

HB220514

# T7 High Yield RNA Synthesis Kit

### **Product Information**

Product Name	Catalog No.	Size
	10623ES50	50 T
T7 High Yield RNA Synthesis Kit	10623ES60	100 T
	10623ES70	500 T

## **Product Description**

T7 High Yield RNA Synthesis Kit optimizes the transcription reaction system. The kit can synthesize the single-stranded RNA efficiently by uses T7 RNA polymerase, the linear double-stranded DNA with the T7 promoter sequence as the template, NTPs as the substrate to control the DNA sequence downstream of the promoter. During transcription, modified nucleotides can be added to the substrate to prepare biotin or dye-labeled RNA.

This kit can synthesize long transcripts and short transcripts, RNA can be produced  $100-200 \mu g$  with  $1 \mu g$  of DNA template input. The RNA synthesized by transcription can be used for various downstream applications, such as RNA structure and function research, RNase protection, probe hybridization, RNAi, microinjection and in vitro translation.

#### **Contents**

		Cat#/Specification		
Contents No.	Name	10623ES50	10623ES60	10623ES70
		(50 T)	(100 T)	(500 T)
10623-A	T7 RNA Polymerase Mix	100 μL	200 μL	1 mL
10623-B	10×Transcription Buffer	100 μL	200 μL	1 mL
10623-C	ATP (100mM)	100 μL	200 μL	1 mL
10623-D	CTP (100mM)	100 μL	200 μL	1 mL
10623-E	GTP (100mM)	100 μL	200 μL	1 mL
10623-F	UTP (100mM)	100 μL	200 μL	1 mL
10623-G	Control DNA Template $(500 ng/\mu L)$	10 μL	20 μL	100μL

## **Applications**

In vitro RNA synthesis

# **Shipping and Storage**

Dry ice transportation. Store at -20°C, valid for two years.

## **Notes:**

- 1. Be careful not to mix RNase in the reaction system.
- 2. Experiment equipment (such as: pipette tip, product tube, etc.) should strictly use RNase Free products.
- 3. For your safety and health, please wear lab coats and disposable gloves.
- 4. For research use only!

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## Synthesis principle



Figure 1: In vitro RNA transcription process

## **Experimental methods**

#### 1. DNA template preparation

Linearized plasmids with double-stranded T7 promoters or PCR amplification products can be used as Hifair<sup>TM</sup> T7 High Yield RNA Synthesis Kit in vitro transcription templates, which can be dissolved in TE buffer or RNase free  $H_2O$ .

T7 promoter sequence: TAATACGACTCACTATAG\*GG (Note: G\* is the first base of RNA transcription) A. Plasmid template

Insert the target DNA to the plasmid vector containing the T7 promoter, and then treated with restriction enzymes, purified after completely linearized.

- Note: 1. The circular plasmids have no effective termination, RNA products of different lengths will be transcribed. In order to obtain a specific length RNA, the plasmid must be completely linearized.
  - 2. The restriction enzyme selected for plasmid linearization needs to be on the right side of the promoter region, downstream of the inserted DNA fragment, and has no recognition site in the inserted DNA fragment. The restriction enzyme should be capable of forming 5' sticky ends or smooth ends.
  - 3. In order to avoid the influence of protein and salt ions on the system, the plasmid is recommended to be purified when used as a template for in vitro transcription after linearization.

## B. PCR product template

The PCR product with T7 promoter can be used as an in vitro transcription template. First, add the T7 promoter sequence (TAATACGA CTCACTATAGGG) to the 5' end of the upstream primer sense strand; next, the T7 promoter DNA template is amplified under the action of high fidelity enzyme; then transcription is performed. PCR products can be used directly as templates without purification, but higher RNA output will be obtained after purification.

- Note: 1. The specificity and concentration of the PCR product must be confirmed by electrophoresis when used as a template. Put 2-5  $\mu$ L of PCR product into the 20  $\mu$ L reaction system.
  - 2. In order to obtain more high-quality RNA, the PCR product should be recovered by gel and used as a template for in vitro transcription.

