

# PCR Using OneTaq<sup>®</sup> DNA Polymerase (NEB #M0480)

## Materials Required but not Supplied

### OneTaq<sup>®</sup> DNA Polymerase

- Deoxynucleotide (dNTP) Solution Mix (NEB #N0447)
- Nuclease-free Water (NEB #B1500)

## Overview

The Polymerase Chain Reaction (PCR) is a powerful and sensitive technique for DNA amplification. *Taq* DNA Polymerase is an enzyme widely used in PCR. OneTaq DNA Polymerase allows for greater amplification sensitivity across a wide variety of amplicons regardless of GC content. The following guidelines are provided to help ensure successful PCR using New England Biolabs' OneTaq DNA Polymerase. These guidelines cover most routine PCR. Specialized applications may require further optimization.

## Protocol

### Reaction setup:

1. Assemble all reaction components on ice.

Each component should be gently mixed before adding to the reaction in a sterile thin-walled PCR tube. Mix the entire reaction to ensure homogeneity. Collect all liquid to the bottom of the tube with a quick spin if necessary. Overlay the sample with mineral oil if using a PCR machine without a heated lid.

| COMPONENT                           | 25 µl reaction | 50 µl reaction | FINAL CONCENTRATION    |
|-------------------------------------|----------------|----------------|------------------------|
| 5X OneTaq Standard Reaction Buffer* | 5 µl           | 10 µl          | 1X                     |
| 10 mM dNTPs (NEB #N0447)            | 0.5 µl         | 1 µl           | 200 µM                 |
| 10 µM Forward Primer                | 0.5 µl         | 1 µl           | 0.2 µM                 |
| 10 µM Reverse Primer                | 0.5 µl         | 1 µl           | 0.2 µM                 |
| OneTaq DNA Polymerase               | 0.125 µl       | 0.25 µl        | 1.25 units/50 µl PCR** |
| Template DNA                        | variable       | variable       | < 1,000 ng             |
| Nuclease-free water                 | to 25 µl       | to 50 µl       |                        |

\*OneTaq GC Reaction Buffer and High GC Enhancer can be used for difficult amplicons.

\*\*For amplicons between 3–6 kb, use 2.5–5 units/50 µl rxn.

Notes: Gently mix the reaction. Collect all liquid to the bottom of the tube by a quick spin if necessary. Overlay the sample with mineral oil if using a PCR machine without a heated lid.

2. Quickly transfer the reactions to a thermocycler preheated to the denaturation temperature (94°C) and begin thermocycling.

#### Thermocycling Conditions for a Routine PCR:

| STEP                 | TEMP    | TIME            |
|----------------------|---------|-----------------|
| Initial Denaturation | 94°C    | 30 seconds      |
| 30 Cycles            | 94°C    | 15-30 seconds   |
|                      | 45-68°C | 15-60 seconds   |
|                      | 68°C    | 1 minute per kb |
| Final Extension      | 68°C    | 5 minutes       |
| Hold                 | 4-10°C  |                 |

\*Use of the [NEB Tm Calculator](#) is highly recommended.

## General Guidelines

### 1. Template:

Use of high-quality, purified DNA templates greatly enhances the success of PCR. Recommended amounts of DNA template for a 50 µl reaction are as follows:

| DNA              | Amount     |
|------------------|------------|
| Genomic          | 1 ng–1 µg  |
| Plasmid or Viral | 1 pg–10 ng |

### 2. Primers:

Oligonucleotide primers are generally 20–40 nucleotides in length and ideally have a GC content of 40–60%. Computer programs such as [Primer3](#) can be used to design or analyze primers. The final concentration of each primer in a reaction may be 0.05–1 µM, typically 0.2 µM.

### 3. Mg<sup>++</sup> and Additives:

Mg<sup>++</sup> concentration of 1.5–2.0 mM is optimal for most PCR products generated with OneTaq DNA Polymerase. The final Mg<sup>++</sup> concentration in 1X OneTaq Standard Reaction Buffer is 1.8 mM. This supports satisfactory amplification of most amplicons. However, Mg<sup>++</sup> can be further optimized in 0.2 mM increments using MgCl<sub>2</sub> ([NEB #B9021](#)).

Amplification of some difficult targets, like GC-rich sequences (50-65% GC), may be improved by the use of OneTaq GC Reaction Buffer. The final Mg<sup>++</sup> concentration in 1X OneTaq GC Reaction Buffer is 2.0 mM. To optimize the Mg<sup>++</sup> concentration of the OneTaq GC Reaction Buffer, MgSO<sub>4</sub> should be used ([NEB #B1003](#)).

For extremely difficult amplicons (>65% GC), 10–20% OneTaq High GC Enhancer can be added to reactions with OneTaq GC Reaction Buffer. The enhancer should not be used alone and typically increases yields when other conditions have failed.

### 4. Deoxynucleotides:

The final concentration of dNTPs is typically 200 µM of each deoxynucleotide.

## 5. **OneTaq DNA Polymerase Concentration:**

Generally, use OneTaq DNA Polymerase at a concentration of 25 units/ml (1.25 units/50 µl reaction) for amplicons up to 3 kb. The optimal concentration of OneTaq DNA Polymerase may range from 5–100 units/ml (0.25–5 units/50 µl reaction). For specialized applications, including 3–6 kb amplicons, 2.5–5 units/50 µl reaction is recommended. Note that in some cases increasing the amount of enzyme in the reaction can be inhibitory.

## 6. **Denaturation:**

An initial denaturation of 30 seconds at 94°C is sufficient for most amplicons from pure DNA templates. For difficult templates such as GC-rich sequences, a longer initial denaturation of 2–4 minutes at 94°C is recommended prior to PCR cycling to fully denature the template. With colony PCR, an initial 5-minute denaturation at 94°C is recommended to lyse cells.

During thermocycling, a 15–30 second denaturation at 94°C is recommended.

## 7. **Annealing:**

The annealing step is typically 15–60 seconds. The annealing temperature is based on the  $T_m$  of the primer pair and is typically 45–68°C. Annealing temperatures can be optimized by doing a temperature gradient PCR starting 5°C below the calculated  $T_m$ . The NEB [T<sub>m</sub> Calculator](#) is recommended to calculate an appropriate annealing temperature.

## 8. **Extension:**

The recommended extension temperature is 68°C. Extension times are generally 1 minute per kb. A final extension of 5 minutes at 68°C is recommended.

## 9. **Cycle Number:**

Generally, 25–35 cycles yield sufficient product. Up to 45 cycles may be required to detect low-copy-number targets.

## 10. **2-step PCR:**

When primers with annealing temperatures above 68°C are used, a 2-step thermocycling protocol (combining annealing and extension into one step) is possible.

### **Thermocycling conditions for a routine 2-step PCR:**

| STEP                 | TEMP   | TIME            |
|----------------------|--------|-----------------|
| Initial Denaturation | 94°C   | 30 seconds      |
| 30 Cycles            | 94°C   | 15-30 seconds   |
|                      | 68°C   | 1 minute per kb |
| Final Extension      | 68°C   | 5 minutes       |
| Hold                 | 4-10°C |                 |

## 11. **PCR Product:**

A significant portion of the PCR products generated using OneTaq DNA Polymerase contain dA overhangs at the 3' end; therefore the PCR products can be ligated to dT/dU-overhang vectors.

## Related Resources

- [T<sub>m</sub> Calculator](#)