

# SARS-CoV-2 S1 RBD3 Indian B.1.617.2 (delta)

Catalogue #	P-372-100
Description:	Protein contains amino acids 319-537, mutations L452R, T478K plus two extra amino acids (AS) in N-terminus and His-6 tag at C-terminus and GSG linker between protein and tag.
Uniprot ID:	P0DTC2
MW:	25.89 kDa
Host:	CHO-based cell line (expressed by QMCF Technology)
Purification:	Purified by Ni-affinity chromatography and gel- filtration from serum-free CHO growth media, sterile filtrated
Concentration:	1 mg/ml
Concentration: Buffer:	1 mg/ml PBS pH 7.4
Buffer:	PBS pH 7.4
Buffer: Endotoxine:	PBS pH 7.4 NA Measured by its binding ability to ACE2 protein
Buffer: Endotoxine: Bioproperties:	PBS pH 7.4 NA Measured by its binding ability to ACE2 protein by OCTET RED96 system. SDS-PAGE, NanoDrop A280, Analytical SEC,

100020000	+D1		
			245 180 140
	-		100
	-		75
	-		60
	÷.,		45
	- 8	8	35
	-		25
	-		20
			15 10
1	2 3	3	

DTT

DTT

**Figure 1.** Simply Blue Safe stained SDS-PAGE analysis of SARS-CoV-2 S1 RBD3 Indian B.1.617.2 (delta). 4-12% gradient gel is used for analysis. Lane 1. 0.8 µg SARS-CoV-2 S1 RBD3 Indian B.1.617.2 (delta) (-DTT). Lane 2. Protein marker (Smobio). Lane 3. 0.8 µg SARS-CoV-2 S1 RBD3 Indian B.1.617.2 (delta) (+DTT).

#### Peak Table

Peak #	RT (min)	Area	Area %
1	10.462	2333.08	100.00

## Chromatogram

ICO944 #174 - DAD1B,Sig=220,16 Ref=of

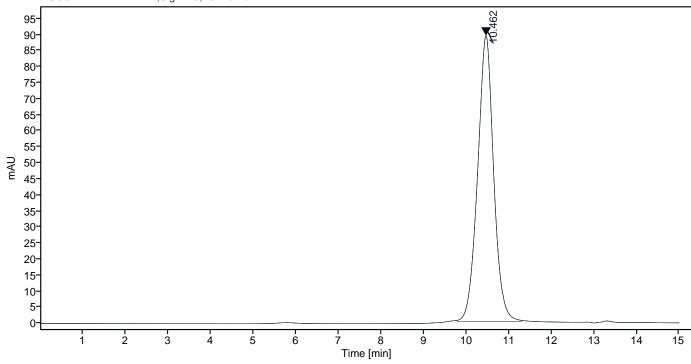


Figure 2. HPLC analytical SEC for final product.

#### Peak Table

Peak #	RT (min)	Area	Area %
1	10.461	2352.93	100.00

## Chromatogram

ICO944 #174 3x - DAD1B,Sig=220,16 Ref=of

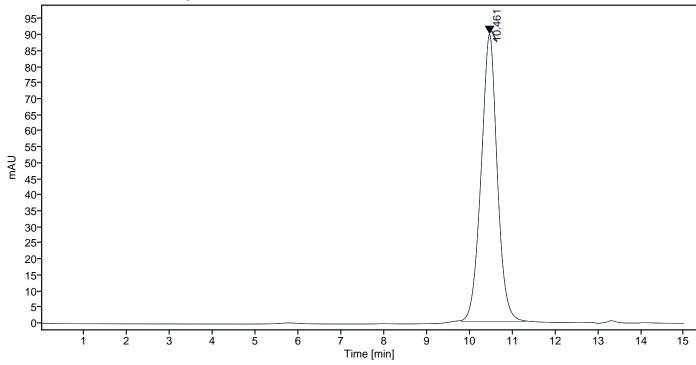


Figure 3. HPLC analytical SEC after 3 freeze-thaw cycles.

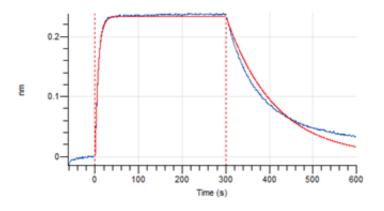


Figure 4. Octet RED96e analysis of SARS-CoV-2 Spike Delta protein binding to the ACE2 receptor.