Code: MORV0003 Code: MORV0003-5 Code: MORV0003-15
Pack Size: 1ml Pack Size: 5x1ml Pack Size: 15x1ml



1. Introduction

Genie Fusion Ultra High-Fidelity DNA Polymerase is a new generation superior enzyme based on Genie Fusion Ultra DNA Polymerase for robust PCR with higher fidelity. The unique extension factor, specificity-promoting factors and plateau-inhibiting factor newly added to Genie Fusion Ultra greatly improve its long-fragment amplification ability, specificity, and PCR yield. Genie Fusion Ultra is capable of amplifying long fragments such as 40 kb λ DNA, 40 kb plasmid DNA, 20 kb genomic DNA and 10 kb cDNA. The amplification error rate of Genie Fusion Ultra is 53-fold lower than that of conventional Tag and 6-fold lower than that of Pfu. In addition, Genie Fusion Ultra has a good resistance to PCR inhibitors and can be used for direct PCR amplifications of bacteria, fungi, plant tissues, animal tissues, and even whole blood samples. Genie Fusion Ultra contains two monoclonal antibodies inhibiting the $5'\rightarrow 3'$ polymerase activity and $3'\rightarrow 5'$ exonuclease activity at room temperature, which enable Genie Fusion Ultra to perform greatlyspecific Hot-Start PCRs. Genie Fusion Ultra High-Fidelity Red 2X Master Mix contains Genie Fusion Ultra High-Fidelity DNA Polymerase, dNTP, an optimized buffer system, and loading dye. The amplification can start only with the addition of primer and template, thereby easing PCR setup and improving reproducibility. Protective agents in the Genie Fusion Ultra High-Fidelity Red 2X Master Mix enable the resistance to repeated freeze-thaw cycles. Amplification will generate blunt-ended products, which are compatible with GenieClone II One Step Cloning Kit Series (Assay Genie #MORV0004).

2. Package Information

Components	MORV0003	MORV0003-5	MORV0003-15
Genie Fusion Ultra High-Fidelity Red 2X Master Mix	1 ml	5 x 1ml	15 x 1 ml

3. Storage

Store at -20°C; avoid repeated freezing and thawing.

4. Unit Definition

One unit (U) is defined as the amount of enzyme that incorporates 10 nmol of whole dNTPs into acid-insoluble products in 30 minutes at 74° C with activated salmon sperm DNA as the template / primer.

5. Quality Control

Residual Endonulease Test: The product is tested in a reaction containing 25 μ l of Genie Fusion Ultra High-Fidelity Red 2X Master Mix and 0.3 μ g of Supercoiled pBR322 DNA. After incubation at 37°C for 4 hours, there is no visually discernible change in DNA bands determined by agarose gel electrophoresis.

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ResidualE.Coli gDNA Test: The residual nucleotide in 25 μ l of Genie Fusion Ultra High-Fidelity Red 2X Master Mix is tested by SYBR® Red qPCR using specific primers of E.Coli gDNA. The residual E.Coli gDNA is lower than 10 copies.

Functional Assay 1: In a 50μ l PCR system with 1U of Genie Fusion Ultra High-Fidelity Red 2X Master Mix, 100 ng of human genomic DNA was used as template. After 35 cycles, 1/10 of PCR products were detected by 1% agarose gel electrophoresis. A single DNA band of 8.2 kb was detected after EB staining.

Functional Assay 2: In a 50μ l PCR system with 1U of Genie Fusion Ultra High-Fidelity Red 2X Master Mix, 10 ng of λ DNA was used as template. After 30 cycles, 1/10 of PCR products were detected by 1% agarose gel electrophoresis. A single DNA band of 15 kb was detected after EB staining.

6. Experimental Processes

6.1 For Conventional PCR

Recommended PCR System

Keep all components on ice during the experiment. All components need to be mixed up thoroughly after thawing and put back to -20°C Immediately for storage after using.

DdH ₂ O	up to 50 μl
Genie Fusion Ultra High-Fidelity Red 2X Master Mix	25 μl
rimer 1 (10 μM)	5 μΙ
Primer 2 (10 μM)	5 μΙ
Template DNA ^a	x μl

The Assay Genie PCR Enhancer is recommended for unsuccessful amplification of fragments with GC content > 60%. a. Optimal reaction concentration varies in different templates. In a 50 μ l system, the recommended template usage is as follows:

Templates	Input Template DNA
Genomic DNA	50 - 400 ng
Plasmid or Virus DNA	10 pg- 30 ng
cDNA	1 - 5 μ l (≤ 1/10 of the total volume of PCR
	system)

Recommended PCR Program

Steps	Temperature	Time	Cycles
Pre-denaturation ^a	95°C	30 sec / 3 min	1
Denaturation	95°C	15 sec	•
Annealing ^b	56-72°C	15 sec	}
Extension ^c	72°C	30 - 60 sec / kb	5 25 - 35
Final Extension	72°C	5 min	1

a. For pre-denaturation, the recommended temperature is 95°C, and the recommended time is 30 sec for plasmid / virus DNA and 3 min for genomic DNA / cDNA.

b. For annealing, the recommended temperature is the Tm of the primers. If the Tm of the primers is higher than 72°C, the annealing step can be removed (two-step PCR). If necessary, annealing temperature can be

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further optimized in a gradient. In addition, the amplification specificity depends directly on the annealing temperature. Raising annealing temperature is helpful to improve poor amplification specificity.

c. Longer extension time is helpful to increase the amplification yield.

