SUPPORTING INFECTIOUS DISEASE RESEARCH

Vector pLVX-EF1α-IRES-Puro Containing the SARS-Related Coronavirus 2, USA-WA1/2020 Non-Structural Protein 13 Gene

Catalog No. NR-52961

Product Description:

Note: The vial label indicates this product contains a TST tag. This nomenclature refers to a 2X Strep tag. This product does not express the Twin-Strep-tag[®] that is commonly referred to as a TST tag. The non-structural protein 13 (nsp13) gene from severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), USA-WA1/2020 (GenBank: MN985325) was codon optimized and modified by the addition of C-terminal 2X Strep tag and cloned into the pLVX-<u>EF1α-IRES-Puro</u> lentiviral expression plasmid. The vector contains an internal ribosomal entry site (IRES) that allows a gene-of-interest and a puromycin resistance gene to be simultaneously co-expressed from a single mRNA transcript. Expression of the transcript is driven by the human elongation factor 1 alpha (EF1α) promoter. The beta-lactamase gene, *bla*, provides transformant selection through ampicillin resistance in *Escherichia coli (E. coli)* and the puromycin resistance gene, *pac*, provides transformant selection through puromycin resistance in eukaryotic cells. The deposited plasmid was transformed into NEB[®] Stable Competent *E. coli* cells (New England Biolabs[®] C3040H), grown in Luria-Bertani broth with ampicillin (100 µg per mL) for 1 day at 30°C in an aerobic atmosphere, extracted using a Plasmid *Plus* Maxi Kit (QIAGEN[®] 12963) and vialed in TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH 8.0).

Lot: 70037728

Manufacturing Date: 22JUL2020

TEST	SPECIFICATIONS	RESULTS
Next-Generation DNA Sequencing	~ 10,720 base pairs	10,719 base pairs ¹
Genotypic Analysis Sequencing of nsp13 insert (~ 1810 base pairs)	≥ 99% sequence identity to depositor's sequence 2X Strep tag sequence confirmed	100% sequence identity to depositor's sequence ² 2X Strep tag sequence confirmed ³
Antibiotic Resistance Ampicillin (encoded by beta-lactamase gene <i>bla</i>) ⁴ Puromycin (encoded by puromycin n-acetyltransferase gene <i>pac</i>)	<i>bla</i> sequence present <i>pac</i> sequence present	<i>bla</i> sequence present <i>pac</i> sequence present
Concentration by PicoGreen® Measurement	≥ 2 µg/mL	0.1 μg in 20 μL per vial (6 μg/mL)
Amount per Vial	Report results	0.1 μg per vial
OD ₂₆₀ /OD ₂₈₀ Ratio (pre-vial)	1.7 to 2.1	1.9
Effective Bacterial Transformation NEB® Stable Competent <i>E. coli</i> cells	≥ 50 colonies per ng	> 500 colonies per ng

¹The sequence was assembled pre-vial using the predicted sequence as the reference sequence. The complete plasmid sequence and map are provided on the BEI Resources webpage.

²The NR-52961 insert was codon optimized but is 100% identical with the SARS-CoV-2, USA-WA1/2020 ORF1ab polyprotein (GenPept QHO60603.1).

³This TST tag is a tandem Strep tag, defined by the sequence N-WSHPQFEKGGGSGGGSGGGSWSHPQFEK-C. For more information, please see Gordon, D. E., et al. "A SARS-CoV-2 Protein Interaction Map Reveals Targets for Drug Repurposing." <u>Nature</u> 583 (2020): 459-468. PubMed: 32353859.

⁴The antibiotic ampicillin degrades quickly during growth. Bacterial stationary phase should be minimized during plasmid expansion to avoid plasmid loss and increased antibiotic concentrations may be necessary.

Certificate of Analysis for NR-52961

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/Heather Couch/ Heather Couch

09 OCT 2020

Program Manager or designee, ATCC Federal Solutions

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