

# *Genomika i molekularna dijagnostika SARS-CoV-2*

Ivan-Christian Kurolt

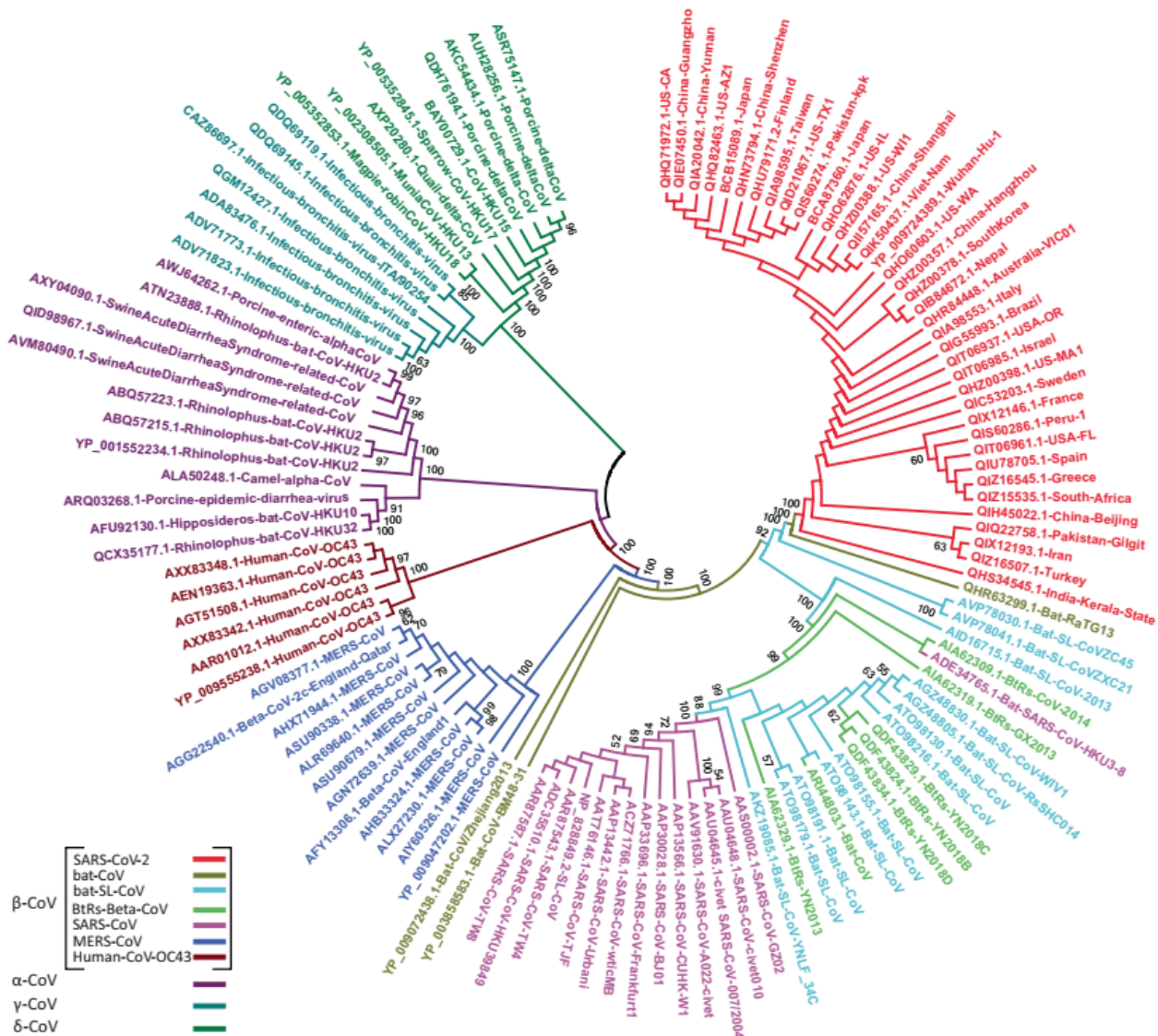
Klinika za infektivne bolesti “Dr. Fran Mihaljević”

