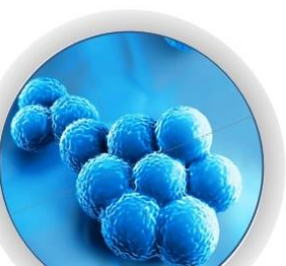




AMR u doba COVID-19 pandemije

Prof. dr Ivana Ćirković

Institut za mikrobiologiju i imunologiju
Medicinski fakultet, Univerzitet u Beogradu



The future if we do not act now

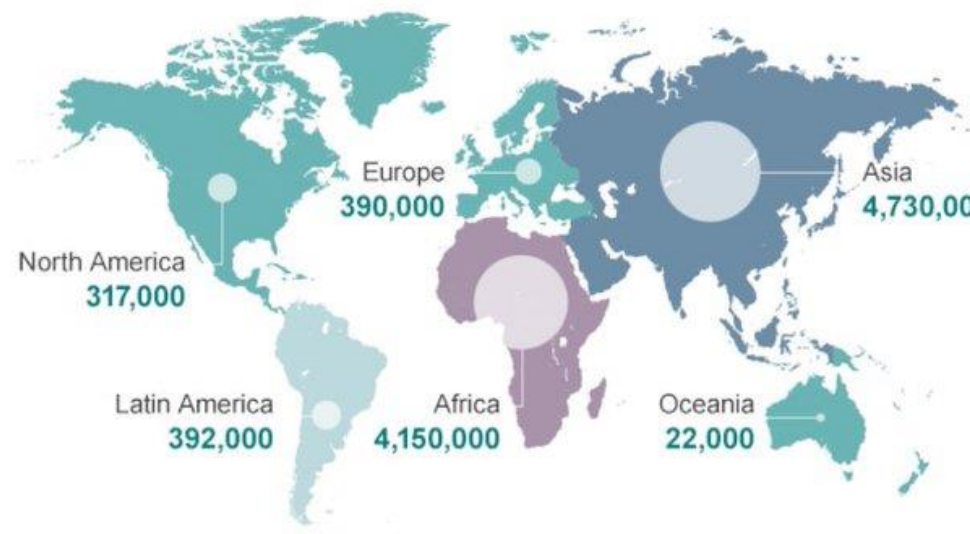
GLOBAL

Addressing the problem of antimicrobial resistance could result in:

10m deaths by 2050
Costing £66 trillion

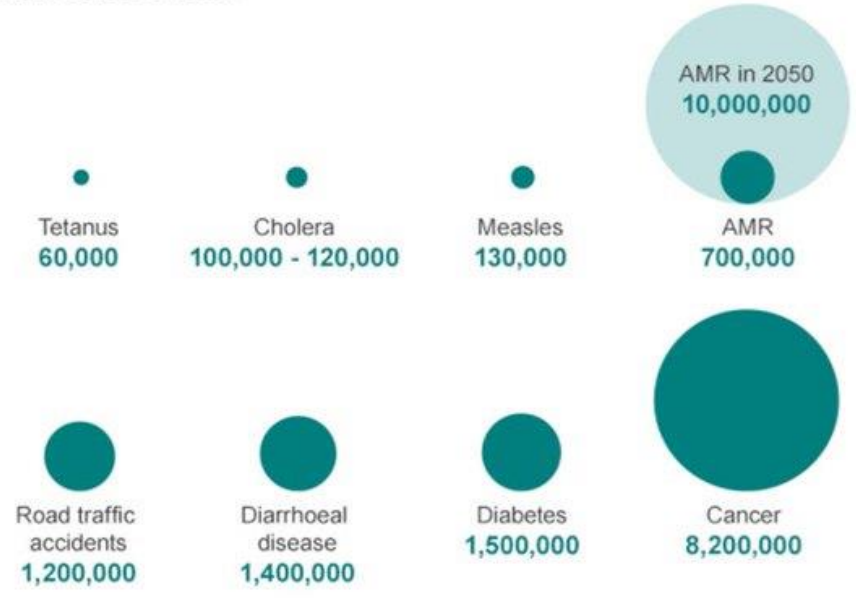


Deaths attributable to antimicrobial resistance every year by 2050



Source: Review on Antimicrobial Resistance 2014

Deaths attributable to antimicrobial resistance every year compared to other major causes of death



Source: Review on Antimicrobial Resistance 2014

By 2050: more deaths from resistant infections compared to e.g. cancer

<http://amr-review.org/>



Tackling drug-resistant infections globally

Could COVID-19 lead to an increase in antimicrobial resistance?

2nd September 2020



Mathew Upton, Professor of Medical Microbiology at the university and a co-author on the research, added: "Antibiotics underpin all of modern medicine, but AMR is an issue that could impact millions of lives in the decades to come. Currently, the COVID-19 pandemic is causing immense suffering and loss of life across the globe, but AMR has been – and will remain – one of the most significant threats to global human health."

AMR & COVID-19

Antimikrobna otpornost (AMR) nastaje kada se mikroorganizmi (kao što su bakterije i virusi) izmene nakon izlaganja antimikrobnim lekovima. Ove promene mogu značiti da postaju otporne na lekove koji se koriste za njihovo lečenje. Postoje različite vrste antimikrobnih sredstava koja deluju protiv različitih vrsta mikroorganizama, npr. antibakterijski ili antibiotici protiv bakterija, antivirusna sredstva protiv virusa, antimikotična sredstva protiv gljivica, itd. **Otpornost na antibiotike** uzrokuje uporno prekomerno korišćenje i zloupotreba antibiotika u zdravlju ljudi i životinja.



Antibiotici ne leče niti sprečavaju viruse, uključujući i one koji prozrokuju COVID-19!



Antibiotici deluju samo protiv **bakterijskih infekcija**. Štaviše, neprimerena upotreba antibiotika povećava rizik od otpornosti na antibiotike, što rizikuje sve čak i od blažih infekcija.

Kada pacijenti sa COVID-19 mogu dobiti antibiotike?



Neki pacijenti sa COVID-19 mogu razviti **ko-bakterijske infekcije**. Ako je to slučaj, onda zdravstveni radnici mogu prepisati antibiotike za lečenje sekundarne bakterijske infekcije kod tih pacijenata.

Tačna dijagnoza je ključna!

Tačna dijagnoza je od vitalnog značaja za lečenje. Testiranje pomaže u razlikovanju virusnih (poput virusa koji uzrokuje COVID-19) od bakterijskih infekcija. Zbog toga je daleko manja verovatnoća da će se antibiotici nepotrebno prepisivati i koristiti, što zauzvrat smanjuje rizik od otpornost na antibiotike i optimizuje negu pacijenta.



Nikada se nemojte sami lečiti antibioticima!

Važno je poslušati savete lekara. Ako se ne osećate dobro, potražite lekarsku pomoć i nemojte se sami dijagnostifikovati i lečiti antibioticima. Zapamtite - **uzimajte antibiotike samo ako su vam prepisani.**



Stalno upražnjavajte redovnu higijenu!

Higijena ruku je presudna u doba COVID-19. Upraznjavajte redovnu higijenu ruku kod kuće i u zdravstvenim ustanovama, tako što ćete redovno prati vaše ruke. Kijajte i kašljite u savijeni lakat ili maramicu koju treba baciti u zatvorenu kantu. Ovo su neki od najefikasnijih načina smanjenja širenja mnogih infekcija, uključujući organizme otporne na antibiotike.



Antibiotici ne leče niti sprečavaju viruse, uključujući i one koji prozrokuju **COVID-19!**



Antibiotici deluju samo protiv **bakterijskih infekcija**. Štaviše, neprimerena upotreba antibiotika povećava rizik od otpornosti na antibiotike, što rizikuje sve čak i od blažih infekcija.

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Stop the **COVID-19** pandemic from becoming an **AMR** catastrophe



Why pay attention to antimicrobial resistance?

Antimicrobial resistance (AMR) occurs when microorganisms change and become resistant to antimicrobial drugs used to treat them. There are different types of antimicrobials which work against different types of microorganisms, such as antibacterials or antibiotics against bacteria, antivirals against viruses, and antifungals against fungi.

Because of antibiotic resistance, the world is running out of effective antibiotics to treat infectious diseases. Unless appropriate action is taken, decades of progress in health and medicine risk being undone.

In May 2015, the World Health Assembly (WHA) endorsed a global action plan on AMR and urged all Member States to develop national action plans. WHA72 (May 2019) called for accelerated implementation.

What has antibiotic use got to do with COVID-19?

Up to 15% of patients with severe COVID-19 infection need antibiotics for so called co-infections (bacterial infections they have alongside COVID-19). However, currently available evidence suggests that 75% of COVID-19 patients get antibiotics, meaning that a majority are prescribed antibiotics unnecessarily without having a diagnosis of bacterial co-infection confirmed.

This overuse of antibiotics will drive the development of resistance unless action is taken. It is vital that prescribers pay close attention to guidelines in order to avoid negative long-term consequences. Both COVID-19 and AMR pose a serious public health threat with long-term negative humanitarian and economic consequences. Failure to systematically include AMR-specific activities in the COVID-19 response undermines the global fight against AMR.

Preventing and controlling AMR and COVID-19 is a complex issue which involves many different sectors and requires a comprehensive approach and international cooperation.

Masovno i prekomereno propisivanje antibiotika kod COVID-19 bolesnika - OPRAVDANJE

- Identifikacija/detekcija bakterijske koinfekcije ili superinfekcije kod COVID-19 je izazov (simptomi slični virusnim infekcijama)
- Uprkos saznanju da je COVID-19 virusna infekcija, standardni refleks je da se započne sa antibiotskom terapijom kod pacijenata sa kašljem, temperaturom i rengenkim infiltratima – pokazatelji bakterijske vanbolničke pneumonije
- Racionalnost koja stoji iza odluke propisati antibiotik COVID-19 pacijentu se zasniva i na iskustvu sa gripom, kod koga je pokazana stopa bakterijskih koinfekcija i superinfekcija u 11-35% slučajeva
- Nepostojanje efikasne antivirusne terapije
- Anksioznost zbog visoke stope mortaliteta

Masovno i prekomereno propisivanje antibiotika kod COVID-19 bolesnika - **NERACIONALNO**

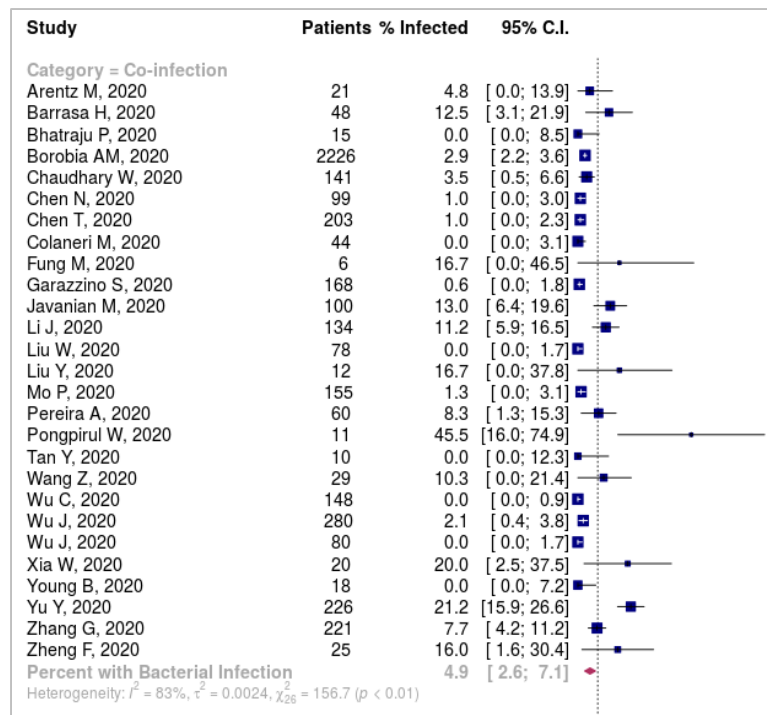
- In a study of 99 patients with COVID-19 pneumonia in Wuhan Jinyintan Hospital, **most patients** were given antibiotic treatment (25% patients were treated with a single antibiotic and 45% patients were given combination therapy); [Chen N *et al.* Lancet 2020; 395: 507–13]
- In a Chinese study involving 41 COVID-19 patients, **all patients** were administered with empirical antibiotic treatment; [Huang C *et al.* Lancet 2020; 395: 497–506]
- In a review of medical literature (18 full texts), the use of broad-spectrum ABs was widely reported, with **72% of COVID-19 cases** receiving AB therapy; [Rawson TM *et al.* CID 2020; <https://doi.org/10.1093/cid/ciaa530>]
- In a study in Dallas on 289 hospitalized COVID-19 patients, antibiotics were given to most patients (**93.8%**); [Crotty MP *et al.* medRxiv 2020]
- In a living meta-analysis and systematic review, **over 70% of COVID-19 patients** received antibiotics [Langford BJ *et al.* <https://www.tarm.org/covid>]

- **MEĐUTIM, studije u prethodnom periodu su pokazale nisku stopu bakterijskih koinfekcija kod COVID-19 bolesnika**

Masovno i prekomereno propisivanje antibiotika kod COVID-19 bolesnika - **NERACIONALNO**

Bakterijske koinfekcija kod **3-5%** COVID-19 bolesnika

Najčešći uzročnici (> 90% slučajeva) - *S. aureus*, *H. influenzae* i *S. pneumoniae*



Revidiran stav o upotrebi antibiotika kod hospitalizovanih COVID-19 bolesnika.

