

## **Certificate of Analysis for NR-3050**

## Genomic DNA from Escherichia coli, Strain CDC (ATCC® 12807™)

Catalog No. NR-3050

**Product Description:** Genomic DNA was extracted from a preparation of *Escherichia coli* (*E. coli*), strain CDC (ATCC® 12807™), serotype O126:K71(B16):H.

Lot<sup>1</sup>: 64204561 Manufacturing Date: 24MAY2016

TEST	SPECIFICATIONS	RESULTS
Genotypic Analysis Sequencing of 16S rRNA gene (~ 880 base pairs)	≥ 99% sequence identity to <i>E. coli</i> type strain (GenBank: X80725)	99.5% sequence identity to <i>E. coli</i> type strain (GenBank: X80725) <sup>2</sup>
Agarose Gel Electrophoresis	High molecular weight chromosomal DNA	High molecular weight chromosomal DNA (Figure 1)
Concentration by PicoGreen® Measurement	0.7 to 1.5 μg in 25 to 100 μL per vial	0.95 μg in 54 μL per vial (17.6 μg/mL)
Functional Activity by PCR Amplification 16S ribosomal RNA gene	~ 1500 base pairs amplicon	~ 1500 base pairs amplicon
Genotypic Analysis of Virulence Markers <sup>3,4</sup> PCR amplification of plasmid markers EAF (pEAF) bfpA (pEAF) elt (pJY11) esth (pCS1) estp (pCS1) invE (pINV) CVD432 (pAA) aggR (pAA) PCR amplification of chromosomal markers eaeA stx1 sx2 astA	Report results Report results Negative Negative Negative Negative Negative Negative Report results Negative Negative Report results Report results Report results	Positive Positive Negative
OD <sub>260</sub> /OD <sub>280</sub> Ratio	1.7 to 2.1	1.8
Bacterial Inactivation 10% of total yield plated on agar <sup>5,6</sup>	No viable bacteria detected	No viable bacteria detected

<sup>&</sup>lt;sup>1</sup>The bacterial preparation used for extraction of genomic DNA was produced by inoculation of BEI Resources NRS-99 (Lot 3663830) into Tryptic Soy broth. After incubation for 24 hours at 37°C, genomic DNA was extracted using proprietary technology.

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<sup>&</sup>lt;sup>2</sup>Also consistent with *Shigella* species.

<sup>&</sup>lt;sup>3</sup>Kimata, K., et al. "Rapid Categorization of Pathogenic *Escherichia coli* by Multiplex PCR." <u>Microbiol. Immunol.</u> 49 (2005): 485-492. PubMed: 15965295.

<sup>&</sup>lt;sup>4</sup>Virulence marker results were obtained from the source organism used to produce the seed lot used for nucleic acid extraction (NR-99, lot 3663825).

<sup>&</sup>lt;sup>5</sup>An extraction procedure was used that has been shown to consistently inactivate 100% of Gram-negative and Gram-positive bacteria.

<sup>&</sup>lt;sup>6</sup>Plates were incubated for 14 days under propagation conditions.

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1 2

Base Pairs

12000
5000

2000
1650
1000
650
400

Figure 1: Agarose Gel Electrophoresis

Lane 1: Invitrogen™ 1Kb Plus DNA Ladder™

Lane 2: 200 ng of NR-3050

28 FEB 2018

Program Manager or designee, ATCC Federal Solutions

ATCC®, on behalf of BEI Resources, hereby represents and warrants that the material provided under this certificate has been subjected to the tests and procedures specified and that the results described, along with any other data provided in this certificate, are true and accurate to the best of ATCC®'s knowledge.

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