Jedinica za znanstvena istraživanja

ikurolt@bfm.hr



# Coronaviridae - Orthocoronavirinae

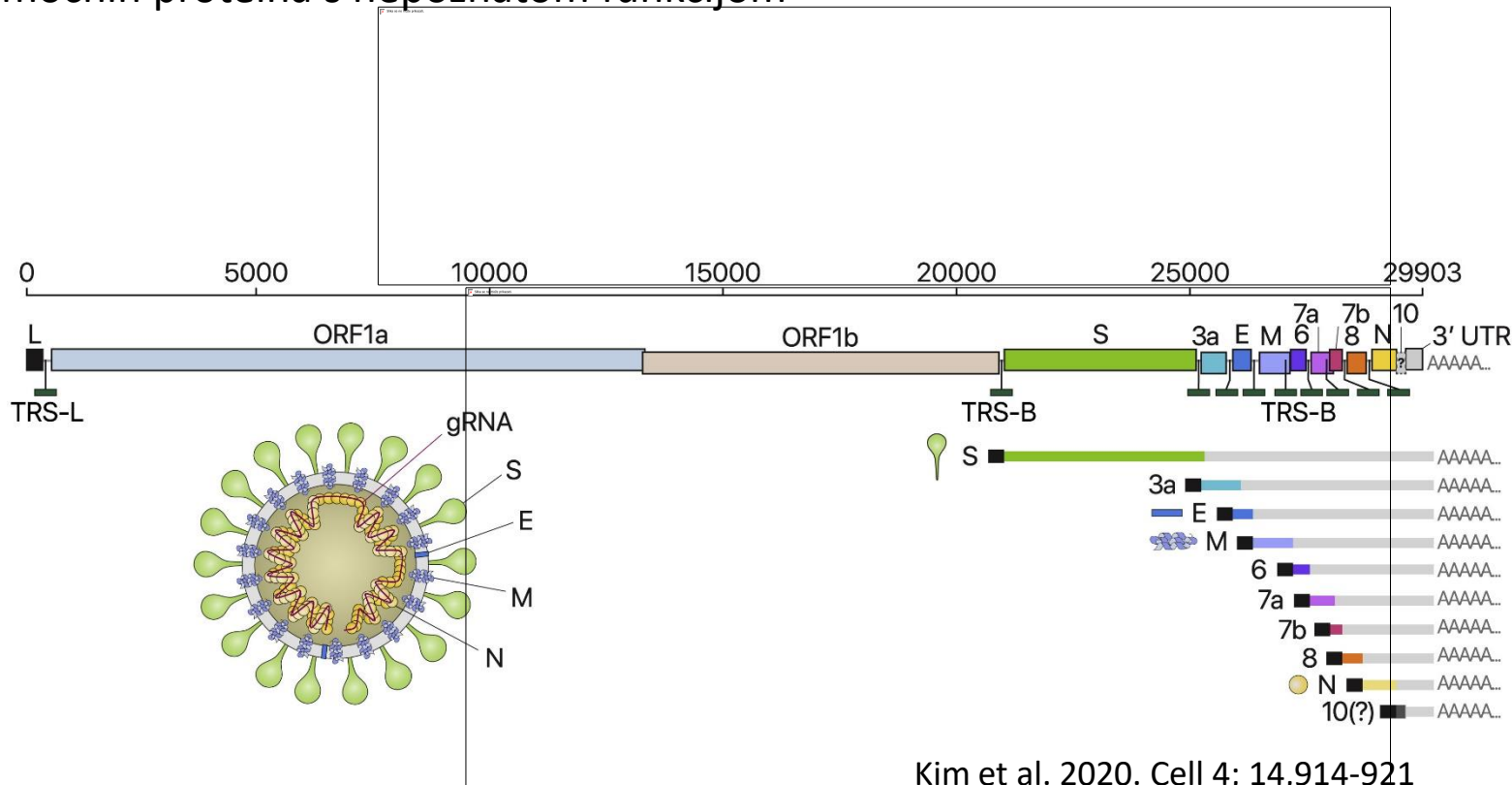


ejiang2013  
J9  
CCDC1  
U5



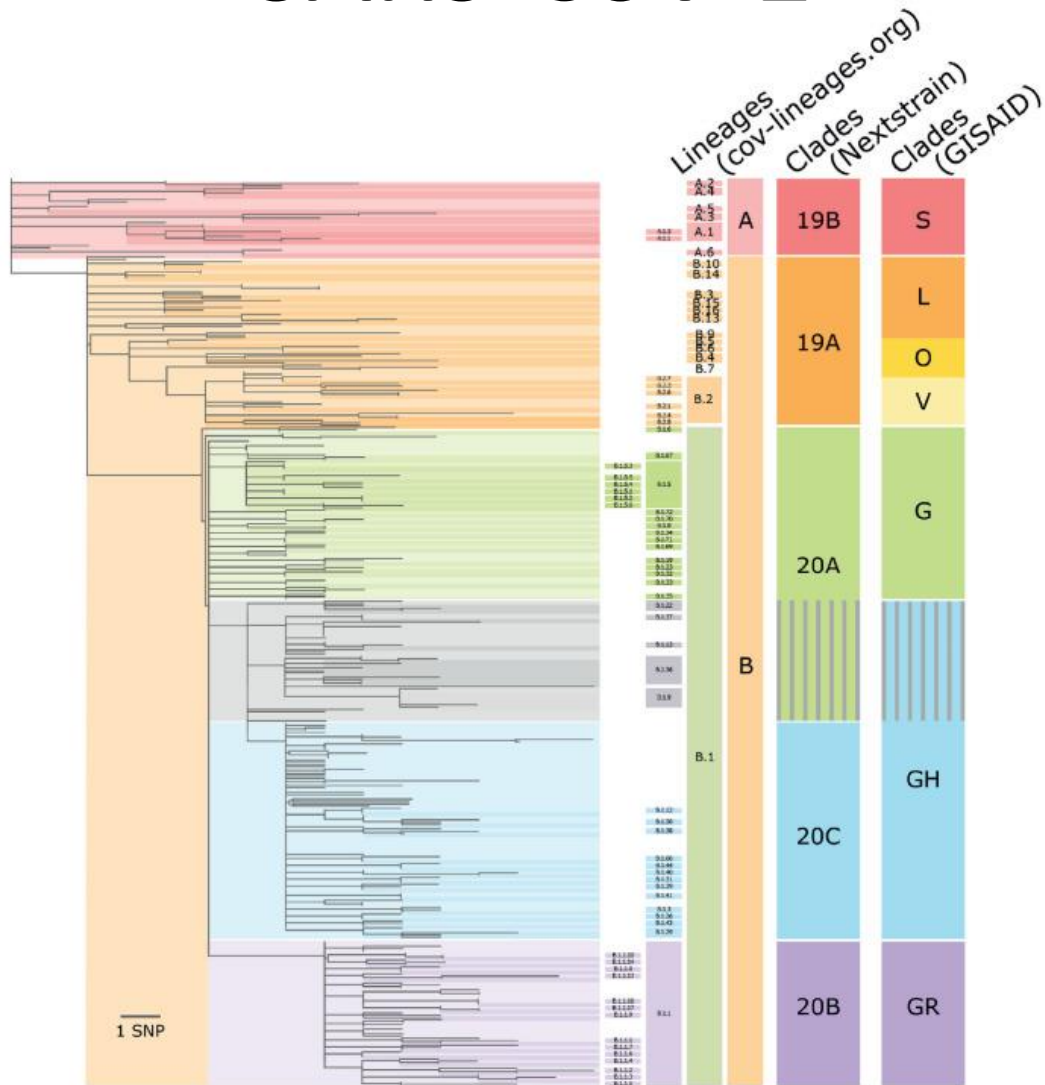
# Coronavirus

- + ss RNA genom, 29,9 kB
- 14 ORF, 26 različitih proteina
  - Strukturnih: Spike (S), Envelope (E), Membrane (M), Nucleocapsid (N)
  - 16 nestrukturiranih proteina – prvenstveno potrebno u replikaciji
  - 6 pomoćnih proteina s nepoznatom funkcijom





# Filogenetska raznolikost SARS-CoV-2





# Implementacija

01.12.2019.: Prvi slučaj COVID-19 u Wuhanu, Kina

21.01.2020.: Odluka o uvođenju testa na COVID-19

27.01.2020.: Isporuka pozitivnih kontrola za COVID-19

29.01.2020.: Isporuka početnica i fluorescentnih proba

30.01.2020.: Zaprimanje prvog uzorka na testiranje COVID-19

WHO PHEIC

14.02.2020.: 1. autohtoni slučaj u Italiji

25.02.2020.: 1. dokazani slučaj u Hrvatskoj



**EVA**g  
European Virus Archive Global

European Virus Archive - GLOBAL



# Molekularni testovi za SARS-CoV-2



World Health Organization

Health Topics ▾

Countries ▾

< Coronavirus disease 2019

< Technical guidance

Laboratory guidance

Early investigations

Patient management

Surveillance and case definitions

Infection prevention and control

Points of entry and mass gatherings

## 2. Mole

Several assays may

In-house de

Some groups will be willing to recommend

Summary t

Country

China

Germany

Hong Kong

## *Coronavirus Test Kits Sent to States Are Flawed, C.D.C. Says*

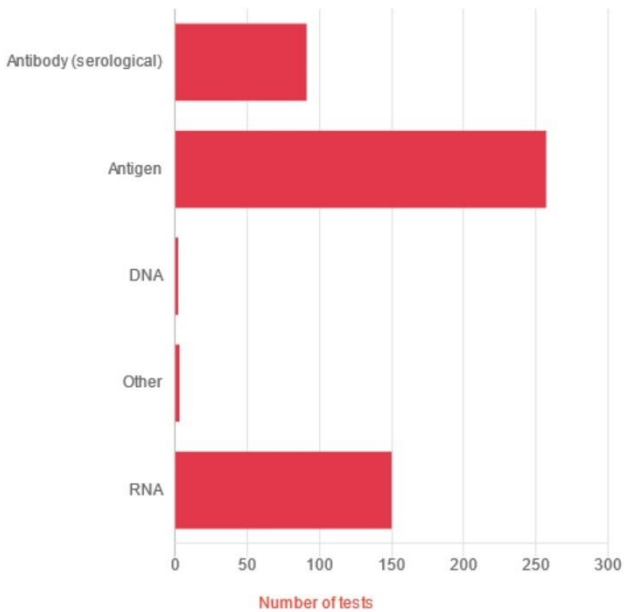
Some tests distributed by the agency deliver “inconclusive” readings. The C.D.C. will need to ship new ingredients, further delaying results.