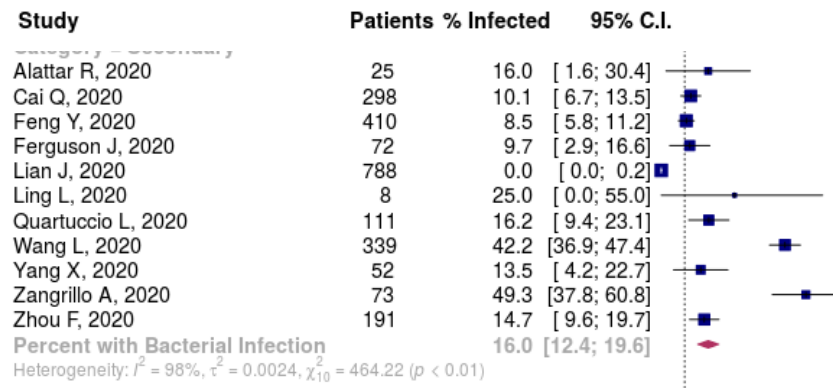
Antibiotska terapija se preporučuje:

- ✓ kod COVID-19 pacijenata kod kojih je radiološki detektovana bakterijska infekcija
- ✓ u JIL
- ✓ kod imunokompromitovanih bolesnika

Masovno i prekomereno propisivanje antibiotika kod COVID-19 bolesnika

Bakterijske **superinfekcija** kod **15-20%** COVID-19 bolesnika

Najčešći uzročnici –
Acinetobacter spp.,
Enterobacterales,
P. aeruginosa i *S. aureus*



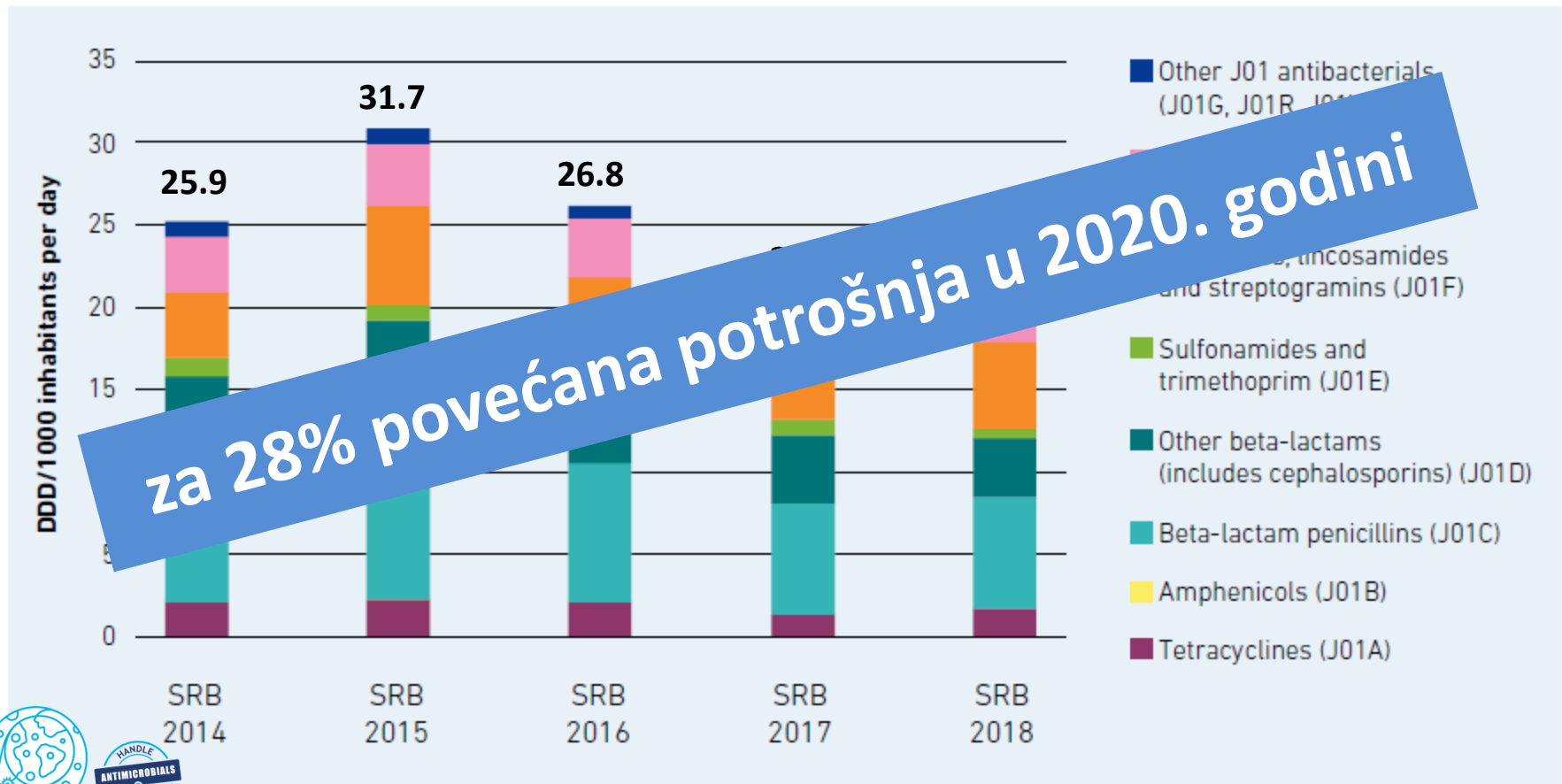
✓ Infekcije krvi česte sekundarne bolničke infekcije kod COVID-19 bolesnika (14,3-36,3%)

“Antibiotici nisu bombone”

Pregled situacije u Srbiji



✓ Potrošnja antibiotika se prati na nacionalnom nivou i u okviru AMC mreže (WHO)



“Antibiotici nisu bombone”

Pregled situacije u EU

Antimicrobial consumption in the EU/EEA (ESAC-Net)

Annual Epidemiological Report for 2020



Key facts

- For 2020, twenty-nine countries (27 European Union (EU) Member States and two European Economic Area (EEA) countries - Iceland and Norway) reported data on antimicrobial consumption. Twenty-five countries reported data for both community and hospital consumption; two countries (Germany and Iceland) reported only community consumption, and two countries (Cyprus and Czechia) reported total consumption for both sectors combined.
- The Anatomical Therapeutic Chemical (ATC) classification index with defined daily doses (DDD) 2021 was used for the analysis of both 2020 data and historical data. Antimicrobial consumption is expressed as DDD per 1 000 inhabitants per day.
- In 2020, the mean total (community and hospital sector combined) consumption of antibacterials for systemic use (ATC group J01) in the EU/EEA was 16.4 DDD per 1 000 inhabitants per day (country range: 8.5–28.9). During the period 2011–2020, a statistically significant decrease was observed for the EU/EEA overall, as well as for eight individual countries. A statistically significant increasing trend was observed for two countries.
- The EU/EEA mean total (community and hospital sector combined) consumption of antivirals for systemic use (ATC group J05) was 2.56 DDD per 1 000 inhabitants per day (country range: 0.59–11.19), with no statistically significant trends in the five-year period between 2016–2020.

“Antibiotici nisu bombone”

Pregled situacije u Srbiji



✓ Rezistencija bakterija na antibiotike se prati na nacionalnom nivou i u okviru CAESAR mreže (WHO)

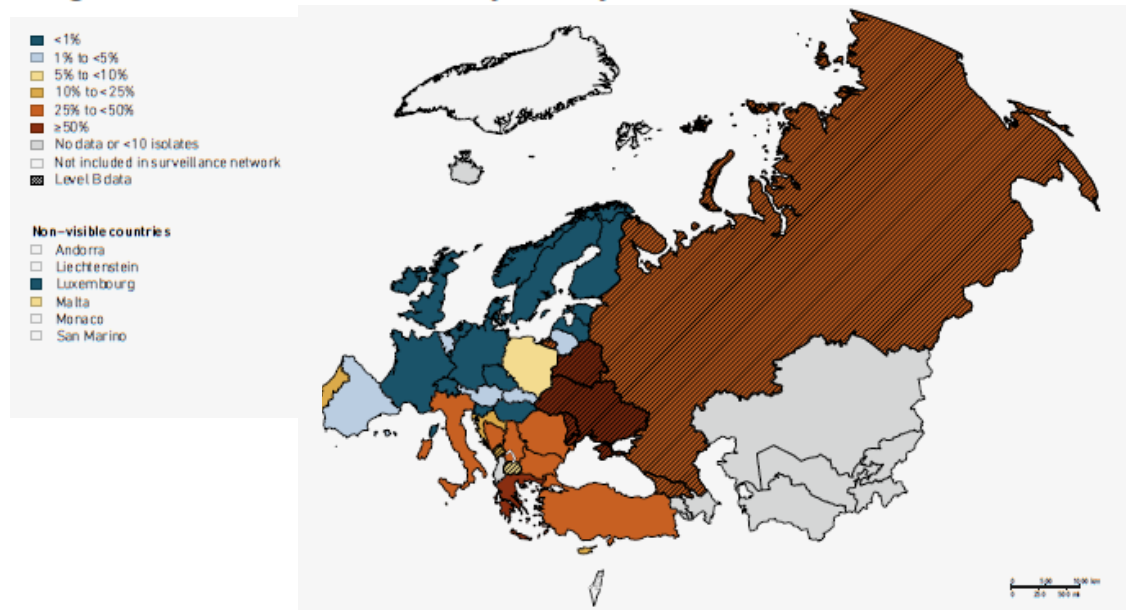


Central Asian and European
Surveillance of Antimicrobial Resistance

Annual report 2020



Fig. 2.5 Percentage of invasive *K. pneumoniae* isolates resistant to carbapenems in the WHO European Region (EARS-Net and CAESAR), by country or area, 2019



“Antibiotici nisu bombone”

