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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: | Q5® Hot Start High-Fidelity DNA Polymerase |
|------------------------|---|
| Catalog Number: | M0493L |
| Concentration: | 2,000 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 74°C. |
| Packaging Lot Number: | 10191817 |
| Expiration Date: | 02/2025 |
| Storage Temperature: | -20°C |
| Storage Conditions: | Proprietary |
| Specification Version: | PS-M0493S/L v1.0 |

| Q5® Hot Start High-Fidelity DNA Polymerase Component List | | | | |
|---|--|------------|----------------------|--|
| NEB Part Number | Component Description | Lot Number | Individual QC Result | |
| M0493LVIAL | Q5® Hot Start High-Fidelity DNA Polymerase | 10178502 | Pass | |
| B9028AVIAL | Q5® High GC Enhancer | 10180077 | Pass | |
| B9027SVIAL | Q5® Reaction Buffer Pack | 10180076 | Pass | |

| Assay Name/Specification | Lot # 10191817 |
|--|----------------|
| Endonuclease Activity (Hot Start, Nicking) A 50 µl reaction in NEBuffer 2 in the presence of 400 µM dNTPs containing 1 µg of supercoiled pUC19 DNA and a minimum of 10 units of Q5® High-Fidelity DNA Polymerase incubated for 4 hours at 37°C results in <10% conversion to the nicked form as determined by agarose gel electrophoresis. | Pass |
| PCR Amplification (20 kb Lambda DNA) A 50 μ I reaction in Q5® Reaction Buffer in the presence of 200 μ M dNTPs and 1.0 μ M primers containing 10 ng Lambda DNA with 1 unit of Q5® Hot Start High-Fidelity DNA Polymerase for 22 cycles of PCR amplification results in the expected 20 kb product. | Pass |
| PCR Amplification (7 kb Human Genomic DNA) A 50 μl reaction in Q5® Reaction Buffer in the presence of 200 μM dNTPs and 0.5 μM primers containing 20 ng Human Genomic DNA with 1 unit of Q5® Hot Start High-Fidelity DNA Polymerase for 30 cycles of PCR amplification results in the expected 7 kb product. | Pass |
| PCR Amplification (Enhancer Dependent, >65% GC-rich) A 50 µl reaction in Q5® Reaction Buffer and Q5® High GC Enhancer in the presence of | Pass |





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| Assay Name/Specification | Lot # 10191817 |
|--|----------------|
| 200 μ M dNTPs and 0.5 μ M primers containing 20 ng Human Genomic DNA with 1 unit of Q5® Hot Start High-Fidelity DNA Polymerase for 30 cycles of PCR amplification results in the enhancer-dependent production of the expected 452 bp product. | |
| PCR Amplification (Hot Start, Human Genomic DNA) A 50 μ I reaction in Q5® Reaction Buffer plus Q5® High GC Enhancer in the presence of 200 μ M dNTPs and 0.5 μ M primers containing 100 ng Human Genomic DNA with 1 unit of Q5® Hot Start High-Fidelity DNA Polymerase for 25 cycles of PCR amplification results in the expected 665 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 |
| Phosphatase Activity (pNPP) A 200 µl reaction in 1M Diethanolamine, pH 9.8, 0.5 mM MgCl2 containing 2.5 mM p-Nitrophenyl Phosphate (pNPP) and a minimum of 100 units Q5® High-Fidelity DNA Polymerase incubated for 4 hours at 37°C yields <0.0001 unit of alkaline phosphatase activity as determined by spectrophotometric analysis. | Pass |
| Protein Purity Assay (SDS-PAGE) Q5® High-Fidelity DNA Polymerase is ≥ 95% pure as determined by SDS-PAGE analysis using Coomassie Blue detection. | Pass |
| RNase Activity (Extended Digestion) A 10 μ I reaction in NEBuffer 4 containing 40 ng of a 300 base single-stranded RNA and a minimum of 1 μ I of Q5® Hot Start High-Fidelity 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 units of Q5® High-Fidelity 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.

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.





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