

Long Taq DNA Polymerase

Catalog No.: P2082 (250U)

| Concentration: | 5U/µl | | |
|----------------|------------------------------|--------|--|
| Contents: | Long Taq DNA Polymerase 50µl | | |
| | 10xLong PCR Buffer I | 1.25ml | |
| | 10xLong PCR Buffer I | 1.25ml | |
| | dNTPs(2.5mM each) | 1 ml | |
| | PCR Enhancer | 500µl | |
| | 6xLoading Buffer | 1ml | |
| | | 0. | |

Note: 10xLong PCR Buffer I and 10xLong PCR Buffer II are Mg^{2+} plus.

Store at -20°C

In total 6 vials.

Description

Long Taq DNA Polymerase, a combination of two thermostable DNA polymerases, Taq and Pfu, is a special formulation designed for amplifying large fragment. This specially formulated Long Taq was shown to amplify long templates from λ phage genome of up to 20 kb. It is also a better choice for amplifying complex template, such as GC-rich template.

Long taq is suitable as a direct replacement for ordinary Taq Polymerase in most applications. Using Long Taq in your PCR reactions results in 3´-dA overhangs PCR products, which can be used in TA clone.

Unit Definition

One unit is defined as the amount of the enzyme required to catalyze the incorporation of 10 nmole of dNTPs into an acid-insoluble form in 30 minutes at 70°C using hering sperm DNA as substrate.

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Storage Buffer

20mM TrisCl (pH8.0), 100mM KCl, 3mM MgCl₂ 1mM DTT, 0.1% NP-40, 0.1% Tween20, 0.2mg/ml BSA, 50% (v/v) glycerol

10X Long PCR Buffer I with Mg²⁺

500mM Tris-HCl pH 8.8, 160mM (NH₄)₂SO₄, 25mM MgCl₂, 1% Triton X-100

PRODUCT USE LIMITATION.

This product is developed, designed and sold exclusively for research purposes and in vitro use only. The product was not tested for use in diagnostics or for drug development, nor is it suitable for administration to humans or animals.

10X Long PCR Buffer II with Mg²⁺

200mM Tris-HCl PH8.8, 100mM KCl, 100mM (NH₄)₂SO₄, 16mM MgSO₄, 1% Tritonx-100

Note:

10xLong PCR Buffer I is classical Long Taq DNA Polymerase buffer, is good for long template especially above 10kb.

10xLong PCR Buffer II is an optimized special buffer. It is for better fidelity but not good at long template above 10kb.

Users could choose suitable buffer for different template.

Features

- High fidelity: three times fidelity of Taq DNA Polymerase.
- Longer fragment: amplify long templates as long as 40kb.
- Amplification of complex template (GC rich or repetitive

sequence).

• Generates 3'-dA and blunt end PCR products.

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Applications

- PCR amplification of complex templates
- PCR amplification of long templates
- DNA sequencing
- PCR for cloning

Basic PCR Protocol

The following basic protocol serves as a general guideline and a starting point for any PCR amplification. Optimal reaction conditions (incubation time and temperature, concentration of Taq DNA Polymerase, primers, Mg²⁺, and template DNA) vary and need to be optimized.

1. Add the following components to a sterile microcentrifuge tube sitting on ice:

| Reagent | Quantity, for 50µl reaction | Final concentration |
|---|-----------------------------|---------------------|
| Sterile deionized water | variable | - |
| 10X Long PCR Buffer (Mg ²⁺ plus) | 5µl | 1X |
| dNTPs (10mM each) | lμl | 0.2 mM each |
| Primer I | variable | 0.4-1µM |
| Primer II | variable | 0.4-1µM |
| Long Taq DNA Polymerase (5U/µl) | 0.25-0.5 μl | 1.25-2.5U/50 μl |
| Template DNA | variable | 10рд-1µд |
| Total | | 50µl |

1.1 Recommended PCR assay with PCR Buffer (Mg^{2+} plus)



BIO KNOWLEDGE

LAB

1.2 Recommended PCR assay with PCR Buffer (Mg²⁺ free)

| Reagent | Quantity, for 50µl reaction | Final concentration |
|---|-----------------------------|---------------------|
| Sterile deionized water | variable | - |
| 10X Long PCR Buffer (Mg ²⁺ free) | 5µl | 1X |
| dNTPs (10mM each) | lμl | 0.2 mM each |
| Primer I | variable | 0.4-1µM |
| Primer II | variable | 0.4-1µM |
| 25mM Mg ²⁺ | variable | 1-4mM |
| Long Taq DNA Polymerase (5U/µl) | 0.25-0.5 μl | 1.25-2.5U/50 μl |
| Template DNA | variable | 10рд-1µд |
| Total | 1 | 50μΙ |