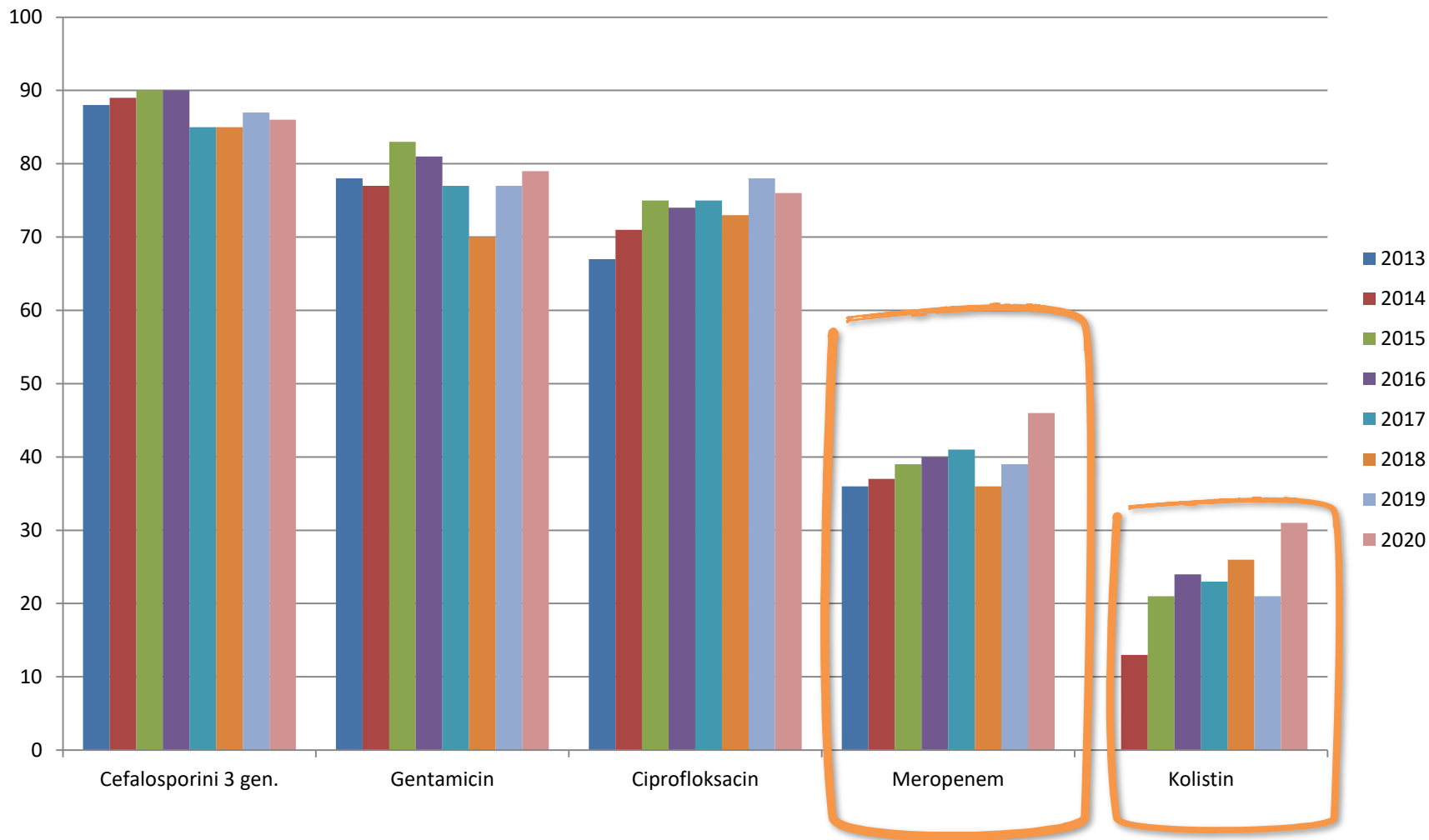
Pregled situacije u Evropi



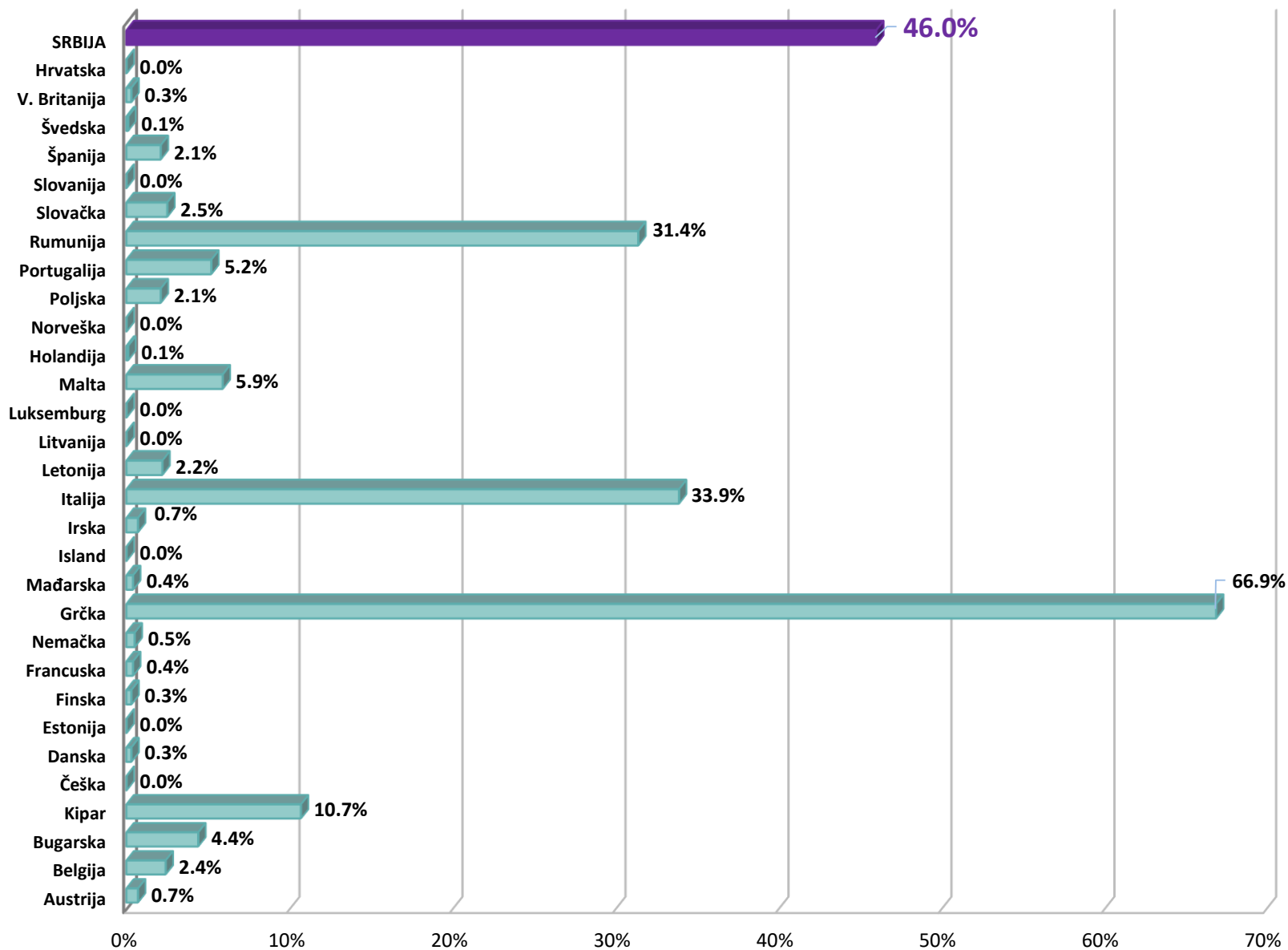
Surveillance of antimicrobial resistance in Europe, 2020 data

1. *Acinetobacter* spp.
2. *Enterococcus faecalis* i *Enterococcus faecium*

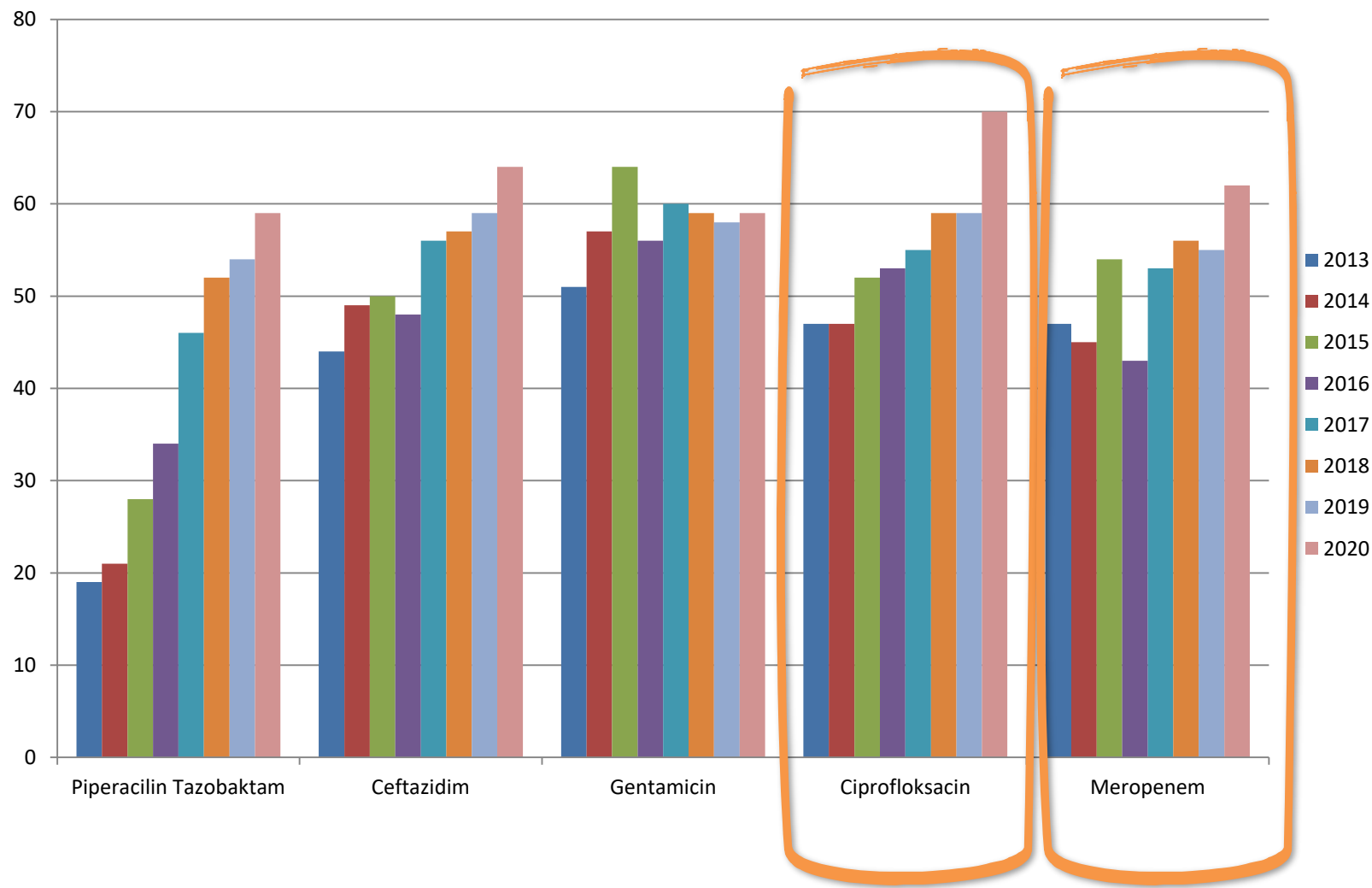
Klebsiella pneumoniae: rezistencija na antibiotike u Srbiji, 2013-2020.



