

LEGEND MAX™

ELISA Kit



SARS-CoV-2 Nucleocapsid Pro-

Cat. No. 448007

ELISA Kit for Accurate Quantitation of SARS-CoV2 Nucleocapsid Protein from Cell Culture Supernatant, Serum, Plasma and Other Biological Fluids

BioLegend, Inc. biolegend.com



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It is highly recommended that this manual be read in its entirety before using this product. Do not use this kit beyond the expiration date.

For Research Purposes Only. Not for use in diagnostic or therapeutic procedures. Purchase does not include or carry the right to resell or transfer this product either as a stand-alone product or as a component of another product. Any use of this product other than the permitted use without the express written authorization of BioLegend is strictly prohibited.

LEGEND MAX™ SARS-CoV2 Nucleocapsid Protein ELISA Kit

Introduction:

The nucleocapsid protein is an integral component of SARS-CoV-2, and is highly conserved among the coronaviruses. It performs several important functions for the virus: interacting with viral RNA, providing structure and making up the viral core and nucleocapsid, and interacting with the membrane protein during assembly. These functions increase the efficiency of viral transmission and assembly.

In SARS and other viral infections, nucleocapsid protein levels in patient serum have been show to be a descriptive marker of diease progression. Nucleocapsid protein can be detected in patient serum early in disease onset, prior to antiviral IgG or viral RNA, and detection levels tend to stay high into the infection.

BioLegend's LEGEND MAX™ SARS-CoV2- Nucleocapsid Protein ELISA kit is a Sandwich Enzyme-Linked Immunosorbent Assay (ELISA) with a 96-well strip plate. The plate is pre-coated with a monoclonal mouse capture antibody. The detection antibody is a biotinylated monoclonal mouse antibody. This kit is specifically designed for the accurate quantitation of SARS-CoV-2 Nucleocapsid Protein from cell culture supernatant, serum, plasma and other biological fluids. It is analytically validated with ready-to-use reagents.

Materials Provided:

Description	Quantity	Volume	Part #
SARS-CoV2 Nucleocapsid Protein Pre-Coated 96-well Strip Microplate	1 plate		750002219
SARS-CoV2 Nucleocapsid Protein Detection Antibody	1 bottle	12 mL	750002164
SARS-CoV2 Nucleocapsid Protein Standard	1 vial	lyophilized	750002917
Avidin HRP	1 bottle	12 mL	77897
Assay Buffer H	1 bottle	25 mL	750002328
Wash Buffer (20x)	1 bottle	50 mL	78233
Substrate Solution F	1 bottle	12 mL	76335
Stop Solution	1 bottle	12 mL	79133
Plate Sealers	4 sheets		78101

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LEGEND MAX™ SARS-CoV2 Nucleocapsid Protein ELISA Kit

Materials to be Provided by the End-User:

- Microplate reader able to measure absorbance at 450 nm
- Adjustable pipettes to measure volumes ranging from 1 μL to 1,000 μL
- · Deionized water
- Wash bottle or automated microplate washer
- Log-Log graph paper or software for data analysis
- Polypropylene tubes to prepare standard dilutions
- Timer
- Plate Shaker
- Polypropylene vials

Storage Information:

Store unopened kit components between 2°C and 8°C. Do not use this kit beyond its expiration date.

O	Opened or Reconstituted Components			
Microplate wells	If not all microplate strips are used, remove the excess strips by pressing up from underneath each strip. Place excess strips back in the foil pouch with the included desiccant pack and reseal. Store between 2°C and 8°C for up to one month.			
Standard	The remaining reconstituted standard stock solution can be aliquoted into polypropylene vials and stored at -70°C for up to one month. Avoid repeated freeze-thaw cycles.			
Detection Antibody				
Avidin-HRP				
Assay Buffer H	Store opened reagents between 2°C and 8°C and use within one month.			
Wash Buffer (20X)				
Substrate Solution F				
Stop Solution				

Health Hazard Warnings:

- 1. Reagents that contain preservatives may be harmful if ingested, inhaled or absorbed through the skin. Refer to the MSDS online at BioLegend's website for details (www.biolegend.com/msds).
- 2. Substrate Solution F is harmful if inhaled or ingested. Avoid skin, eye and clothing contact.

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- To reduce the likelihood of blood-borne transmission of infectious agents, handle all serum, plasma and other biological fluids in accordance with NCCLS regulations.
- 4. Stop Solution contains strong acid. *Wear eye, hand, and face protection.*
- 5. Before disposing of the plate, rinse it with an excess amount of tap water.

