

# Taq Plus DNA Polymerase

Catalog # P201



Version 5.1

Vazyme biotech co., ltd.

## Introduction

Taq Plus DNA polymerase is a mixture of Taq DNA polymerase and an enzyme containing 3'→5' exonuclease activity. The fidelity of Taq Plus is 6 times greater than that of Taq DNA Polymerase. Compared with Taq DNA Polymerase, Taq Plus DNA polymerase has stronger amplification performance, higher sensitivity and yield, and is more tolerant of impurities within 5 kb amplifying range. The obtained PCR products are compatible with ClonExpress II One Step Cloning Kit series (Vazyme, #C112, #C113). The PCR products contain A at the 3'-end and can be directly cloned into T-Vectors.

## Package Information

Components	P201-01 250 U	P201-d1 250 U	P201-02/d2 1,000 U	P201-03/d3 3,000 U
10× Taq Plus Buffer (Mg <sup>2+</sup> plus)	1 ml	1 ml	4 ml	
dNTP Mix (10 mM each)	-----	200 μl	± 800 μl	
Taq Plus DNA Polymerase (5 U/μl)	50 μl	50 μl	200 μl	P201-02/d2 × 3

## Storage

Store at -20°C.

## Unit Definition

One unit (U) is defined as the amount of enzyme that incorporates 10 nmol of dNTPs into acid-insoluble products in 30 min at 74°C, with activated salmon sperm DNA as the template / primer.

## Quality Control

**Exonuclease Activity:** The product is tested in a reaction containing 10 U of Taq DNA Polymerase and 0.6 μg of λ-Hind III. After incubation at 37°C for 16 hours, there is no visually discernible change to DNA bands determined by agarose gel electrophoresis.

**Endonuclease Activity:** The product is tested in a reaction containing 10 U of Taq DNA Polymerase and 0.6 μg of Supercoiled pBR322. After incubation at 37°C for 4 hours, there is no visually discernible change to DNA band determined by agarose gel electrophoresis.

**Functional Assay:** The human α-1-antitrypsin gene is amplified for 30 cycles in a 50 μl system using 1.25 U of Taq DNA Polymerase and 100 ng human genomic DNA as template. A single DNA band of 360 bp is detected by 1% agarose gel electrophoresis.

## Protocol

### 1. General reaction mixture for PCR:

ddH <sub>2</sub> O	to 50 μl
10× Taq Plus Buffer (Mg <sup>2+</sup> plus)	5 μl
25 mM MgCl <sub>2</sub> <sup>a</sup>	optional
dNTP Mix (10 mM each)	1 μl
Template DNA <sup>b</sup>	optional
Primer 1 (10 μM)	2 μl
Primer 2 (10 μM)	2 μl
Taq Plus DNA Polymerase (5 U/μl) <sup>c</sup>	0.5 μl

a. The final concentration of Mg<sup>2+</sup> of this mixture is 2 mM, as for most PCR reactions, the optimized final concentration of Mg<sup>2+</sup> is 1.5 mM- 2 mM. However, if necessary, the concentration of Mg<sup>2+</sup> can be increased by adding 25 mM MgCl<sub>2</sub>.

b. The recommended amount of DNA template for a 50 μl reaction system is as follows:

Human Genomic DNA	0.1 - 1 μg
Bacterial Genomic DNA	10 - 100 ng
λ DNA	0.5 - 5 ng
Plasmid DNA	0.1 - 10 ng

c. The amount of Taq DNA Polymerase can be adjusted between 0.25 μl and 1 μl. Generally, higher level of Taq will increase the yield of PCR products but may decrease the specificity of PCR amplification.



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## 2. Thermocycling conditions for a routine PCR:

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94°C	5 min (Pre-denaturation)	
94°C	30 sec	} 30 - 35 cycles
55°C*	30 sec	
72°C	60 sec / kb	
72°C	7 min (Final extension)	

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\*The optimal annealing temperature should be 1-2°C lower than the  $T_m$  of the primers used.

### Handling Notes

Taq DNA Polymerase also shows polymerase activity at room temperature. Therefore, it is recommended to set up reaction systems on ice and then immediately start the reaction in a PCR amplifier, so as to reduce nonspecific amplification during preparation and get better PCR results.

### Primers Designing Notes

1. Choose C or G as the last base of the 3'-end of the primer;
2. Avoid continuous mismatching at the last 8 bases of the 3'-end of the primer;
3. Avoid hairpin structure at the 3'-end of the primer;
4.  $T_m$  of the primers should be within the range of 55°C - 65°C;
5. Additional sequence should not be included when calculating  $T_m$  of the primers;
6. GC content of the primers should be within the range of 40% - 60%;
7.  $T_m$  and GC content of forward and reverse primers should be as similar as possible.