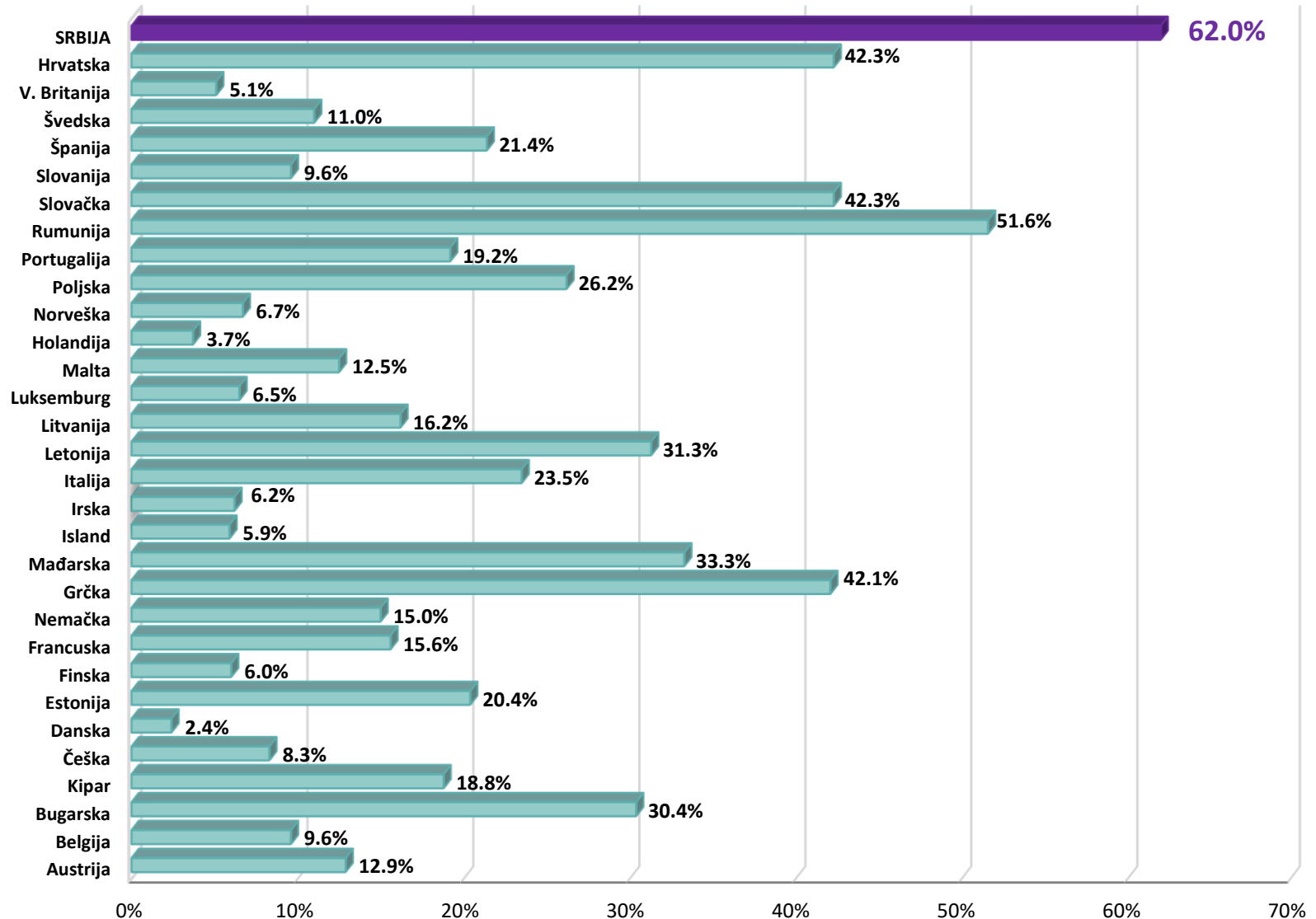
Proporcije karbapenem-rezistentnih izolata *Klebsiella pneumoniae* u Evropi



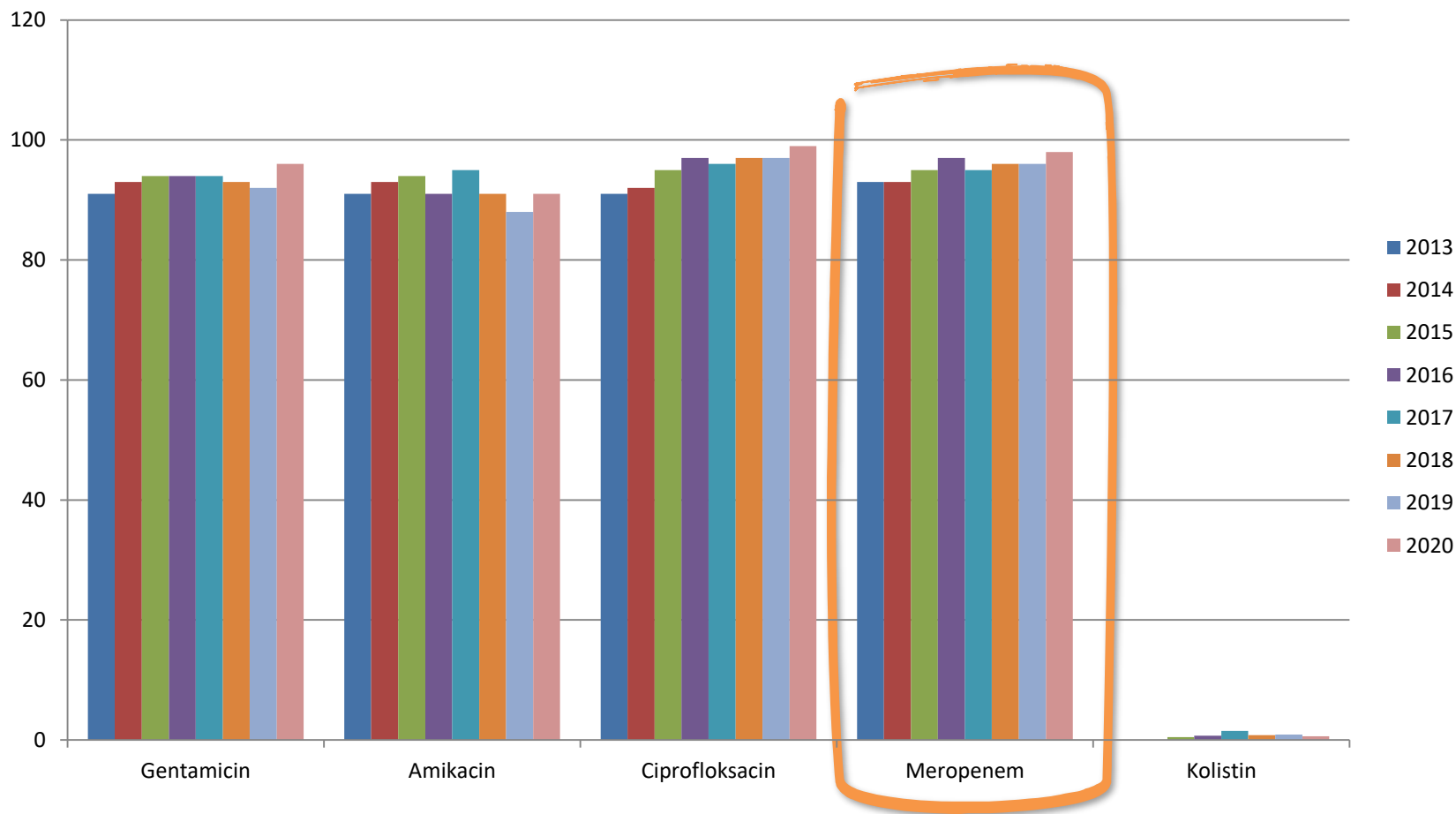
Pseudomonas aeruginosa: rezistencija na antibiotike u Srbiji, 2013-2020.



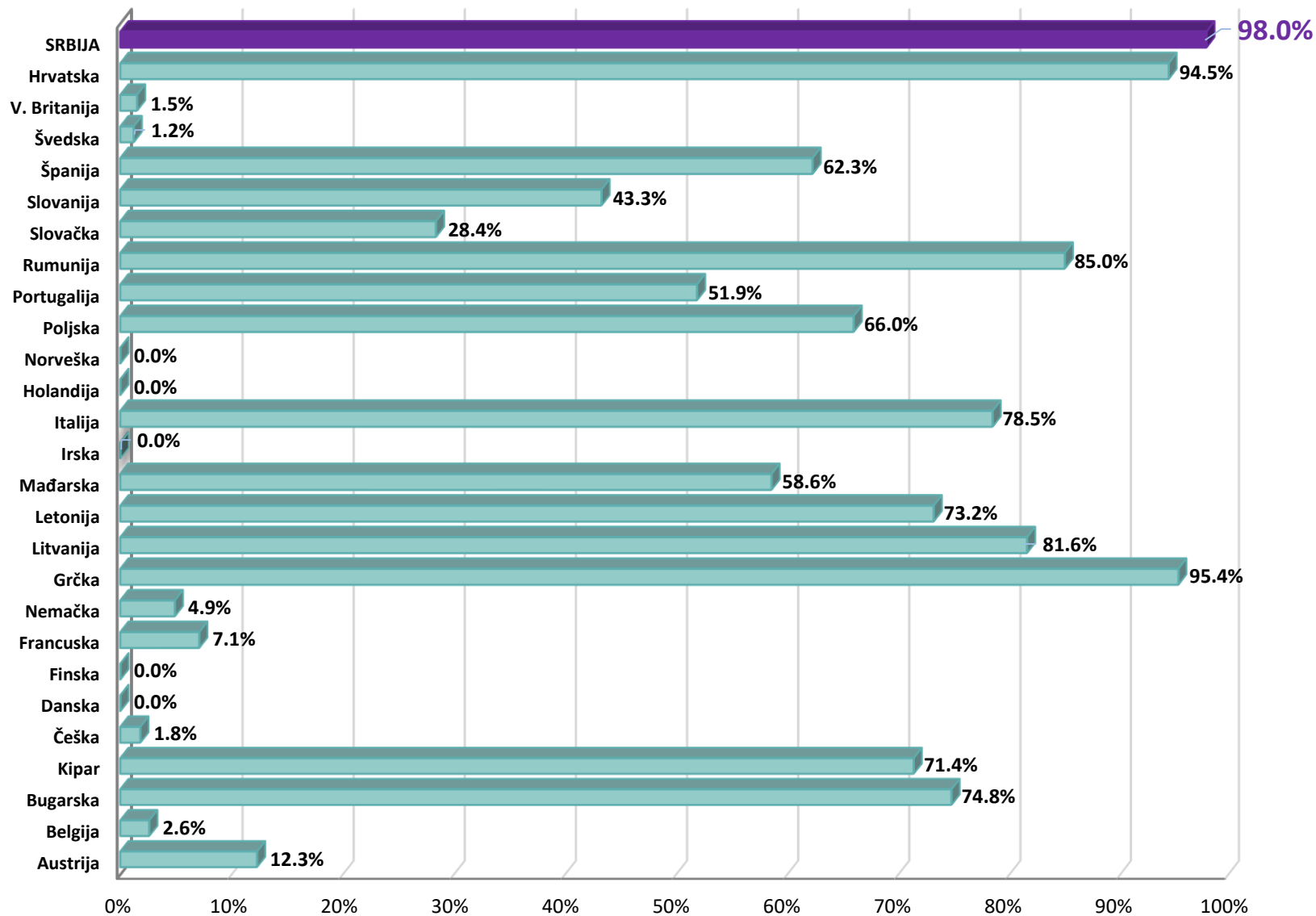
Proporcije karbapenem-rezistentnih izolata *Pseudomonas aeruginosa* u Evropi