Specimen Collection and Handling:

Specimens should be clear and non-hemolyzed. If possible, unknown samples should be run at a number of dilutions to determine the optimal dilution factor that will ensure accurate quantitation.

<u>Cell Culture Supernatant</u>: If necessary, centrifuge all samples to remove debris prior to analysis. It is recommended that samples be stored at < -70°C. Avoid repeated freeze-thaw cycles.

<u>Serum:</u> Use a serum separator tube and allow clotting for at least 30 minutes, then centrifuge for 10 minutes at 1,000 x g. Remove serum layer and assay immediately or store serum samples at < -70°C. Avoid repeated freeze-thaw cycles.

<u>Plasma:</u> Collect blood samples in citrate, heparin or EDTA containing tubes. Centrifuge for 10 minutes at 1,000 x g within 30 minutes of collection. Assay immediately or store plasma samples at < -70°C. Avoid repeated freeze-thaw cycles.

Reagent and Sample Preparation:

Note: All reagents should be diluted immediately prior to use.

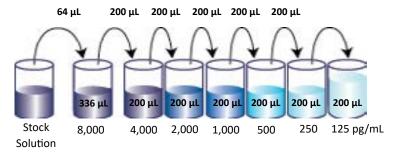
- Dilute the 20X Wash Buffer to 1X with deionized water. For example, make 1 liter of 1X Wash Buffer by adding 50 mL of 20X Wash Buffer to 950 mL of deionized water. If crystals have formed in the 20X Wash Buffer, bring to room temperature and vortex until dissolved.
- Reconstitute the lyophilized SARS-CoV-2 Nucleocapsid Protein Standard by adding the volume of Assay Buffer H to make the 50,000 pg/mL standard stock solution (Refer to LEGEND MAX Kit Lot-Specific Certificate of Analysis/ LEGEND MAX Kit Protocol). Allow the reconstituted standard to sit at room temperature for 15-20 minutes, then briefly vortex to mix completely.
- 3. In general, no dilution is recommended for healthy serum and citrate plasma samples, but samples can be diluted in Assay Buffer H to fit within the range of the assay as determined by the end user. A 2-fold dilution is recommended for EDTA plasma and a 4-fold dilution is recommended for heparin plasma.

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Assay Procedure:

Note: Do not mix reagents from different kits or lots. Reagents and/or antibodies from different manufacturers should not be used with this kit.

- 1. Bring all reagents to room temperature prior to use. It is strongly recommended that all standards and samples be run in duplicate or triplicate. A standard curve is required for each assay.
- 2. If not all microplate strips will be used, remove the excess strips by pressing up from underneath each strip. Place excess strips back in the foil pouch with the included desiccant pack and reseal.
- 3. Prepare 400 μL of the 8,000 pg/mL top standard by adding 64 uL of the 50,000 pg/mL standard stock solution in 336 μL Assay Buffer H. Perform six two-fold serial dilutions of the 8,000 pg/mL top standard in separate tubes using Assay Buffer H as the diluent. Thus, the SARS-CoV-2 Nucleocapsid Protein standard concentrations in the tubes are 8,000 pg/mL, 4,000 pg/mL, 2,000 pg/mL, 1,000 pg/mL, 500 pg/mL, 250 pg/mL, and 125 pg/mL, respectively. Assay Buffer H serves as the zero standard (0 pg/mL).



- 4. Wash the plate 4 times with at least 300 μ L of 1X Wash Buffer per well; blot any residual buffer by firmly tapping the plate upside down on abs bent paper. All subsequent washes should be performed similarly.
- 5. Add 50 μL of Assay Buffer H to each well that will contain either standard dilutions or samples. Then add 50 μL of standard dilutions or samples to the appropriate wells.
- 6. Seal the plate with a Plate Sealer included in the kit and incubate the plate for 2 hours at room temperature with shaking.
- 7. Discard the contents of the plate into a sink, then wash the plate 4 times with 1X Wash Buffer as in step 4.
- 8. Add 100 μL of SARS-CoV-2 Nucleocapsid Protein Detection Antibody solution

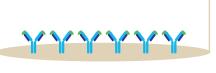
to each well, seal the plate and incubate at room temperature for 1 hour while shaking.