Table for selection of 25 mM MgCl₂ solution volume in

50 μ l reaction mix :

| Final Mg ²⁺ conc. | 1.0mM | 1.5mM | 2.0mM | 2.5mM | 3mM | 4mM |
|------------------------------|-------|-------|-------|-------|-----|-----|
| Mg ²⁺ Stock | 2µl | 3μl | 4µl | 5µl | 6µl | 8μl |

Recommendations with Template DNA in a 50μ l reaction volume

| Human genomic DNA | 0.1 µg-1 µg |
|--------------------|--------------|
| Plasmid DNA | 0.5 ng-5 ng |
| Phage DNA | 0.1 ng-10 ng |
| E.coli genomic DNA | 10 ng-100 ng |

2. Mix contents of tube. Cap tubes and centrifuge briefly to collect the contents to the bottom.

When using a thermal cycler that does not contain a heated lid, overlay the reaction mixture with 25 μl mineral oil.



3. Perform 25-35 cycles of PCR amplification as follows:

| Initial Denaturation | 94°C | 3 minutes |
|----------------------|---------|--------------|
| 25-35 Cycles | 94°C | 30 seconds |
| | 55-68°C | 30 seconds |
| | 72°C | 1-10 minutes |
| Final Extension | 72°C | 10 minutes |

4. Incubate for an additional 10 min at 72° C and maintain the reaction at 4° C. The samples can be stored at -20° C until use.

5. Analyze the amplification products by agarose gel

electrophoresis and visualize by ethidium bromide staining. Use appropriate molecular weight standards.

Notes on cycling conditions

- 4. Initial denaturation can be performed over an interval of 1-5 min at 95°C depending on the GC content of template.
- 5. Denaturation for 30 sec to 2 min at 94-95°C is sufficient. If the amplified DNA has a very high GC content, denaturation time may be increased up to 3-4 min.
- 6. Optimal annealing temperature is 5°C lower than the melting temperature of primer-temperature DNA duplex. If nonspecific PCR products are obtained optimization of annealing temperature can be performed by increasing temperature stepwise by 1-2°C.
- 7. The number of PCR cycles depends on the amount of template DNA in the reaction mix and on the expected yield of the PCR product. 25-35 cycles are usually sufficient for the majority PCR reaction. Low amounts of starting template may require 40 cycles.
- 8. The time of the final extension step can be extended for amplicons that will be cloned into T/A vectors.



Guidelines for preventing contamination of PCR reaction

During PCR more than 10 million copies of template DNA are generated. Therefore, care must be taken to avoid contamination with other templates and amplicons that may be present in the laboratory environment. General recommendations to lower the risk of contamination are as follows:

- Prepare your DNA sample, set up the PCR mixture, perform thermal cycling and analyze PCR products in separate areas.
- Set up PCR mixtures in a laminar flow cabinet equipped with an UV lamp.
- Wear fresh gloves for DNA purification and reaction set up.
- Use reagent containers dedicated for PCR. Use positive displacement pipettes, or use pipette tips with aerosol filters to prepare DNA samples and perform PCR set up.
- Always perform "no template control" (NTC) reactions to check for contamination

Quality Control

The absence of endodeoxyribonucleases, exodeoxyribonucleases and ribonucleases is confirmed by appropriate quality tests. Functionally tested in amplification of a single-copy gene from human genomic DNA. Endodeoxyribonuclease Assay No detectable conversion of covalently closed circular DNA to a nicked DNA was observed after incubation of 10U Long Taq Polymerase with 1 μ g pBR322 DNA in 50 μ l for 4 hours at 37°C and 70°C.

Exodeoxyribonuclease Assay

No detectable degradation of lambda DNA-HindIII fragments was observed after incubation of 10U Long Taq Polymerase with $1\mu g$ digested DNA for 4 hours at 37°C and 70°C.

Ribonuclease Assay

0% of the total radioactivity was released into trichloroacetic acid-soluble fraction after incubation of 10U Long Taq Polymerase with 1μg E.coli [3H]-RNA (40000cpm/μg) for 4 hours at 37°C and 70°C.