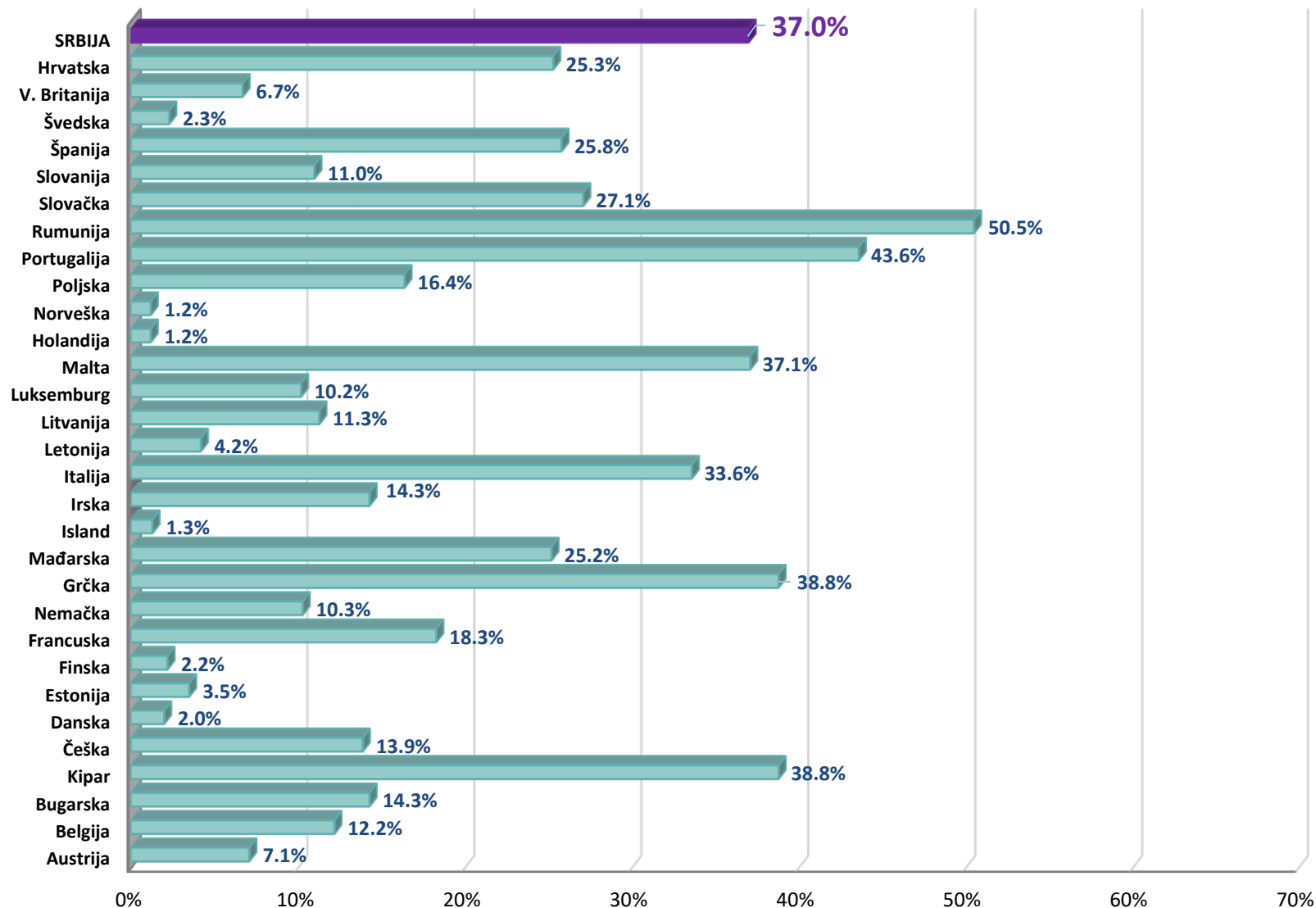
Acinetobacter spp.: rezistencija na antibiotike u Srbiji, 2013-2020.



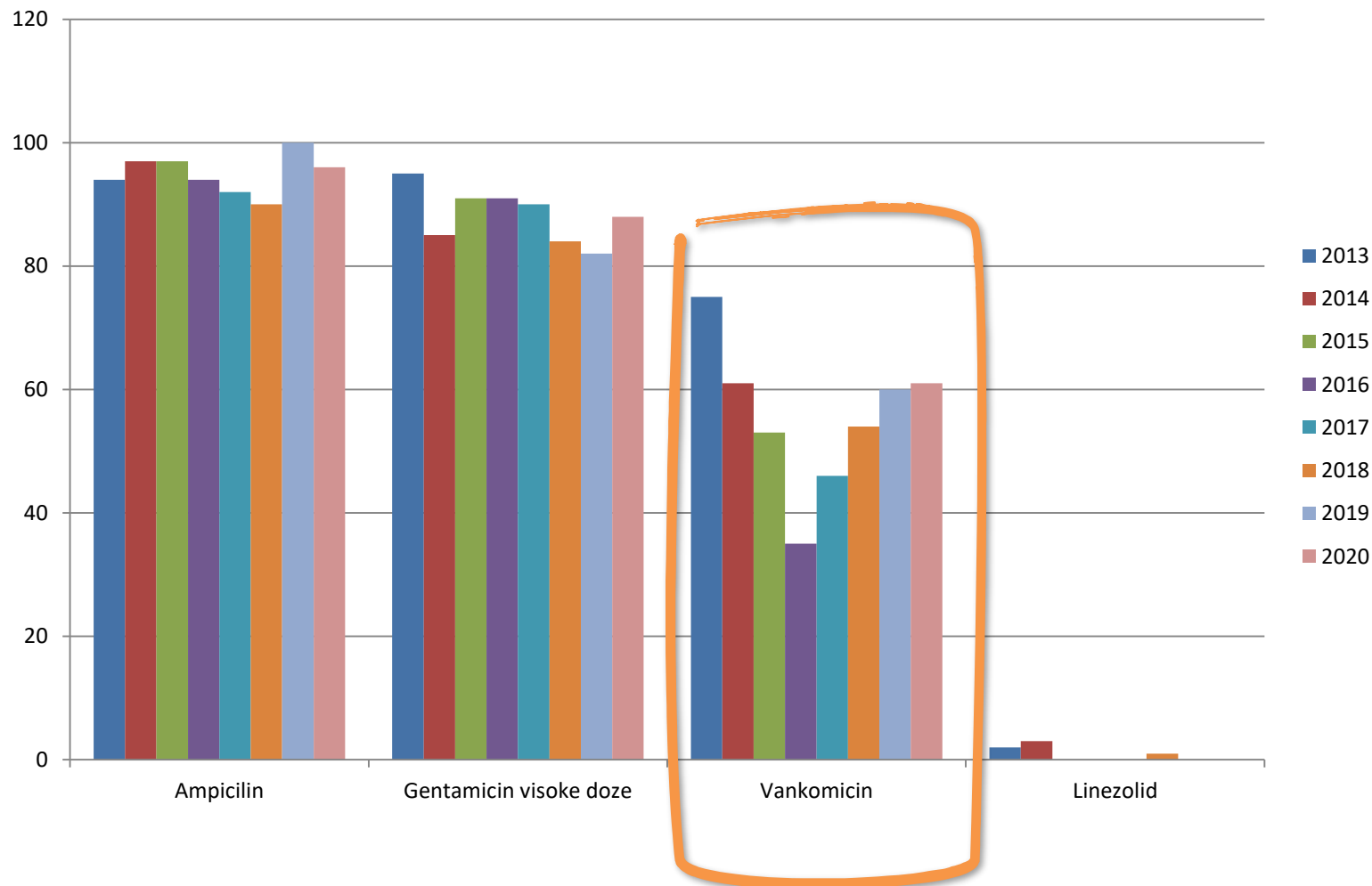
Proporcije karbapenem-rezistentnih izolata *Acinetobacter* spp. u Evropi



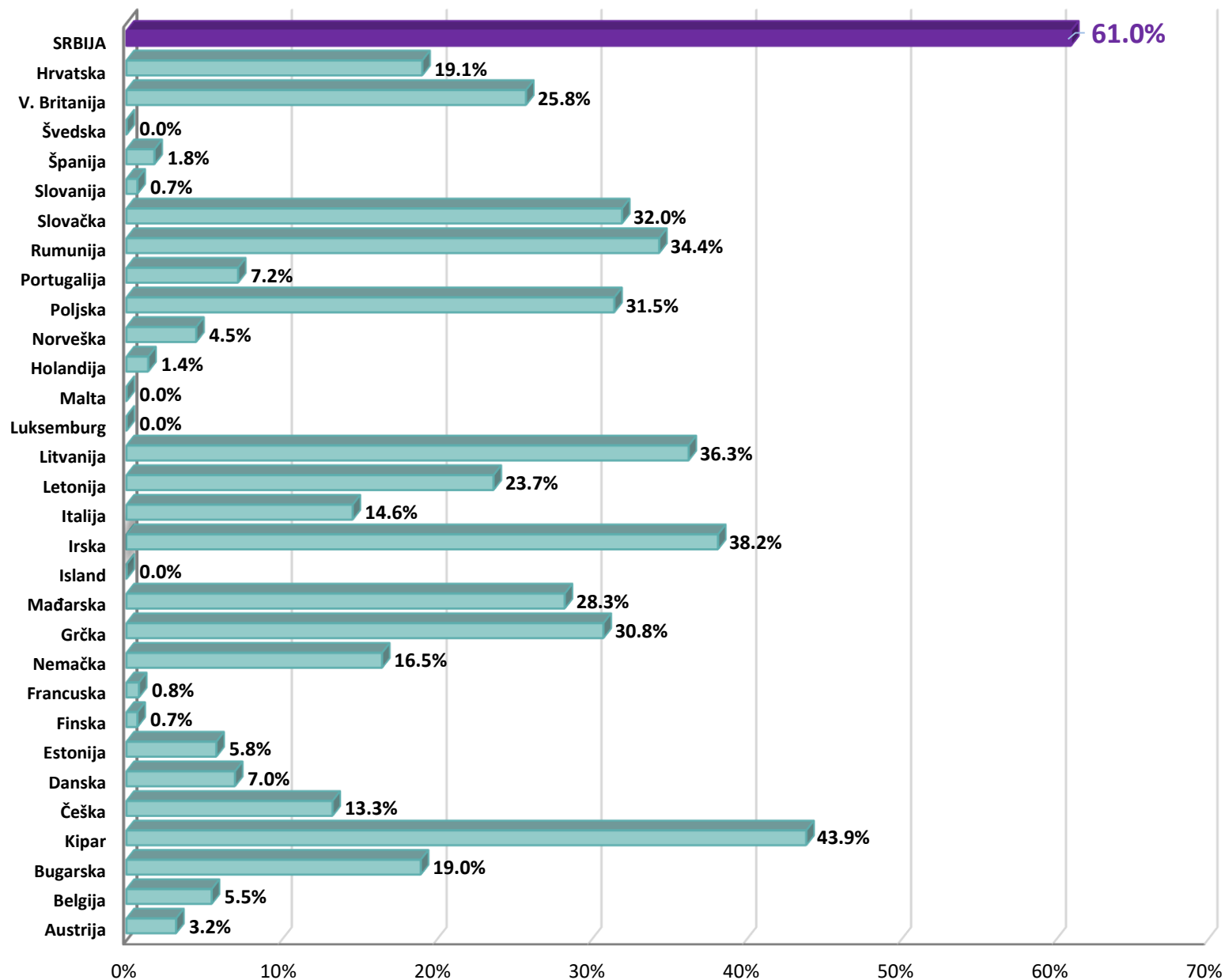
Proporcije izolata meticilin-rezistentnih *Staphylococcus aureus* (MRSA) u Evropi



Enterococcus faecium: rezistencija na antibiotike u Srbiji, 2013-2020.



