

**Vector pCAGGS Containing the SARS-Related Coronavirus 2, Wuhan-Hu-1 Spike Glycoprotein Gene (soluble, stabilized)**

**Catalog No. NR-52394**

This reagent is the tangible property of the U.S. Government.

**Product Description:**

The vector for the spike (S) glycoprotein gene from severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), Wuhan-Hu-1 (GenBank: [MN908947](#)) was designed for expression of a soluble S glycoprotein with a polybasic cleavage site deletion (RRAR to A; residues 682 to 685) and stabilizing mutations (K986P and V987P, wild type numbering) with a C-terminal thrombin cleavage site, T4 foldon trimerization domain and hexa-histidine tag. The S sequence was codon optimized for mammalian expression and subcloned into the [pCAGGS](#) mammalian expression vector. NR-52394 contains the beta-lactamase gene, *bla*, to provide transformant selection through ampicillin resistance in *Escherichia coli* (*E. coli*). Lot 70038122 was produced from a preparation of glycerol stock (NRC-52394 lot 70034724), grown in Luria-Bertani broth with ampicillin (50 µg per mL) for 1 day at 37°C in an aerobic atmosphere, extracted using a Plasmid *Plus* Maxi Kit (QIAGEN® 12963) and vialled in TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH 8.0).

**Lot: 70038122**

**Manufacturing Date: 07AUG2020**

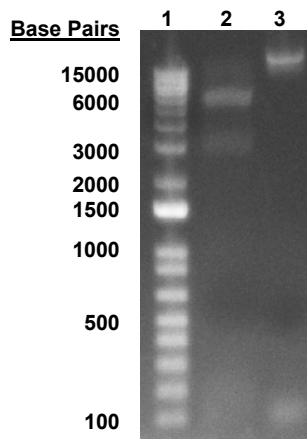
TEST	SPECIFICATIONS	RESULTS
<b>Next-Generation DNA Sequencing<sup>1</sup></b>	~ 8530 base pairs	8537 base pairs
<b>Genotypic Analysis<sup>1</sup></b> Sequencing of S glycoprotein insert (~ 3770 base pairs)	≥ 99% sequence identity to depositor's sequence Thrombin protease site sequence confirmed T4 foldon trimerization domain sequence confirmed His <sub>6</sub> tag sequence confirmed	100% sequence identity to depositor's sequence <sup>2</sup> Thrombin protease site sequence confirmed T4 foldon trimerization domain sequence confirmed His <sub>6</sub> tag sequence confirmed
<b>Antibiotic Resistance</b> Ampicillin (encoded by beta-lactamase gene <i>bla</i> ) <sup>3</sup>	<i>bla</i> sequence present	<i>bla</i> sequence present
<b>Agarose Gel Electrophoresis</b> Digestion with <i>ScaI</i>	~ 6 kb and ~ 3 kb	~ 6 kb and ~ 3 kb (Figure 1)
<b>Concentration by Picogreen® Measurement</b>	≥ 2 µg/mL	0.1 µg in 20 µL per vial (6.8 µg/mL)
<b>Amount per Vial</b>	Report results	0.1 µg per vial
<b>OD<sub>260</sub>/OD<sub>280</sub> Ratio (pre-vial)</b>	1.7 to 2.1	2.0
<b>Effective Bacterial Transformation</b> Invitrogen™ One Shot™ TOP10 <i>E. coli</i>	≥ 50 colonies per ng	> 500 colonies per ng

<sup>1</sup>This test was performed pre-vial on a previous lot of extracted material (NR-52394 lot 70034717). The complete plasmid sequence and map are provided on the BEI Resources webpage.

<sup>2</sup>The NR-52394 insert was codon optimized for mammalian expression with mutations for stability and solubility, but otherwise is consistent with the SARS-CoV-2, Wuhan-Hu-1 S protein (GenPept: QHD43416).

<sup>3</sup>The 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.

**Figure 1: Agarose Gel of Undigested and Restriction Enzyme Digested NR-52394**



Lane 1: Invitrogen™ TrackIt™ 1 Kb Plus DNA Ladder  
 Lane 2: NR-52394 digested  
 Lane 3: NR-52394 undigested

/Heather Couch/

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27 AUG 2020

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