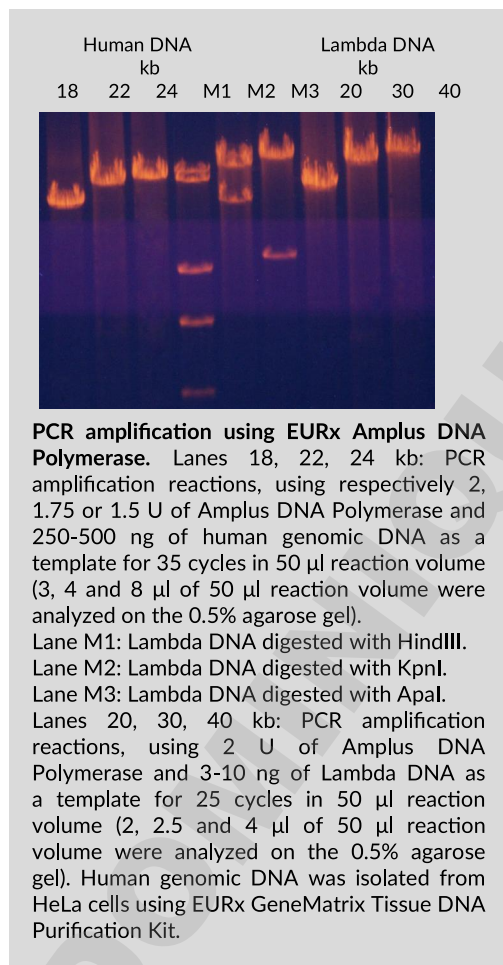


Amplus DNA Polymerase

Cat. No.	size
EK2900-01	100 units
EK2900-02	500 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 MgCl₂, 200 µM each of dATP, dCTP, dGTP, dTTP (a mix of unlabeled and [³H]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.



References:

- Cline, J., Braham, J. and Hogrefe, H. (1996) *Nucleic Acids Res.* 24, 3546.

Mixture of thermophilic DNA polymerases, a polymerase-enhancing factor and a unique polymerase buffer enabling long-range PCR amplification of genomic targets longer than 25 kb and episomal targets up to 40 kb.

Description:

- Amplus DNA Polymerase is a modified and optimized thermostable enzymes blend containing *Thermus sp.* DNA polymerase, *Pyrococcus sp.* DNA polymerase and the polymerase-enhancing factor.
- Ultrapure, recombinant enzymes are used to prepare Amplus DNA Polymerase.
- A unique composition of the amplification buffer enables effective buffering of pH at high temperatures and allows to achieve robust yield from long genomic targets.
- The polymerase amplifies efficiently large genomic DNA fragments from 3 kb to over 25 kb and episomal fragments up to 40 kb. It also provides reliable amplification of smaller fragments (as small as 0.1 kb).
- The enzymes blend is especially suitable for eukaryotic genome amplification, cloning and analysis.
- Ideal for genome mapping and sequencing by greatly simplifying contig assembly from large amplification products.
- Enables characterization of cloned sequences in lambda phages and cosmids.
- Improves PCR results with critical templates, such as containing GC-rich regions.
- Amplus DNA Polymerase exhibits the 3'→5' proofreading activity, resulting in considerably higher PCR fidelity and processivity than possible with unmodified Taq DNA polymerase (1).
- Maintains the 5'→3' exonuclease activity.
- Adds extra A at the 3' ends.

Storage Buffer:

20 mM Tris-HCl (pH 8.0 at 22°C), 100 mM KCl, 0.1 mM EDTA, 1 mM dithiothreitol, 50% glycerol and stabilizers.

10 x Reaction Buffer:

10 x Amplus Buffer:

The buffer contains 29 mM MgSO₄ and is optimized for use with Eurx dNTP mix.