- 9. Discard the contents of the plate into a sink, then wash the plate 4 times with 1X Wash Buffer as in step 4.
- 10. Add 100 μ L of Avidin-HRP solution to each well, seal the plate and incubate at room temperature for 30 minutes while shaking.
- 11. Discard the contents of the plate into a sink, then wash the plate 5 times with 1X Wash Buffer as in step 4. For this final wash, soak wells in 1X Wash Buffer for 30 seconds to 1 minute for each wash. This will help minimize background.
- 12. Add 100 μL of Substrate Solution F to each well and incubate for 10 minutes in the dark. Wells containing SARS-CoV-2 Nucleocapsid Protein should turn blue in color with an intensity proportional to its concentration. It is not necessary to seal the plate during this step.
- 13. Stop the reaction by adding 100 μ L of Stop Solution to each well. The solution color should change from blue to yellow.
- 14. Read absorbance at 450 nm within 30 minutes. If the reader is capable of reading at 570 nm, the absorbance at 570 nm can be subtracted from the absorbance at 450 nm.

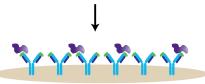
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Assay Procedure Summary

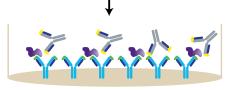
1. Wash 4 times. Add 50 μL Assay Buffer H to standard wells and sample wells



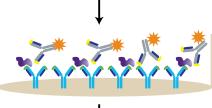
2. Add 50 μL of standard or sample, incubate 2 hr, RT, shaking



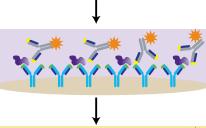
Wash 4 times
 Add 100 μL of Detection Antibody
 solution
 Incubate 1 hr, RT, shaking



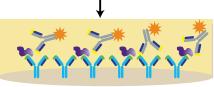
Wash 4 times
 Add 100 μL Avidin-HRP solution
 Incubate 30 min, RT, shaking



5. Wash 5 times Add 100 μL Substrate Solution F Incubate 10 min, RT, in the dark



Add 100 μL Stop Solution



7. Read absorbance at 450 nm and 570 nm

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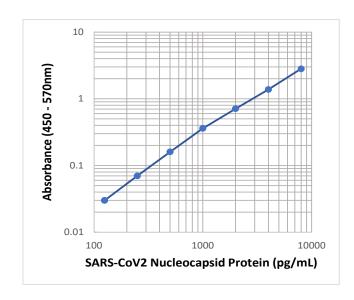
Calculation of Results:

The data can be best calculated with computer-based curve-fitting software using a 5- or 4-parameter logistics curve-fitting algorithm. If an appropriate software is not available, use log-log graph paper to determine sample concentrations. Determine the mean absorbance for each set of duplicate or triplicate standards, controls, and samples. Plot the standard curve on log-log graph paper with cytokine concentration on the X-axis and absorbance on the Y-axis. Draw a best fit line through the standard points. To determine the unknown cytokine concentrations, find the mean absorbance value of the unknown concentration on the Y-axis and draw a horizontal line to the standard curve. At the point of intersection, draw a vertical line to the X-axis and read the cytokine concentration.

If samples were diluted, multiply the concentration by the appropriate dilution factor. If a test sample's absorbance value falls outside the linear portion of the standard curve, the test sample needs to be re-analyzed at a higher (or lower) dilution as appropriate.

Typical Data:

This standard curve was generated at BioLegend for demonstration purposes only. A standard curve must be run with each assay.



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LEGEND MAX™ SARS-CoV2 Nucleocapsid Protein ELISA Kit Performance Characteristics:

<u>Specificity:</u> This kit recognizes natural and recombinant SARS-CoV-2 Nucleocapsid Protein as well as recombinant SARS-CoV Nucleocapsid Protein. No cross reactivity was observed when this kit was used to analyze recombinant MERS-CoV Nucleocapsid protein at 50 ng/mL.

<u>Sensitivity:</u> The average minimum detectable concentration of SARS-CoV-2 Nucleocapsid Protein is 21.4 pg/mL (n=6). The range of calculated minimum detectable concentrations was 11.00 - 32.76 pg/mL.

<u>Recovery:</u> Recombinant or natural SARS-CoV-2 Nucleocapsid Protein at 3 different concentrations was spiked into different human samples of Serum, Citrate Plasma, EDTA Plasma, and Heparin Plasma. Sample recovery was then analyzed with the LEGEND MAX™ SARS-CoV-2 Nucleocapsid Protein ELISA Kit.

Sample Type	N	% Recovery
Serum	5	103.3
Citrate Plasma	5	106.8
EDTA Plasma	5	115.5
Heparin Plasma	4	83.1
Cell Culture Supernatant	3	102.1

<u>Linearity:</u> Recombinant or natural SARS-CoV-2 Nucleocapsid Protein was spiked into human samples, then diluted 2 fold in serial to produce samples within the dynamic range of the kit. Samples were then assayed to determine the dilutional linearity.

