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# Opti*Taq* PCR Master Mix (2x)

2x Opti*Taq* PCR Master Mix

*Taq* DNA Polymerase *Pyrococcus sp.* DNA Polymerase

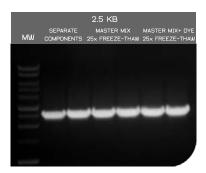
# Cat. No. Size E2910-01 100 reactions 50 µl each

E2910-02 200 reactions 50 μl each E2910-03 500 reactions 50 μl each

## 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 min at 74°C. The reaction conditions are: 50 mM Tris-HCI (pH 9.0 at 25°C), 50 mM NaCI, 5 mM MgCl<sub>2</sub>, 200  $\mu$ M each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and I<sup>3</sup>H1dTTP), 10  $\mu$ g activated calf thymus DNA and 0.1 mg/ml BSA in a final volume of 50  $\mu$ l.

**Storage Conditions:** Store at -20°C for long- term storage (more than 12 months) or at 4°C for up to 2 months.



# Figure 1: PCR amplification using EURx Opti*Taq* DNA Polymerase.

A 2.5 kb PCR amplicon was amplified with Opti*Taq* DNA Polymerase either in stand-alone or in master mix format, respectively.

Lane MW: molecular size marker-Perfect 1 kb DNA Ladder (Cat. No. E3130).

Lanes POL (1,2): PCR amplification reactions using 1.25 U Opti*Taq* DNA Polymerase, Pol Buffer B and dNTPs Lanes mM (3,4): PCR amplification reactions using Opti*Taq* PCR Master Mix (2x), after 25 freeze-thaw cycles.

Lanes MM+COL (5,6): PCR amplification reactions using Opti*Taq* PCR Master Mix (2x) and 10 x Color Load, after 25 freezethaw cycles. Opti*Taq* DNA Polymerase Master mix, with stable and reproducible high performance even after more than 25 freeze-thaw cycles or more than 12 months of storage.

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## Description:

- → Opti*Taq* PCR Master Mix (2x) is a ready-to-use solution containing Opti*Taq* DNA Polymerase, optimized reaction buffer, MgCl<sub>2</sub> and dNTPs.
- → Use of Opti*Taq* PCR Master Mix (2x) saves time, increases reproducibility (due to minimizing calculation and pipetting errors) and reduces contamination risk (due to fewer pipetting steps) during PCR set-up.
- → Opti*Taq* DNA Polymerase is a modified and optimized thermostable enzymes blend containing *Thermus aquaticus* DNA polymerase, *Pyrococcus sp.* DNA polymerase and enhancing factors.
- → The enzymes blend exhibits  $3^{\prime} \rightarrow 5^{\prime}$  proofreading activity, resulting in considerably higher PCR fidelity, processivity and yield than possible with proofreading-deficient *Taq* DNA polymerase (1).
- ➔ Ultrapure, recombinant enzymes are used to prepare Opti*Taq* DNA Polymerase.
- → Opti Taq PCR Master Mix is stable with respect to multiple cycles of freezing and thawing. Even after more than 25 freeze-thaw cycles, no decline in performance is detected.
- → Same performance as standalone OptiTaq DNA Polymerase (Cat. No. E2600). Additionally, aliquots of clean nuclease free water are supplied, allowing the setup of PCR reactions without the risk of introducing unwanted DNA through contaminated water.
- ➔ For optional use, a 10 x Color load buffer is supplied. The Color Load buffer allows to directly load PCR products to agarose gels.
- → The enzyme replicates DNA at 74°C and exhibits a half-life of 40 min at 95°C (1,2).
- → Maintains the 5' $\rightarrow$ 3' exonuclease activity.
- → Adds an extra A overhang to some, but not all 3'-ends. PCR amplicons are suitable for both, blunt and TA-cloning.
- → Opti*Taq* PCR Master Mix (2x) is recommended for use in PCR and primer extension reactions at elevated temperatures to obtain a wide range of DNA products up to 20 kb.

## Taq PCR Master Mix (2x) Package Contents:

- 1. Opti*Tag* PCR Master Mix (2x)
- 2. Water, nuclease free
- 3. 10 x Color Load Buffer

## Opti*Taq* PCR Master Mix (2x):

Supplied in 2 x Pol Buffer B containing 3 mM MgCl<sub>2</sub> and 0.4 mM of each dNTP. Final concentrations: 1.5 mM MgCl<sub>2</sub> and 0.2 mM of each dNTP.