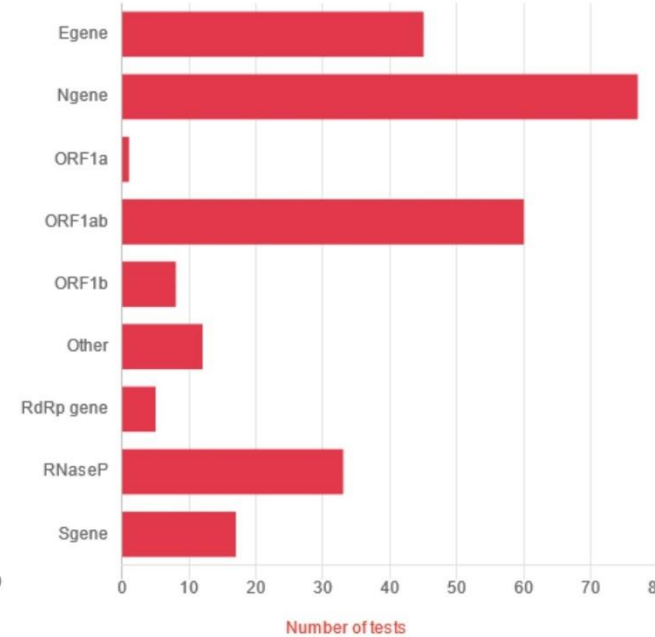


# Molekularni testovi za SARS-CoV-2

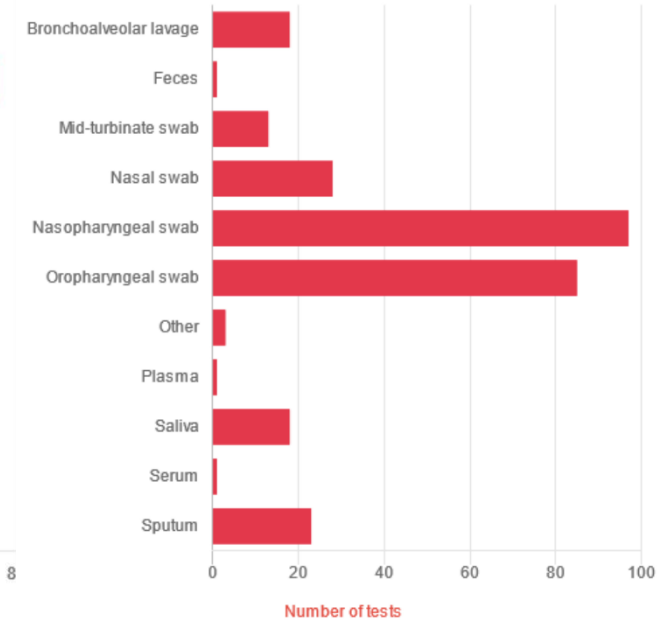
Assay target



Target analyte

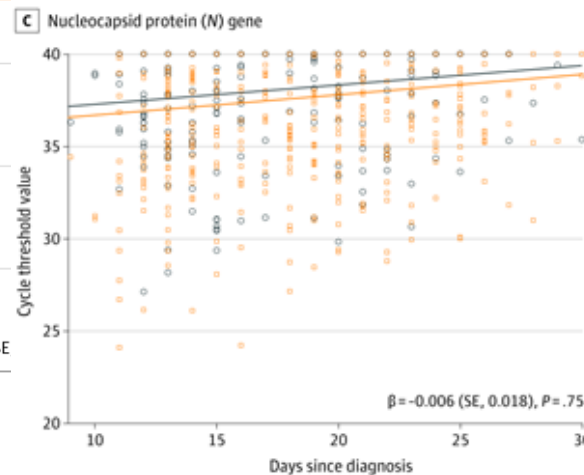
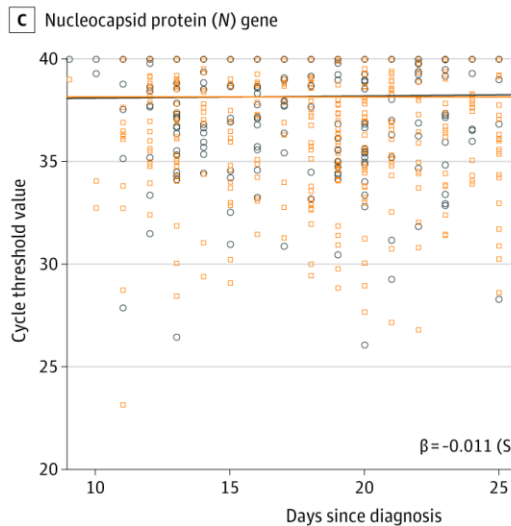
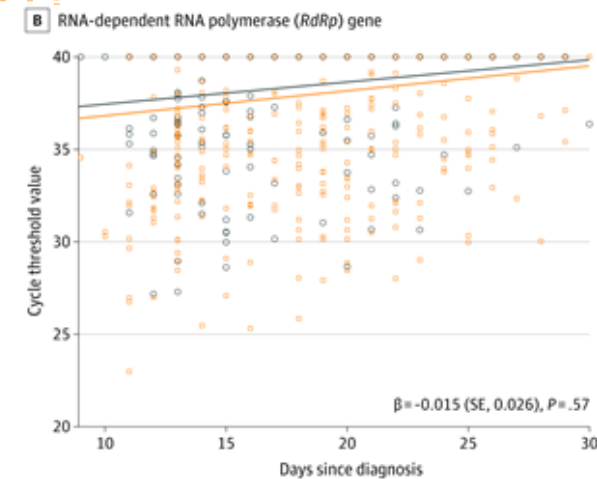
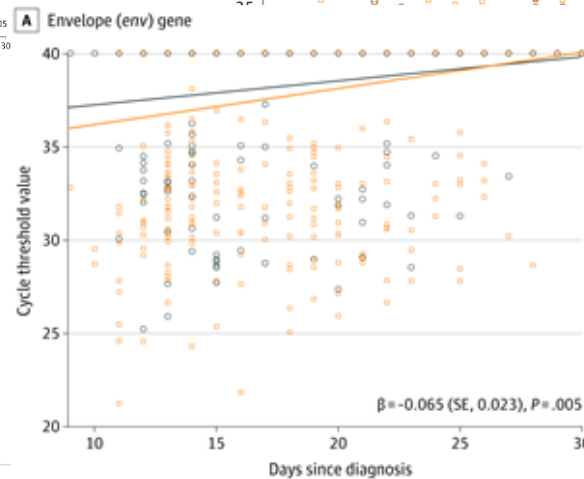
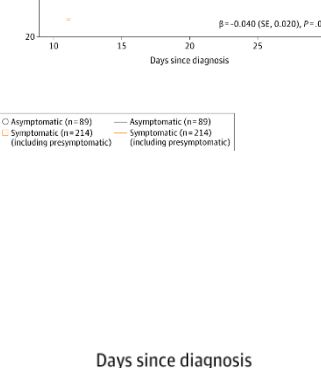
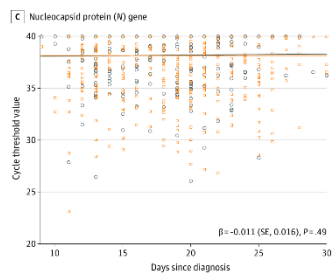
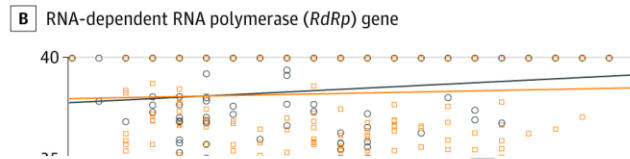
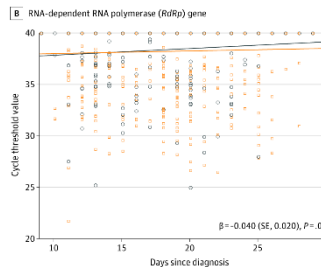
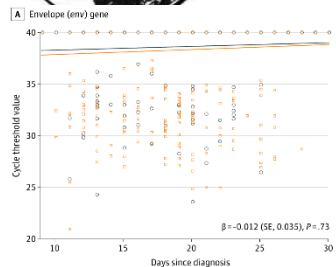


Validated sample types





# SARS-CoV-2 virusno opterećenje



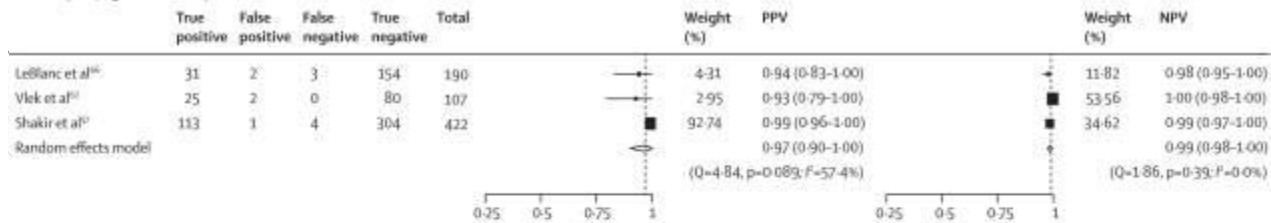
○ Asymptomatic (n = 89)    — Asymptomatic (n = 89)  
□ Symptomatic (n = 214)    — Symptomatic (n = 214)  
(including presymptomatic)    (including presymptomatic)



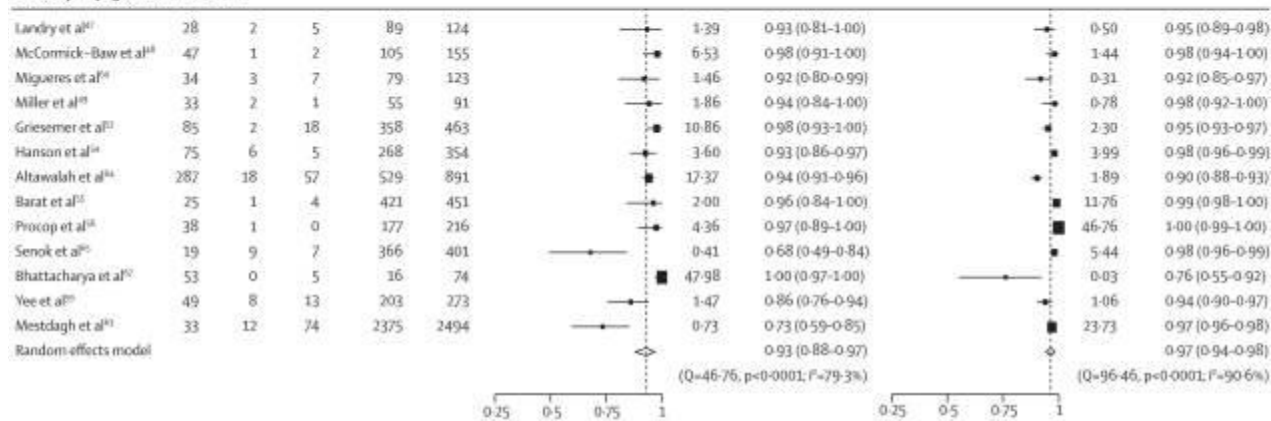


# SARS-CoV-2 virusno opterećenje

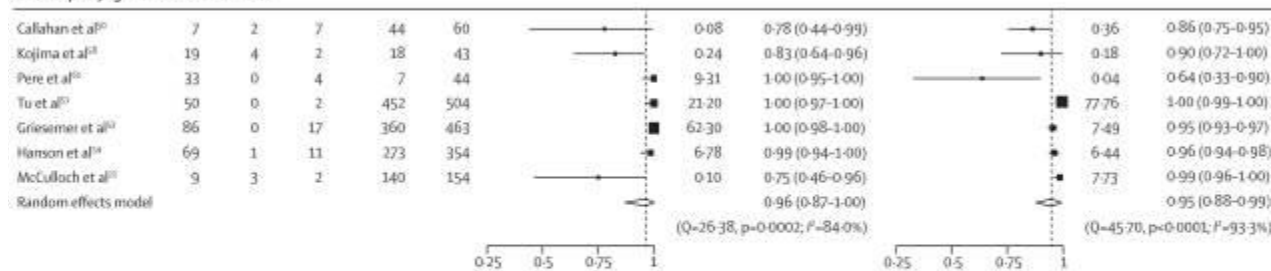
## A Nasopharyngeal swab and pooled nasal and throat swab



