

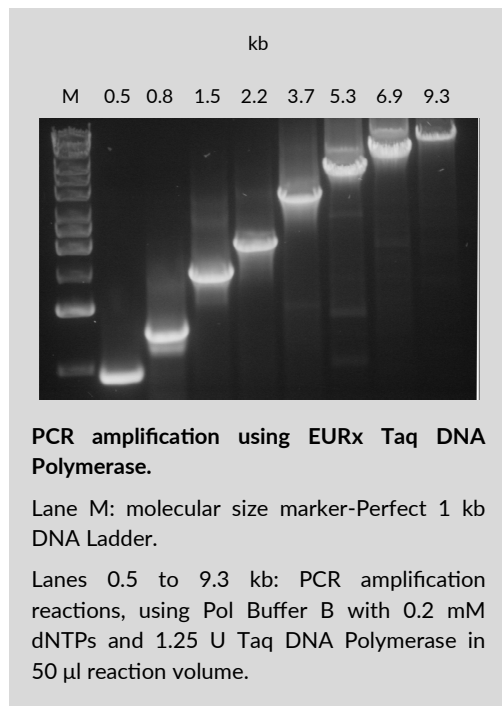
Taq DNA Polymerase

(*Thermus aquaticus*)

| Cat. No. | size |
|----------|------------|
| E2500-01 | 200 units |
| E2500-04 | 500 units |
| E2500-02 | 1000 units |
| E2500-03 | 5000 units |

Unit Definition: One unit is defined as the amount of enzyme required to catalyze the incorporation of 10 nmoles of dNTP into acid-insoluble material in 30 minutes at 74°C. The reaction conditions are: 50 mM Tris-HCl (pH 9.0 at 25°C), 50 mM NaCl, 5 mM MgCl₂, 200 μM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]dTTP), 10 μg activated calf thymus DNA and 0.1 mg/ml BSA in a final volume of 50 μl.

Storage Conditions: Store at -20°C.



Stable thermophilic DNA polymerase, suitable for applications requiring high temperature synthesis of DNA.

Description:

- Taq DNA Polymerase is a thermostable enzyme of approximately 94 kDa from *Thermus aquaticus*.
- Ultrapure, recombinant protein.
- The enzyme replicates DNA at 74°C and exhibits a half-life of 40 min at 95°C (1,2).
- Catalyzes the polymerization of nucleotides into duplex DNA in the 5'→3' direction in the presence of magnesium ions.
- Maintains the 5'→3' exonuclease activity.
- Lacks the 3'→5' exonuclease activity.
- Adds extra A at the 3' ends.
- Taq DNA Polymerase is recommended for use in PCR and primer extension reactions at elevated temperatures to obtain a wide range of DNA products up to 10 kb.

Storage Buffer:

20 mM Tris-HCl (pH 8.0 at 22°C), 100 mM KCl, 0.1 mM EDTA, 1 mM dithiothreitol, 50% glycerol and stabilizers

10 x Reaction Buffers:

10 x Pol Buffer A (optimization buffer without MgCl₂):

The buffer allows to optimize MgCl₂ concentration.

10 x Pol Buffer B (general application, up to 10 kb):

The buffer contains 15 mM MgCl₂ and is optimized for use with 0.2 mM of each dNTP.

10 x Pol Buffer C (colored):

10 x Pol Buffer B enriched with two gel tracking dyes and a gel loading reagent. The buffer enables direct loading of PCR products onto an agarose gel.

Quality Control:

All preparations are assayed for contaminating endonuclease, 3'-exonuclease, and nonspecific single- and double-stranded DNase activities. Typical preparations are greater than 95% pure, as judged by SDS polyacrylamide gel electrophoresis.

References:

1. Chien, A., Edgar, D.B. and Trela, J.M. (1976) *J. Bacteriol.* 127, 1550.
2. Kaledin, A.S., Sliusarenko, A.G. and Gorodetskii, S.I. (1980) *Biokhimiya* 45, 644.

This product is developed, designed and sold exclusively for research purposes and in vitro use only.

