SeqAmp[™] DNA Polymerase Protocol-At-A-Glance

SeqAmp DNA Polymerase (Cat. Nos. 638504 and 638509) is a high fidelity PCR enzyme with hot start capabilities that is well-suited for use with specific SMARTer® kits for next-generation sequencing (NGS). This optimized PCR enzyme has been shown to perform well even with challenging templates containing GC-rich and AT-rich regions. When using SeqAmp DNA Polymerase with a SMARTer kit, follow the protocol in the particular SMARTer kit user manual. For other applications use the protocol below.

1. Set up the reaction by adding each component indicated in Table 1. To determine how much template to use, refer to Table 2. After adding all the components, mix by tapping the bottom of the tube, then centrifuge briefly.

NOTE: Keep enzymes and reagents on ice.

 Table 1. Recommended Reaction Set-Up

Reagent	Volume	Final Conc.
2X SeqAmp PCR Buffer (includes Mg ²⁺ , dNTPs)	25 µl	1X ¹
Primer 1	10–15 pmol	0.2–0.3 μM ²
Primer 2	10–15 pmol	0.2–0.3 µM ²
Template	< 500 ng	
SeqAmp DNA Polymerase	1 µl	1.25 U/50 µl
Sterilized distilled water	up to 50 µl	
Total volume per reaction	50.0 µl	

¹ Mg²⁺ (1 mM); dNTPs (200 µM each).

²When amplifying fragments \geq 10 kb, use 0.2 μ M.

Table 2. Recommended Amounts of Purified DNA and cDNA Templates

DNA Source	Recommended Amount (standard)	Recommended Amount (long targets)
Human genomic DNA	5–500 ng	100–500 ng
<i>E. coli</i> genomic DNA	100 pg–200 ng	10–200 ng
Plasmid DNA	10 pg–10 ng	1–10 ng
cDNA	25–750 ng	250–750 ng

NOTE: This product is incompatible with bisulfite-treated DNA or other uracil-containing templates.

2. Program your thermal cycler with one of the following sets of cycling conditions:

a. 3-Step PCR

94°C	1 min ¹	
98°C	10 sec	
55 or 60°C ²	15 sec	30 cycles
68°C ³	30 sec/kb ⁴	

b. 2-Step PCR

98°C 10 sec 68°C 30 sec/kb⁴

30 cycles

¹When amplifying a GC-rich region or a long target, perform the initial denaturation at 94°C for 1 min.

² Set the annealing temperature to 60°C for primers with a T_m (determined by the following equation) > 55°C, and to 55°C for primers with a T_m ≤ 55 °C. **NOTE:** T_m (°C) = [(nA + nT) x 2] + [(nC + nG) x 4] - 5

³ For 3-step PCR, set the extension temperature to 68°C.

⁴When amplifying from crude samples, set the extension time to 1 min/kb.

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