Phusion® High-Fidelity PCR Master Mix with HF Buffer



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100 reactions (50 μl vol) Lot: 0091206 RECOMBINANT Store at -20°C Exp: 6/13

Description: High Fidelity DNA Polymerases are important for applications in which the DNA sequence needs to be correct after amplification. Manufactured and quality controlled at New England Biolabs, Thermo Scientific® Phusion High-Fidelity DNA Polymerase offers both high fidelity and robust performance, and thus can be used for all PCR applications. Its unique structure, a novel Pyrococcus-like enzyme fused with a processivity-enhancing domain, increases fidelity and speed. Phusion DNA Polymerase is an ideal choice for cloning and can be used for long or difficult amplicons. With an error rate 50-fold lower than that of *Tag* DNA Polymerase and 6-fold lower than that of Pyrococcus furiosus DNA Polymerase (1), Phusion is one of the most accurate thermostable polymerases available. Phusion DNA Polymerase possesses $5' \rightarrow 3'$ polymerase activity, $3 \rightarrow 5$ exonuclease activity and will generate blunt-ended products.

Phusion High-Fidelity PCR Master Mix offers robust performance and can be used for all PCR applications. It is a 2X master mix consisting of Phusion DNA Polymerase, deoxynucleotides and reaction buffer that has been optimized and includes MgCl₂. All that is required is the addition of template, primers and water.

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

Applications:

- PCR
- Cloning
- · Long or Difficult Amplification
- High-throughput PCR

Reagents Supplied with Master Mix: 100% DMSO (500 µl)

Reaction Conditions: 1X Phusion High-Fidelity PCR Master Mix with HF Buffer, DNA template, 0.5 μ M primers and 3% DMSO (optional) in a total reaction volume of 50 μ l.

Phusion 1X Master Mix:

20 units/ml Phusion DNA polymerase 0.2 mM each dNTP 1X Phusion HF Buffer

Unit Definition: One unit is defined as the amount of enzyme that will incorporate 10 nmol of dNTP into acid insoluble material in 30 minutes at 74°C.

Unit Assay Conditions: 25 mM TAPS-HCl (pH 9.3 @ 25°C), 50 mM KCl, 2 mM MgCl $_2$, 1 mM β -mercaptoethanol, 200 μ M dNTPs including [³H]-dTTP and 400 μ g/ml activated Calf Thymus DNA.

Heat Inactivation: No

Quality Control Assays

7.5 kb Genomic DNA PCR: 30 cycles of PCR amplification in a 50 μ l reaction containing 50 ng genomic DNA with 1X Phusion High-Fidelity PCR Master Mix with HF Buffer and 1.0 μ M primers results in the expected 7.5 kb product.

20 kb Lambda DNA PCR: 22 cycles of PCR amplification in a 50 µl reaction containing 10 ng Lambda DNA with 1X Phusion High-Fidelity PCR Master Mix with HF Buffer and 1.0 µM primers results in the expected 20 kb product.

Note: Product specifications for individual components in the Phusion High-Fidelity PCR Master Mix with HF Buffer are available separately.

PCR

The following guidelines are provided to ensure successful PCR using Phusion Master Mixes. These guidelines cover routine PCR reactions. Amplification of templates with high GC content, high secondary structure, low template concentrations or long amplicons may require further optimization.

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 and centrifuged prior to use. It is important to add Phusion Master Mix last in order to prevent any primer degradation caused by the $3' \rightarrow 5'$ exonuclease activity. **Please note that**

protocols with Phusion DNA Polymerase may differ from protocols with other standard polymerases. As such, conditions recommended below should be used for optimal performance.

COMPONENT	25 μl REACTION	50 μl REACTION	FINAL CONCENTRATION
10 μM Forward Primer	1.25 μl	2.5 μ1	0.5 μΜ
10 μM Reverse Primer	1.25 μl	2.5 μl	0.5 μΜ
DMSO (optional)	(0.75 µl)	(1.5 µl)	(3%)
2X Phusion Master Mix	12.5 μl	25 μ1	1X
Template DNA	variable	variable	<250 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 from ice to a PCR machine with the block preheated to 98°C and begin thermocycling:

Thermocycling Conditions for a Routine PCR:

STEP	ТЕМР	TIME
Initial Denaturation	98°C	30 seconds
	98°C	5–10 seconds
25–35 Cycles	45–72°C	10-30 seconds
	72°C	15-30 seconds/kb
Final Extension	72°C	5–10 minutes
Hold	4°C	

General Guidelines:

1. Template:

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

DNA	AMOUNT	
Genomic	50 ng-250 ng	
Plasmid or Viral	1 pg-10 ng	

If the template DNA is obtained from a cDNA synthesis reaction, the volume added should be less than 10% of the total reaction volume.

