

# Expand Long Template PCR System

Deoxynucleoside-triphosphate: DNA deoxynucleotidyltransferase, *E.C. 2.7.7.7*

**Cat. No. 11 681 834 001** 150 U  
**Cat. No. 11 681 842 001** 720 U (2 × 360 U)  
**Cat. No. 11 759 060 001** 3,600 U (10 × 360 U)

 **Version 25**

Content version: February 2017

Store the kit at -15 to -25°C

## 1. What this Product Does

### Number of Tests

The kit is designed for

- approx. 38 reactions (Cat. No. 11 681 834 001)
- approx. 190 reactions (Cat. No. 11 681 842 001)
- approx. 950 reactions (Cat. No. 11 759 060 001)

with a final reaction volume of 50 µl each.

### Kit Contents

| Vial | Label                                | Contents   |
|------|--------------------------------------|--|
| 1    | Expand Long Template Enzyme mix      | <ul style="list-style-type: none"> <li>• 30 µl (150 U pack size)</li> <li>• 2 × 72 µl (720 U pack size)</li> <li>• 10 × 72 µl (3,600 U pack size)</li> </ul> Enzyme storage buffer: 20 mM Tris-HCl, pH 7.5 (25°C), 100 mM KCl, 1 mM dithiothreitol (DTT), 0.1 mM EDTA, 0.5% Nonidet P40 (v/v), 0.5% Tween 20 (v/v), 50% glycerol (v/v) |
| 2    | Expand Long Template buffer <b>1</b> | 10 × conc. with 175 mM MgCl <sub>2</sub> <ul style="list-style-type: none"> <li>• 1 ml (150 U pack size)</li> <li>• 2 × 1 ml (720 U pack size)</li> <li>• 10 × 1 ml (3,600 U pack size)</li> </ul>   |
| 3    | Expand Long Template buffer <b>2</b> | 10 × conc. with 275 mM MgCl <sub>2</sub> <ul style="list-style-type: none"> <li>• 1 ml (150 U pack size)</li> <li>• 2 × 1 ml (720 U pack size)</li> <li>• 10 × 1 ml (3,600 U pack size)</li> </ul>   |
| 4    | Expand Long Template buffer <b>3</b> | 10 × conc. with 275 mM MgCl <sub>2</sub> and detergents <ul style="list-style-type: none"> <li>• 1 ml (150 U pack size)</li> <li>• 2 × 1 ml (720 U pack size)</li> <li>• 10 × 1 ml (3,600 U pack size)</li> </ul>  |

### Storage and Stability

Store the kit at -15 to -25°C. When properly stored, the kit is stable until the expiration date printed on the label.

⚠ Always thaw and equilibrate all buffers at +37°C to +56°C before use. Vortex thoroughly. If crystals have formed, incubate at +37°C to +56°C until they are dissolved.

Once the kit is opened, store the kit components as described in the following table:

| Vial | Label                                | Storage  |
|------|--------------------------------------|--|
| 1    | Expand Long Template Enzyme mix      | <ul style="list-style-type: none"> <li>• Aliquot and store at -15 to -25°C.</li> <li>• Avoid repeated freezing and thawing!</li> </ul> |
| 2    | Expand Long Template buffer <b>1</b> | Store at -15 to -25°C.   |
| 3    | Expand Long Template buffer <b>2</b> |  |
| 4    | Expand Long Template buffer <b>3</b> |  |

### Additional Equipment and Reagents Required

- Thermal block cycler (*e.g.*, Applied Biosystems GeneAmp PCR System 9600)
- 0.2 ml thin-walled PCR tubes
- Sterile reaction (Eppendorf) tubes for preparing master mixes and dilutions

### Application

Expand Long Template PCR System is a special enzyme mix that contains thermostable Taq DNA polymerase and Tgo DNA polymerase (1), a thermostable DNA polymerase with proofreading activity. This powerful polymerase mixture produces a high yield of PCR product from genomic DNA. Expand Long Template PCR System is optimized to efficiently amplify large genomic DNA fragments around 20 kb long. Due to the inherent 3'-5' exonuclease ("proofreading") activity of Tgo DNA polymerase, Expand Long Template PCR System copies DNA three times as accurately than Taq DNA polymerase.

