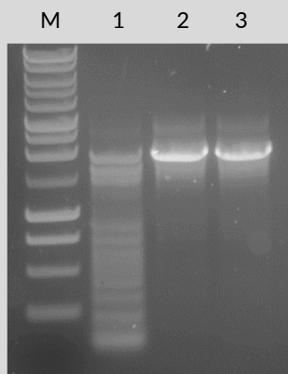


**onTaq**  
**DNA Polymerase** HOT START

| Cat. No. | size       |
|----------|------------|
| E2713-01 | 200 units  |
| E2713-04 | 500 units  |
| E2713-02 | 1000 units |
| E2713-03 | 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 MgCl<sub>2</sub>, 200 µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [<sup>3</sup>H]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.



**PCR amplification using EURx onTaq DNA Polymerase.**

A 2 kb amplicon of the human  $\beta$ -globin gene was amplified using EURx onTaq DNA Polymerase, 10 x Pol Buffer B and 0.2 mM dNTPs in 50 µl reaction volume.

Lane M: molecular size marker-Perfect Plus 1 kb DNA Ladder.

Lane 1: PCR amplification reaction using 1.25U Taq DNA Polymerase. Reaction was incubated 30 min at 25°C before PCR.

Lane 2, 3: PCR amplification reactions using 1.25 U onTaq DNA Polymerase. Reactions were incubated 30 min at 25°C before PCR.

**Description:**

- onTaq DNA Polymerase is a modified "hot start" enzyme which is blocked at moderate temperatures and allows room temperature reaction setup.
- The polymerase activity is restored during 10 min initial denaturation step.
- onTaq DNA Polymerase provides very tight inhibition of the polymerase activity at moderate temperatures.
- Use of the onTaq DNA Polymerase allows for the increase of PCR specificity, sensitivity and yield in comparison to the conventional PCR assembly method.
- Both increased specificity and reduced mispriming improve multiplex PCR.
- Thermostable onTaq DNA Polymerase replicates DNA at 72°C and exhibits a half-life of 40 min at 95°C.
- Catalyzes the polymerization of nucleotides into duplex DNA in the 5'→3' direction in the presence of magnesium ions.
- Contains the 5'→3' exonuclease activity.
- Lacks the 3'→5' exonuclease activity.
- Adds extra A at the 3' ends.
- onTaq 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 9.0 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 MgCl<sub>2</sub>):**

The buffer allows to optimize MgCl<sub>2</sub> concentration.

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

The buffer contains 15 mM MgCl<sub>2</sub> 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.

## Preparation of PCR Reaction:

| Component   | Volume/reaction  | Final concentration |
|---|--|---------------------|
| 10 x Pol Buffer A or<br>10 x Pol Buffer B or<br>10 x Pol Buffer C | 5 µl   | 1x                  |
| 25 mM MgCl <sub>2</sub>   | 2-10 µl when using<br>10 x Pol Buffer A                        | 1-5 mM              |
|   | 0-7 µl when using<br>10 x Pol Buffer B or<br>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          |
| onTaq DNA<br>Polymerase, 2.5 U/µl                                 | 0.5 µl   | 1.25 U              |
| Template DNA  | Variable   | <0.5 µg/50 µl       |
| Sterile double-distilled<br>water                                 | Variable   | -                   |
| Total volume  | 50 µl  | -                   |

## 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 at room temperature. Use of onTaq DNA Polymerase allows room temperature reaction setup. Mix well.
3. Reactions can be placed in a room temperature thermal cycler.
4. Standard concentration of MgCl<sub>2</sub> in PCR reaction is 1.5 mM (as provided with 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 MgCl<sub>2</sub>.
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. 1.25U of onTaq 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 10<sup>4</sup> 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 10<sup>11</sup> molecules, 1 µg of *E. coli* genomic DNA equals 2 x 10<sup>8</sup> molecules, 1 µg of human genomic DNA equals 3 x 10<sup>5</sup> molecules).

## Thermal Cycling Conditions

| Step                 | Temperature | Time       | Number of Cycles |
|----------------------|-------------|------------|------------------|
| Initial Denaturation | 95°C        | 10 min     | 1                |
| Denaturation         | 94°C        | 30 s       | 25-35            |
| Annealing            | 50-68°C     | 30 s       |                  |
| Extension            | 72°C        | 1 min/1 kb |                  |
| Final Extension      | 72°C        | 7 min      | 1                |
| Cooling              | 4°C         | Indefinite | 1                |

## Notes:

1. onTaq is activated by a 10-minute initial denaturation step at 95°C.
2. Annealing temperature should be optimized for each primer set based on the primer T<sub>m</sub>. Optimal annealing temperatures may be above or below the estimated T<sub>m</sub>. As a starting point, use an annealing temperature 5°C below T<sub>m</sub>.
3. When amplifying long PCR products (over 5 kb) use an elongation temperature of 68°C instead of 72°C.

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

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