

*be* INSPIRED *drive* DISCOVERY *stay* GENUINE

240 County Road Ipswich, MA 01938-2723 Tel 978-927-5054 Fax 978-921-1350 www.neb.com info@neb.com

## New England Biolabs Certificate of Analysis

| Product Name:          | LongAmp® Hot Start Taq 2X Master Mix  |
|------------------------|---|
| Catalog Number:        | M0533S  |
| Concentration:         | 2 X Concentrate   |
| Packaging Lot Number:  | 10174026  |
| Expiration Date:       | 03/2024   |
| Storage Temperature:   | -20°C   |
| Specification Version: | PS-M0533S/L v2.0  |
| Composition (1X):      | 60 mM Tris-SO4 (pH 9.1 @ 25°C), 20 mM (NH4)2SO4, 2 mM MgSO4, 0.3 mM<br>dATP, 0.3 mM dCTP, 0.3 mM dGTP, 0.3 mM dTTP, 3 % Glycerol, 0.06 %<br>IGEPAL® CA-630, 0.05 % Tween® 20, 125 units/ml LongAmp® Hot Start<br>Taq DNA Polymerase |

| LongAmp® Hot Start Taq 2X Master Mix Component List |                                      |            |                      |  |
|---|--------------------------------------|------------|----------------------|--|
| <b>NEB Part Number</b>                              | Component Description                | Lot Number | Individual QC Result |  |
| M0533SVIAL  | LongAmp® Hot Start Taq 2X Master Mix | 10164435   | Pass                 |  |

| Assay Name/Specification  | Lot # 10174026 |
|---|----------------|
| <b>PCR Amplification (30 kb Lambda DNA, Master Mix)</b><br>A 25 μl reaction in 1X LongAmp® Hot Start Taq Master Mix and 0.4 μM primers<br>containing 1 ng Lambda DNA for 28 cycles of PCR amplification results in the<br>expected 30 kb product.   | Pass           |
| <b>PCR Amplification (30 kb Human Genomic DNA, Master Mix)</b><br>A 25 µl reaction in 1X LongAmp® Hot Start Taq Master Mix and 0.4 µM primers<br>containing 500 ng Human Genomic DNA for 28 cycles of PCR amplification results in<br>the expected 30 kb product.   | Pass           |
| <b>Non-Specific DNase Activity (16 hour, Buffer)</b><br>A 50 µl reaction in 1X LongAmp® Hot Start Taq Master Mix containing 1 µg of T3 or T7<br>DNA in addition to a reaction containing Lambda-HindIII DNA incubated for 16 hours<br>at 37°C results in a DNA pattern free of detectable nuclease degradation as<br>determined by agarose gel electrophoresis.                                     | Pass           |
| PCR Amplification (Hot Start, Human Genomic DNA, Master Mix)<br>A 50 µl reaction in 1X LongAmp® Hot Start Taq Master Mix and 0.2 µM primers<br>containing 2 ng Human Genomic DNA for 35 cycles of PCR amplification results in the<br>expected 306 bp product and a decrease in non-specific genomic bands after<br>pre-incubation at room temperature for 1 hour, when compared to a non-hot start | Pass           |





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| Assay Name/Specification   | Lot # 10174026 |
|--|----------------|
| control reaction.  |                |
| Inhibition of Primer Extension (Hot Start, Radioactivity Incorporation)<br>A 50 µl primer extension assay in ThermoPol® Reaction Buffer in the presence of 200<br>µM dNTPs including [ <sup>3</sup> H]-dTTP, containing 15 nM primed single-stranded M13mp18 with<br>10 units of LongAmp® Hot Start Taq DNA Polymerase incubated for 16 hours at 25°C<br>yields >95% inhibition when compared to a non-hot start control reaction. | Pass           |
| <b>RNase Activity (Extended Digestion)</b><br>A 10 µl reaction in NEBuffer 4 containing 40 ng of a 300 base single-stranded RNA<br>and a minimum of 1 µl of LongAmp® Hot Start Taq 2X Master Mix is incubated at 37°C.<br>After incubation for 4 hours, >90% of the substrate RNA remains intact as determined<br>by gel electrophoresis using fluorescent detection.  | Pass           |
| <b>qPCR DNA Contamination (E. coli Genomic)</b><br>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 $\leq$ 1 E. coli genome. | Pass           |

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

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mail & non

Trinh Nguyen Production Scientist 03 Oct 2022

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