### Enzyme Properties

|   |   |
|---|---|
| Volume activity                         | 5 U/µl  |
| Error rate <sup>1</sup>                 | three times more accurate than Taq DNA polymerase   |
| Optimal enzyme concentration            | varies from 0.5 to 5.0 U per 50 µl reaction   |
| Standard enzyme concentration           | 3.75 U per 50 µl reaction   |
| Optimal elongation temperature          | 68°C  |
| Standard Mg <sup>2+</sup> concentration | 1.75 mM (as MgCl <sub>2</sub> ) when using 350 µM of each dNTP and 2.75 mM (as MgCl <sub>2</sub> ) when using 500 µM of each dNTP |
| PCR product size                        | around 20 kb  |
| Repair of mismatched primers at 3' end  | yes, due to the 3'-5' exonuclease activity of the proofreading polymerase   |
| Incorporation of modified nucleotides   | accepts modified nucleotides like DIG-dUTP, Biotin-dUTP and Fluorescein-dUTP  |
| Incorporation of dUTP                   | No  |

<sup>1</sup> Relative fidelity determined by the lacl assay (2).

## 2. How To Use this Product

### 2.1 Before You Begin

#### General considerations

The optimal conditions (incubation times and temperatures, concentrations of enzyme, template DNA, Mg<sup>2+</sup> concentration) depend on the system used and must be determined for each system. In particular, to ensure optimal reaction efficiency, you should titrate the Mg<sup>2+</sup> concentration and the amount of enzyme used per assay. If you are developing a new assay, you should test all three amplification systems to find the one that gives the best results.

As a starting point for developing your assays, use the following guidelines:

- Optimal enzyme concentration: 0.5 to 5.0 U/μl. The recommended starting concentration is 3.75 U (0.75 μl).
- The quality of the template has a tremendous effect on the success of the PCR.
- dNTP concentration: Always use balanced concentrations of all four dNTPs. The final concentration of each dNTP should be between 350 and 500 μM.
- ⚠ If you increase the concentration of dNTP, also increase the Mg<sup>2+</sup> concentration.
- The optimal buffer for dilution of the template DNA is either Water, PCR Grade\* or 5 to 10 mM Tris (pH 7-8). Avoid dissolving the template in TE buffer because EDTA chelates Mg<sup>2+</sup>.
- Reactions do not usually require additives. Nevertheless, in some cases you can improve yield by adding up to 100 μg/ml bovine serum albumin (BSA), 0.1% Tween 20 (v/v) or 1 to 2% DMSO.
- Use 0.2 ml thin-walled PCR tubes.
- Keep denaturation steps as short as possible and denaturation temperature as low as possible.

#### Sample Material

Template DNA, *e.g.* human genomic DNA\*

- ⚠ The quality of the template has a tremendous effect on the success of the PCR.

### 2.2 Setting up the Experiment

#### Preparation of Reaction Mixes

- Always thaw and equilibrate all buffers at +37°C to +56°C before use. Vortex thoroughly.
- Briefly vortex and centrifuge all reagents before starting.
- Set up the reaction components in a sterile microfuge tube (on ice).
- After pipetting the last reaction component, start the reactions immediately. Do not store complete reaction mixes on ice.
- If the initial reaction produces too many primer dimers, try using two separate master mixes. (Master Mix 1 contains dNTPs, primers, and template DNA; Master Mix 2 contains buffer and enzyme. Final volume of each mix: 25 μl. For final conc. of each component, see below.) Add these two to the tube on ice, then just before starting the reaction, vortex the tubes to produce a homogeneous reaction mix.