### 10 x Color Load:

10 x Color Load contains two gel tracking dyes and a gel loading reagent. It 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. Cline J. et al. (1996) Nucleic Acid Res. 24 (18) 3546-3551.

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# Opti*Taq* PCR Master Mix (2x) PCR PROTOCOL (1)

#### PCR Reaction Assembly

## Preparation of PCR Reaction:

| Component                               | Volume / Reaction | Final Concentration                                |  |
|---|-------------------|--|--|
| Opti <i>Taq</i> PCR master<br>Mix (2 x) | 25 µl             | 1.25 U Opti <i>Taq</i> DNA<br>Polymerase           |  |
|   |                   | l x Reaction Buffer<br>(1.5 mM MgCl <sub>2</sub> ) |  |
|   |                   | 0.2 mM of each dNTP                                |  |
| Upstream primer                         | Variable          | 0.1-0.5 μM   |  |
| Downstream primer                       | Variable          | 0.1-0.5 μM   |  |
| Optional:<br>10 x Color Load            | 5 µl              | 1 x  |  |
| Template DNA                            | Variable          | <0.5 µg/50 µl                                      |  |
| Sterile double-<br>distilled water      | Το 50 μl          | -  |  |
| Total volume                            | 50 µl             | -  |  |

General formula for calculating total gene  $\prime$  genome copy numbers from the total amount of template DNA:

Template DNA copy number [molecules] =

$$\frac{DNA \ amount \ [ng] \cdot 6.022 x 10^{23} \ [molecules \ mol^{-1}]}{Genomic \ DNA \ length \ [kb] \cdot 616 \ [g \ mol^{-1} \ bp^{-1}]} \cdot \frac{10^{-3} \ [kb \ bp^{-1}]}{10^{9} \ [ng \ g^{-1}]}$$

Optimum: 10<sup>4</sup> template DNA copies Maximum: 0.5 µg template DNA or less

(MW per bp: siehe Dolezel et al. Cytometry, 2003, Vol. 51A, 2, 127-8)

#### Notes

- First Law of PCR: PCR is a sort of homeopathic process. It works best, as long as *all* components are assembled in homeopathic doses only.
- . **Concentration Differences.** Thaw, gently vortex and centrifuge Opti*Taq* PCR Master Mix (2x) and primers to avoid localized differences in salt concentration.
- On Ice. PCR reactions should be set up on ice.

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- Primer Mix. Primers can be either added separately or as primer mix prepared separately.
- Mix Template. Vortex the samples and briefly spin down.
- . Preheat Cycler. Place reactions in a thermal cycler that has been preheated to  $94-95^{\circ}C$ .
- .  $MgCl_2$  Standard concentration of  $MgCl_2$  in PCR reactions is 1.5 mM (as provided in the 1 x Opti *Taq* Master Mix) when using 0.2 mM dNTP (each). In most cases this concentration will produce satisfactory results. However, if higher MgCl\_2 concentrations are required, prepare a 25 mM MgCl\_2 stock solution (or request us to ship an aliquot along with your order) and add an appropriate amount to the reaction. Adding 1  $\mu$ I of a 25 mM MgCl\_2 solution to a total reaction volume of 50  $\mu$ I will add 25 nmol MgCl\_2 and thus increase total MgCl\_2 reaction concentration in 0.5 mM.
- 8. Color Load. Use of the 10 x Color Load allows PCR reactions to be loaded directly onto an agarose gel without prior addition of a gel loading buffer. The 10 x Color Load contains a gel loading reagent and two gel tracking dyes (a red dye and an 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 to enzymatic manipulation.
- 9. PCR additives / PCR Enhancers. In most cases there is no need to add any 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% [v/v]. The recommended starting DMSO concentration (if required) is 3% [v/v].
- 10. **Template Copies.** As a general guide for how much template DNA to use, start with a minimum  $10^4$  copies of the target sequence to obtain a signal in 25-35 cycles (i.e.  $1 \mu g$  of 1 kb ds DNA equals  $9.8 \times 10^{11}$  molecules,  $1 \mu g$  of *E. coli* genomic DNA equals  $2 \times 10^{9}$  molecules,  $1 \mu g$  of human genomic DNA equals  $3 \times 10^{5}$  molecules).