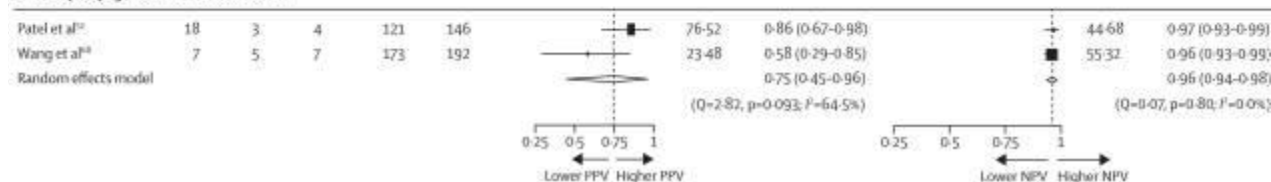
## B Nasopharyngeal swab and saliva



## C Nasopharyngeal swab and nasal swab

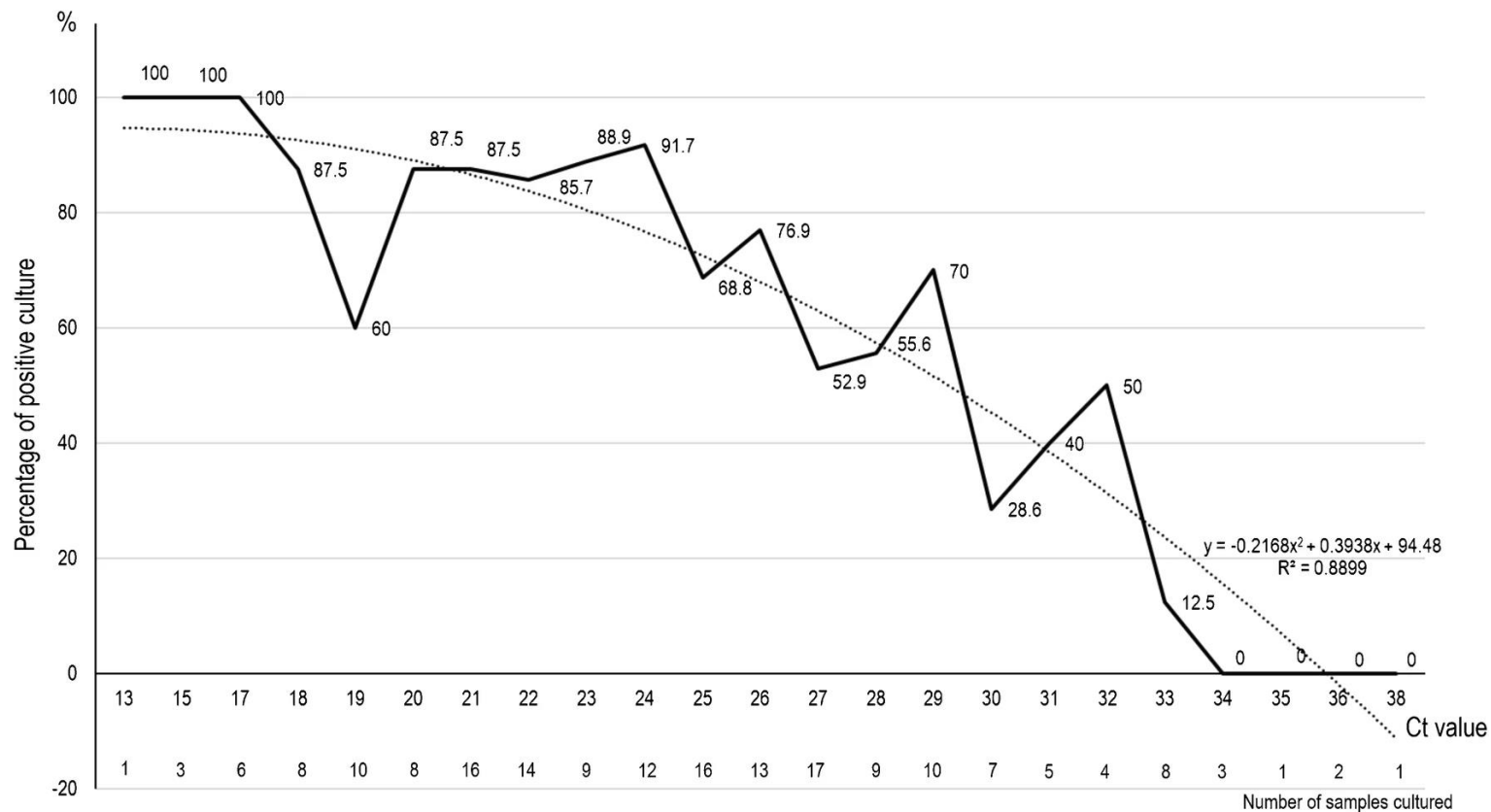


## D Nasopharyngeal swab and throat swab



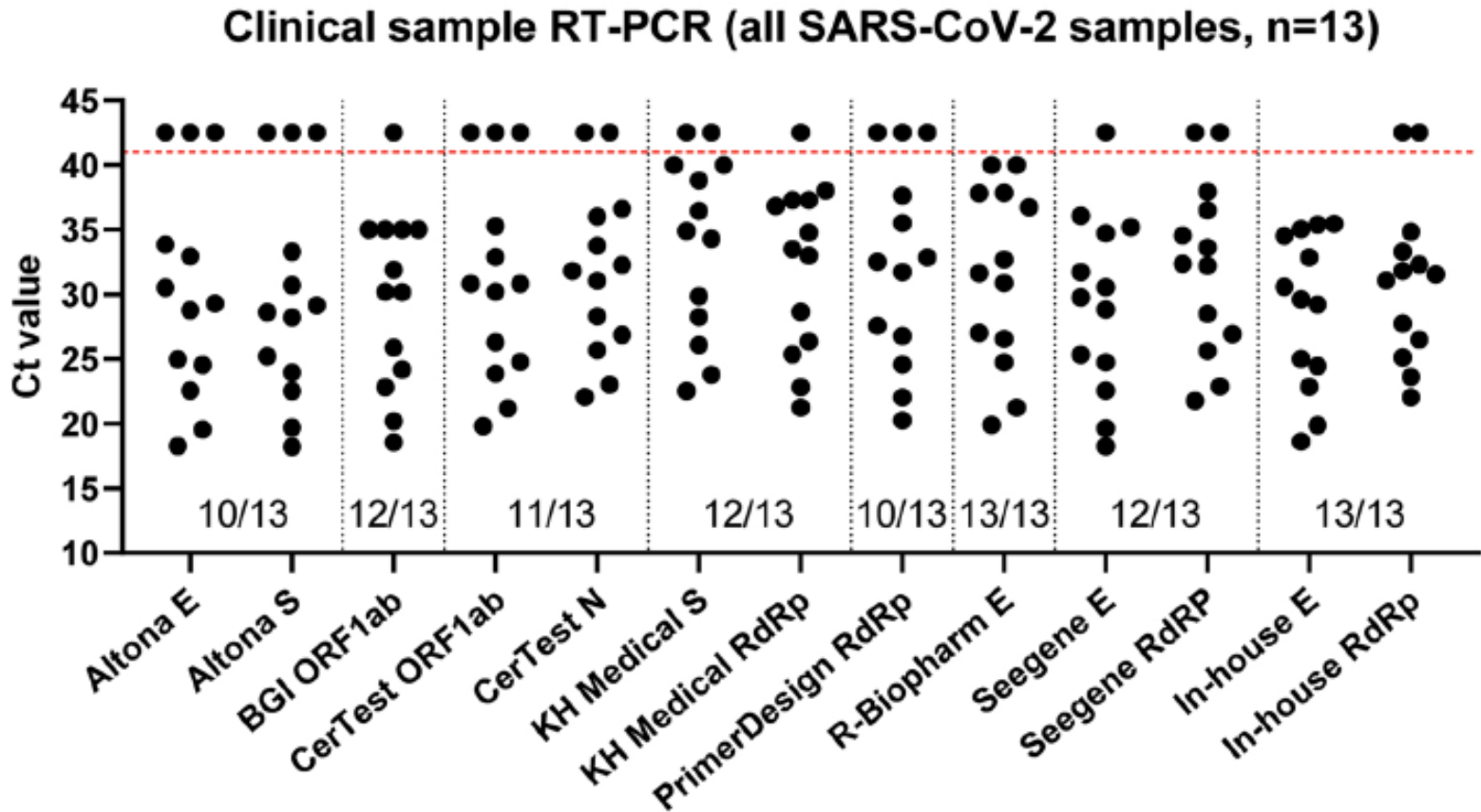


# SARS-CoV-2 virusno opterećenje





# SARS-CoV-2 virusno opterećenje





# SARS-CoV-2 virusno opterećenje

Panel 2	Ref	20	28	Influenza B	RSV	30	34	neg kontrola	21	Influenza A	31	26
Ustanova 1	Rezultat	poz	poz	neg	neg	poz	neg	neg	poz	neg	neg	poz
	gen E	24,5	33,5	-	-	34,5	38,7	-	25,8	-	-	29,2
	gen RDRp	26,3	34,5	-	-	35,3	-	-	27,4	-	-	32
Ustanova 2	Rezultat	poz	neod.	neg	neg	neod.	neg	neg	poz	neg	neod.	poz
	gen E	26	34	-	-	35,5	-	-	28,1	-	35,6	32,8
	gen RdRP	30	-	-	-	-	-	-	33	-	-	38,5
Ustanova 3	Rezultat	poz	poz	neg	neg	poz	neg	neg	poz	neg	neg	poz
	gen E	23	33	-	-	31	38	-	26	-	-	31
	gen RdRp	28	38	-	-	35	-	-	32	-	-	35
Ustanova 4	Rezultat	poz	poz	neg	neg	poz	poz	neg	poz	neg	poz	poz
	gen E	20,54	29,57	-	-	30,40	36,24	-	22,45	-	33,08	27,01
	gen RdRp	22,26	30,97	-	-	32,53	38,60	-	24,44	-	33,90	28,67
Ustanova 5	Rezultat	POZ	POZ	neg	neg	repeat	repeat	neg	POZ	neg	repeat	POZ
	gen: E	25,47	34,56	-	-	36,46	37,24	-	27,48	-	37,19	30,29
	gen: N	24,68	34,00	-	-	36,43	37,26	-	26,72	-	37,08	29,54
	gen: RdRP	26,57	31,48	-	-	-	-	-	28,23	-	-	30,34
Ustanova 6	Rezultat	POZ	GR	NEG	NEG	GR	NEG	NEG	POZ	NEG	GR	POZ
	gen E	22,14	32,21	/	/	34,39	/	/	25,31	/	37,31	28,69
	gen RdRp	23,69	/	/	/	/	/	/	26,76	/	/	29,36