- ⚠ If the amplification system recommended for a given fragment length does not give satisfactory amplification, repeat the experiment using one of the other two possible amplification systems.

| Amplification of human gen. DNA       | 0.5 – 9 kb System 1 |                           | 9 – 12 kb System 2 |                           | > 12 kb System 3 |                           |
|---------------------------------------|---------------------|---------------------------|--------------------|---------------------------|------------------|---------------------------|
|                                       | Vol.                | Final conc.               | Vol.               | Final conc.               | Vol.             | Final conc.               |
| add PCR grade water                   | 50 μl               |                           | 50 μl              |                           | 50 μl            |                           |
| dATP*10 mM                            | 1.75 μl             | 350 μM                    | 2.5 μl             | 500 μM                    | 2.5 μl           | 500 μM                    |
| dCTP*10 mM                            | 1.75 μl             | 350 μM                    | 2.5 μl             | 500 μM                    | 2.5 μl           | 500 μM                    |
| dGTP*10 mM                            | 1.75 μl             | 350 μM                    | 2.5 μl             | 500 μM                    | 2.5 μl           | 500 μM                    |
| dTTP*10 mM                            | 1.75 μl             | 350 μM                    | 2.5 μl             | 500 μM                    | 2.5 μl           | 500 μM                    |
| Downstream primer                     | x μl                | 300 nM                    | x μl               | 300 nM                    | x μl             | 300 nM                    |
| Upstream primer                       | y μl                | 300 nM                    | y μl               | 300 nM                    | y μl             | 300 nM                    |
| Components                            | Vol.                | Final conc.               | Vol.               | Final conc.               | Vol.             | Final conc.               |
| 10x PCR buffer with MgCl <sub>2</sub> | 5 μl                | 1.75 mM MgCl <sub>2</sub> | 5 μl               | 2.75 mM MgCl <sub>2</sub> | 5 μl             | 2.75 mM MgCl <sub>2</sub> |
| template DNA                          | z μl                | up to 500 ng genom. DNA   | z μl               | up to 500 ng genom. DNA   | z μl             | up to 500 ng genom. DNA   |
| Expand Long Template Enzyme mix       | 0.75 μl             |                           | 0.75 μl            |                           | 0.75 μl          |                           |

\* Instead of using single dNTP solutions, you may use the PCR Grade Nucleotide Mix.

#### Thermal Cycling

Place samples in the thermal block cycler, and cycle according to the thermal profile below. Gradually increasing extension time ensures a higher yield of amplification products.

|                      | Temperature               | Time  | Cycles  |
|----------------------|---------------------------|---|---------|
| Initial Denaturation | +92 to +94°C              | 2 min   | 1       |
| Denaturation         | +92 to +94°C              | 10 s  | 10      |
| Annealing            | +45 to +65°C <sup>a</sup> | 30 s  |         |
| Elongation           | 68°C                      | 45 s – 30 min <sup>b</sup>  |         |
| Denaturation         | +92 to +94°C              | 15 s  | 15 – 25 |
| Annealing            | +45 to +65°C <sup>a</sup> | 30 s  |         |
| Elongation           | 68°C                      | 45 s – 30 min <sup>b</sup> + 20 s cycle elongation for each successive cycle <sup>c</sup> |         |
| Final Elongation     | 68°C                      | 7 min   | 1       |
| Cooling              | 4°C                       | unlimited time  |         |

<sup>a</sup> Optimal annealing temperature depends on the melting temperature of the primers and the system used.

<sup>b</sup> Elongation time depends on fragment length: Use 2 min for up to 3 kb, 4 min for 6 kb, 8 min for 10 kb, 15 min for 20 kb, 20 min for 30 kb.

<sup>c</sup> For example, cycle no. 11 is 20 s longer than cycle 10, cycle no. 12 is 40 s longer than cycle 10, cycle no. 13 is 60 s longer than cycle 10, etc.

⚠ The thermal profile above was developed for the Applied Biosystems GeneAmp PCR System 9600. Other thermal block cyclers may require a different profile. Long range PCR in general is sensitive to even minute differences between ramping or heat transfer rates of different thermal block cyclers. Therefore, always develop and run your Expand Long Template PCR experiment on the same thermal block cycler. If you switch to a different thermal block cycler, adjust the reaction conditions and thermal profile.