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Preparation of PCR Reaction:

| Component | Volume/reaction | Final concentration |
|-------------------------------------------------------------------|----------------------------------------------------------------|---------------------|
| 10 x Pol Buffer A or 10 x Pol Buffer B or 10 x Pol Buffer C | 5 µl | 1x |
| 25 mM MgCl ₂ | 2-10 µl when using 10 x Pol Buffer A | 1-5 mM |
| | 0-7 µl when using 10 x Pol Buffer B or 10 x Pol Buffer C | 1.5-5 mM |
| dNTP mix (5 mM each) | 2 µl | 0.2 mM of each dNTP |
| Upstream primer | Variable | 0.1-0.5 µM |
| Downstream primer | Variable | 0.1-0.5 µM |
| Taq DNA Polymerase, 5 U/µl | 0.25 µl | 1.25 U |
| Template DNA | Variable | <0.5 µg/50 µl |
| Sterile double-distilled water | Variable | - |
| Total volume | 50 µl | - |

Thermal Cycling Conditions:

| Step | Temperature | Time | Number of Cycles |
|-------------------------|-------------|------------|------------------|
| Initial Denaturation | 94-95°C | 2-5 min | 1 |
| Denaturation | 94-95°C | 15-60 s | 25-35 |
| Annealing | 50-68°C | 30-60 s | |
| Extension | 72°C | 1 min/1 kb | |
| Final Extension | 72°C | 7 min | 1 |
| Cooling | 4°C | Indefinite | 1 |

Notes:

1. Completely thaw and mix thoroughly all components of PCR reaction before use to avoid localized differences in salt concentration. It is especially important for magnesium solutions, because they form concentration gradient when frozen.
2. Prepare reaction mixes on ice, mix well.
3. Place reactions in a thermal cycler that has been preheated to 94-95°C.
4. Standard concentration of MgCl₂ in PCR reaction is 1.5 mM (as provided by the 1 x Pol Buffer B or the 1 x Pol Buffer C) when using 0.2 mM dNTP (each). In most cases this concentration will produce satisfactory results. However, in some cases, reaction may be improved by determining optimal concentration of MgCl₂.
5. The 10 x Pol Buffer C allows PCR reactions to be loaded directly onto an agarose gel without prior addition of a gel loading buffer. The buffer contains a gel loading reagent and two gel tracking dyes (a red dye and a yellow dye) that separate during electrophoresis. In a 1% agarose gel, the red dye migrates at the same rate as 600 bp DNA fragment and the yellow dye migrates faster than 20 bp. The dyes do not interfere with most downstream enzymatic applications, however it is recommended to purify PCR products prior enzymatic manipulation.
6. 1.25 U of Taq DNA Polymerase is recommended concentration of the enzyme per 50 µl amplification reaction. For most applications, enzyme will be in excess and will produce satisfactory results. Increased amounts of enzyme may generate artifacts like as smearing of bands, etc.
7. In most cases there is no need to add additives to the PCR reaction. For some difficult targets such as: GC-rich sequences, sequences with complex secondary structures additives such as DMSO can be included to improve amplification. Use DMSO in concentrations of 2-8%. The recommended starting DMSO concentration (if needed) is 3%.
8. As a general guide for how much template DNA to use, start with a minimum 10⁴ copies of the target sequence to obtain a signal in 25-35 cycles (i.e. 1 µg of 1 kb ds DNA equals 9.1 x 10¹¹ molecules, 1 µg of *E. coli* genomic DNA equals 2 x 10⁸ molecules, 1 µg of human genomic DNA equals 3 x 10⁵ molecules).

Notes:

1. Annealing temperature should be optimized for each primer set based on the primer T_m. Optimal annealing temperatures may be above or below the estimated T_m. As a starting point, use an annealing temperature 5°C below T_m.
2. When amplifying long PCR products (over 5 kb):
 - a. initial denaturation should be 2 min at 94°C
 - b. cycle denaturation should be 15-20 s at 94°C
 - c. use an elongation temperature of 68°C instead of 72°C.