500 μl</b> of <b>Fenozol Plus.</b>
5.	Vortex the sample for <b>10 s</b> and incubate for <b>5 min</b> at <b>50 °C</b> .
6.	Add <b>170 μl</b> of <b>ultrapure water</b> . Vortex the sample for <b>10 s.</b> <b>Note</b> . Adding water results in DNA precipitation.
7.	Centrifuge the sample for <b>10 min</b> at <b>12 000 RPM</b> .
8.	Attention. In the isolation protocol, use the supernatant as the sample. Take the filtrate starting from the top so that the precipitate is not carried over to the next purification step. The step is important for RNA purity. Follow point 1. of the protocol.

#### Solid tissue

1. Homogenize tissue sample in liquid nitrogen. Transfer the sample of tissue to 1.5 ml Eppendorf tube (not included). Note. For soft tissues, use mechanical lysis. Transfer the sample to a tube containing zirconia beads (A&A Biotechnology cat. # 106-50Z). Place the tube in the Beadbeater and perform the shaking process. 2. Add 500 µl of Fenozol Plus. 3. Vortex the sample for 10 s and incubate for 5 min at 50 °C. 4. Add 170 µl of ultrapure water. Vortex the sample for **10 s**. Note. Adding water results in DNA precipitation. 5. Centrifuge the sample for 10 min at 12 000 RPM. Attention. In the isolation protocol, use the supernatant as the sample. Take the filtrate starting from the top so that the 6. precipitate is not carried over to the next purification step. The step is important for RNA purity. Follow point 1. of the protocol.

### Protocol

1. Place XS-R strips in the rack.



- 2. Remove the foil from the XS-R strips.
- 3. Add 600 µl of samples to the well 1 (first from the left) on the XS-R strip.

Note. The wells are numbered on the side of the strip.

Attention: If no pellet is visible at the bottom of the tube, take the filtrate starting from the top so as not to mix the whole after centrifugation. This step is intended to separate the fraction containing DNA.

- 4. Place the rack in the extraction device.
- 5. Place the appropriate number of **tip combs 8** in the extraction device.

#### 6.

#### Run the program according to the table below:

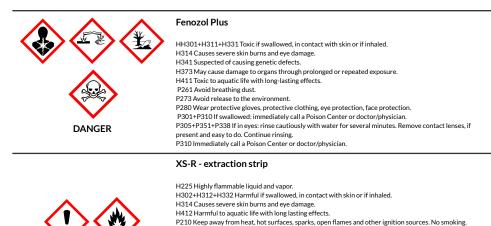
Step	Well	Name	Mix Time (min)	Magnet (s)	Wait Time (min)	Volume (µl)	Mix Speed (1-10)	Temp. (°C)
1	1	MIX	5.0	0	0.0	900	4	OFF
2	5	BEADS	0.5	30	0.0	600	5	OFF
3	1	BIND	5.0	60	0.0	900	4	OFF
4	2	WASH1	2.0	30	0.0	600	6	OFF
5	3	WASH2	1.0	30	0.0	600	6	OFF
6	4	WASH3	1.0	30	5.0	600	6	OFF
7	6	ELUTION	6.0	60	0.0	100	8	50
8	3	DROP	1.0	0	0.0	600	5	OFF

After the program is over, remove the combs and then remove the rack from the extraction device and transfer RNA located in well **6** (first from the right) on the **XS-R** strip into sterile tubes (not included).

Note. Store extracted material at -80 °C.

### Safety information

DANGER



P273 Avoid release to the environment.

if present and easy to do. Continue rinsing.

or shower.

P280 Wear protective gloves/protective clothing/eye protection/face protection/hearing protection.

P303+P361+P353 If on skin (or hair): Take off immediately all contaminated clothing. Rinse skin with water

P301+P312+P330 If swallowed: Call a poison center/doctor/ if you feel unwell.

P304+P340 If inhaled: Remove person to fresh air and keep comfortable for breathing. P305+P351+P338 If in eyes: rinse cautiously with water for several minutes. Remove contact lenses,

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