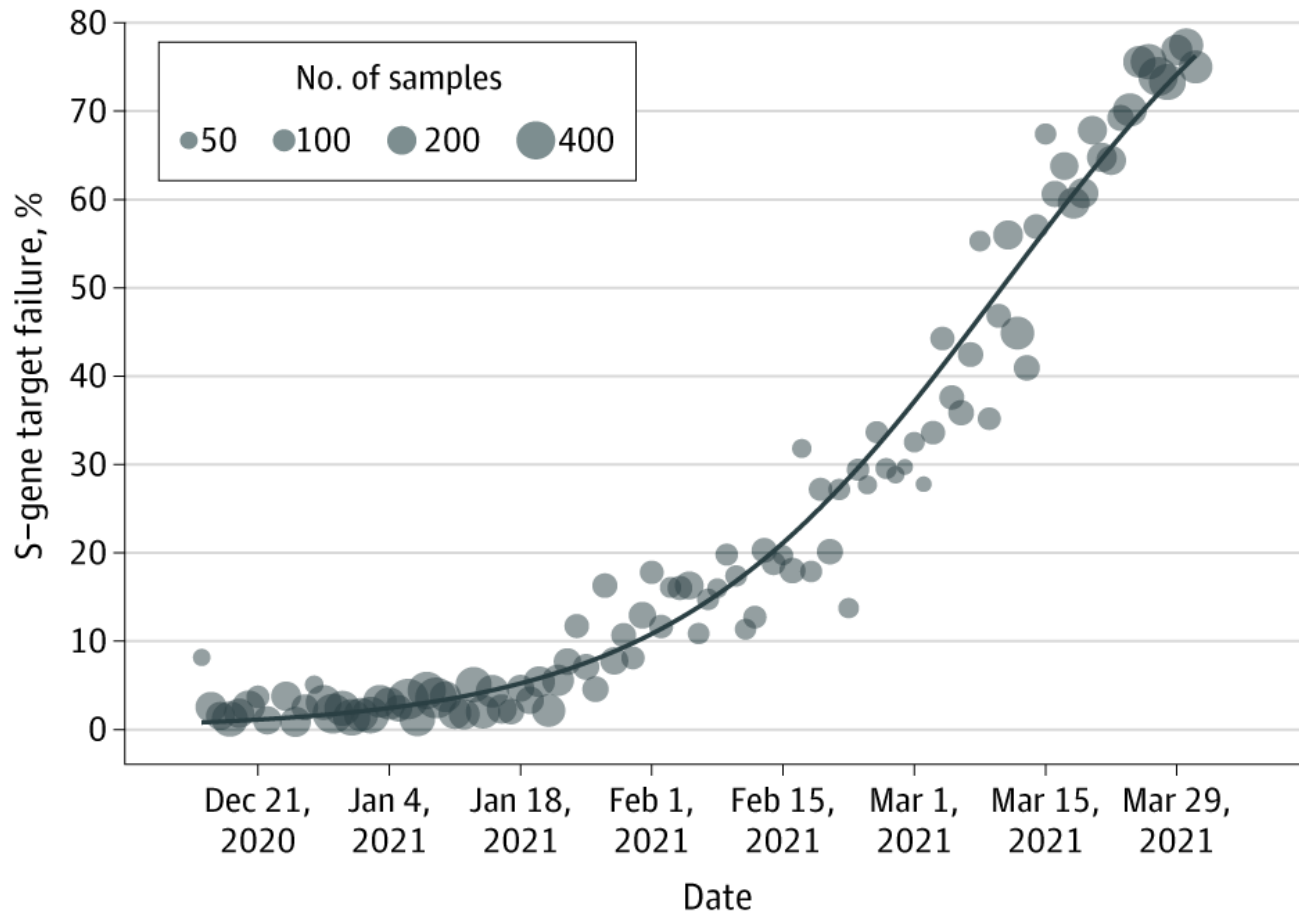


# SARS-CoV-2 virusno opterećenje

Barkod Hanks + iClean		Rez Hanks + iClean		Barkod Gongdong + iClean		Rez Gongdong + iClean		Razlika
0204*577		17,4		0204*649		23,5		6,1
0204*574		16,7		0204*647		24,3		7,6
Barkod Hanks + iClean		17,6		Barkod Gongdong mediji + iClean		21,7		4,1
0204*581		17,6		0204*646		21,7		4,1
Gondong štapić		18		0604*335		23,7		5,7
0704*210		22,6		0704*322		27,9		5,3
0704*211		21,4	26,2	0604*349		32,9		6,7
0704*241		23,8	21,6	0604*327		31,7		10,3
0704*310		19,3	18,4	0604*611		26,5		4,9
0704*352		35,1	20	0704*318		26,4		2,6
0704*448		29,3	16,5	0604*626		23,5		5,1
0704*458		20,5	16,2	0704*433		26,1		6,8
0704*510		15,5	22,5	0604*729		23,5		3,5
0704*513		20,3	25,8	0704*435		37		1,9
0704*514		18,1	16,5	0604*738		25,2		8,7
0604*526		16,2		0704*534		29,9		0,6
0604*541		22,5		0604*741		24,4		8,2
0604*483		25,8		0704*530		24,9		4,4
				0604*742		33,2		10,7
				0704*536		21,4		5,9
				0604*744		28		2,2
				0704*538		29,8		9,5
				0604*738		25,2		8,7
				0704*538		23,4		5,3
				0604*741		24,4		8,2
				0604*742		33,2		10,7
				0604*745		28		2,2



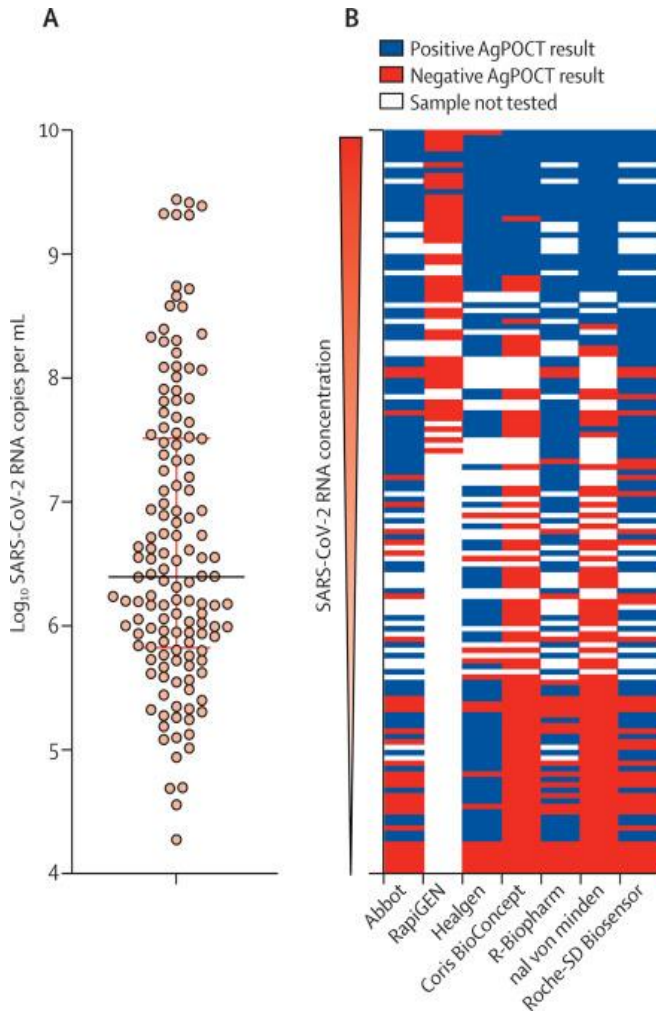
# SARS-CoV-2 virusno opterećenje



Brown et al. JAMA. Published online April 8, 2021



# Antigeniski testovi



**Table 2** Antigen-based detection methods utilized for COVID-19 diagnostics

Ref.	Detection method	Antigen	Sample type	Num. of samples	Days since symptom onset (days)	Sensitivity (%) / LOD	Specificity (%)
[106]	Fluorescence immunochromatographic assay	Nucleocapsid protein	NPS and urine	239	3	68	100
[115]	Fluorescence immunochromatographic assay	Nucleocapsid protein	Saliva	103	9	11.7	N/A
[116]	Chemiluminescence enzyme immunoassay	Nucleocapsid protein	NPS	313	N/A	55.2	99.6
[117]	Fluorescence immunochromatographic assay	Nucleocapsid protein (Genscript Cat #Z03488 & Genemedi GMP-V-2019nCoV-N002)	Non-clinical samples (in PBS buffer)	N/A	N/A	Genemedi – 0.65 ng/mL, Genscript – 3.03 ng/mL	N/A
[118]	Fluorescence immunochromatographic assay	SARS-CoV-2 antigen	NPS	19	N/A	N/A (low)	N/A
[114]	Fluorescence immunochromatographic assay	SARS-CoV-2 antigen	NPS and OPS	127	<7 for 93.7% of samples	93.9	100
[119]	GICA	Nucleoprotein	NPS	138	N/A	50	100
[120]	GICA	Nucleoprotein	NPS	148	Median: 4, mean: 6.6, range: 0–34	30.2	100
[121]	GICA	Nucleoprotein	NPS	328	N/A	57.6	99.5