### 3. Troubleshooting

#### Little or no PCR product

| Possible Cause                               | Recommendation   |  |
|--|--|--|
| Difficult template                           | Try amplification system 2 or 3 first, even if you have short amplicons.   |  |
| Poor DNA template quality                    | Check quality and concentration of template: <ul style="list-style-type: none"> <li>Analyze an aliquot on an agarose gel to check for possible degradation.</li> <li>Include a control reaction using a known template under established PCR conditions.</li> <li>Check or repeat purification of template.</li> </ul> |  |
| Enzyme concentration too low                 | Increase the amount of enzyme mix in 0.5 U steps.  |  |
| MgCl <sub>2</sub> concentration too low      | Increase the MgCl <sub>2</sub> concentration in 0.25 mM steps. (Minimum concentration is 1.75 mM MgCl <sub>2</sub> .)  |  |
| Cycle conditions not optimal                 | <ul style="list-style-type: none"> <li>Reduce annealing temperature.</li> <li>Increase number of cycles.</li> <li>Be sure to perform the final elongation step.</li> </ul>   |  |
| Primer design not optimal                    | Design alternative primers.  |  |
| Primer concentration not optimal             | <ul style="list-style-type: none"> <li>Both primers must be present in the reaction at the same concentration.</li> <li>Titrate primer concentration (0.3 – 0.6 μM).</li> </ul>  |  |
| Annealing temperature too high               | <ul style="list-style-type: none"> <li>Reduce annealing temperature. (Minimum annealing temperature is 45°C.)</li> <li>Determine the optimal annealing temperature by touch-down PCR.</li> </ul>   |  |
| Primer quality or storage problems           | <ul style="list-style-type: none"> <li>If you are using an established primer pair, check their performance under established PCR conditions (with a control template).</li> <li>Make sure primers are not degraded.</li> <li>Always store primers at –15 to –25°C.</li> </ul>   |  |
| Multiple bands or background smear           | Annealing temperature too low  | Increase annealing temperature according to primer length.   |
|  | Primer design or concentration not optimal   | <ul style="list-style-type: none"> <li>Review primer design.</li> <li>Titrate primer concentration (0.3 – 0.6 μM).</li> <li>Both primers must be present in the reaction at the same concentration.</li> <li>Perform nested PCR with nested primers.</li> <li>Primers should have similar melting temperatures.</li> </ul> |
| PCR products in negative control experiments | DNA template problems  | Use serial dilutions of template.  |
|  | Carry-over contamination   | <ul style="list-style-type: none"> <li>Replace all reagents, especially water.</li> <li>Use aerosol-resistant pipette tips.</li> <li>Set up PCR reactions in an area separate from that used for PCR product analysis.</li> </ul>  |
| Specific problems in RT-PCR                  | No product, additional bands, background smear   | <ul style="list-style-type: none"> <li>The volume of cDNA template (from RT reaction) should not exceed 10% of the final PCR reaction volume.</li> <li>Increase MgCl<sub>2</sub> in 0.25 mM steps.</li> <li>Follow troubleshooting suggestions above.</li> </ul>   |

### 4. Additional Information on this Product

#### References

- Hopfner, K.P. et al. (1999) *Proc. Natl Acad. Sci.* **96**, 3600-3605.
- Frey, B. & Suppmann, B. (1995) *Biochemica* **2**, 8-9.
- Lin, Z. et al (2000) Polymorphisms of human SP-A, SP-B, and SP-D genes: association of SP-B Thr131Ile with ARDS. *Clinical Genetics* **58**, 181.
- van Beilen; J.B. et al (2001) Analysis of *Pseudomonas putida* alkane-degradation gene clusters and flanking insertion sequences: evolution and regulation of the alk genes. *Microbiology* **147**, 1621-1630.



#### Quality Control

Each lot of Expand Long Template PCR System is function tested in PCR. Routinely, Expand Long Template PCR is used to amplify a human genomic DNA with primers that are specific for 9 kb, 12 kb and 15 kb fragments. According to our Quality Control procedures the enzyme mix is tested for the absence of any contaminating activities, including endo- or exonucleases and nicking activity.

### 5. Supplementary Information

#### 5.1 Symbols

In this document the following symbols are used to highlight important information:

| Symbol  | Description  |
|---|--|
|  | Information Note:<br>Additional information about the current topic or procedure.              |
|  | Important Note:<br>Information critical to the success of the procedure or use of the product. |

#### Changes to previous version

Editorial Changes.

