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New England Biolabs Certificate of Analysis

Product Name: T4 Gene 32 Protein

Catalog Number: M0300L
Concentration: 10 mg/ml
Packaging Lot Number: 10121337
Expiration Date: 08/2023
Storage Temperature: -20°C

Storage Conditions: 20 mM Tris-HCl, 100 mM NaCl, 0.5 mM DTT, 1 mM EDTA, 50% Glycerol,

(pH 8.0 @ 25°C)

Specification Version: PS-M0300S/L v1.0

| T4 Gene 32 Protein Component List | | | | |
|-----------------------------------|-----------------------|------------|----------------------|--|
| NEB Part Number | Component Description | Lot Number | Individual QC Result | |
| M0300LVIAL | T4 Gene 32 Protein | 10117898 | Pass | |
| B7004SVIAL | NEBuffer™ 4 | 10115023 | Pass | |

| Assay Name/Specification | Lot # 10121337 |
|---|----------------|
| Exonuclease Activity (Radioactivity Release) A 50 µl reaction in NEBuffer 4 containing 1 µg of a mixture of single and double-stranded [³H] E. coli DNA and a minimum of 10 µg of T4 Gene 32 Protein incubated for 4 hours at 37°C releases <0.1% of the total radioactivity. | Pass |
| Protein Purity Assay (SDS-PAGE) T4 Gene 32 Protein is ≥ 99% pure as determined by SDS-PAGE analysis using Coomassie Blue detection. | Pass |
| Protein Concentration (A280) The concentration of T4 Gene 32 Protein is 10 mg/ml +/- 10% as determined by UV absorption at 280 nm. Protein concentration is determined by the Pace method using the extinction coefficient of 39,670 and molecular weight of 33,506 daltons for T4 Gene 32 Protein (Pace, C.N. et al. (1995) Protein Sci., 4, 2411-2423). | Pass |
| RNase Activity Assay (2 Hour Digestion) A 10 µl reaction in NEBuffer 4 containing 40 ng of fluorescein labeled RNA transcript and a minimum of 10 µg of T4 Gene 32 Protein incubated for 2 hours at 37°C results in no detectable degradation of the RNA as determined by gel electrophoresis using fluorescent detection. | Pass |



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| Assay Name/Specification | Lot # 10121337 |
|---|----------------|
| RNase Activity (Extended Digestion) A 10 µl reaction in NEBuffer 4 containing 40 ng of fluorescein labeled RNA transcript and a minimum of 10 µg of T4 Gene 32 Protein 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 |
| Single Stranded DNase Activity (FAM-Labeled Oligo) A 50 µl reaction in CutSmart® Buffer containing a 20 nM solution of a fluorescent internal labeled oligonucleotide and a minimum of 10 µg of T4 Gene 32 Protein incubated for 16 hours at 37°C yields <5% degradation as determined by capillary electrophoresis. | Pass |
| Functional Testing (Single Stranded DNA Binding - FAM Labeled Oligo) A 20 μl reaction in NEBuffer 4 containing 20 μM FAM-labeled 50-mer and a maximum of 80 μg of T4 Gene 32 Protein incubated for 30 minutes at 37°C produces a mobility shift in >95% of the starting material as determined by TBE gel electrophoresis and UV imaging. | Pass |
| Phosphatase activity (FAM Labeled Oligo) A 50 ul reaction in CutSmart® Buffer containing a 20 nM solution of a fluorescent internal labeled oligonucleotide with a 5' phosphate and a minimum of 10 µg of T4 Gene 32 Protein incubated for 16 hours at 37°C yields <5% degradation as determined by capillary electrophoresis. | Pass |
| Non-Specific DNase Activity (16 Hour) A 50 μl reaction in NEBuffer 4 containing 1 μg of Lambda-HindIII DNA and a minimum of 30 μg of T4 Gene 32 Protein incubated for 16 hours at 37°C results in a DNA pattern free of detectable nuclease degradation as determined by agarose gel electrophoresis. | Pass |
| Endonuclease Activity (Nicking) A 50 μl reaction in NEBuffer 4 containing 1 μg of supercoiled PhiX174 DNA and a minimum of 10 μg of T4 Gene 32 Protein incubated for 4 hours at 37°C results in <10% conversion to the nicked form as determined by agarose gel electrophoresis. | Pass |
| qPCR DNA Contamination (E. coli Genomic) A minimum of 10 μg of T4 Gene 32 Protein 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.



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BioLabs inc.

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Bo Wu

Production Scientist 21 Sep 2021 Michael Tonello

Packaging Quality Control Inspector

21 Sep 2021

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