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:

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

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Component	Volume/reaction	Final concentration
10 x Amplus Buffer contains 29 mM MgSO ₄	5 µl	1x
dNTP mix (5 mM each)	5 µl	0.5 mM of each dNTP
Upstream primer	Variable	0.3-0.4 µM
Downstream primer	Variable	0.3-0.4 µM
Amplus DNA Polymerase, 5 U/µl	0.3-0.4 µl	1.5-2U
Template DNA	Variable	<0.5 µg/50 µl
Sterile double-distilled 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 on ice, mix well.
3. Place reactions in a thermal cycler that has been preheated to denaturation temperature.
4. For Amplus DNA Polymerase-based PCR, optimal concentration of MgSO₄ in PCR reaction is 2.9 mM (as provided with the 1 x Amplus Buffer) when using Eurx 0.5 mM dNTP (each) and recommended amount of DNA template (see point 7). In most cases this concentration will produce satisfactory results. If increased Mg²⁺ concentration is needed use 25 mM MgSO₄ provided and increase Mg²⁺ concentration in increments of 0.1 mM up to 3.5 mM.
5. 1.75 U of Amplus DNA Polymerase is recommended starting concentration of the enzyme per 50 µl amplification reaction. For most applications, enzyme will be in excess and will produce satisfactory results. Too much enzyme may generate artifacts like as smearing of bands, etc. In some cases it may be necessary to optimize the enzyme concentration in range 1.5–2 U to achieve its best performance (try 0.3, 0.35, 0.4 µl of the enzyme per 50 µl PCR reaction).
6. The recommended concentration of primers is 0.3-0.4 µM. This concentration enables to achieve satisfying yield while not compromising amplification specificity from genomic targets.
7. For long range PCR use: 100-500 ng of human genomic DNA, 0.1-10 ng of bacterial DNA, 1-50 ng of phage DNA and 1-20 ng of plasmid DNA. If lower than recommended amount of DNA is used (when amplifying shorter, standard PCR targets) it may be necessary to increase/optimize Mg²⁺ concentration in increments of 0.1 mM up to 3.5 mM.
8. The quality of the template influences dramatically the performance of PCR. Ensure that a 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). Template DNA should be prepared carefully, applying gentle methods that do not shear the template in order to receive high molecular weight DNA of high purity.
9. Complex genomic DNA should be stored at 2-8°C and should not have been frozen. Avoid vortexing the genomic DNA. Avoid freeze-thaw steps.
10. Use only thin-walled 0.2 ml tubes performing long PCR amplification.
11. In most cases there is no need to add additives to PCR reaction. For some difficult targets such as GC-rich sequences and very long targets over 30 kb, additives such as DMSO may be included to improve amplification.

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Step	Temperature	Time	Number of Cycles
Initial Denaturation	93-94°C	2 min	1
Denaturation	93-94°C	10 s	25-35
Annealing	60-68°C	30 s	
Extension	68°C	1 min/1 kb	
Final Extension	68°C	7 min	1
Cooling	4°C	Indefinite	1

Thermal Cycling Conditions for Products 0.1-10 kb

Thermal Cycling Conditions for Products over 10 kb

Step	Temperature	Time	Number of Cycles
Initial Denaturation	92-93°C	2 min	1
Denaturation	92-93°C	10 s	10
Annealing	60-68°C	30 s	
Extension	68°C	1 min/1 kb	
Denaturation	92-93°C	10 s	15-25
Annealing	60-68°C	30 s	
Extension	68°C	1 min/1 kb +20 s in each additional cycle	
Final Extension	68°C	7 min	1
Cooling	4°C	Indefinite	1

Notes:

- Annealing temperature should be optimized for each primer set based on the primer T_m . Optimal annealing temperatures may be above or below the estimated T_m . As a starting point, use an annealing temperature 5°C below T_m .
- Typical primers for long PCR amplification have a length of 22-34 and should have annealing temperatures above 60°C to enhance reaction specificity.
- When amplifying long PCR products keep denaturation temperature as low as possible. **For genomic DNA over 15 kb denaturation temperature of 92°C is strongly recommended** (genomic DNA is much more liable to breaks than episomal DNA, especially at denaturation temperatures). Sometimes fragments with high GC-content need higher denaturation temperatures, but keep in mind that the yield increases when denaturation temperature/duration is decreased.
- For PCR products between 5 and 10 kb elongation of extension step (+20 s in each additional cycle starting from 11th cycle) is optional and allows to achieve better yield. For PCR products over 10 kb the elongation of extension step is necessary to achieve satisfying results due to loss of processivity of the enzymes blend.

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