

Product information

Product Name: 2X Omni KlenTaq 2 & LA Mix

Catalog No: VN1900M, VN1901M, VN1903M, VN1904M Packing Size: $500 \times 25 \,\mu$ l rxns and $250 \times 25 \,\mu$ l rxns

Shipping Condition: Ice pack Storage Condition: -20°C Product Description:

Omni KlenTaq 2 or Omni KlenTaq 2 LA Mix is a pre-made 2X mixture containing Omni KlenTaq 2 or Omni KlenTaq 2 LA DNA polymerase, salts, magnesium, and dNTPs, which simplifies PCR reactions. You only need to add the template, specific primers, and water to save time and minimize prinetting errors

2X Omni KlenTaq 2 or Omni KlenTaq 2 LA Mix is recommended for standard PCR applications. The mix consists of Omni KlenTaq 2 or OmniKlenTaq 2 LA polymerase (for amplifying longer fragments) and a novel buffer system that delivers high yield PCR amplification across a wide range of templates. It has been designed to produce more robust amplification than other commonly used mixes, making it suitable for challenging templates and for use in the presence of some PCR inhibitors or direct PCR without DNA extraction.

Protocol:

PCR setting for a 25 µl reaction

1 OK Setting for a 20 pri reaction		
Reagent	Volume	Final Concentration
2X Omni KlenTaq 2 / LA PCR Mix	12.5 µl	1x
Left Primer	Variable	0.2-0.4 μM
Right Primer	Variable	0.2-0.4 μM
DNA Template [†]	Variable	0.1-100 ng
De-ionized Distilled H ₂ O	Adjust final volume to 25 µl	=

[†]DNA amount depends mostly on genome size and target gene copy number.

Typical Cycling Parameters

Typical Cycling Larameters			
Initial Denaturation	95°C	2-5 min (for purified DNA)	
		5-15 min (for crude samples)	
25-40 cycles			
Denaturation	94°C	20 to 30 sec	
Annealing	50°C to 68°C	20 to 60 sec	
Extension	70°C	2 min / 1kb target	
Final Extension	70°C	5 min	
Hold	4ºC		

Troubleshooting guide

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No PCR products	Please check your PCR settings to ensure that all necessary components are included in the PCR mix. It's also recommended to perform a gradient annealing temperature experiment to determine the optimal temperature for your specific target. This can help to improve PCR efficiency and specificity, resulting in more reliable and consistent results.	
The bands in agarose gel are smear	It's important to check for DNA and primer degradation as degraded DNA or primers can also negatively impact the PCR results.	
Low yield of products	If you are using crude samples, it's important to initially denature the samples for 5-15 minutes to ensure proper release of DNA. Additionally, you can try the following strategies to improve PCR performance: Conduct an enzyme titration experiment to optimize the enzyme concentration for your specific target. Increase the extension time to ensure complete amplification of the target DNA. Try a gradient annealing temperature experiment to identify the optimal annealing temperature for your primers. Consider redesigning your primers if the amplification is not specific or efficient enough.	
Non-specific products are observed	If you are observing non-specific products in your PCR reaction, there are several strategies you can try: Perform a gradient annealing temperature experiment to identify the optimal temperature for your primers, which can help to reduce non-specific products. Check the GC content of your target sequence. If it's above 65%, it may be necessary to use our PEC-GC or other PCR enhancers. Evaluate your primers and redesign them if necessary to improve specificity and reduce non-specific products.	

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