Jayamohan et al. Anal Bioanal Chem. 2021;413(1):49-71



# Serologija

**Table 3** Antibody-based tests utilized for COVID-19 diagnostics

Ref.	Detection method	Antibody	Sample type	Num. of samples	Seroconversion (days)	Sensitivity	Specificity
[129]	GICA	IgG and IgM	Serum/whole blood	134	7	96.8 <sup>1</sup>	N/A
[130]	GICA	IgG and IgM	Serum/whole blood	525	N/A	88.66 <sup>2</sup>	90.63 <sup>2</sup>
[131]	GICA	IgG and IgM	Serum	814	5	86.89 <sup>2</sup>	99.39 <sup>2</sup>
[132]	GICA	IgG and IgM	Serum	179	8	95.10 <sup>2</sup>	91 <sup>2</sup>
[125]	CLIA	IgG and IgM	Serum	285	13	100 <sup>1</sup>	N/A
[123]	CLIA	IgG and IgM	Serum	159	14	91.14 <sup>2</sup>	80 <sup>2</sup>
[124]	CLIA	IgG and IgM for nucleocapsid protein	Serum	222	4	81.5 <sup>2</sup>	96.6 <sup>2</sup>
[133]	ELISA	IgG and IgM	Serum	238	11	81.3 <sup>2</sup>	N/A
[134]	ELISA	IgG and IgM for nucleocapsid and spike protein	Serum	214	10	82.2	N/A
[135]	ELISA	IgG and IgM	Serum	15	5	N/A	N/A
[136]	ELISA	IgA, IgM, and IgG	Serum	208	5	85.4	N/A
[126]	ELISA	IgG, IgA for spike protein	Serum	61	N/A	N/A	N/A

<sup>1</sup> Highest sensitivity among samples tested

<sup>2</sup> Sensitivity reported as a mean of all samples tested

N/A. data not reported or not relevant in the context of the referenced publication





# Varijante

## VoC – Variants of Concern

WHO label	Lineage + additional mutations	Country first detected (community)	Spike mutations of interest	Year and month first detected	Evidence for impact on transmissibility	Evidence for impact on immunity	Evidence for impact on severity	Transmission in EU/EEA
Alpha	<b>B.1.1.7</b>	United Kingdom	N501Y, D614G, P681H	September 2020	Yes (v) [1]	No	Yes (v) [3, 4]	Dominating
	<b>B.1.1.7+ E484K</b>	United Kingdom	E484K, N501Y, D614G, P681H	December 2020	Yes (v) [1]	Neutralisation (v) [2, 5]	Yes (v) [3]	Outbreaks
Beta	<b>B.1.351</b>	South Africa	K417N, E484K, N501Y, D614G, A701V	September 2020	Yes (v) [6]	Escape (v) [7, 8]	Yes (v) [4, 9]	Community
Gamma	<b>P.1</b>	Brazil	K417T, E484K, N501Y, D614G, H655Y	December 2020	Yes (v) [10]	Neutralisation (v) [11]	Yes (v) [4]	Community
Delta	<b>B.1.617.2</b>	India	L452R, T478K, D614G, P681R	December 2020	Yes (v) [12-14]	Escape (v) [15]		Community



# Varijante

## Vol – Variants of Interest

WHO label	Lineage + additional mutations	Country first detected (community)	Spike mutations of interest	Year and month first detected	Evidence for impact on transmissibility	Evidence for impact on immunity	Evidence for impact on severity	Transmission in EU/EEA
Eta	<b>B.1.525</b>	Nigeria	E484K, D614G, Q677H	December 2020		Neutralisation (m) [5]		Community
Epsilon	<b>B.1.427/B.1.429</b>	USA	L452R, D614G	September 2020	Unclear [12]	Neutralisation (v) [12]		Sporadic/Travel
Theta	<b>P.3</b>	The Philippines	E484K, N501Y, D614G, P681H	January 2021	Yes (m) [1]	Neutralisation (m) [5]		Sporadic/Travel
	<b>B.1.616</b>	France	V483A, D614G, H655Y, G669S	February 2021	Detection (c) [17]			Single outbreak
Kappa	<b>B.1.617.1</b>	India	L452R, E484Q, D614G, P681R	December 2020	Yes (v) [18]	Neutralisation (v) [15, 17]		Outbreaks
	<b>B.1.620</b>	Unclear (b)	S477N, E484K, D614G, P681H	February 2021		Neutralisation (m) [5, 14]		Outbreaks
	<b>B.1.621</b>	Colombia	R346K, E484K, N501Y, D614G, P681H	January 2021	Yes (m) [1]	Neutralisation (m) [5]		Sporadic/



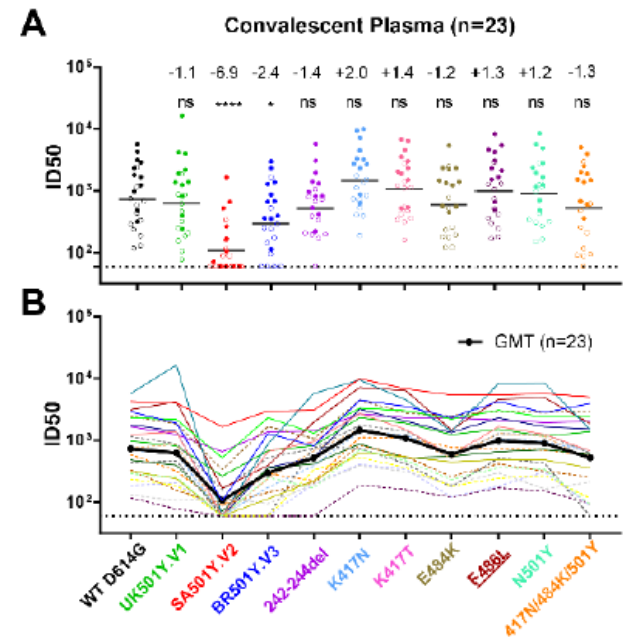
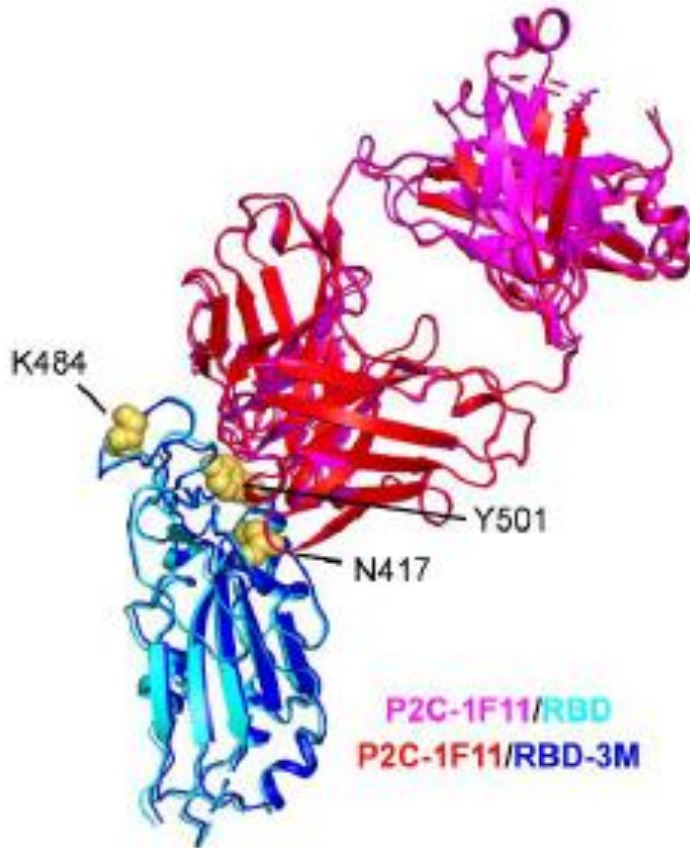
# Varijante

# Vum – Variants under monitoring

WHO label	Lineage + additional mutations	Country first detected (community)	Spike mutations of interest	Year and month first detected	Evidence for impact on transmissibility	Evidence for impact on immunity	Evidence for impact on severity	Transmission in EU/EEA
	<b>B.1.617.3</b>	India	L452R, E484Q, D614G, P681R	February 2021	Yes (m) [1]	Neutralisation (m) [5, 12]		Not detected
	<b>B.1.214.2</b>	Unclear (b)	Q414K, N450K, ins214TDR, D614G	December 2020				Detected (a)
	<b>A.23.1+ E484K</b>	United Kingdom	V367F, E484K, Q613H	December 2020		Neutralisation (m) [5]		Detected (a)
	<b>A.27</b>	Unclear (b)	L452R, N501Y, A653V H655Y	December 2020	Yes (m) [1]	Neutralisation (m) [12]		Detected (a)
	<b>A.28</b>	Unclear (b)	E484K, N501T, H655Y	December 2020		Neutralisation (m) [5]		Detected (a)
	<b>C.16</b>	Unclear (b)	L452R, D614G	October 2020		Neutralisation (m) [5]		Detected (a)
	<b>C.37</b>	Peru	L452Q, F490S, D614G	December 2020				Detected (a)
	<b>B.1.351+P384L</b>	South Africa	P384L, K417N, E484K, N501Y, D614G, A701V	December 2020	Yes (v) [6]	Escape (v) [7, 8]	Unclear [9]	Detected (a)
	<b>B.1.351+E516Q</b>	Unclear (b)	K417N, E484K, N501Y, E516Q, D614G, A701V	January 2021	Yes (v) [6]	Escape (v) [7, 8]	Unclear [9]	Detected (a)
	<b>B.1.1.7+L452R</b>	United Kingdom	L452R, N501Y, D614G, P681H	January 2021	Yes (v) [1]	Neutralisation (m) [12]	Yes (v) [3]	Detected (a)
	<b>B.1.1.7+S494P</b>	United Kingdom	S494P, N501Y, D614G, P681H	January 2021	Yes (v) [1]	Neutralisation (m) [15]	Yes (v) [3]	Detected (a)
	<b>C.36+L452R</b>	Egypt	L452R, D614G, Q677H	December 2020		Neutralisation (m) [12]		Detected (a)
	<b>AT.1</b>	Russia	E484K, D614G, N679K, ins679GIAL	January 2021		Neutralisation (m) [5]		Detected (a)
Iota	<b>B.1.526</b>	USA	E484K, D614G, A701V	December 2020		Neutralisation (m) [5]		Detected (a)
	<b>B.1.526.1</b>	USA	L452R, D614G	October 2020		Neutralisation (m) [12]		Detected (a)
	<b>B.1.526.2</b>	USA	S477N, D614G	December 2020				Detected (a)
	<b>B.1.1.318</b>	Unclear (b)	E484K, D614G, P681H	January 2021		Neutralisation (m) [5]		Detected (a)
Zeta	<b>P.2</b>	Brazil	E484K, D614G	January 2021		Neutralisation (m) [5]		Detected (a)
	<b>B.1.1.519</b>	Mexico	T478K, D614G	November 2020		Neutralisation (m) [12]		Detected (a)
	<b>AV.1</b>	United Kingdom	N439K, E484K, D614G, P681H	March 2021		Neutralisation (m) [5]		Detected (a)
	<b>P.1+P681H</b>	Italy	D614G, E484K, H655Y, K417T, N501Y, P681H	February 2021		Unclear [24, 25]		