## 2. In vitro RNA transcription

#### A. Thawing reagents

Centrifuge the T7 RNA Polymerase Mix briefly and place on ice. Thaw 10× Transcription Buffer and ribonucleotides (ATP, CTP, GTP, UTP), mix and centrifuge to the bottom of the tube, place 10× Transcription Buffer at room temperature, and place 4 types of ribonucleotides on ice.

# B. Assembly transcription reaction at room temperature

Prepare the reaction system according to the following system:

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Components	Volume (µL)	Final concentration
RNase free H <sub>2</sub> O	Up to 20	-
10×Transcription Buffer	2	1×
CTP / GTP/ ATP/ UTP (100 mM each)	2 each	10 mM each
Template DNA	1 μg	-
T7 RNA Polymerase Mix	2	-

- Note: 1. The reaction is configured at room temperature. Since 10× Transcription Buffer contains spermidine, the concentration of spermidine too high will cause DNA template precipitation at low temperature.
  - 2. Short transcript (<100 nt), 2 μg template can be used, transcription time increased to 4-8 hs.
  - 3. For long transcripts (>1000 nt), recommended to use linearized plasmid templates for transcription.
  - 4. Perform the reaction in a PCR machine with the hot lid open to prevent the reaction solution from evaporating for a long time.
  - 5. The reaction product may have a white precipitate. This is free pyrophosphate and magnesium ions produce the magnesium pyrophosphate in the reaction, won't affect the subsequent experiments. You can add some EDTA to clear it. If the addition of EDTA affects subsequent experiments, the supernatant can also be recovered by centrifugation.
  - 6. The reagents and containers must without RNase contamination.

#### C. Incubate at 37°C for 2 hours

Mix the above reaction solution, briefly centrifuge to the bottom of the tube, and incubate at 37°C for 2 hs. If the transcript length is less than 100 nt, increase the reaction time to 4-8 hs.

#### D. DNase I treatment (optional)

After the reaction is complete, add 2  $\mu$ L of DNase I (RNase free) to each tube and incubate at 37°C for 15 mins to remove the template DNA.

### 3. Product purification

## A. RNA Cleaner magnetic bead

Take out RNA clean beads from  $4^{\circ}$ C in advance, equilibrate to room temperature (about 30 minutes), dilute the transcription product to 50  $\mu$ L with RNase free H<sub>2</sub>O.

- ① Invert or vortex to mix the magnetic beads thoroughly, add  $2^{\times}$  magnetic beads (100  $\mu$ L) to the RNA sample (50  $\mu$ L), pipette 6 times to mix thoroughly. Incubate for 5 min at room temperature, allow RNA to bind to the magnetic beads.
- ②Place the sample on the magnetic stand for 5 mins. After the solution is clear, remove the supernatant carefully.
- 3Keep the sample on the magnetic stand, add  $200~\mu$ L of freshly prepared 80% ethanol to rinse the magnetic beads, incubate at room temperature for 30~s, carefully remove the supernatant. Repeat this operation once again.

(Note: The 80% ethanol used for rinsing needs to be freshly prepared with RNase free  $H_2O$  to prevent the introduction of RNase enzyme from causing RNA degradation.)

(4) Keep the sample on the magnetic stand, open the lid for 5 mins to dry the beads.

(Note: Avoid excessive drying. If the magnetic beads are cracked, it means that the magnetic beads are too dry and the elution efficiency of RNA will be reduced).

- $\odot$ Remove the sample from the magnetic stand, add 22  $\mu$ L RNase free H<sub>2</sub>O, pipette 6 times to mix well, incubate for 5 mins at room temperature.
- 6 Place the sample on the magnetic stand for 5 mins. After the solution clarifies, carefully transfer 20  $\mu$ L of the supernatant to a new RNase free PCR tube.

Note: Leave 2-3  $\mu$ L of liquid when transferring the supernatant, not attract the magnetic beads and affect the subsequent experiments. The obtained RNA is extremely unstable, please proceed to the next step as soon as possible. If you want to save, please store at

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-80°C.)