Sample Type	N	% Recovery
Serum	4	101.1
Citrate Plasma	4	110.1
EDTA Plasma	4	102.8
Heparin Plasma	4	97.8
Cell Culture Supernatant	3	102,34

<u>Intra-Assay Precision:</u> Two samples containing different SARS-CoV-2 Nucleocapsid Protein concentrations were tested on one plate with 12 replicates.

Concentration	Sample 1	Sample 2
Number of Assays	12	12
Mean Concentration (pg/mL)	654.79	2735.41
Standard Deviation	14.30	173.74
%CV	2.18	6.35

<u>Inter-Assay Precision:</u> Two samples containing different SARS-CoV-2 Nucleocapsid Protein concentrations were tested in ten independent assays.

Concentration	Sample 1	Sample 2
Number of Assays	10	10
Mean Concentration (pg/mL)	849.8	3309.0
Standard Deviation	67.5	283.8
%CV	7.94	8.58

<u>Biological Samples:</u> 293E cells were transfected with PBL-909 (100 X 10⁶ cells) overnight and harvested along with control, untransfected 293E cells. Cells were lysed with Triton-X 100 lysis buffer (50 mM Tris, 150 mM NaCl, 1% TX-100, 5 mM EDTA). Cell culture supernatant and cell lysate were assayed for natural SARS-CoV-2 Nucleocapsid Protein.

	Untransfected Cell Culture	Transfected Cell Culture
	Supernatant	Supernatant
Concentration	Non-detectable	66,239.3
(ng/mL)		

	Untransfected Cell Lysis	Transfected Cell Lysis
Concentration	Non-detectable	9,437.5
(ng/mL)		

Troubleshooting Guide:

Problem	Probable Cause	Solution
High Background	Background wells were contaminated	Avoid cross-well contamination by using the provided plate sealers. Use multichannel pipettes and change tips between pipetting samples and reagents.
	Insufficient washes	Increase number of washes. Increase soaking time between washes prior to addition of substrate solution.
	TMB Substrate Solution was contaminated	TMB Substrate Solution should be clear and colorless prior to addition to wells. Use a clean container prior to pipetting substrate solution into wells.
No or poor signal	Detection Antibody, Avidin-HRP or Substrate solution were NOT added	Down the essey and follow the protocol
	Wrong reagent or reagents were added in wrong sequential order	Rerun the assay and follow the protocol.
	Insufficient plate agitation	The plate should be agitated during all incubation steps using a plate shaker at a speed where solutions in wells are within constant motion without splashing.
	The wash buffer contains Sodium Azide (NaN3)	Avoid Sodium Azide contamination in the wash buffer as it inhibits HRP activity.
	Incubations were done at an inappropriate temperature, timing or without agitation	Rerun the assay and follow the protocol.
Low or poor standard curve	The standard was incorrectly reconstituted or diluted	Adjust the calculations and follow the protocol.
signal	Standard was inappropriately stored	Store the reconstituted standard stock solution in polypropylene vials at -70°C. Avoid repeated freeze-thaw cycles.
	Reagents added to wells with incorrect concentrations	Check for pipetting errors and the correct reagent volume.

Problem	Probable Cause	Solution		
Signal is high, standard curves have saturated	Standard reconstituted with less volume than required	Reconstitute new lyophilized standard with the correct volume of solution recommended in the protocol.		
signal	Standards/samples, detection antibody, Avidin-HRP or substrate solution were incubated for too long	Rerun the assay and follow the protocol.		
Sample readings	Samples contain no or below detectable levels of the analyte	If samples are below detectable levels, it may be possible to use a larger sample volume. Contact technical support for appropriate protocol modifications.		
are out of range	Samples contain analyte concentrations greater than highest standard point	Samples may require dilution and analysis.		
	Multichannel pipette errors	Confirm that pipette calibrations are accurate.		
High variation in samples and/or	Plate washing was not adequate or uniform	Ensure pipette tips are tightly secured. Ensure uniformity in all wash steps.		
standards	Non-homogenous samples	Thoroughly mix samples before assaying.		
	Samples may have high particulate matter	Remove particulate matter by centrifugation.		
	Cross-well contamination	Do not reuse plate sealers.		
		Always change tips for reagent additions. Ensure that pipette tips do not touch the reagents on the plate.		

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