6.2 For Long-fragment PCR

Genie Fusion Ultra High-Fidelity DNA Polymerase can extraordinarily perform a long-fragment amplification with high specificity and yields. If the recommended program is failure to work, the following Touch Down two-step PCR may be helpful:

Steps	Temperature	Time	Cycles
Pre-denaturation	95°C	3 min	1
Denaturation	92°C	15 sec	} 5
Extension	74°C	60 sec/kb	
Denaturation	95℃	15 sec	} 5
Extension	72°C	60 sec/kb	
Denaturation	95℃	15 sec	} 5
Extension	70°C	60 sec/kb	
Denaturation	95℃	15 sec	} 25
Extension	68°C	60 sec/kb	
Final Extension	68°C	5 min	1

It is recommended to use high-quality templates and long primers . Increasing the input of template DNA may be helpful to improve the amplification yield.

6.3 For PCR Using Crude Material as Template

Genie Fusion Ultra Master Mix have a good resistance to PCR inhibitors and can be used for direct PCR amplifications of bacteria, fungi, plant tissues, animal tissues, and even whole blood samples. Crude materials that have been successfully amplified with Genie Fusion Ultra Master Mix are as follows:

Sample Type	Amplification	Temperature Recommendation
	Method	(for a 50 μl PCR system)
Whole Blood	Direct PCR	1 - 5 μΙ
Filter Paper Dry Blood	Direct PCR	1 - 2 mm² filter paper
Cultured Cells	Direct PCR	Low amounts of cells
Yeast	Direct PCR	Single clone or 1 µl suspension
Bacteria	Direct PCR	Single clone or 1 µl suspension
Mod	Direct PCR	Low amount of sample
Sperm	Direct PCR	Low amount of sample
Plankton	Direct PCR	Low amount of sample
Plant Tissue	Direct PCR	1 - 2 mm² tissue
Mouse Tail	PCR with lysate	1 - 5 μl lysate*
Food	PCR with lysate	1 - 5 μl lysate*

^{*}Lysate Preparation:

Genie Fusion Ultra High-Fidelity Red 2X Master Mix Code: MORV0003 Code: MORV0003-5 Code: MORV0003-15 Pack Size: 1ml Pack Size: 5x1ml Pack Size: 15x1ml Animal Submerge little Mix well and spin 60°C 10 **Tissues** mount of samples in at room or Food lysis buffer with final min temperature. 95°C 10 samples concentration 200 Collect the μg/ml of Proteinase min supernatant as K (self-provide). lysate.

Lysis Buffer: 20 mM of Tris-HCl, 100 mM of EDTA, 0.1% SDS, pH 8.0 (not included in this kit).

7. Attention

- 1. Use high-quality templates.
- 2. DO NOT use dUTP or any primers or templates that contain uracil.
- 3. The Genie Fusion Ultra High-Fidelity DNA Polymerase has strong proofreading activity. Therefore, the PCR products must be purified before adding A-Tailing when TA cloning.
- 4. Primers design notes:
- * Choose C or G as the last base of the 3'-end of the primer.
- * Avoid continuous mismatching at the last 8 bases of the 3'-end of the primer.
- * Avoid hairpin structure at the 3'-end of the primer.
- * Tm of the primers should be within the range of 55°C 65°C (recommend to calculate in Primer Premier 5), and the Tm difference between F and R primers should be less than 1°C.
- * Additional sequence should not be included when calculating Tm of the primers.
- * GC content of the primers should be within the range of 40% 60%.
- * The general distribution of A, G, T, C in the primers should be uniform, and avoid using regions with rich GC and rich AT.
- * Keep complementary sequence less than 5 bases within the primers or between two primers, and complementary sequence less than 3 bases at the 3'-end of the primers.
- * Please search the specificity of the designed primers by NCBI BLAST to avoid non-specific amplification.

8. Troubleshooting

Template Input

No or Low Yield of PCR Products

Primers	Optimize primer design
Annealing Temperature	Set gradient annealing temperature to find out the optimal
	one
Concentration of Primers	Optimize the concentration of primers
Extension Time	Optimize the extension time to 30 sec/kb-1 min/kb
Cycle Numbers	Increase cycle numbers to 35 - 40
Purity of Templates	Use high-purity templates

Refer to the recommended reaction system and increase the

input

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Unspecific or Smear Bands in Electrophoresis

Primers Optimize primer design

Annealing Temperature Try to improve annealing temperature and set gradient

annealing temperature to optimize

Concentration of Primers Decrease the concentration of primers to final concentration

as 0.2 μM

Extension Time Appropriately decrease the extension time when bands longer

than target bands appear

Cycle Numbers Decrease cycle number to 25 - 30
PCR Programs Use Two-Step PCR or Touch-down PCR

Purity of Templates Use high purity templates

Template Input Modify or decrease templates input referring to the

recommended reaction system

Enzymes Input Appropriately adjust or decrease the input of high-fidelity

polymerase

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