



## MBT060A Components

# Taq Polymerase (Recombinant) (5 units/µl)

**MBT060A Reagents provided** 1000 Units 5000 Units 500 Units 20 x 500 Units Tag Polymerase (5 U/ $\mu$ l) 100 µl 200 µl 1 ml 2 ml (20 x 100 µl) 10X HiBuffer A (With 25mM MgCl<sub>2</sub>) 2 ml 4 ml 20 ml 40 ml (20 x 2 ml) 10X HiBuffer S (With 17.5 mM MgCl<sub>2</sub>) 2 ml 20 ml 40 ml (20 x 2 ml) 4 ml 50mM MgCl<sub>2</sub> 1 ml 2 ml 10 ml 20 ml (20 x 1 ml)

## Description:

Taq DNA Polymerase is a thermostable DNA polymerase of thermophilic bacterium *Thermus aquaticus*. The enzyme catalyzes 5'-3' synthesis of DNA, has no detectable 3'-5' exonuclease (proof-reading) activity, but maintains the 5'-3' exonuclease activity. In addition, Taq DNA Polymerase exhibits deoxynucleotidyl transferase activity, resulting in addition of extra adenines at 3'-end of PCR products.

### Features:

- Thermostable enzyme of approximately 94 kDa from *Thermus aquaticus*
- Ultra pure recombinant protein
- Replicates DNA at 74°C and exhibits a half-life of 40 minutes at 95°C
- Generates mostly 3' dA overhang PCR products which are suitable for TA cloning
- DNA sequencing and labeling
- Incorporates modified nucleotides (E.g.: biotin, deoxygenin, fluroscently-labelled nucleotides)
- Supplied with two buffers- **10X Hibuffer A** with KCl and **10X Hibuffer S** with (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> [(NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> allows for PCR at a wide range of Mg<sup>2+</sup> concentrations and decrease unspecific priming].

Concentration: 5 U/µl

Source: E.coli cells with a pol gene from Thermus aquaticus

Molecular weight: 94 kDa monomer

#### **Unit Definition:**

1U is defined as amount of enzyme that is required to catalyze the incorporation of 10 nmoles of dNTP into acid-insoluble material in 30 minutes at  $74^{\circ}$ C.

# **Reaction Buffer:**

# 10X HiBuffer A (With 25mM MgCl<sub>2</sub>):

**500mM KCl**, 100mM Tris-HCl (pH 9.1 at 20°C) and 0.1% Triton X-100. The buffer is optimized for use with 0.1 - 0.2mM of each dNTP.

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#### 10X HiBuffer S:

**160mM**  $(NH_4)_2SO_4$ , 500mM Tris-HCl (pH 9.2 at 22°C), **17.5mM MgCl<sub>2</sub>** and 0.1% Triton X-100. The buffer is optimized for use with 0.35mM of each dNTP.

#### Storage Buffer:

200 mM Tris-HCl (pH 8.0 at 22°C), 100 mM KCl, 0.5% Tween 20, 0.5% Nonidet-P40, 0.1 mM EDTA, 1mM DTT and 50% Glycerol. Store at  $-20^{\circ}$ C.

#### Guidelines for PCR optimization using HiMedia's Taq Polymerase:

#### DNA Template

- 1. Use high quality, purified DNA templates
- 2. Approximately 10<sup>4</sup> copies are required to detect the amplification in 25-30 PCR cycles
- 3. Use higher DNA concentration when few PCR cycles are desired

#### • Primers

- 1. Generally 20-30 bp in size
- 2. GC content between 40-60% ideally
- 3. Melting temperatures should be between 42-65°C
- 4. Final concentration to be used  $0.1-0.5\mu M$  of each primer

#### Magnesium Concentration

- 1. Ideal for Taq Polymerase is 1.5-2.0mM
- 2. Optimum concentration depends on template, buffer and dNTPs
- 3. Higher than optimal concentration yields undesired products and if concentration is too low the concentration, no amplification products are detected.

#### • dNTPs

- 1. Typical concentration to be used is 200µM
- 2. Higher than optimal concentration of dNTPs yields higher yield but low fidelity

## • Taq Polymerase

1. Typical concentration to be used is 0.5 to 2 units per  $50\mu$ l of reaction

#### • PCR reaction

- 1. Thaw all reaction components on ice
- 2. To PCR reaction, add Taq Polymerase at the end
- 3. Once the reaction is set, immediately transfer the tubes to pre-heated thermal cycler.
- 4. Start the reaction with desired cycling conditions with annealing temperature set to 5°C difference of melting temperature between forward and reverse primers.

#### **Quality Control:**

All preparations are assayed for contaminating endonuclease, exonuclease, and non-specific DNase activities. Functionally tested in DNA amplification.

#### Buffers recommended for different sizes of template DNA

| Buffers           | Size of template DNA to be amplified |         |          |
|-------------------|--------------------------------------|---------|----------|
|                   | 100bp-5kb                            | 5kb-8kb | 8kb-20kb |
| HiBuffer S (1X)   | -                                    | +       | +        |
| HiBuffer A (1X)   | +                                    | -       | -        |
| MgCl <sub>2</sub> | +                                    | +       | +        |

Key: + Indicates recommended buffer

#### Inhibition and Inactivation:

- Inhibitors: ionic detergents (deoxycholate, sarkosyl and SDS) at concentrations higher than 0.06, 0.02, and 0.01% respectively.
- Inactivated by phenol/chloroform extraction.

#### NOTE:

The error rate for Taq Polymerase, which lacks proof-reading activity is approximately 1 to 2  $X10^{-5}$  errors (or mutation frequency) per nucleotide per duplication. Accordingly, the accuracy of PCR is 4.5 X  $10^{4}$ . Accuracy is an inverse of the error rate and shows an average number of correct nucleotides incorporated before an error occurs.

**Storage conditions:** The Taq Polymerase should be stored at -20°C. When stored under the recommended conditions, the product is stable for 2 years.

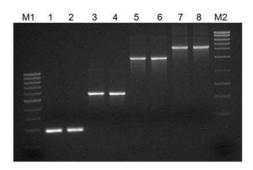


Figure representing amplification of different amplicon sizes using Taq Polymerase with HiBuffer A and HiBuffer S.

#### **Technical Assistance**

At HiMedia, we pride ourselves on the quality and availability of our technical support. For any kind of technical assistance, mail at <u>mb@himedialabs.com</u>.

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