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EntiLink™ 1st Strand cDNA Synthesis Kit

Catalog No.	Specification	Storage/Shelf life
EQ003	100 rxn	-20°C/1.5 years

Introduction

The EntiLink™ 1st Strand cDNA Synthesis Kit is a complete system for the efficient synthesis of first-strand cDNA .

The EntiLink™ Reverse Transcriptase has dramatically improved thermal stability and can withstand reaction temperatures up to 50°C, making it suitable for reverse transcription of RNA templates with complex secondary structures. The EntiLink™ Reverse Transcriptase also has an increased affinity for templates and is suitable for reverse transcription of small amounts of templates and low-copy genes. The EntiLink™ Reverse Transcriptase also has an improved ability to synthesize full-length cDNAs, which can be amplified up to 10 kb in length.

The kit contains all the components needed to synthesize high-quality first-strand cDNA from total RNA or mRNA, and provides two primers for cDNA synthesis: Random Primers N6 and oligo (dT)18. The synthesized single-stranded cDNA product can be used directly in subsequent PCR or qPCR reactions.

Kit Components

Component	Quantity
RNase-Free ddH ₂ O	2×1 mL
5× Buffer	400 µL
EntiLink™ Enzyme Mix	200 µL
Oligo (dT)18 (50 µM)	100 µL
Random Primers N6 (50 µM)	100 µL
User Manual	1 copy

Note: 1) 5× Buffer contains dNTPs. 2) EntiLink™ Enzyme Mix contains RNase inhibitor



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Advantage

1. Can efficiently synthesize full-length first-strand cDNA up to 10kb.
2. Can withstand reaction temperatures up to 50 ° C.
3. Fully provide all the components needed for the RT reaction.

Kit application

1. cDNA library construction.
2. RT-qPCR reaction and RT-PCR reaction.
3. Primer extension.
3. RNA sequencing.

Notes

- 1.All operations should be performed on ice, and RNase contamination should be avoided.
- 2.For your safety and health, please wear lab coat and disposable gloves to operate.
- 3.This product is for research use ONLY!

Self supplied Reagents and items

- 1.RNase-free 200μL microcentrifuge tube.
2. Pipettes and tips (to avoid RNase contamination, RNase-free pipette tips with filter cartridges must be used).
3. Disposable gloves, masks and other protective equipment.
4. Constant temperature water bath.
5. In RNase-free laboratory operations: Because of the RNase in saliva and skin, wear latex . gloves and a mask during the whole process of RNA extraction.



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Operation steps

1. Reverse transcription reaction system preparation (20 μ L system)

Components	Volume
Total RNA	1 ng -5 μ g*
or mRNA	1 ng-500 ng*
5 \times Buffer	4 μ L
Random Primers N6 (50 μ M)	1 μ L
or Oligo (dT)18 (50 μ M)	or 1 μ L
or Gene Specific Primers (2 μ M)	or 1 μ L
EntiLink™ Enzyme Mix	2 μ L
RNase-free H ₂ O	To 20 μ L

[Note]: *If the subsequent experiment is qPCR, it is recommended that the amount of Total RNA or mRNA input does not exceed 1 μ g or 100 ng, and if the expression abundance of the target gene is very low, up to 5 μ g of Total RNA or 500 ng of mRNA can be input.

** If the subsequent experiment is PCR, for complex templates, the RNA, H₂O, and reverse transcription primers can be incubated at 65°C for 5 min and then quickly cooled on ice before adding the EntiLink™ Enzyme Mix; if the subsequent experiment is qPCR, the incubation step at 65°C can be omitted and the EntiLink™ Enzyme Mix can be added directly into the system.

2. Reaction program

Reaction Temperature	Reaction Time
25°C	5 min
42°C	30 min
85°C	5 min

[Note]: 1) Fluorescence quantification experiments can be performed using only Random Primers N6; they can also be mixed 1:1 with Oligo (dT)₁₈ for better results.

2) Reverse transcription temperature: 42°C is recommended. For high GC content templates or complex templates, the reverse transcription temperature can be increased to 50°C.

3) Reverse transcription products can be stored at -20°C for a short period of time, if long-term storage is required, it is recommended to store them at -80 °C after dispensing to avoid repeated freezing and thawing.



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3. Primer selection

- 1) If the template is of eukaryotic origin, it is recommended to choose Oligo (dT)₁₈, which pairs with the 3' Poly A tail of the eukaryotic mRNA for the highest yield of full-length cDNA.
- 2) For reverse transcription of prokaryotic RNA, use Random Primers N6 or gene-specific primers.
- 3) Random Primers N6 is widely applicable. mRNA, rRNA, tRNA, small RNA and LncRNA templates can be reverse transcribed with Random Primers N6.
- 4) For cDNA synthesis of less than 2 kb, use 1-2 μ L of Random primers N6; for cDNA synthesis of more than 2 kb, use 0.4-1 μ L of Random primers N6.