



# Taq DNA Polymerase with Opti Buffer

Cat#  PCR001  PCR002  PCR001-S

Pack Size:  200U  1000U  100U recommended to use 1.25 U per reaction of 25 ul

## Storage and Handling:

- Store at -20°C in a frost-free freezer. Taq DNA Polymerase with Opti Buffer is stable for up to 1 years if stored at -20°C. Avoid repeated freeze-thaw cycles. Prior to use, thaw the Opti Buffer on ice. After opening, store at -20°C

## Kit contents:

Components	PCR001	PCR002	PCR001-S
Taq Polymerase 5U/μL	40 μL	200 μL	20 μL
10X Opti Buffer	1mL	3mL	1mL

## Introduction

Taq DNA Polymerase with Opti Buffer is a recombinant enzyme derived from the thermophilic bacterium *Thermus aquaticus*. It catalyzes the polymerization of nucleotides into DNA, and has a high degree of processivity and fidelity, making it a popular choice for PCR applications. The Opti Buffer provided with the Taq DNA Polymerase is optimized for efficient amplification of DNA targets, with enhanced specificity and yield. These guidelines apply to standard PCR. For amplification of templates with high GC content, high secondary structure, low template concentrations, or amplicons up to 5.5 kb. The enzyme lacks a 3'→5' exonuclease activity (no proofreading ability). Taq DNA Polymerase leaves an A' overhang, which makes the enzyme ideal for TA cloning. Additional optimization may be required depending on the sample type.

**10X Opti Buffer with MgCl<sub>2</sub>:** The reaction buffer has been optimized to work with wide rand of template. It contains, Tris-HCl (pH 8.8 at 25°C), KCl and 20mM MgCl<sub>2</sub>.

**Note:** 10X reaction buffer can be supplied without MgCl<sub>2</sub> on demand. Concentration gradients may form in frozen products and should be dispersed upon thawing. Mix well prior to use.

**Source:** An *E. coli* strain that carries the Taq DNA Polymerase gene.

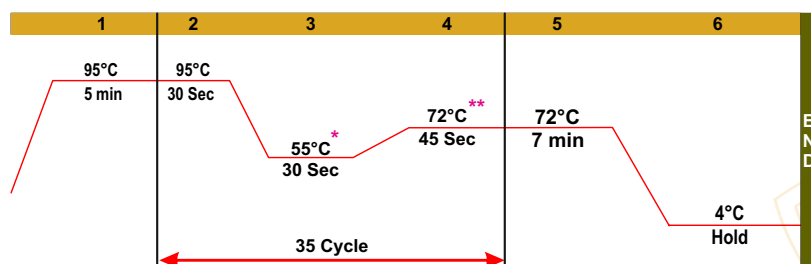
**Storage Buffer:** 50mM Tris-HCl (pH 8.0 at 25°C), 0.1mM EDTA, 2mM DTT, 50% glycerol.

**Unit Definition** One unit of Taq DNA Polymerase with Opti Buffer is defined as the amount of enzyme required to incorporate 10 nmol of dNTPs into acid-insoluble form in 30 minutes at 72°C using poly(dA)-oligo(dT) as a template-primer.

## Standard PCR Reaction mix

Component	For 25ul reaction	For 50ul reaction	Final Concentration
Template (DNA)	Variable	Variable	<1,000 ng
10 μM Forward Primer	0.5 μl	1 μl	0.2 μM (0.05-1μM)
10 μM Reverse Primer	0.5 μl	1 μl	0.2 μM (0.05-1μM)
10 mM dNTP each	0.5 μl	1 μl	200μM
10X Opti Buffer	2.5 μl	5 μl	1X
Taq DNA Polymerase	0.25 μl	0.5 μl	1.25 units /25μl PCR reaction
Nuclease-Free Water	to 25 μl	to 50 μl	

## General PCR Condition for 1 kb amplification



\* Depends on the primer T<sub>m</sub> and template used (Need to be optimized)

\*\* Allow approximately 1 minutes for every 1kb to be amplified.

## Result

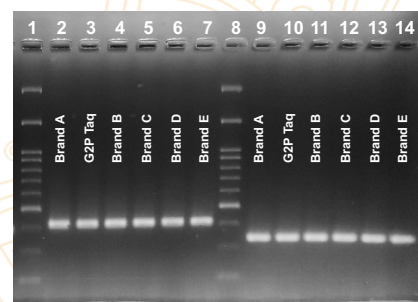


Fig: Different Gene was amplified from plant genome and loaded on 2% agarose gel. Lane 3&11 represent the amplification using Taq DNA Polymerase from Gene to protein Pvt Ltd.

