



Taq DNA Pol 5 U/μl

Cat. #
11EPTQX025

Pack size
250 U

Lot # 1528M0078
Expiration date: Dec-2015

Reference contents:

- Taq DNA Polymerase 5 U/μl
- PCR buffer XD 10xC with MgCl₂

Store at -20°C.

For research use only.

Taq DNA Polymerase is a highly purified thermostable recombinant polymerase, with a 5'-3' exonuclease activity, but no 3'-5' exonuclease activity.

Reaction conditions:

PCR buffer XD diluted to 1xC: 20 mM Tris-HCl pH 8.3 (25°C); 50 mM KCl; 1.5 mM MgCl₂
Taq DNA polymerase is supplied in a formulation containing 50% glycerol.

Reagent	20 μl Reaction	50 μl Reaction	Final Conc.
Sterilized water	Add to 20μl	Add to 50μl	
PCR buffer XD 10xC	2 μl	5 μl	1xC
10 mM dNTPs each	0.4 μl	1 μl	200 μM each
Primer A	x μl	x μl	0.2 to 0.5 μM
Primer B	x μl	x μl	0.2 to 0.5 μM
DNA Template *	x μl	x μl	
Taq DNA Pol 5 U/μl	0.2 μl to 0.3 μl	0.2 μl to 0.3 μl	1 to 1.5 U/rxn **

* Suggested template amount: 100 pg plasmid or phage DNA; 10 - 100 ng genomic DNA; 0.2 - 30 ng cDNA.
** If PCR inhibitors remain in the reaction mix, higher amounts of Taq DNA Pol may be necessary (2-3 U/rxn) to ensure optimal PCR results. Same advice for amplification of fragments up to 3 - 4 kb.

Guidelines for PCR program:

Cycle step	Temperature	Time	Cycles
Initial denaturation (1)	93-95°C	2-4 min	1
Denaturation	93 °C	30 - 60 sec	20 -40
Annealing (2)	(T _m - 5°C)	30 - 60 sec	
Elongation (3)	72°C	1 min /kb	
Final elongation (4)	72°C	10 min	1
End of PCR assay	+ 4°C / -20°C	Hold / Store	

(1) For GC rich templates, extend to 10 min, but add Taq DNA Pol after initial denaturation or keep maximum 5 min at 95°C by adding up to 10% DMSO to reaction mix.

(2) The annealing temperature should be placed 5°C below the lowest T_m of the pair of primers. Avoid annealing temperatures under 48°C.

(3) For amplifications less than 1kb, 15 to 30 sec could be enough.

(4) A final elongation step is recommended to fill-in the incompletely amplified fragments and to add the A residues to the 3'ends of PCR products, (useful for T/A cloning protocols).

Quality control:

Unit assay conditions:

One unit of DNA polymerase is the amount of enzyme required to catalyse the incorporation of 10 nanomoles of nucleosides into a DE81 adsorbable product within 30 min at 74°C under assay conditions.

Absence of contamination: No nickase, endonuclease, 3' exonuclease, ribonuclease activities detected.

PCR assay on human genomic DNA: Specific PCR products of 400 bp using human β-globin gene as a template and decreasing amounts of both DNA template and Taq DNA polymerase are obtained in all assays.

Specificity of PCR buffer XD:

- The buffer contains no BSA and no detergents like Triton.
- That buffer allows optimal results when working with vegetal or beef material or using further dHPLC.

Other PCR buffers available from MP Biomedicals:

- **Standard PCR buffer**, with or without MgCl₂, used in a broad range of PCR conditions.
- **(NH₂)SO₄ PCR buffer** containing ammonium sulphate to improve efficiency and specificity of some application
- **Direct loading buffer** with a densifying agent and a red purple dye allowing direct loading after cycling without new handle.

Troubleshooting:

1. Little or no amplification observed:

- Increase the elongation time on the basis of 60 sec/kb
- Check the T_m of the primers, their quality and their length (> 18 bases for animal material and > 20-22 bases for plants)
- Avoid primer secondary structures, like dimers, hairpins
- Avoid any mismatch or point mutations in the 3' region of the primers
- Avoid 3 G or 3C residues in the 3'end region of the primer
- Lower the annealing temperature by 2°C to 5°C steps
- If annealing temperature (T_a) is close to the elongation temperature, perform PCR by decreasing "T_a" every cycle
- Increase the number of cycles by 5 (if < 35)
- If templates are GC rich or contain difficult secondary structures, add DMSO up to 10% by 2% steps
- Increase the quantity of the template, and/or the primers
- Consider the presence of inhibitory factors

2. Multiple bands or smearing observed:

- Check the T_m of the primers, their design, their length, their quality
- Check the homologous sequences of the primers to the template
- Verify the absence of secondary structures, like hairpins
- Increase the annealing temperature by 2°C to 5°C steps
- Lengthen the homologous sequence of the primer to the template
- Reduce the number of cycles by 5
- Reduce the concentration of DNA template and/or primers

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