Proporcije vankomicin-rezistentnih izolata *Enterococcus faecium* u Evropi



Brzi molekularni test Respiratorni panel

nazofaringealni bris

23 patogena – 1 test – **45 minuta**

19 virusa

Adenovirus

Coronavirus 229E (Koronavirus 229E)

Coronavirus HKU1 (Koronavirus HKU1)

Coronavirus NL63 (Koronavirus NL63)

Coronavirus OC43 (Koronavirus OC43)

Middle East Respiratory Syndrome Coronavirus (Bliskoistočni respiratorni sindrom koronavirus) (MERS-CoV)

Severe Acute Respiratory Syndrome Coronavirus 2 (Teški akutni respiratorni sindrom – koronavirus 2) (SARS-CoV-2)

Human Metapneumovirus (Humani metapneumovirus)

Human Rhinovirus/Enterovirus (Humani rinovirus/enterovirus)

Influenza A (Influenza A), sa podtipovima H1, H3 i H1-2009

Influenza B (Influenza B)

Parainfluenza Virus 1 (Virus parainfluenza 1)

Virus parainfluenza tipa 2 (Virus parainfluenza 2)

Virus parainfluenza tipa 3 (Virus parainfluenza 3)

Virus parainfluenza tipa 4 (Virus parainfluenza 4)

Respiratorni sincicijalni virus (Respiratorni sincicijalni virus)

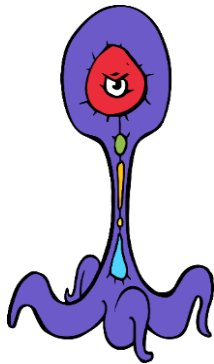
4 bakterije

Bordetella parapertussis

Bordetella pertussis

Chlamydia pneumoniae

Mycoplasma pneumoniae





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Contents lists available at ScienceDirect

Journal of Clinical Virology

journal homepage: www.elsevier.com/locate/jcv



Clinical evaluation of the BioFire® Respiratory Panel 2.1 and detection of SARS-CoV-2



Hannah M. Creager^a, Barbara Cabrera^a, Andy Schnaubelt^a, Jesse L. Cox^a, Allison M. Cushman-Vokoun^a, Salika M. Shakir^b, Keith D. Tardif^b, Meei-Li Huang^c, Keith R. Jerome^{c,d}, Alexander L. Greninger^c, Daria Drobysheva^e, Usha Spaulding^e, Margarita Rogatcheva^e, Kevin M. Bourzac^e, S.H. Hinrichs^a, M.J. Broadhurst^{a,1}, P.D. Fey^{a,*,1}

A B S T R A C T

We evaluated the performance of the BioFire® Respiratory Panel 2.1 (RP2.1) in the detection of SARS CoV-2 in comparison against three other SARS CoV-2 EUA assays. In these studies, the RP2.1 panel had 98 % positive percent agreement (48/49) and 100 % negative percent agreement (49/49). Since 30 % of nasopharyngeal swab specimens have a SARS CoV-2 Ct > 30 and thus detection of virus in low titers is clinically relevant, a sample with a high titer was diluted and each 10 fold dilution was tested in triplicate and compared against 6 other EUA approved SARS CoV-2 assays. These data suggested that the BioFire® RP2.1 panel, along with four other SARS CoV-2 assays (Roche cobas, Cepheid Xpert Xpress, BioFire® Defense COVID19, and NECoV19), consistently detected viral RNA at the 10 – 7 dilution. Overall, these studies suggest that the BioFire® RP2.1 assay can be used to detect acute cases of SARS CoV2 in addition to patients with low viral titer later in disease presentation.

Brzi molekularni test Respiratorni panel

23 patogena – 1 test – **45 minuta**

19 virusa

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Virus parainfluenza tipa 4 (Virus parainfluenza 4)

Respiratorni sincicijalni virus (Respiratorni sincicijalni virus)

4 bakterije

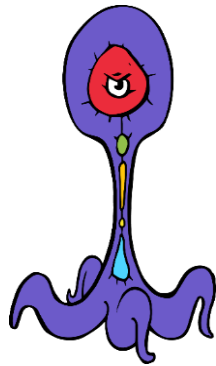
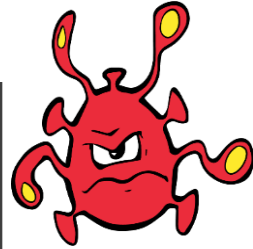
Bordetella parapertussis

Bordetella pertussis

Chlamydia pneumoniae

Mycoplasma pneumoniae

koinfekcije



Respiratorni panel 2.1 plus (RP2.1plus) – naše iskustvo



10 February 2021

BioFire® Respiratory Panel 2.1 plus

Run Summary
Sample ID: Surla Goran 3110954172652 Run Date: 10 Feb 2021 9:57 AM
Detected: Coronavirus NL63 Controls: Passed
Equivalocal: None

Result Summary

Viruses	
Not Detected	Adenovirus
Not Detected	Coronavirus 229E
Not Detected	Coronavirus HKU1
Detected	Coronavirus NL63
Not Detected	Coronavirus OC43
Not Detected	Middle East Respiratory Syndrome Coronavirus (MERS-CoV)
Not Detected	Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2)
Not Detected	Human Metapneumovirus
Not Detected	Human Rhinovirus/Enterovirus
Not Detected	Influenza A
Not Detected	Influenza B
Not Detected	Parainfluenza Virus 1
Not Detected	Parainfluenza Virus 2
Not Detected	Parainfluenza Virus 3
Not Detected	Parainfluenza Virus 4
Not Detected	Respiratory Syncytial Virus

Bacteria	
Not Detected	<i>Bordetella parapertussis</i> (IS1001)
Not Detected	<i>Bordetella pertussis</i> (ptxP)
Not Detected	<i>Chlamydia pneumoniae</i>
Not Detected	<i>Mycoplasma pneumoniae</i>

Run Details
Pouch: RP2.1plus v1.0 Protocol: NPS2 v3.2
Run Status: Completed Operator: Institut Dedinje (LabIKVBD)
Serial No.: 37098155 Instrument: 2FA06341
Lot No.: 107P29



telemikrobiologija

10:57

Mozete li nam pomoći 10:57

Hvala 10:57



14 October 2021

BioFire® Respiratory Panel 2.1 plus

Run Summary
Sample ID: stojanovic milan 1906953710490 Run Date: 13 Oct 2021 10:28 AM
Detected: Coronavirus OC43 Controls: Passed
Equivalocal: None

Result Summary

Viruses	
Not Detected	Adenovirus
Not Detected	Coronavirus 229E
Not Detected	Coronavirus HKU1
Not Detected	Coronavirus NL63
Detected	Coronavirus OC43
Not Detected	Middle East Respiratory Syndrome Coronavirus (MERS-CoV)
Not Detected	Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2)
Not Detected	Human Metapneumovirus
Not Detected	Human Rhinovirus/Enterovirus
Not Detected	Influenza A
Not Detected	Influenza B
Not Detected	Parainfluenza Virus 1
Not Detected	Parainfluenza Virus 2
Not Detected	Parainfluenza Virus 3
Not Detected	Parainfluenza Virus 4
Not Detected	Respiratory Syncytial Virus

Bacteria	
Not Detected	<i>Bordetella parapertussis</i> (IS1001)
Not Detected	<i>Bordetella pertussis</i> (ptxP)
Not Detected	<i>Chlamydia pneumoniae</i>
Not Detected	<i>Mycoplasma pneumoniae</i>

Run Details
Pouch: RP2.1plus v1.0 Protocol: NPS2 v3.2
Run Status: Completed Operator: Institut Dedinje
Serial No.: 43731006 Instrument: 2FA06341
Lot No.: 203F21



Kako biste Vi protumacili Hvala

06:35

Brzi molekularni test Pneumonija panel

Sputum, endotrahealni aspirat, BAL,...

27 patogena + 7 gena rezistencije – 1 test – **60 minuta**

18 bakterija + 9 virusa + 7 gena rezistencije

Bakterije prijavljene sa binovima od 10^4 , 10^5 , 10^6 , ili $\geq 10^7$ kopija/mL

<i>Acinetobacter calcoaceticus-baumannii</i> kompleks	<i>Klebsiella oxytoca</i>	<i>Serratia marcescens</i>
<i>Enterobacter cloacae</i> kompleks	<i>Klebsiella pneumoniae</i> grupa	<i>Staphylococcus aureus</i>
<i>Escherichia coli</i>	<i>Moraxella catarrhalis</i>	<i>Streptococcus agalactiae</i>
<i>Haemophilus influenzae</i>	Proteus vrste	<i>Streptococcus pneumoniae</i>
<i>Klebsiella aerogenes</i>	<i>Pseudomonas aeruginosa</i>	<i>Streptococcus pyogenes</i>

Atipične bakterije

<i>Chlamydia pneumoniae</i>	<i>Legionella pneumophila</i>	<i>Mycoplasma pneumoniae</i>
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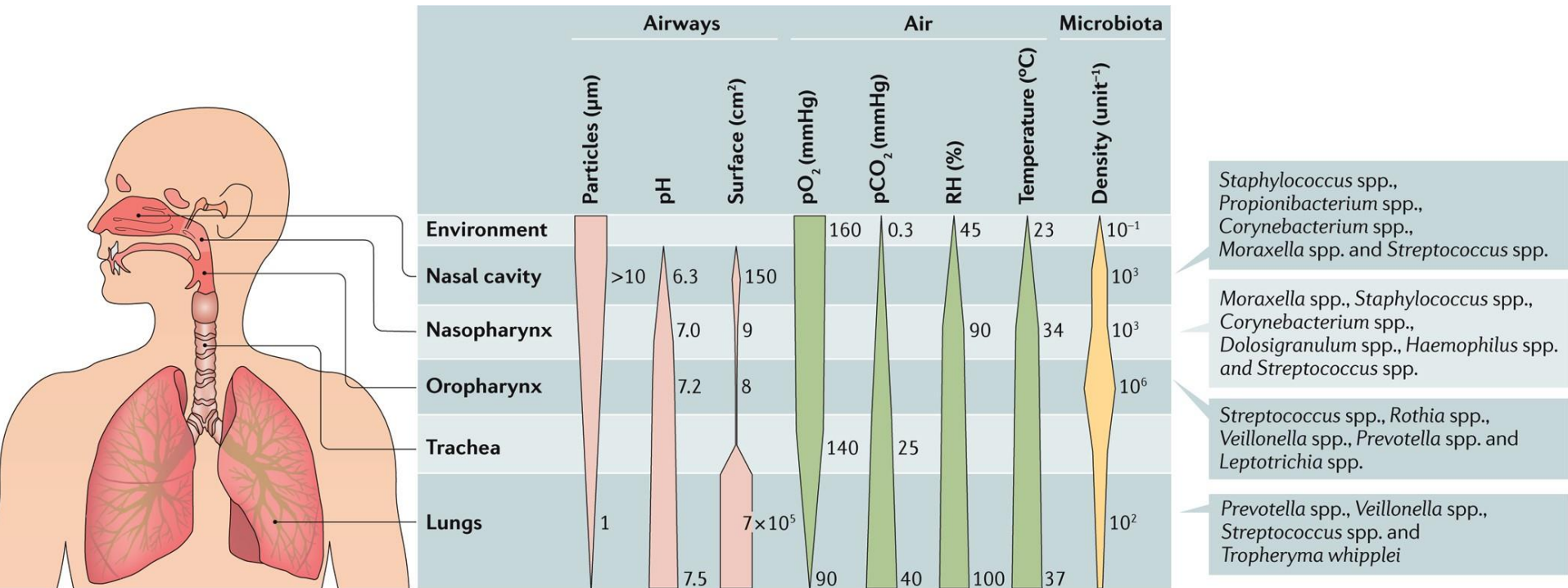
Virusi

Adenovirus	Humani rinovirus/enterovirus	Bliskoistočni respiratorni sindrom – koronavirus (MERS-CoV)
Koronavirus	Influenca A	Virus parainfluenze
Humani metapneumovirus	Influenca B	Respiratorni sincicijalni virus

Geni antimikrobne rezistencije

CTX-M	NDM	<i>mecA/C</i> i MREJ
IMP	OXA-48-like	
KPC	VIM	

Mikrobiota respiratornog trakta



Brzi molekularni test Pneumonija panel

27 patogeni + 7 gena rezistencije – 1 test – **60 minuta**

18 bakterija + 9 virusa + 7 gena rezistencije

Bakterije prijavljene sa binovima od 10^4 , 10^5 , 10^6 , ili $\geq 10^7$ kopija/mL

