



SCIENTIFIC VETERINARY INSTITUTE "NOVI SAD"
21000 NOVI SAD, Rumenčki put 20

☎ 021/4895-308; Fax: 021/518-544; e-mail: niv@niv.ns.ac.rs

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Научни институт за ветеринарство „НОВИ САД“
У Новом Саду са потпуном одговорношћу

29 DEC 2020
19381-150
20 год
НОВИ САД

REPORT OF ANALYSIS

No: 19010 from 05.11.2020.

Owner: „REAL S doo“, Omladinskih brigada 86J/15, 11070, Belgrade, Serbia
Ordering party: Owner of the material

Sample : Nano-silver impregnated canvas for the production of protective masks:
“ALBO nanosilver filter”

Origin of the sample: owner
No. and sign of the sample: 19010
Place and date of sampling: Novi Sad, Serbia, 15.09.2020.
Number of samples: 1
Sampled by: Owner and ordering party

Analysis: Proving of antiviral activity of nano-silver impregnated canvas “ALBO nanosilver filter” against SARS-CoV-2 virus with molecular technique real-time RT-PCR (advanced “viability PCR” method)

Anamnesis:

Condition of the sample upon receiving: Appropriate
Date of receiving the sample: 15.09.2020.
Examination completed: 28.12.2020.

Remark if any:

*The method marked with * are beyond the scope of accreditation.*

HEAD OF DEPARTMENT FOR
epizootiology, clinical diagnostic and
laboratory analysis



Sava Lazić, PhD, DVM, Principal Research Fellow

Distribute to:

1. Owner – Ordering party 2 copies
2. Archive

STATEMENT:

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3. Laboratory NIV-NS includes a disclaimer in the report for the information and sample received from the customer and report refers only to the sample examined, as it has been received.

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DEPARTMENT FOR EPIZOOTIOLOGY, CLINICAL EXAMINATION
AND LABORATORY ANALYSIS

Laboratory for Virology

Report No.: 19010

RESULTS OF ANALYSIS

Examined characteristics	Examination method	Referral value
Analysis of antiviral activity *	Molecular technique <i>real-time RT-PCR</i> (advanced "Viability PCR" metod)	Positive / negative

Description of samples used and procedure:

- Working virus stock in the experiment was the isolate of *SARS-CoV-2* virus, cultivated on VERO cell culture
- Test samples 1.1 P – 1.6 P are six samples of material (*ALBO nanosilver filter*) exposed to (soaked with) *SARS-CoV-2* virus isolate for 60 minutes. Median values of results after testing in duplicates are shown in table.
- Control samples PC 1-P – PC 3-P are three samples of *SARS-CoV-2* virus isolate that were used for testing. Median values of results after testing in duplicates are shown in table.
- Water as a negative test control and NTC as a mixture of test reagents.
- IC - internal control of the reaction whose detection indicates that there was no inhibition of the reaction.

Findings in samples:

Label of the sample	Test samples		Internal controls (IC)	Control samples		Internal controls (IC)
	Label	Ct value SARS-CoV-2	Ct values of IC	Label	Ct value SARS-CoV-2	Ct values of IC
ALBO nanosilver filter incubation with virus 60 minute	1.1 P	37.65	32.44	PC 1-P	28.33	31.76
	1.2 P	37.67	32.32	PC 2-P	28.49	31.41
	1.3 P	38.59	32.04	PC 3-P	28.33	31.59
	1.4 P	38.63	31.95	Water	Negative	31.30
	1.5 P	38.50	32.03	Water	Negative	31.39
	1.6 P	39.00	32.02	Water	Negative	31.52
* used method are beyond the scope of accreditation				NTC	Negative	Not done
				NTC	Negative	Not done
				NTC	Negative	Not done

Comment:

If we compare the values obtained by "viability real-time RT-PCR" method, which detects the RNA genome of only viable (complete / infectious) virus particles, we conclude that the ALBO nanosilver filter showed significant antiviral effect on SARS-CoV-2 virus which was reflected in a significant reduction in number of detectable viral genomes relative to the virus being tested. The mean Ct value determined by incubating the virus with an ALBO nanosilver filter for 60 minutes was Ct 38.34, and the mean Ct value of the virus tested was Ct 28.38. The maximum difference between the control and test samples was ≥ 10.67 Ct, and the smallest ≥ 9.16 Ct values (mean $\geq 38.34 - 26.38 = \geq 9.96$ Ct). Translated into the number of viral particles, i.e. detectable genomes of viable virus particles SARS-CoV-2, the difference between the tested material ALBO nanosilver filter with virus and control samples (virus tested), i.e. the reduction in the number of viable virus particles was from $\geq 2.75 \log_{10}$ up to $\geq 3.20 \log_{10}$, with an average of $\geq 2.99 \log_{10}$ number of detectable viable viral genomes (about 1,000 times less viral genomes in treated samples)

Novi Sad, 28.12.2020.

ANALYSED BY

dr sci. vet. med. Tamaš Petrović



VERIFIED BY:

dr sci. vet. med. Tamaš Petrović