General formula for calculating total gene /genome copy number from template DNA mass:

copy number [molecules] = (DNA amount [ng] x  $6.022 \times 10^{23}$ [molecules mol<sup>-1</sup>]) / (length [bp] x  $1x10^9$  [ng g<sup>-1</sup>] x 616 [g bp<sup>-1</sup>]) (MW per bp: see Dolezel et al. Cytometry, 2003, Vol. 51A, 2, 127-8)

- 11. Template DNA for Long Range PCR. For long range PCR use: 50-500 ng of human genomic DNA, 0.1-10 ng of bacterial DNA, phage DNA or plasmid DNA.
- High Quality Template DNA: Ensure that template DNA is of sufficiently high quality. Use only high-molecular-weight DNA, when amplifying long PCR targets (over 20-50 kb, depending on the amplicon length).
- 13. Storage of High Molecular Weight DNA: Complex genomic DNA should be stored at 2-8°C. Avoid vortexing genomic DNA.
- 14. Thin Walled Reaction Cups: Use only thin-walled 0.2 ml tubes performing long PCR amplification.

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# Opti*Taq* PCR Master Mix (2x) PCR PROTOCOL (2)

#### Thermal Cycling Conditions

#### Thermal Cycling Conditions for Products 0.1-10 kb in Size:

| Step                    | Tempera-<br>ture | Time       | Number of<br>Cycles |
|-------------------------|------------------|------------|---------------------|
| Initial<br>Denaturation | 93-95°C          | 2-5 min    | 1                   |
| Denaturation            | 93-95°C          | 15-30 s    | 25-35               |
| Annealing               | 50-68°C          | 30 s       |                     |
| Extension               | 72°C or<br>68°C  | l min/l kb |                     |
| Final Extension         | 72°C or<br>68°C  | 7 min      | 1                   |
| Cooling                 | 4°C              | Indefinite | 1                   |

### Thermal Cycling Conditions for Products Larger Than 10 kb in Size:

| Step                    | Tempera-<br>ture     | Time   | Number of<br>Cycles |
|-------------------------|----------------------|--|---------------------|
| Initial<br>Denaturation | 92-94 <sup>°</sup> C | 2 min  | 1                   |
| Denaturation            | 92-94°C              | 10-15 s  | 10                  |
| Annealing               | 60-68°C              | 30 s   |                     |
| Extension               | 68°C                 | 1 min/1 kb                                     |                     |
| Denaturation            | 92-94°C              | 10-15 s  | 15-25               |
| Annealing               | 60-68°C              | 30 s   |                     |
| Extension               | 68°C                 | 1 min/1 kb<br>+20 s per<br>additional<br>cycle |                     |
| Final Extension         | 68°C                 | 7 min  | 1                   |
| Cooling                 | 4°C                  | Indefinite                                     | 1                   |

#### Notes

- 1. Annealing: Annealing temperature should be optimized for each primer set based on the primer  $T_{\rm m}$ . Optimal annealing temperatures may be above or below the estimated  $T_{\rm m}$ . As a starting point, use an annealing temperature  $5\,^\circ\text{C}$  below  $T_{\rm m}$ .
- Long PCR Primer Requirements: Typical primers for long PCR amplification have a length of 22-34 bp and should have annealing temperatures above 60°C to enhance reaction specificity.
- 3. Long PCR Short Denaturation Steps: When amplifying long PCR products, keep denaturation steps as short as possible and denaturation temperature as low as possible (try not to exceed 2 min at 94°C during initial denaturation and 10-15 s at 94°C during cycle denaturation step). Sometimes fragments with high GC-content need higher denaturation temperatures, but keep in mind that the yield increases when denaturation temperature / duration is decreased.
- Long PCR Low Elongation Temperature: For PCR products exceeding 5 kb in size, an elongation temperature of 68°C is strongly recommended.
- Long PCR Extended Elongation Period: For PCR products exceeding 10 kb in length, an elongation of the extension step (+20 s in each additional cycle, starting from the 11<sup>th</sup> cycle) is strongly recommended due to loss of processivity of the enzymes blend.
- 6. Minimize Time for Cooling Step (+4°C). Overly prolonged cooling time to 4°C puts a hard strain on your PCR cycler's Peltier elements. To prevent an all too early wearout, minimize the time for the last cooling step at 4°C and, upon completion of the PCR reaction, remove PCR samples from the cycler as soon as possible. Alternately, choose +8°C as final cooling temperature, but

Alternately, choose +8°C as final cooling temperature, but take care to remove the PCR samples as soon as possible for minimizing PCR amplicon exposure time to 3'-5' exonuclease activity of *Pyrococcus sp.* DNA polymerase.

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