<i>Acinetobacter calcoaceticus-baumannii</i> kompleks	<i>Klebsiella oxytoca</i>	<i>Serratia marcescens</i>
<i>Enterobacter cloacae</i> kompleks	<i>Klebsiella pneumoniae</i> grupa	<i>Staphylococcus aureus</i>
<i>Escherichia coli</i>	<i>Moraxella catarrhalis</i>	<i>Streptococcus agalactiae</i>
<i>Haemophilus influenzae</i>	Proteus vrste	<i>Streptococcus pneumoniae</i>
<i>Klebsiella aerogenes</i>	<i>Pseudomonas aeruginosa</i>	<i>Streptococcus pyogenes</i>

Atipične bakterije

<i>Chlamydia pneumoniae</i>	<i>Legionella pneumophila</i>	<i>Mycoplasma pneumoniae</i>
-----------------------------	-------------------------------	------------------------------

Virusi

Adenovirus	Humani rinovirus/enterovirus	Bliskoistočni respiratorni sindrom – koronavirus (MERS-CoV)
Koronavirus	Influenca A	Virus parainfluenze
Humani metapneumovirus	Influenca B	Respiratorni sincicijalni virus

Geni antimikrobne rezistencije

CTX-M	NDM	<i>mecA/C</i> i MREJ
IMP	OXA-48-like	
KPC	VIM	

Prevalence and outcomes of co-infection and superinfection with SARS-CoV-2 and other pathogens: A systematic review and meta-analysis

Published: May 6, 2021

Results

Of 6639 articles screened, 118 were included in the random effects meta-analysis. The pooled prevalence of co-infection was 19% (95% confidence interval [CI]: 14%-25%, $I^2 = 98%$) and that of superinfection was 24% (95% CI: 19%-30%). Pooled prevalence of pathogen type stratified by co- or superinfection were: viral co-infections, 10% (95% CI: 6%-14%); viral superinfections, 4% (95% CI: 0%-10%); bacterial co-infections, 8% (95% CI: 5%-11%); bacterial superinfections, 20% (95% CI: 13%-28%); fungal co-infections, 4% (95% CI: 2%-7%); and fungal superinfections, 8% (95% CI: 4%-13%). Patients with a co-infection or superinfection had higher odds of dying than those who only had SARS-CoV-2 infection (odds ratio = 3.31, 95% CI: 1.82–5.99). Compared to those with co-infections, patients with

Table 2. All identified organisms as a proportion of total number of organisms per pathogen.

Pathogen type	Co-infection (N = 1910) No. (%)	Superinfection (N = 480) No. (%)
Bacteria		
<i>Staphylococcus aureus</i>	148 (7.7)	13 (2.7)
<i>Haemophilus influenza</i>	127 (6.6)	6 (1.3)
<i>Mycoplasma pneumoniae</i>	82 (4.3)	6 (1.3)
<i>Acinetobacter spp.</i>	78 (4.1)	107 (22.3)
<i>Escherichia coli</i>	73 (3.8)	33 (6.9)
<i>Stenotrophomonas maltophilia</i>	10 (0.5)	18 (3.8)
<i>Klebsiella pneumoniae</i>	189 (9.9)	28 (5.8)
<i>Streptococcus pneumoniae</i>	156 (8.2)	4 (0.8)
<i>Chlamydia pneumoniae</i>	29 (1.5)	
<i>Bordetella</i>	3 (0.2)	
<i>Moraxella catarrhalis</i>	32 (1.7)	
<i>Pseudomonas</i>	67 (3.5)	
<i>Enterococcus faecium</i>	14 (0.7)	22 (4.6)
Viruses		
Non-SARS-CoV-2 ^a coronavirus strains	38 (2.0)	9 (1.9)
Human influenza A	426 (22.3)	0 (0)
Human influenza B	73 (3.8)	0 (0)
Respiratory syncytial virus	72 (3.8)	2 (0.4)
Parainfluenza	17 (0.9)	0 (0)
Human metapneumovirus	20 (1.0)	9 (1.9)
Rhinovirus	68 (3.6)	11 (2.3)
Adenovirus	35 (1.8)	2 (0.4)
Fungi		
<i>Mucor</i>	6 (0.3)	1 (0.2)
<i>Candida spp.</i>	19 (1.0)	90 (18.8)
<i>Aspergillus</i>	128 (6.7)	65 (13.5)

respiratorne
koinfekcije

FilmArray® Pneumonia Panel plus – naše iskustvo

FilmArray® Pneumonia Panel plus - IVD
 A BIOMERIEUX COMPANY
 www.BioFireDx.com

Run Information

Sample ID	[REDACTED]	Run Date	05 Feb 2021 11:03 AM
Protocol	SPUTUM v3.3	Serial No.	30889230
Pouch Type	Pneumoplus v2.0	Lot No.	376220
Controls	Passed	Operator	Institut Dedinje (LabIKVBD)
Run Status	Completed	Instrument	2FA06341

Result Summary

Bacteria

	Bin (copies/mL)	Bin (copies/mL)			
		10 ⁴	10 ⁵	10 ⁶	≥10 ⁷
Not Detected	<i>Acinetobacter calcoaceticus-baumannii</i> complex				
Not Detected	<i>Enterobacter cloacae</i> complex				
Not Detected	<i>Escherichia coli</i>				
Not Detected	<i>Haemophilus influenzae</i>				

Antimicrob

☒ N/A	CTX-M
☒ N/A	IMP
☒ N/A	KPC
☑ Detected	<i>mecA/C</i> and MREJ
☒ N/A	NDM
☒ N/A	OXA-48-like
☒ N/A	VIM

Atypical Bacteria

Not Detected	<i>Chlamydia pneumoniae</i>
Not Detected	<i>Legionella pneumophila</i>
Not Detected	<i>Mycoplasma pneumoniae</i>

Viruses

Not Detected	Adenovirus
Not Detected	Coronavirus
Not Detected	Human Metapneumovirus
Not Detected	Human Rhinovirus/Enterovirus
Not Detected	Influenza A
Not Detected	Influenza B
Not Detected	Middle East Respiratory Syndrome Coronavirus (MERS-CoV)
Not Detected	Parainfluenza Virus
Not Detected	Respiratory Syncytial Virus

FilmArray® Pneumonia Panel plus - IVD
 A BIOMERIEUX COMPANY
 www.BioFireDx.com

Run Information

Sample ID	113355 BORZANOVIC MILORAD	Run Date	13 Dec 2020 7:11 PM
Protocol	BAL v3.3	Serial No.	30889242
Pouch Type	Pneumoplus v2.0	Lot No.	376220
Controls	Passed	Operator	lab lab (Lab IKVBD)
Run Status	Completed	Instrument	2FA06341

Result Summary

Bacteria

	Bin (copies/mL)	Bin (copies/mL)			
		10 ⁴	10 ⁵	10 ⁶	≥10 ⁷
☑ Detected	≥10 ⁷	////	////	////	////
Not Detected	<i>Acinetobacter calcoaceticus-baumannii</i> complex				
Not Detected	<i>Enterobacter cloacae</i> complex				
Not Detected	<i>Escherichia coli</i>				
Not Detected	<i>Haemophilus influenzae</i>				
Not Detected	<i>Klebsiella aerogenes</i>				
Not Detected	<i>Klebsiella oxytoca</i>				
☑ Detected	10 ⁵	////	////	////	////
Not Detected	<i>Klebsiella pneumoniae</i> group				
Not Detected	<i>Moraxella catarrhalis</i>				
Not Detected	<i>Proteus</i> spp.				
Not Detected	<i>Pseudomonas aeruginosa</i>				
Not Detected	<i>Serratia marcescens</i>				
Not Detected	<i>Staphylococcus aureus</i>				
Not Detected	<i>Streptococcus agalactiae</i>				
Not Detected	<i>Streptococcus pneumoniae</i>				
Not Detected	<i>Streptococcus pyogenes</i>				

Antimicrobial Resistance Genes

Not Detected	CTX-M
Not Detected	IMP
☑ Detected	KPC
☒ N/A	<i>mecA/C</i> and MREJ
☑ Detected	NDM
Not Detected	OXA-48-like
Not Detected	VIM

Atypical Bacteria

Not Detected	<i>Chlamydia pneumoniae</i>
Not Detected	<i>Legionella pneumophila</i>
Not Detected	<i>Mycoplasma pneumoniae</i>

Viruses

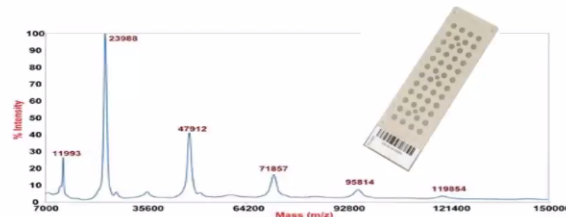
Not Detected	Adenovirus
Not Detected	Coronavirus
Not Detected	Human Metapneumovirus
Not Detected	Human Rhinovirus/Enterovirus
Not Detected	Influenza A
Not Detected	Influenza B
Not Detected	Middle East Respiratory Syndrome Coronavirus (MERS-CoV)
Not Detected	Parainfluenza Virus
Not Detected	Respiratory Syncytial Virus

Infekcije krvi: direktna detekcija iz hemokulture

The PATIENT in ICU



hands-on time = 2-3 min



K. pneumoniae

Bitte die Packungseinlage für Details beachten.

Probe	bk2056		Prozeß Abgeschlossen	01-31-16, 01:16 p.m.
Test	BC-GN	?	Analyse Abgeschlossen	01-31-16, 02:08 p.m.
Kartusche	02148436			

Übersicht

Detail	
Acinetobacter	Nicht detektiert
Enterobacter	Nicht detektiert
E. coli	Nicht detektiert
K. oxytoca	Nicht detektiert
VIM	Nicht detektiert
CTX-M	Detektiert
NDM	Nicht detektiert
S. marcescens	Nicht detektiert
Citrobacter	Nicht detektiert
Proteus	Nicht detektiert
P. aeruginosa	Nicht detektiert
K. pneumoniae	Detektiert
OXA	Nicht detektiert
KPC	Nicht detektiert
IMP	Nicht detektiert

Information Verbrauchsmaterial

Kartusche	REF 20-006-021	2016-05-09	LOT 111115021A
Extraktionsplatte	REF 20-009-021	2016-05-09	LOT 111115021B
Reagenzienplatte	REF 20-011-021	2016-05-09	LOT 111115021C

Prozessor-Qualitätskontrollstatus

Prozessor Modul	A:1
Prozessierungszeit	113 Minuten
Prozessierungstemperatur	38,5 °C
Interne Kontrollen	Bestanden

RESULT: after 1,5-h

- Patient: Mr. J.-C. H. (78)
 - Patient with COPD + diabetes mellitus, already hospitalized for 3 weeks because of COVID-19.
 - **LAST NIGHT around 21:30 patient was transferred from intermediate-care to ICU because of high BT, CRP, and leukocytes.**
 - **The patient was without AB *Th.* and the BC were collected**

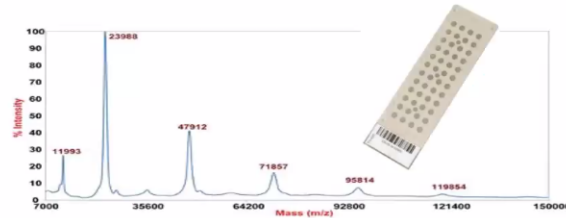
CEFTAZIDIME + MOXIFLOXACIN were administrated

Infekcije krvi: direktna detekcija iz hemokulture/direktna detekcija iz krvi

The PATIENT in ICU



hands-on time = 2-3 min



K. pneumoniae

Bitte die Packungseinlage für Details beachten.

