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New England Biolabs Certificate of Analysis

Product Name: LongAmp® Hot Start Taq DNA Polymerase

Catalog Number: M0534L
Concentration: 2,500 U/ml

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 75°C.

Packaging Lot Number: 10240724
Expiration Date: 12/2025
Storage Temperature: -20°C

Storage Conditions: 10 mM Tris-HCl, 100 mM KCl, 1 mM DTT, 0.1 mM EDTA, 0.5 % Tween®

20 , 0.5 % IGEPAL® CA-630 , 50 % Glycerol, (pH 7.4 @ 25°C)

Specification Version: PS-M0534S/L v2.0

| LongAmp® Hot Start Taq DNA Polymerase Component List | | | | |
|--|---------------------------------------|------------|----------------------|--|
| NEB Part Number | Component Description | Lot Number | Individual QC Result | |
| M0534LVIAL | LongAmp® Hot Start Taq DNA Polymerase | 10221162 | Pass | |
| B0323SVIAL | LongAmp® Taq Reaction Buffer | 10222171 | Pass | |

| Assay Name/Specification | Lot # 10240724 |
|--|----------------|
| Inhibition of Primer Extension (Hot Start, Radioactivity Incorporation) A 50 µl primer extension assay in ThermoPol® Reaction Buffer in the presence of 200 µM dNTPs including [³H]-dTTP, containing 15 nM primed single-stranded M13mp18 with 10 units of LongAmp® Hot Start Taq DNA Polymerase incubated for 16 hours at 25°C yields >95% inhibition when compared to a non-hot start control reaction. | Pass |
| Non-Specific DNase Activity (16 Hour) A 50 µl reaction in NEBuffer 2 containing 1 µg of T3 or T7 DNA in addition to a reaction containing Lambda-HindIII DNA and a minimum of 2.5 units of LongAmp® Hot Start Taq DNA Polymerase incubated for 16 hours at 37°C results in a DNA pattern free of detectable nuclease degradation as determined by agarose gel electrophoresis. | Pass |
| PCR Amplification (30 kb Human Genomic DNA) A 25 μl reaction in LongAmp® Taq Reaction Buffer in the presence of 300 μM dNTPs and 0.4 μM primers containing 500 ng Human Genomic DNA with 2.5 units of LongAmp® Hot Start Taq DNA Polymerase for 28 cycles of PCR amplification results in the expected 30 kb product. | Pass |



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| Assay Name/Specification | Lot # 10240724 |
|--|----------------|
| PCR Amplification (30 kb Lambda DNA) A 25 µl reaction in LongAmp® Taq Reaction Buffer in the presence of 300 µM dNTPs and 0.4 µM primers containing 1 ng Lambda DNA with 2.5 units of LongAmp® Hot Start Taq DNA Polymerase for 28 cycles of PCR amplification results in the expected 30 kb product. | Pass |
| PCR Amplification (Hot Start, Human Genomic DNA) A 50 µl reaction in LongAmp® Taq Reaction Buffer in the presence of 200 µM dNTPs and 0.2 µM primers containing 2 ng Human Genomic DNA with 5 units of LongAmp® Hot Start Taq DNA Polymerase for 35 cycles of PCR amplification results in the expected 306 bp product, and a decrease in non-specific genomic bands after pre-incubation at room temperature for 1 hour, when compared to a non-hot start control reaction. | Pass |
| RNase Activity (Extended Digestion) A 10 µl reaction in NEBuffer 4 containing 40 ng of a 300 base single-stranded RNA and a minimum of 1 µl of LongAmp® Hot Start Taq DNA Polymerase is incubated at 37°C. After incubation for 16 hours, >90% of the substrate RNA remains intact as determined by gel electrophoresis using fluorescent detection. | Pass |
| qPCR DNA Contamination (E. coli Genomic) A minimum of 2.5 units of LongAmp® Hot Start Taq DNA Polymerase is screened for the presence of E. coli genomic DNA using SYBR® Green qPCR with primers specific for the E. coli 16S rRNA locus. Results are quantified using a standard curve generated from purified E. coli genomic DNA. The measured level of E. coli genomic DNA contamination is ≤ 1 E. coli genome. | Pass |

This product has been tested and shown to be in compliance with all specifications.

One or more products referenced in this document may be covered by a 3rd-party trademark. Please visit www.neb.com/trademarks for additional information.

Lea Antonopoulos

31 Jan 2024

Michael Tonello

Packaging Quality Control Inspector

05 Jun 2024



M0534L / Lot: 10240724

Page 2 of 2