

# Protocol for Q5® High-Fidelity 2X Master Mix

Protocols.io also provides an [interactive version of this protocol](#) where you can discover and share optimizations with the research community.

**Please note that protocols with Q5 High-Fidelity DNA Polymerase may differ from protocols with other polymerases. Conditions recommended below should be used for optimal performance.**

## Reaction Setup:

We recommend assembling all reaction components on ice and quickly transferring the reactions to a thermocycler preheated to the denaturation temperature (98°C). All components should be mixed prior to use.

| Component                      | 25 µl Reaction | 50 µl Reaction | Final Concentration |
|--------------------------------|----------------|----------------|---------------------|
| Q5 High-Fidelity 2X Master Mix | 12.5 µl        | 25 µl          | 1X                  |
| 10 µM Forward Primer           | 1.25 µl        | 2.5 µl         | 0.5 µM              |
| 10 µM Reverse Primer           | 1.25 µl        | 2.5 µl         | 0.5 µM              |
| Template DNA                   | variable       | variable       | < 1,000 ng          |
| Nuclease-Free Water            | to 25 µl       | to 50 µl       |                     |

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.

Transfer PCR tubes to a PCR machine and begin thermocycling.

## Thermocycling Conditions for a Routine PCR:

| STEP | TEMP | TIME |
|------|------|------|
|      |      |      |

|                      |          |                  |
|----------------------|----------|------------------|
| Initial Denaturation | 98°C     | 30 seconds       |
| 25–35 Cycles         | 98°C     | 5–10 seconds     |
|                      | *50–72°C | 10–30 seconds    |
|                      | 72°C     | 20–30 seconds/kb |
| Final Extension      | 72°C     | 2 minutes        |
| Hold                 | 4–10°C   |                  |

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

**Protocol changes as of 7/14 (after talking with Airola lab lady)**

**use 1 uL of plasmid and maybe 5 uL of 10mM primers  
make pcr final extension time 5 min instead of 2 min  
annealing temp 62 C should work.**

**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    |
|-------------|-----------|
| DNA Genomic | 1 ng–1 µg |

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 best results are typically seen when using each primer at a final concentration of 0.5  $\mu$ M in the reaction.

3.  $Mg^{++}$  and additives:

The Q5 High-Fidelity Master Mix contains 2.0 mM  $Mg^{++}$  when used at a 1X concentration. This is optimal for most PCR products generated with this master mix.

4. Deoxynucleotides:

The final concentration of dNTPs is 200  $\mu$ M of each deoxynucleotide in the 1X Q5 High-Fidelity Master Mix. Q5 High-Fidelity DNA Polymerase cannot incorporate dUTP and is not recommended for use with uracil-containing primers or templates.

5. Q5 High-Fidelity DNA Polymerase concentration:

The concentration of Q5 High-Fidelity DNA Polymerase in the Q5 High-Fidelity 2X Master Mix has been optimized for best results under a wide range of conditions.

6. Denaturation:

An initial denaturation of 30 seconds at 98°C is sufficient for most amplicons from pure DNA templates. Longer denaturation times can be used (up to 3 minutes) for templates that require it.

During thermocycling, the denaturation step should be kept to a minimum. Typically, a 5–10 second denaturation at 98°C is recommended for most templates.

7. Annealing:

Optimal annealing temperatures for Q5 High-Fidelity DNA Polymerase tend to be higher than for other PCR polymerases. The [NEB Tm Calculator](#) should be used to determine the annealing temperature when using this enzyme. Typically use a 10–30 second annealing step at 3°C above the  $T_m$  of the lower  $T_m$  primer. A temperature gradient can also be used to optimize the annealing temperature for each primer pair.

For high  $T_m$  primer pairs, two-step cycling without a separate annealing step can be used (see note 10).

8. Extension:

The recommended extension temperature is 72°C. Extension times are generally 20–30 seconds per kb for complex, genomic samples, but can be reduced to 10 seconds per kb for simple templates (plasmid, *E. coli*, etc.) or complex templates < 1 kb. Extension time can be increased to 40 seconds per kb for cDNA or long, complex templates, if necessary.

A final extension of 2 minutes at 72°C is recommended.

9. Cycle number:

Generally, 25–35 cycles yield sufficient product. For genomic amplicons, 30–35 cycles are recommended.

10. 2-step PCR:

When primers with annealing temperatures  $\geq 72^\circ\text{C}$  are used, a 2-step thermocycling protocol (combining annealing and extension into one step) is possible.

11. Amplification of long products:

When amplifying products > 6 kb, it is often helpful to increase the extension time to 40–50 seconds/kb.

12. PCR product:

The PCR products generated using Q5 High-Fidelity 2X Master Mix have blunt ends. If cloning is the next step, then blunt-end cloning is recommended. If T/A-cloning is preferred, the DNA should be purified prior to A-addition, as Q5 High-Fidelity DNA Polymerase will degrade any overhangs generated.