

# Hotstart KnTaq Polymerase

## #P1221, 250U

**Concentration:** 5U/ $\mu$ l

**Contents:**

Hotstart KnTaq Polymerase (5U/ $\mu$ l)	50 $\mu$ l
10 $\times$ Taq Buffer (Mg <sup>2+</sup> Plus)	1.25 ml

**Store at -20°C**

For research use only.

In total 2 vials.

### **Description**

Hotstart KnTaq Polymerase is an N-truncated Taq DNA polymerase with a stable hot-start system and no 5'-3' exonuclease activity. It is optimized for excellent differentiation of error extensions, and can be used for SNP analysis, genotyping and allele-specific PCR.

### **Unit Definition**

One unit is defined as the amount of the enzyme required to catalyze the incorporation of 10 nmole of dNTPs into an acid-insoluble form in 30 minutes at 70°C using hering sperm DNA as substrate.

### **Applications**

- SNP analysis and genotyping
- Allele-specific PCR
- Multiplexing
- Arbitrarily primed PCR
- Formulation of dried-down amplification reagents

### **Features**

- Thermostable: half-life over 40 min at 95°C incubation
- Generates 3'-dA overhangs PCR products

### **Basic PCR Protocol**

The following basic protocol serves as a general guideline and a

starting point for any PCR amplification. Optimal reaction conditions (incubation time and temperature, concentration of Hotstart Kntaq Polymerase, primers and template DNA) vary and need to be optimized.

### 1. Add the following components to a sterile microcentrifuge tube sitting on ice:

Reagent	50- $\mu$ l rxn	Final conc.
10 $\times$ Taq Buffer (Mg <sup>2+</sup> plus)	5 $\mu$ l	1 $\times$
dNTPs (10mM each) <sup>[1]</sup>	1 $\mu$ l	0.2mM each
Primer I	variable	0.4-1 $\mu$ M <sup>[2]</sup>
Primer II	variable	0.4-1 $\mu$ M <sup>[2]</sup>
Hotstart Kntaq Polymerase (5U/ $\mu$ l) <sup>[2]</sup>	0.5-1.0 $\mu$ l	2.5-5U/50 $\mu$ l <sup>[3]</sup>
Template DNA	variable	10pg-1 $\mu$ g
Nuclease-free water	To 50 $\mu$ l	-

[1] Recommended range of final primer concentration: 0.1-1 $\mu$ M. The concentration can be reduced when the specificity is poor, and the concentration can be increased when the efficiency is low.

[2] Adjust the amount of DNA polymerase according to the difficulty of the amplification.

### 2. Mix contents in tube. Cap tubes and centrifuge briefly to collect the contents to the bottom.

When using a thermal cycler that does not contain a heated lid, overlay the reaction mixture with 25  $\mu$ l mineral oil.

### 3. Perform 25-35 cycles of PCR amplification as follows:

Initial Denaturation	94°C	2 minutes
25-35 Cycles	94°C	30 seconds
	55°C <sup>[1]</sup>	30 seconds

	72°C <sup>[2]</sup>	Variable
Final Extension	72°C	5~10 minutes

[1] The annealing temperature should be set according to primers with lower T<sub>m</sub> values.

[2] The optimal extension time is 2min/kb (up to 20s/kb for simple templates).

### 4. Incubate for an additional 10 min at 72°C and maintain the reaction at 4°C. The samples can be stored at -20°C until use.

### 5. Analyze the amplification products by agarose gel electrophoresis and visualize by nucleic acid dye staining. Use appropriate molecular weight standards.

#### Notes on Cycling Conditions

- Hotstart Kntaq Polymerase adopts improved antibody modification technology, so it relies on temperature to activate the polymerase activity, which can effectively inhibit non-specific bindings, and the reaction system can be configured at room temperature.
- The half-life of enzyme is >40 minutes at 95°C.
- Hotstart Kntaq Polymerase accepts modified nucleotides (e.g. biotin-, digoxigenin-, fluorescent-labeled nucleotides) as substrates for the DNA synthesis.
- The number of PCR cycles depends on the amount of template DNA in the reaction mix and on the expected yield of the PCR product. 25-35 cycles are usually sufficient for the majority PCR reaction. Low amounts of starting template may require 40 cycles.

### **Guidelines for Preventing Contamination of PCR Reaction**

During PCR more than 10 million copies of template DNA are generated. Therefore, care must be taken to avoid contamination with other templates and amplicons that may be present in the laboratory environment. General recommendations to lower the risk of contamination are as follows:

- Prepare your DNA sample, set up the PCR mixture, perform thermal cycling and analyze PCR products in separate areas.
- Set up PCR mixtures in a laminar flow cabinet equipped with an UV lamp.
- Wear fresh gloves for DNA purification and reaction set up.
- Use reagent containers dedicated for PCR. Use positive displacement pipettes, or use pipette tips with aerosol filters to prepare DNA samples and perform PCR set up.
- Always perform “no template control” (NTC) reactions to check for contamination.

### **Quality Control**

The absence of endodeoxyribonucleases, exodeoxyribonucleases and ribonucleases is confirmed by appropriate quality tests. Functionally tested in amplification of a single-copy gene from human genomic DNA.

### **Endodeoxyribonuclease Assay**

No detectable conversion of covalently closed circular DNA to a nicked DNA was observed after incubation of 10U Hotstart Kntaq Polymerase with 1µg pBR322 DNA for 4 hours at 37°C and 70°C.

### **Exodeoxyribonuclease Assay**

No detectable degradation of lambda DNA-HindIII fragments was observed after incubation of 10U Hotstart Kntaq Polymerase with 1µg digested DNA for 4 hours at 37°C and 70°C.

### **Ribonuclease Assay**

0% of the total radioactivity was released into trichloroacetic acid-soluble fraction after incubation of 10U *Super* Hotstart Taq Polymerase with 1µg E.coli [3H]-RNA (40000cpm/µg) for 4 hours at 37°C and 70°C.

### **PRODUCT USE LIMITATION.**

This product is developed, designed and sold exclusively for *research purposes and in vitro use only*. The product was not tested for use in diagnostics or for drug development, nor is it suitable for administration to humans or animals.