# Varijante



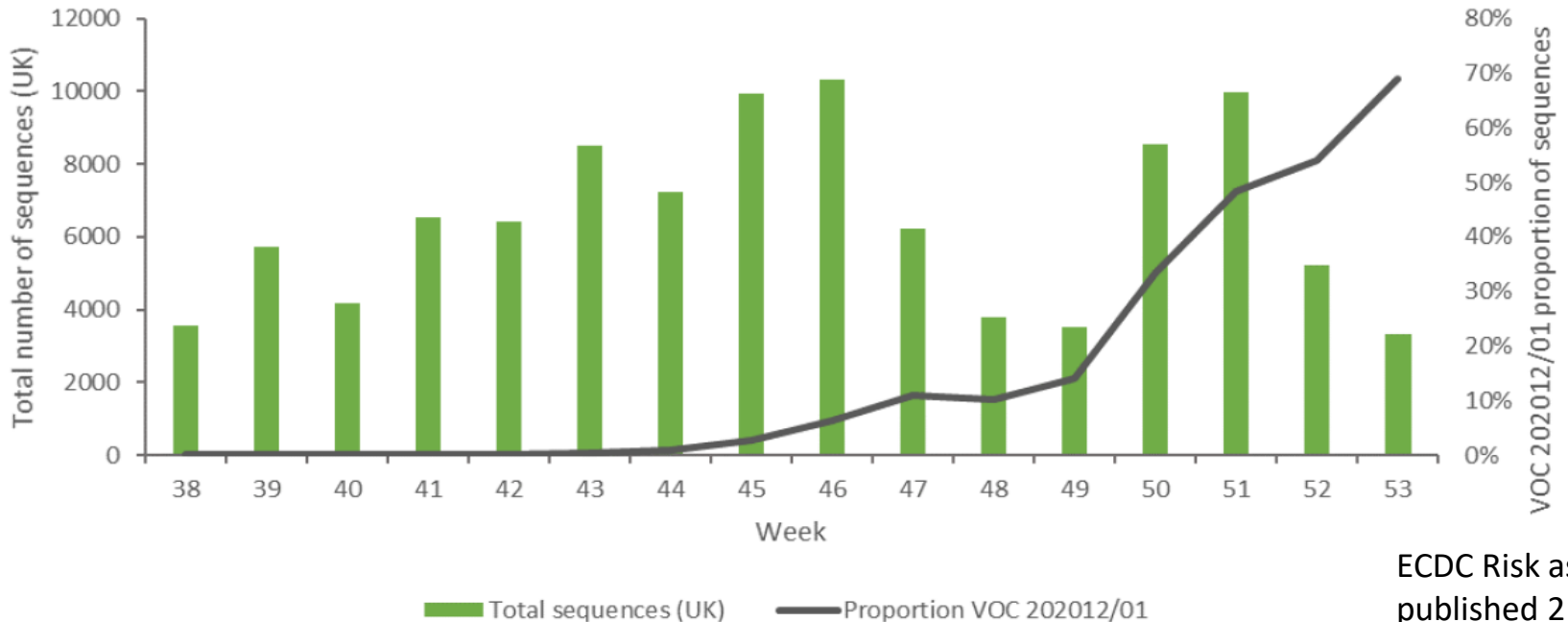
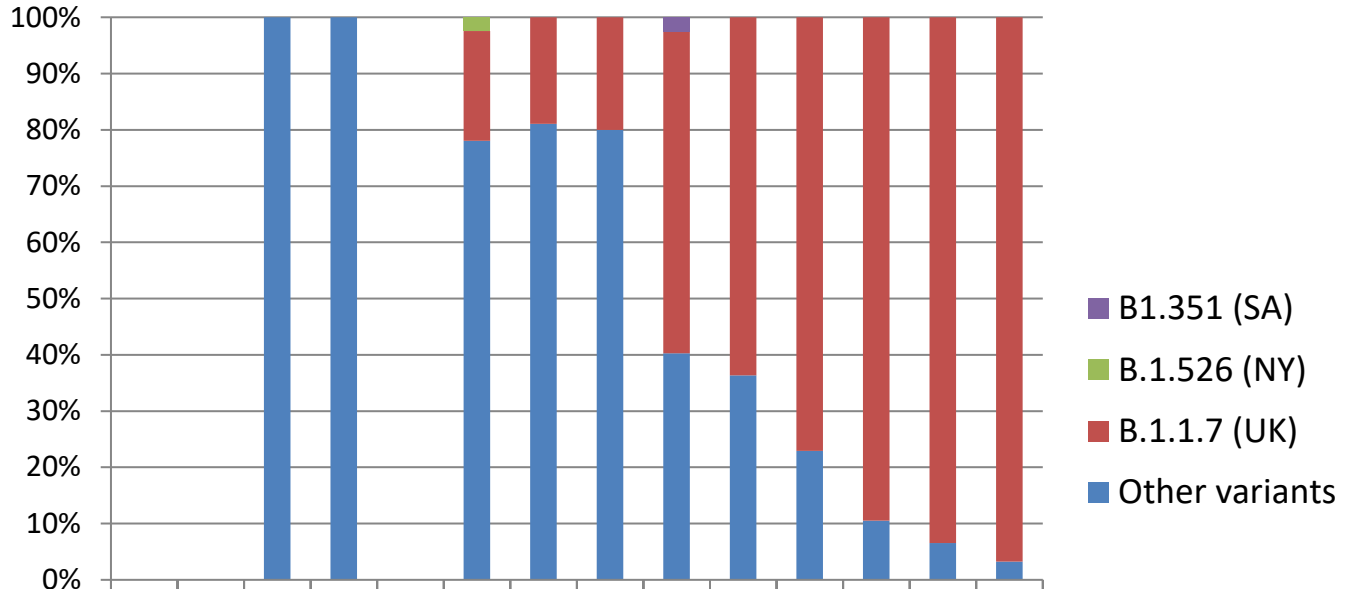
Fold changes in ID50	WT	Variants				NTD		RBD			
C-Plasma1	+1.0	-1.5	-27.6	-2.3	-3.6	+1.5	+1.2	-1.2	+1.5	-1.0	+1.4
C-Plasma2	+1.0	-1.1	-2.6	-1.4	-1.4	+2.3	+1.6	+1.3	+1.3	+1.3	+1.2
C-Plasma3	+1.0	-1.2	-3.7	-1.4	-1.2	+2.4	+1.9	+1.3	+1.6	+1.2	+1.4
C-Plasma4	+1.0	-1.3	-2.5	-1.2	-1.2	+1.6	+1.4	+1.4	+1.3	+1.2	+1.2
C-Plasma5	+1.0	-1.8	-6.7	-2.0	-2.7	+1.9	+1.9	+1.3	+1.3	+1.2	-1.0
C-Plasma6	+1.0	-1.3	-7.2	-2.2	-1.6	+1.5	+1.2	-2.0	+1.1	+1.1	-1.9
C-Plasma7	+1.0	-1.3	-7.2	-1.5	-2.3	+1.8	+1.1	+1.0	+1.2	+1.2	+1.2
C-Plasma8	+1.0	-1.3	BDL	-5.0	-3.4	+1.8	+1.1	-1.3	+1.3	-1.1	-1.2
C-Plasma9	+1.0	-1.5	BDL	BDL	BDL	+1.6	+1.4	+1.0	+1.4	+1.3	-1.2
C-Plasma10	+1.0	+1.3	-18.7	-8.0	-1.6	+2.3	+2.0	-2.1	+1.5	+1.6	-1.7
C-Plasma11	+1.0	-1.2	-6.7	-1.4	-1.1	+1.8	+1.1	+1.1	+1.4	+1.5	+1.3
C-Plasma12	+1.0	+1.1	BDL	-3.1	-1.1	+2.0	+1.5	-1.4	+1.7	+1.5	-1.6
C-Plasma13	+1.0	-1.1	-4.4	-1.0	-1.8	+1.4	+1.3	-1.0	+1.3	+1.0	+1.1
C-Plasma14	+1.0	-1.1	BDL	-3.9	-1.8	+2.0	+1.8	-1.2	+1.5	+1.4	-1.0
C-Plasma15	+1.0	+1.1	BDL	BDL	+1.1	+2.2	+1.8	-1.5	-1.0	+1.7	-2.1
C-Plasma16	+1.0	+1.1	BDL	BDL	+1.8	+1.7	-1.0	-2.5	+1.2	-1.0	BDL
C-Plasma17	+1.0	-1.3	BDL	BDL	+1.4	+2.3	+1.5	-1.3	+1.0	+1.1	-2.0
C-Plasma18	+1.0	-2.1	-3.7	-2.3	-1.6	+2.5	+1.6	-1.3	+1.3	-1.0	-1.3
C-Plasma19	+1.0	-1.2	-1.1	-1.4	+1.5	+3.0	+2.4	+1.5	+2.3	+1.3	+1.6
C-Plasma20	+1.0	-1.3	BDL	-2.8	-1.4	+2.0	+1.7	+1.4	+1.5	+1.5	+1.1
C-Plasma P#2	+1.0	+2.9	BDL	-8.5	+1.0	+1.6	-1.3	-4.1	+1.4	+1.5	-3.8
C-Plasma P#5	+1.0	+1.1	BDL	-1.9	-1.3	+2.2	+1.2	-2.2	+1.4	+1.0	-1.8
C-Plasma P#22	+1.0	-1.1	BDL	-1.1	-1.7	+2.4	+1.8	-1.2	+1.3	+1.2	-3.1
C-Plasma Standard	+1.0	BDL	BDL	+4.3	BDL	+1.5	+1.1	+1.3	+2.0	+1.1	+1.0

