

# DNA Polymerase I, Large (Klenow) Fragment

pH7.9 @ 25°C



**Catalog:** RK20525

**Size:** 200 U / 1,000 U / 5,000 U

**Concentration:** 5,000 U/ml

**Components:**

DNA Polymerase I, Large (Klenow) Fragment (5,000 U/ml)	RM20515
10X ABuffer B	RM20126

## Product Description

DNA Polymerase I, Large (Klenow) Fragment (about 68 kD) is a proteolytic product of *E. coli* DNA Polymerase I, which retains polymerization and 3'→5' exonuclease activity, but has lost 5'→3' exonuclease activity. Klenow retains the polymerization fidelity of the holoenzyme without degrading 5' termini.

It is applicable to DNA sequencing by the Sanger dideoxy method, fill-in of 5' overhangs to form blunt ends, removal of 3' overhangs to form blunt ends, second-strand cDNA synthesis and second-strand synthesis in mutagenesis protocols.

### Product Source:

An *E. coli* strain that contains the *E. coli* polA gene that has had its 5'→3' exonuclease domain removed.

### Unit Definition:

One unit is defined as the amount of enzyme that will incorporate 10 nmol of dNTP into acid insoluble material in 30 minutes at 37°C.

### Storage Conditions:

25 mM Tris-HCl, 1 mM DTT, 0.1 mM EDTA, 50% Glycerol, pH 7.4 @25°C

**Storage Temperature:** -20°C

**Reaction Conditions:** 1X ABuffer B, Incubate at 25°C

### 1X ABuffer B:

10 mM Tris-HCl, 50 mM NaCl, 10 mM MgCl<sub>2</sub>, 1 mM DTT,

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**Heat Inactivation:** 75°C for 20 min

**Molecular Weight:** Theoretical 68000 daltons

**5' - 3' Exonuclease:** No

**3' - 5' Exonuclease:** Yes

**Strand Displacement:** +

**Error Rate:** ~ 18x10<sup>-6</sup> bases

## Instructions

### Protocol for blunting ends by 3' overhang removal and fill-in of 3' recessed (5' overhang) ends using DNA Polymerase I, Large (Klenow) Fragment

1. DNA should be dissolved in 1X ABuffer A/B/C/S or T4 DNA Ligase Reaction buffer supplemented with 33 µM each dNTP.
2. Add 1 unit of Klenow per microgram DNA.
3. Incubate for 15 minutes at 25°C.
4. Stop reaction by adding EDTA to a final concentration of 10 mM and heating for 20 minutes at 75°C.

### Notes:

- **CAUTION:** Elevated temperatures, excessive amounts of enzyme, failure to supplement with dNTPs, or long reaction times may result in recessed ends due to the 3' → 5' exonuclease activity of the enzyme.
- When DNA Polymerase I, Large (Klenow) Fragment is used to sequence DNA using the dideoxy method of Sanger *et al.*, 1 unit/5 µl reaction volume is recommended.
- DNA Polymerase I, Large (Klenow) Fragment is also active in 1X ABuffer A/B/C/S and T4 DNA Ligase Reaction Buffer when supplemented with dNTPs.

### QC Process:

- ◆ Purity is above 95% detected by SDS-PAGE.
- ◆ No nuclease, RNase contamination.

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◆ No residual host genomic DNA is detected by PCR.

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