## COMPONENTS OF THE REACTION MIXTURE

**Template DNA;** For both plasmid and phage DNA, the ideal amounts of template DNA in the 50- $\mu$ L reaction volume are **0.1–1 ng**, and for genomic DNA, they are 0.1–1  $\mu$ g. greater number of templates raises the possibility of non-specific PCR product production. The accuracy of the amplification is decreased with less template. The preparation of templates can be done using any standard DNA purification technique, such as G2P Genomic DNA Purification Kits (PUR11/12/13/14/15/16/17/18/19/20). Taq DNA polymerases can be inhibited by traces of some chemicals used for DNA purification, including phenol, EDTA, and proteinase K. Trace impurities are often removed from DNA samples using ethanol precipitation and repeated washing with 70% ethanol on the DNA pellet.

**Concentration of MgCl<sub>2</sub>;** Because Mg<sup>2+</sup> binds to dNTPs, primers, and DNA templates, Mg<sup>2+</sup> concentration must be adjusted. The concentration range indicated is 1-4 mM. The yield of the PCR product may be lowered if the Mg<sup>2+</sup> content is too low. If the Mg<sup>2+</sup> concentration is too high, non-specific PCR products may arise and PCR fidelity may be lowered. Taq DNA Polymerase is supplied with **20mM of MgCl<sub>2</sub>**, which is optimal in most of the case.

**dNTPs;** Each dNTP should be used at a final concentration of 0.2 mM. Higher dNTP concentrations are preferred in certain PCR applications. Because Mg<sup>2+</sup> binds to dNTPs, the MgCl<sub>2</sub> concentration must be adjusted correspondingly. The reaction mixture must contain equal amounts of all four nucleotides (dATP, dCTP, dGTP, and dTTP). Use the following dNTP mix volumes to achieve a **0.2 mM concentration of each dNTP** in the PCR mixture.

**Primers;** The PCR primers' suggested concentration range is **0.1-1  $\mu$ M**. Excessive primer concentrations raise mispriming and the production of non-specific PCR products are both possible. Higher primer concentrations in the region of 0.3-1 M are frequently preferred for defective primers.

## Troubleshooting

	Possible Cause	Recommendation
Little or no PCR Product	Difficult template e.g., GC-rich templates	<ul style="list-style-type: none"> <li>• Add DMSO (final concentration, 8%) and reduce enzyme concentration (e.g., use as little as 0.5 U per reaction).</li> </ul>
	DNA template problems	Check quality and concentration of template: <ul style="list-style-type: none"> <li>• Analyze an aliquot on an agarose gel to check for possible degradation.</li> <li>• Test the template with an established primer pair or PCR system.</li> <li>• Check or repeat template purification.</li> </ul>
	Enzyme concentration too low	<ul style="list-style-type: none"> <li>• Increase enzyme concentration to 2 U Taq DNA Polymerase per 50 <math>\mu</math>l reaction.</li> </ul>
	MgCl <sub>2</sub> concentration too low	Increase the MgCl <sub>2</sub> concentration in 0.25 mM steps; the minimal acceptable concentration is 1.5 mM MgCl <sub>2</sub> .
	Cycle conditions not optimal	<ul style="list-style-type: none"> <li>• Decrease annealing temperature.</li> <li>• Increase cycle number.</li> <li>• Make sure that the final elongation step is included in the program.</li> </ul>
	Primer design not optimal	Design alternative primers.
	Primer concentration not optimal	<ul style="list-style-type: none"> <li>• Both primers must have the same concentration.</li> <li>• Titrate primer concentration (0.1 to 0.6 M).</li> </ul>
	Primer quality or storage problems	<ul style="list-style-type: none"> <li>• If you use an established primer pair, check performance in an established PCR system (e.g., with a control template).</li> <li>• Make sure that the primers are not degraded.</li> <li>• Always store primers at -15 to -25°C.</li> </ul>
Multiple bands or background smear	Annealing temperature too low	Increase annealing temperature (Longer primers have higher annealing temperatures).
	Primer design or concentration not optimal	<ul style="list-style-type: none"> <li>• Review primer design.</li> <li>• Titrate primer concentration (0.1 to 0.6 M).</li> <li>• Both primers must have the same concentration.</li> <li>• Perform nested PCR with nested primers.</li> </ul>
	DNA template problems	Use serial dilution of template.
PCR products in negative control experiments	Carryover contamination	<ul style="list-style-type: none"> <li>• Replace all reagents, especially water.</li> <li>• Use aerosol-resistant pipette tips.</li> <li>• Set up PCR reactions in an area separate from that used for PCR product analysis.</li> <li>• To eliminate carryover contaminants: Use dUTP* (600mM) instead of dTTP (200 M) and thermolabile UNG* (1 U/50 l reaction); also, increase Mg<sup>2+</sup> concentration (to a maximum of 4 mM) to compensate for higher dNTP conc</li> </ul>