WT D614G  
UK501Y.V1  
SA501Y.V2  
BR501Y.V3  
242-244del  
K417N  
K417T  
E484K  
E486L  
N501Y  
417N/484K/501Y

Wang et al. 2021. SARS-CoV-2 variants resist antibody neutralization and broaden host ACE2 usage



# Varijante



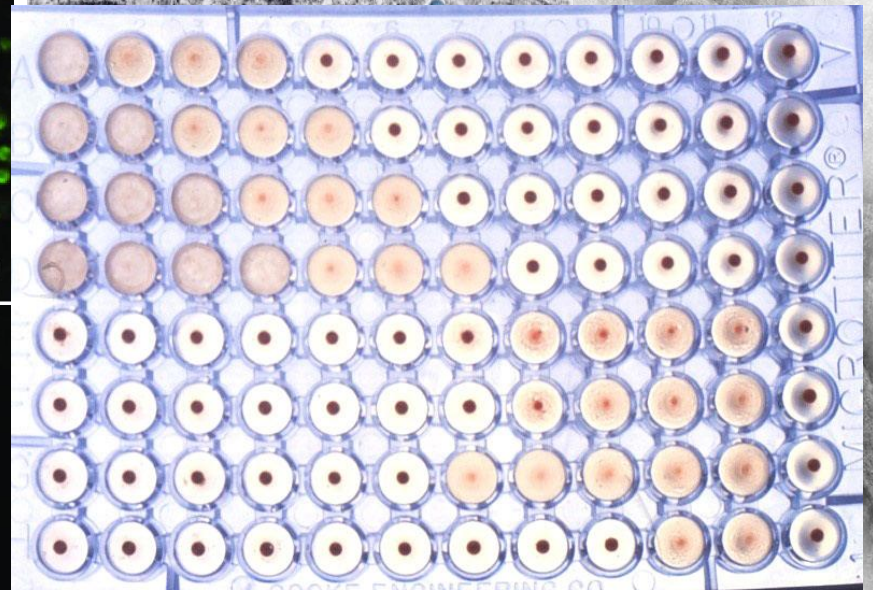
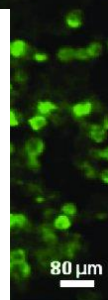
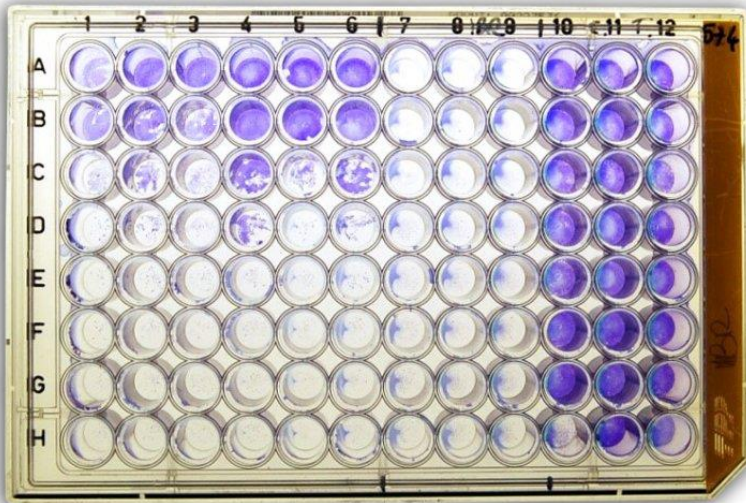
ECDC Risk assessment published 21.01.21.



# Otkrivanje novih virusa – klasična virusologija

Serijska karakterizacija uzročnika:

1. Test na stanice, oplodena jaja ili miševu
2. Teste karakterizacije
  - CPE / smrt ploda / smrt miša
  - Imunofluorescencija / elektronska mikroskopija



B

80 μm

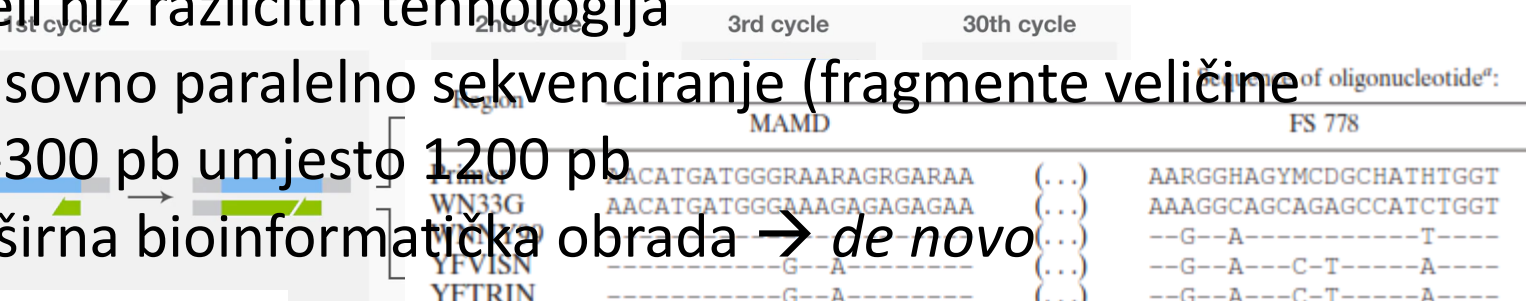
CDC/ F.A. Murphy



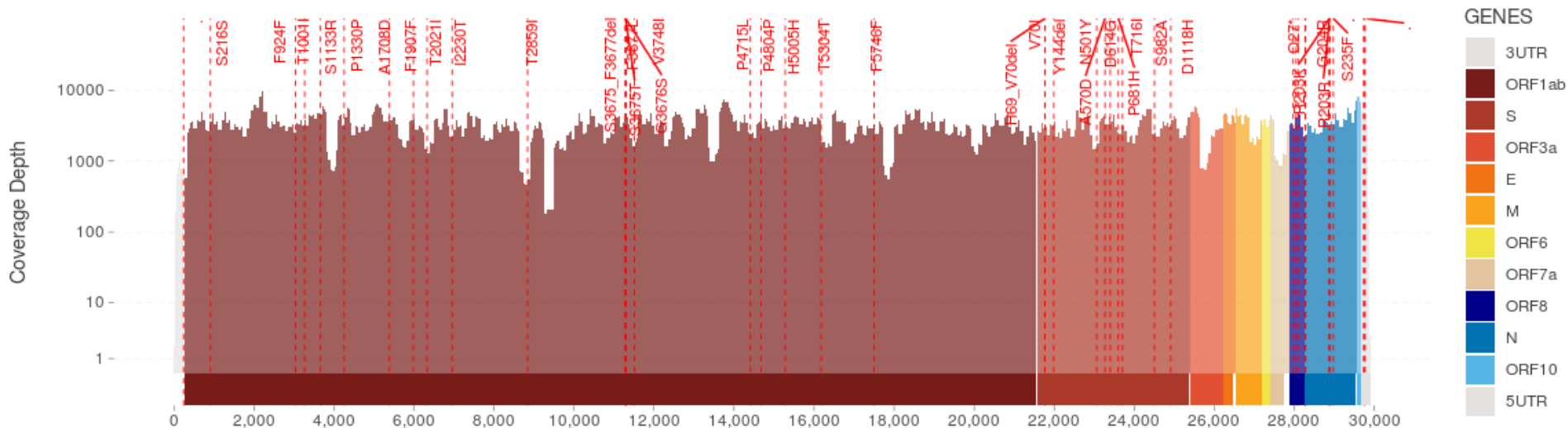
# Otkrivanje novih virusa – molekularne metode

Next-Gen PCR vs. konvencionalne PCR za otkrivanje grupe uzročnika

- Cijeli niz različitih tehnologija
- Masovno paralelno sekvenciranje (fragmente veličine 75-300 pb umjesto 1200 pb)
- Opširna bioinformatička obrada → *de novo*



SARS-CoV-2 Genome Coverage (MN908947: Reference Wuhan-Hu-1)





**Hvala na pažnji!**

[ikurolt@bfm.hr](mailto:ikurolt@bfm.hr)

[www.bfm.hr](http://www.bfm.hr)