

Klentaq Pol Hot Start
Heat-activatable DNA polymerase for high specificity, chemically modified
Thermus aquaticus, recombinant, E.coli

|  | Cat. ${ }^{\circ}$. | Amount |
| :---: | :---: | :---: |
| $\square$ | POL-128XS | 100 units |
| $\square$ | POL-128S | 250 units |
| $\square$ | POL-128M | 500 units |
| $\square$ | POL-128L | 1.000 units |
| $\square$ | POL-128XL | $2 \times 1.000$ units |

## Unit Definition:

One unit is defined as the amount of the enzyme required to catalyze the incorporation of 10 nmoles of dNTPs into an acidinsoluble form in 30 minutes at $74^{\circ} \mathrm{C}$.

## Concentration:

5 units/ $\mu \mathrm{L}$

## Shipping:

Shipped on blue ice

## Storage Conditions:

Store at $-20^{\circ} \mathrm{C}$

## For in vitro use only!

## Additional Storage Conditions:

Avoid freeze/thaw cycles

## Shelf Life:

12 months

## Kit contents:

## Klentaq Polymerase Hot Start(blue cap)

5 units/ $\mu$ l Klentaq Pol Hot Start in Tris-HCl pH $9.0\left(25^{\circ} \mathrm{C}\right), \mathrm{KCl}$, EDTA, 50\% (v/v) Glycerol and stabilizers.
Klentaq Pol HS Reaction Buffer complete (red cap) - 10x conc. Tris- HCl pH $8.0\left(25^{\circ} \mathrm{C}\right)$ and stabilizers.

## Description:

Klentaq Pol Hot Start provides improved specificity and sensitivity when amplifying low-copy-number targets in complex backgrounds or when prolonged room-temperature set up is required. This ultra-pure enzyme, in addition to its hot-start capabilities, reduces false positives, amplifies a wide range of DNA sequence contexts. Klentaq Pol is purified by an additional separation process to reduce contaminating bacterial DNA sequences from the enzyme preparation The polymerase activity is chemically blocked at ambient temperature and switched on automatically at the onset of the initial denaturation. The thermal activation prevents the extension of nonspecifically annealed primers and primer-dimer formation at low temperatures during PCR setup. The enzyme catalyzes the polymerization of nucleoside into duplex DNA in $5^{\prime} \quad 3^{\prime}$ direction in the presence of magnesium. Klentaq lacks the 5' 3' exonuclease activity.

## Activation step:

Klentaq Hot Start Pol requires a prolonged heating or denaturing step. The chemical modification of the polymerase is reversed by the increased temperature of the hot start cycle.

## PCR Reaction Setup

The PCR procedure below shows appropriate volumes for a single $50 \mu \mathrm{~L}$ reaction. For multiple reactions, prepare a master mix of components common to all and then dispense appropriate volumes into each PCR reaction tube prior to adding template DNA and primers.
Thaw, mix, and briefly centrifuge each component before use. Add the following components to a microcentrifuge tube:

## 1. Prepare PCR master mix

Note: Consider the volumes for all components listed next steps to determine the correct amount of water required to reach your final reaction volume.

| Components | $\mathbf{5 0} \boldsymbol{\mu L} \mathbf{r x n}$ | [ final ] |
| :--- | :---: | :---: |
| Water, grade PCR | To $50 \boldsymbol{\mu}$ |  |
| 10x Reaction Buffer | $5 \mu \mathrm{l}$ |  |
| dNTP (Mix 10 mM$)$ | $1 \mu \mathrm{l}$ | $200 \mu \mathrm{M}$ |
| Klentaq Pol HS $(5 \mathrm{U} / \mu \mathrm{l})$ | $0,5 \mu \mathrm{l}$ | $2,5 \mathrm{U} /$ reaction |

Mix and briefly centrifuge the components.

## 2. Add template DNA and primers

| Components | $\mathbf{5 0} \boldsymbol{\mu \mathrm { L }} \mathbf{~ r x n}$ | [ final ] |
| :--- | :---: | :---: |
| Foward primer $(10 \mu \mathrm{M})$ | $0,5-2,5 \mu \mathrm{l}$ | $0,1-0,5 \mu \mathrm{M}$ |
| Reverse primer $(10 \mu \mathrm{M})$ | $0,5-2,5 \mu \mathrm{l}$ | $0,1-0,5 \mu \mathrm{M}$ |
| DNA template |  | $10 \mathrm{pg}-1 \mu \mathrm{~g} * *$ |

**genomic DNA: $1 \mathrm{ng}-1 \mu$ g; plasmidial or viral DNA: $1 \mathrm{pg}-1 \mathrm{ng}$ Cap each tube, mix, and briefly centrifuge the content.
3. Incubate reactions in a thermal cycler. Recommended cycling conditions:

| Step | Temp. | Time |  |
| :--- | :---: | :---: | :---: |
| Initial denaturation | $95^{\circ} \mathrm{C}$ | 10 min |  |
|  | Denaturation | $95^{\circ} \mathrm{C}$ | $15-30 \mathrm{sec}$ |
| $25-40$ | Annealing ${ }^{1}$ | $45-68^{\circ} \mathrm{C}$ | $15-30 \mathrm{sec}$ |
| cycles | Elongation ${ }^{2}$ | $68^{\circ} \mathrm{C}$ | $1 \mathrm{~min} / \mathrm{kbp}$ |
|  | $68^{\circ} \mathrm{C}$ | $1-2 \mathrm{~min}$ |  |
| Final extension (optional) | $4-8{ }^{\circ} \mathrm{C}$ |  |  |
| Hold |  |  |  |

1)The annealing temperature depends on the melting temperature of the primers used.
2)The elongation time depends on the length of the fragments to be amplified. A time of $2 \mathrm{~min} / \mathrm{kb}$ is recommended.

For optimal specificity and amplification an individual optimization of the recommended parameters may be necessary for each new template DNA and/or primer pair.