#### 5.2 Ordering Information

|   | Product  | Pack Size                    | Cat. No.       |
|---|--|------------------------------|----------------|
| High fidelity PCR                               | Expand High Fidelity PCR System                  | 100 U                        | 11 732 641 001 |
|   |  | 2 × 250 U                    | 11 732 650 001 |
|   |  | 10 × 250 U                   | 11 759 078 001 |
|   | High Fidelity PCR Master                         | 1 kit                        | 12 140 314 001 |
|   | FastStart High Fidelity PCR System               | 125 U                        | 03 553 426 001 |
|   |  | 2 × 250 U                    | 03 553 400 001 |
| 10 × 250 U                                      |  | 03 553 361 001               |                |
| Expand High Fidelity <sup>PLUS</sup> PCR System | 125 U  | 03 300 242 001               |                |
|   | 2 × 250 U  | 03 300 226 001               |                |
|   | 10 × 250 U                                       | 03 300 234 001               |                |
| Long template PCR                               | Expand Long Range dNTPack                        | 175 U                        | 04 829 034 001 |
|   |  | 700 U                        | 04 829 042 001 |
|   |  | 3500 U                       | 04 829 069 001 |
|   | Expand 20 kb <sup>PLUS</sup> PCR System          | 200 U                        | 11 811 002 001 |
|   | Expand 20 kb <sup>PLUS</sup> PCR System, dNTPack | 200 U                        | 04 743 814 001 |
|   | Difficult templates & challenging assays         | FastStart Taq DNA Polymerase | 50 U           |
| 100 U   |  |                              | 12 032 902 001 |
| 500 U   |  |                              | 12 032 929 001 |
| 4 × 250 U                                       |  |                              | 12 032 937 001 |
| 10 × 250 U                                      |  |                              | 12 032 945 001 |
| 20 × 250 U                                      |  |                              | 12 032 953 001 |
| FastStart High Fidelity PCR System              |  | 125 U                        | 03 553 426 001 |
|   |  | 2 × 250 U                    | 03 553 400 001 |
|   |  | 10 × 250 U                   | 03 553 361 001 |
|   |  | GC-RICH PCR System           | 100 U          |

|                    | <b>Product</b>                  | <b>Pack Size</b>   | <b>Cat. No.</b> |
|--------------------|---------------------------------|--------------------|-----------------|
| <b>Nucleotides</b> | dATP, PCR Grade                 | 25 µmol, 250 µl    | 11 934 511 001  |
|                    |                                 | 125 µmol, 1,250 µl | 11 969 013 001  |
|                    |                                 | 4 × 125 µmol,      | 03 732 681 001  |
|                    |                                 | 4 × 1,250 µl       |                 |
|                    | dCTP, PCR Grade                 | 25 µmol, 250 µl    | 11 934 520 001  |
|                    |                                 | 125 µmol, 1,250 µl | 11 969 021 001  |
|                    |                                 | 4 × 125 µmol,      | 03 732 690 001  |
|                    |                                 | 4 × 1,250 µl       |                 |
|                    | dGTP PCR Grade                  | 25 µmol, 250 µl    | 11 934 538 001  |
|                    |                                 | 125 µmol, 1,250 µl | 11 969 030 001  |
|                    |                                 | 4 × 125 µmol,      | 03 732 703 001  |
|                    |                                 | 4 × 1,250 µl       |                 |
|                    | dTTP PCR Grade                  | 25 µmol, 250 µl    | 11 934 546 001  |
|                    |                                 | 125 µmol, 1,250 µl | 11 969 048 001  |
|                    |                                 | 4 × 125 µmol,      | 03 732 711 001  |
|                    |                                 | 4 × 1,250 µl       |                 |
|                    | dUTP PCR Grade                  | 25 µmol, 250 µl    | 11 934 554 001  |
|                    |                                 | 125 µmol, 1,250 µl | 11 969 056 001  |
|                    |                                 | 4 × 125 µmol,      | 03 732 720 001  |
|                    |                                 | 4 × 1,250 µl       |                 |
|                    | PCR Nucleotide Mix<br>PCR Grade | 200 µl             | 11 581 295 001  |
|                    |                                 | 2,000 µl           | 11 814 362 001  |
|                    | High Fidelity PCR<br>Master     | 10 × 500 µl        | 12 140 314 001  |
|                    | PCR Master                      | 10 × 500 µl        | 11 636 103 001  |

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