Probe	bk2056		Prozeß Abgeschlossen	01-31-16, 01:16 p.m.
Test	BC-GN	?	Analyse Abgeschlossen	01-31-16, 02:08 p.m.
Kartusche	02148436			

Übersicht

Detail	
Acinetobacter	Nicht detektiert
Enterobacter	Nicht detektiert
E. coli	Nicht detektiert
K. oxytoca	Nicht detektiert
VIM	Nicht detektiert
CTX-M	Detektiert
NDM	Nicht detektiert
S. marcescens	Nicht detektiert
Citrobacter	Nicht detektiert
Proteus	Nicht detektiert
P. aeruginosa	Detektiert
OXA	Nicht detektiert
KPC	Nicht detektiert
IMP	Nicht detektiert

Information Verbrauchsmaterial

Kartusche	REF 20-006-021	2016-05-09	LOT 111115021A
Extraktionsplatte	REF 20-009-021	2016-05-09	LOT 111115021B
Reagenzienplatte	REF 20-011-021	2016-05-09	LOT 111115021C

Prozessor-Qualitätskontrollstatus

Prozessor Modul	A:1
Prozessierungszeit	113 Minuten
Prozessierungstemperatur	38,5 °C
Interne Kontrollen	Bestanden

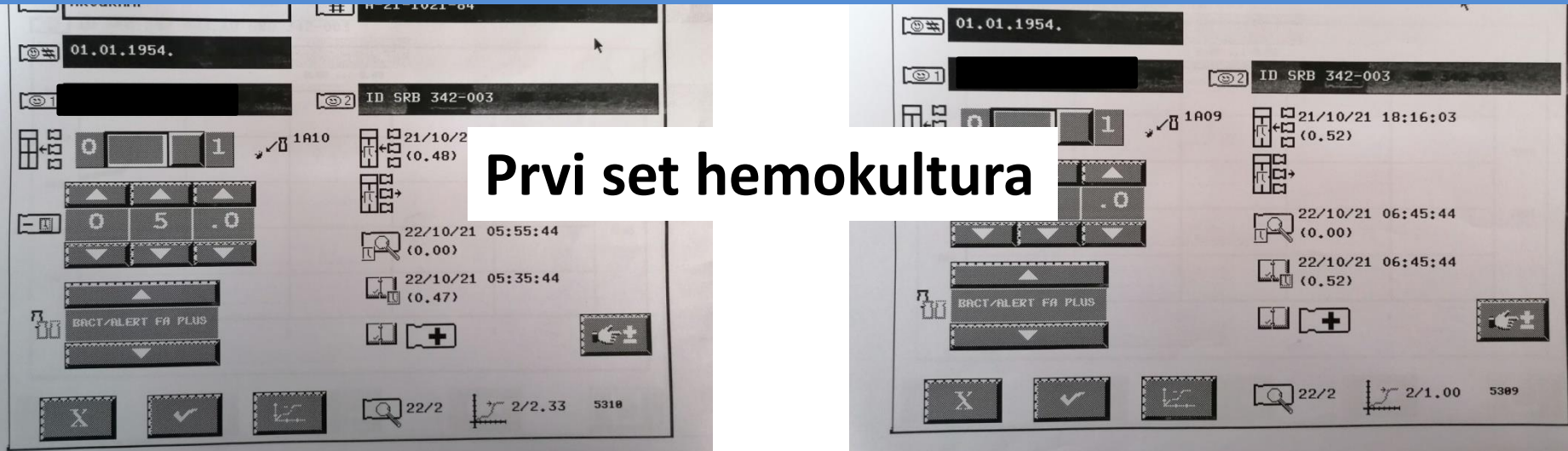
RESULT: after 1,5-h

- Patient: Mr. J.-C. H. (78)
 - Patient with COPD + diabetes mellitus, already hospitalized for 3 weeks because of COVID-19.
 - **LAST NIGHT around 21:30 patient was transferred from intermediate-care to ICU because of high BT, CRP, and leukocytes.**
 - **The patient was without AB *Th.* and the BC were collected**

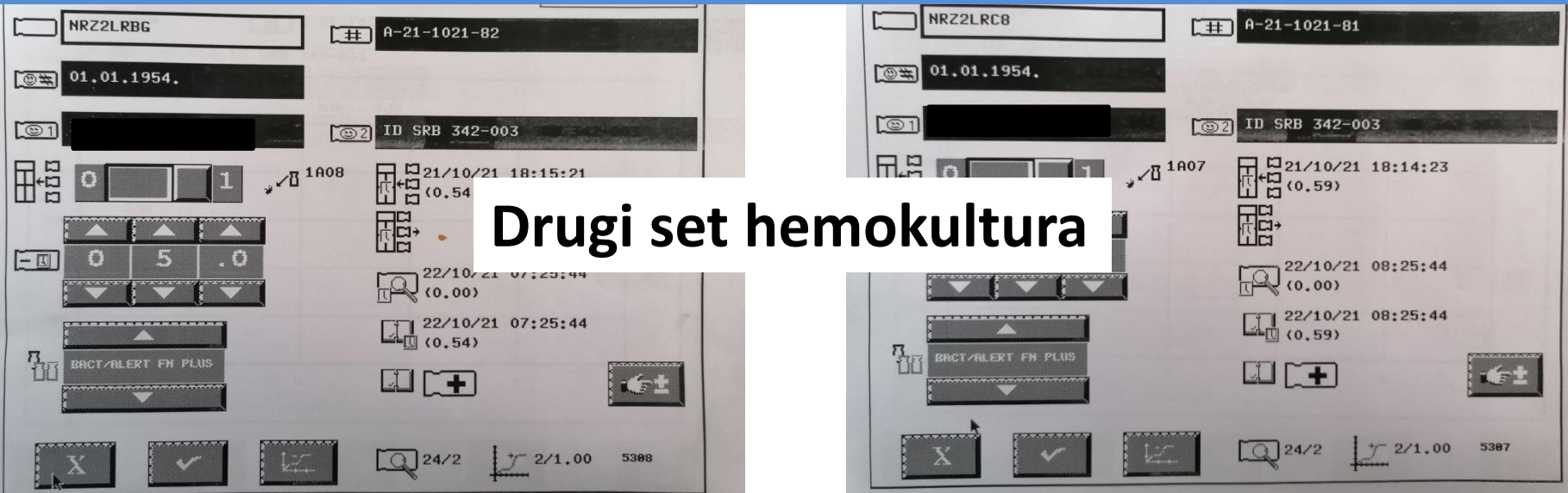
CEFTAZIDIME + MOXIFLOXACIN were administrated

Therapy was escalated to MEROPENEM

Naše iskustvo



U dva seta hemokultura (4 bočice) – detektovan *Staphylococcus epidermidis*, meticilin osetljiv (MSSE)



Brzi molekularni test

Blood culture (hemokultura) panel

Pozitivne hemokultura

43 targeta – 1 test – 60 minuta

GRAM-NEGATIVE BACTERIA

Acinetobacter calcoaceticus-
baumannii complex

Bacteroides fragilis

Enterobacterales

Enterobacter cloacae cp.

Escherichia coli

Klebsiella aerogenes

Klebsiella oxytoca

Klebsiella pneumoniae group

Proteus

Salmonella

Serratia marcescens

Haemophilus influenzae

Neisseria meningitidis

Pseudomonas aeruginosa

Stenotrophomonas maltophilia

GRAM-POSITIVE BACTERIA

Enterococcus faecalis

Enterococcus faecium

Listeria monocytogenes

Staphylococcus

Staphylococcus aureus

Staphylococcus epidermidis

Staphylococcus lugdunensis

Streptococcus

Streptococcus agalactiae

Streptococcus pneumoniae

Streptococcus pyogenes

YEAST

Candida albicans

Candida auris

Candida glabrata

Candida krusei

Candida parapsilosis

Candida tropicalis

Cryptococcus

neoformans/gattii

ANTIMICROBIAL RESISTANCE GENES

Carbapenemases

IMP

KPC

NDM

OXA-48-like

VIM

Colistin Resistance

mcr-1

ESBL

CTX-M

Methicillin Resistance

mecA/C

***mecA/C* and MREJ (MRSA)**

Vancomycin Resistance

vanA/B



ATC

03-011

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ENISO 15189

<http://www.konzilijum.rs>

info@konzilijum.rs

Ime pacijenta: [REDACTED]

Datum prijema: [REDACTED]

Pol: Ž Datum rođenja: 05.10.1943

Datum izdavanja: [REDACTED]

IZVEŠTAJ O ISPITIVANJU

Broj protokola [REDACTED]

Vrsta uzorka: **Brisevi kože i površinskih rana/bris rane**

Analiza: **Bakteriološka kultura-izolovanje, identifikacija i ispitivanje osetljivosti na antibiotike klinički značajnih bakterija**

REZULTAT

ANTIBIOGRAM

Metoda: 3/1

Pseudomonas aeruginosa - Veliki broj

Metoda: 1/1, 2/1

Piperacilin	I
Piperacilin + tazo baktam	I
Ceftazidim	I
Cefepim	I
Imipenem	I
Meropenem	S
Amikacin	S
Tobramicin	S
Ciprofloksacin	I
Levofloksacin	I

Napomena: Za terapiju sistemskih infekcija aminoglikozidne antibiotike primenjivati isključivo u kombinaciji sa drugim aktivnim antimikrobnim lekom.

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REZULTAT	ANTIBIOGRAM <small>Metoda: 3/1</small>
----------	--

Pseudomonas aeruginosa - Veliki broj

Metoda: 1/1, 2/1

Piperacilin	I
Piperacilin + tazo baktam	I
Ceftazidim	I
Cefepim	I
Imipenem	I
Meropenem	S
Amikacin	S
Tobramicin	S
Ciprofloksacin	I
Levofloksacin	I

Aminoglikozidi – samo kombinovana terapija

Napomena: Za terapiju sistemskih infekcija aminoglikozidne antibiotike primenjivati isključivo u kombinaciji sa drugim aktivnim antimikrobnim lekom.

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IZVEŠTAJ O ISPITIVANJU

Broj protokola [REDACTED]

Vrsta uzorka: **Brisevi kože i površinskih rana/bris rane**

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REZULTAT	ANTIBIOGRAM <small>Metoda: 3/1</small>
----------	--

Pseudomonas aeruginosa - Veliki broj

Metoda: 1/1, 2/1

Ceftazidim-avibaktam
Ceftolozan-tazobaktam
Meropenem-vaborbaktam
Imipenem-relebaktam
Kolistin

Piperacilin	I
Piperacilin + tazo baktam	I
Ceftazidim	I
Cefepim	I
Imipenem	I
Meropenem	S
Amikacin	S
Tobramicin	S
Ciprofloksacin	I
Levofloksacin	I

Aminoglikozidi – samo kombinovana terapija

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Broj protokola [REDACTED]

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Analiza: **Bakteriološka kultura-izolovanje, identifikacija i ispitivanje osetljivosti na antibiotike klinički značajnih bakterija**

REZULTAT

ANTIBIOGRAM

Metoda: 3/1

Pseudomonas aeruginosa - Veliki broj

Metoda: 1/1, 2/1

Piperacilin	I
Piperacilin + tazo baktam	I
Ceftazidim	I
Cefepim	I
Imipenem	I
Meropenem	S
Amikacin	S
Tobramicin	S
Ciprofloksacin	I
Levofloksacin	I

Napomena: Za terapiju sistemskih infekcija aminoglikozidne antibiotike primenjivati isključivo u kombinaciji sa drugim aktivnim antimikrobnim lekom.

SIR - нове дефиниције 2019. год.





ATC

03-011

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Napomena: Za terapiju sistemskih infekcija aminoglikozidne antibiotike primenjivati isključivo u kombinaciji sa drugim aktivnim antimikrobnim lekom.

00:09:00

**Bacteria
growth**

With penicillin

Without penicillin

4 um