#### B. Phenol/chloroform

- ①Add 115μL RNase free H<sub>2</sub>O and 15 μL 3M sodium acetate (pH 5.2) to 20 μL reaction mixture, mix well.
- ②Extract once with an equal volume of phenol/chloroform (1:1), and then extract twice with an equal volume of chloroform. Collect the supernatant and transfer it to a new RNase free EP tube.
- ③ Add 2 volumes of absolute ethanol to precipitate RNA. Mixing uniformly and place it at -20°C for at least 30 mins, centrifuge at 4°C for 15 mins at maximum speed, and collect the precipitate.
- 4) Add 500 µL of ice-cold 70% ethanol to wash the RNA pellet.
- ⑤Dissolve the RNA pellet with 20 μL RNase free H<sub>2</sub>O. Store the purified RNA solution at -80°C.

## C. Lithium chloride precipitation

The lithium chloride precipitation method demands the RNA length must be greater than 300 nt, and the concentration must not be less than 100 ng/ $\mu$ L.

- ①Add 30  $\mu$ L RNase free H<sub>2</sub>O and 30  $\mu$ L 7.5M lithium chloride to 20  $\mu$ L reaction mixture.
- ② After mixing uniformly, place it at -20°C for at least 30 mins, centrifuge at 4°C for 15 mins at maximum speed, and collect the precipitate.
- ③Add 500 μL of ice-cold 70% ethanol to wash the RNA pellet.
- 4 Dissolve the RNA pellet with 20 μL RNase free H<sub>2</sub>O. The purified RNA solution is stored at -80°C.

#### D. Column purification

Before purification, add  $80\mu L$  RNase free  $H_2O$  to dilute the product to  $100~\mu L$ , and then follow the column purification instructions for purification.

# 4. RNA quantification

A. Ultraviolet absorption

## B. Dye method

Use RiboGreen dye to quantify RNA, free nucleotides won't affect quantification, purified or unpurified RNA in reaction products can be accurately quantified.

#### 5. RNA size and quality detection

## A. Agarose electrophoresis

In order to determine the size, integrity and quality of RNA, agarose gel electrophoresis or polyacrylamide gel electrophoresis is required for detection.

#### B. Agilent 2100 Bioanalyzer detection

Agilent 2100 Bioanalyzer can be used to evaluate the integrity and quality of RNA. It only requires a small amount of RNA for analysis. High-quality RNA should show obvious and sharp peaks on the electrogram.

## **Frequently Asked Questions:**

#### 1. Low transcript yield

The quality of template is closely related to the yield. The yield of experimental group is significantly lower than the control group. The possible reasons are: ① the experimental template contains inhibitory components; ② The template has something wrong. Suggestions: ① Re-purify the template; ② Determine the template quantification and its integrity; ③ Extend the reaction time; ④ Increase the amount of template input; ⑤ Try other promoters and RNA polymerases.

#### 2. Low yield of short transcripts

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Short transcription initiation fragment will inhibit the reaction. When the transcription product is less than 100 nt, extending the reaction time to 4-8 hs or increasing the amount of template to 2 µg will increase RNA yield.

### 3. RNA transcription length is greater than expected

If the electrophoresis shows that the product band is larger than the expected size, the possible reasons: ①The plasmid template may not be completely linearized; ②The 3' end of the sense strand has a prominent structure; ③The RNA has a secondary structure that is not completely denatured.

Suggestions: ①Check whether the template is completely linearized, and if necessary, perform additional linearization; ②Select a suitable restriction enzyme to avoid 3' overhangs, or use Klenow Fragment /T4 DNA polymerase to complete the transcription before proceeding; ③Use denatured gel to detect RNA products.

## 4. RNA transcription length is less than expected

If the electrophoresis shows that the product band is smaller than the expected size, the possible reasons: ①The template contains a termination sequence similar to T7 RNA polymerase; ②The GC content in the template is high.

Suggestions: ① Lower the reaction temperature (for example, 30°C). Sometimes lowering the temperature can increase the transcription length, but it will reduce the yield. Or try different RNA polymerases for transcription; ②If the template GC content is high, use 42°C to transcript, or add SSB to increase the yield and transcription length.

## 5. Electrophoresis tailing of transcription products

There is tailing phenomenon during electrophoresis. Possible reasons: ①Contaminated by RNase during experimental operation; ② Contaminated DNA template by RNase.

Suggestions: ①Use RNase-free pipette tips and EP tubes, wear disposable latex gloves and masks, and all reagents are prepared with RNase free  $H_2O$ . ②Re-purify the template DNA.

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