Primers:

Oligonucleotide primers are generally 20–40 nucleotides in length and ideally have a GC content of 40–60%. Computer programs such as Primer3 (http://frodo.wi.mit.edu/primer3) can be used to design or analyze primers. The final concentration of each primer in a PCR reaction using the Phusion Master Mix may be 0.2–1 µM, while 0.5 µM is recommended.

Mg⁺⁺, deoxynucleotides and additives:
 At 1X concentration, Phusion Master Mix provides 1.5 mM MgCl₂ and 200 μM of each dNTP in the final reaction. Phusion cannot incorporate dUTP and is not recommended for use with uracil-containing primers or template.

Amplification of difficult targets, such as those with GC-rich sequences or secondary structure, may be improved by the presence of additives such as DMSO (included). A final concentration of 3% DMSO is recommended, although concentration can be optimized in 2% increments. It is important to note that if a high concentration of DMSO is used, the annealing temperature must be lowered as it decreases the primer T_m (2). Phusion DNA polymerase is also compatible with other additives such as formamide or glycerol.

4. Phusion DNA Polymerase Concentration: The concentration of Phusion DNA Polymerase in the Phusion PCR Master Mix has been optimized for best results under a wide range of conditions. If reactions are set up according to recommendations listed, the final concentration of Phusion DNA Polymerase in the reaction is 1 unit/50 µl or 0.4 units/20 µl.

5. Buffers

This Master Mix contains Phusion HF Buffer. HF buffer is recommended as the default buffer for high-fidelity amplification. For difficult templates, such as GC-rich templates or those with secondary structure, Phusion High-Fidelity PCR Master Mix with GC Buffer (NEB #M0532) can improve reaction performance. GC buffer should be used in experiments where HF buffer does not work.

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:

Annealing temperatures required for use with Phusion tend to be higher than with other PCR

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polymerases. The NEB T_m calculator (www.neb.com/TmCalculator) should be used to determine the annealing temperature when using Phusion. Typically, primers greater than 20 nucleotides in length anneal for 10–30 seconds at 3°C above the T_m of the lower T_m primer. If the primer length is less than 20 nucleotides, an annealing temperature equivalent to the T_m of the lower primer should be used. A temperature gradient can also be used to optimize the annealing temperature for each primer pair. For two-step cycling, the gradient can be set as high as the extension temperature.

For high T_m primer pairs, two-step cycling without a separate annealing step can be used.

8. Extension:

The recommended extension temperature is 72°C. Extension times are dependent on amplicon length and complexity. Generally, an extension time of 15 seconds per kb can be used. For complex amplicons, such as genomic DNA, an extension time of 30 seconds per kb is recommended. Extension time can be increased to 40 seconds per kb for cDNA templates, if necessary.

 Cycle number: Generally, 25–35 cycles yields sufficient product.

10. 2-step PCR:

When primers with annealing temperatures $\geq 72^{\circ}\text{C}$ are used, a 2-step thermocycling protocol is recommended.

Thermocycling Conditions for a Routine 2-Step PCR:

STEP	ТЕМР	TIME
Initial Denaturation	98°C	30 seconds
25–35 Cycles	98°C	5–10 seconds
	72°C	15–30 seconds/kb
Final Extension	72°C	5–10 minutes
Hold	4°C	

11. PCR product:

The PCR products generated using Phusion DNA Polymerase have blunt ends; if cloning is the next step, then blunt-end cloning is recommended. If T/A-cloning is preferred, then DNA should be purified prior to A-addition, as Phusion DNA Polymerase will degrade any overhangs generated.

Addition of an untemplated -dA can be done with *Taq* DNA Polymerase (NEB #M0267) or Klenow exo⁻ (NEB #M0212).

References:

- 1. Frey, B. and Suppman, B. (1995) *BioChemica*, 2, 34–35.
- 2. Chester, N. and Marshak, D.R. (1993) *Analytical Biochemistry*, 209, 284–290.

Companion Products Sold Separately:

Phusion HF Buffer Pack

#B0518S 6.0 ml

Phusion GC Buffer Pack

#B0519S 6.0 ml

Detergent-free Phusion HF Buffer Pack

#B0520S 6.0 ml

Detergent-free Phusion GC Buffer Pack

#B0521S 6.0 ml

Magnesium Chloride (MgCl₂) Solution #B9021S 6.0 ml

Phusion High-Fidelity PCR Kit

#E0553S 50 reactions

#E0553L 200 reactions

Phusion High-Fidelity DNA Polymerase

#M0530S 100 units

#M0530L 500 units

Phusion High-Fidelity PCR Master Mix

with GC Buffer

#M0532S 100 reactions

#M0532L 500 reactions

Deoxynucleotide Solution Set #N0446S 25 umol of each

Deoxynucleotide Solution Mix

#N0447S 8 µmol of each #N0447L 40 µmol of each Phusion® DNA Polymerase was developed by Finnzymes Oy, now a part of Thermo Fisher Scientific. This product is manufactured by New England Biolabs, Inc. under agreement with, and under the performance specifications of Thermo Fisher Scientific.

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