

Oltre l'epidemia da SARS-CoV-2: l'impatto su vita e lavoro ed i patogeni da monitorare nel prossimo futuro

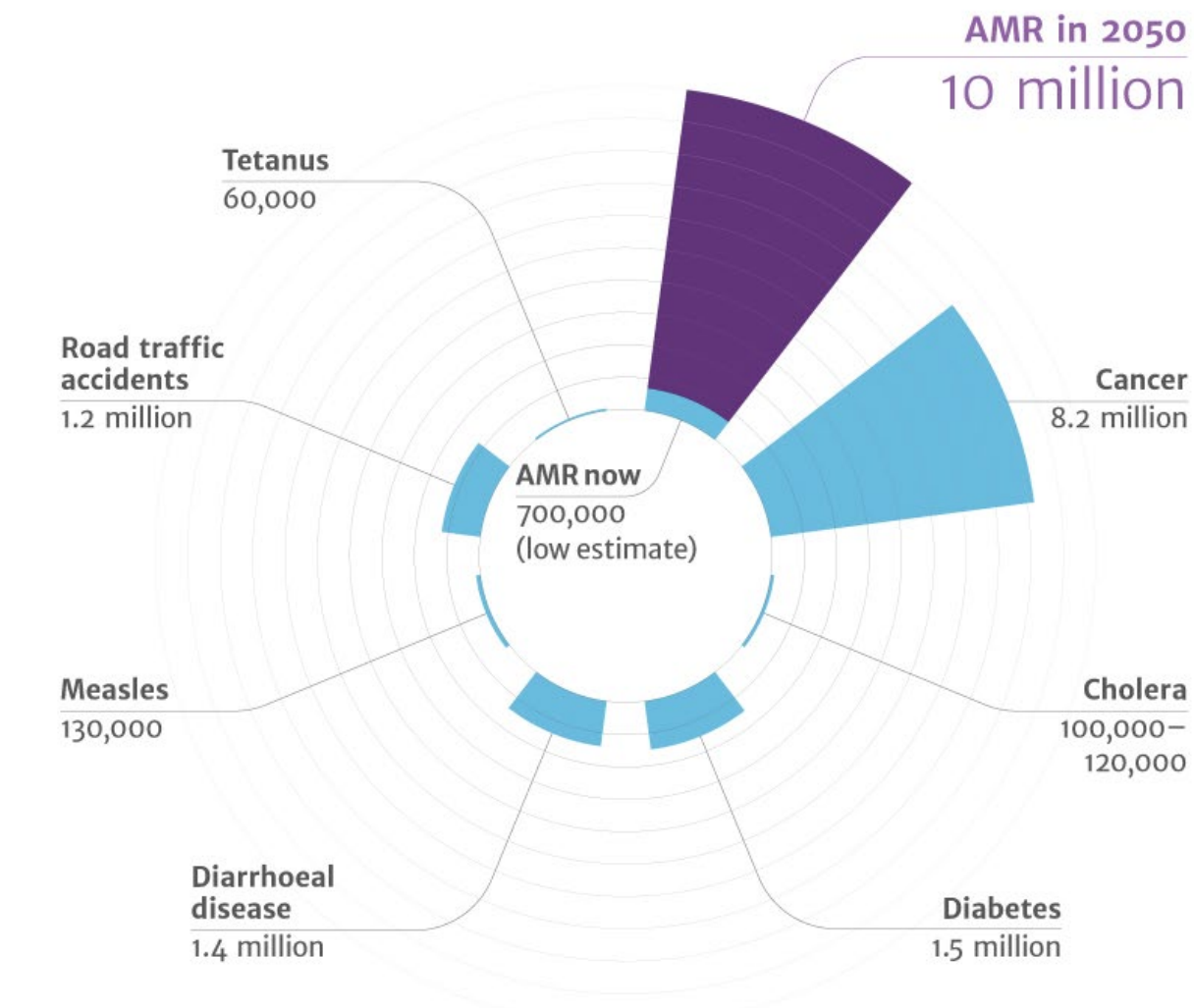
WEBINAR ECM **Crediti ECM: 4**

26 ottobre 2023 ore 9.30-14.00

L'andamento dell'antibiotico-resistenza in Toscana nel 2022 (Rete SMART)

Silvia LM Forni
ARS Toscana

L'AMR è la più grande minaccia alla salute globale e alla medicina moderna



Sources:

Diabetes: www.who.int/mediacentre/factsheets/fs312/en/ Cancer: www.who.int/mediacentre/factsheets/fs297/en/
Cholera: www.who.int/mediacentre/factsheets/fs107/en/ Diarrhoeal disease: www.sciencedirect.com/science/article/pii/S0140673612617280
Measles: www.sciencedirect.com/science/article/pii/S0140673612617280 Road traffic accidents: www.who.int/mediacentre/factsheets/fs358/en/
Tetanus: www.sciencedirect.com/science/article/pii/S0140673612617280

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



Articles

Global burden of bacterial antimicrobial resistance in 2019: a systematic analysis



Antimicrobial Resistance Collaborators*

Summary

Background Antimicrobial resistance (AMR) poses a major threat to human health around the world. Previous publications have estimated the effect of AMR on incidence, deaths, hospital length of stay, and health-care costs for specific pathogen–drug combinations in select locations. To our knowledge, this study presents the most comprehensive estimates of AMR burden to date.



Lancet 2022; 399: 629–55
Published Online
January 20, 2022
[https://doi.org/10.1016/S0140-6736\(21\)02724-0](https://doi.org/10.1016/S0140-6736(21)02724-0)

Associated deaths

Most inclusive estimate of AMR burden

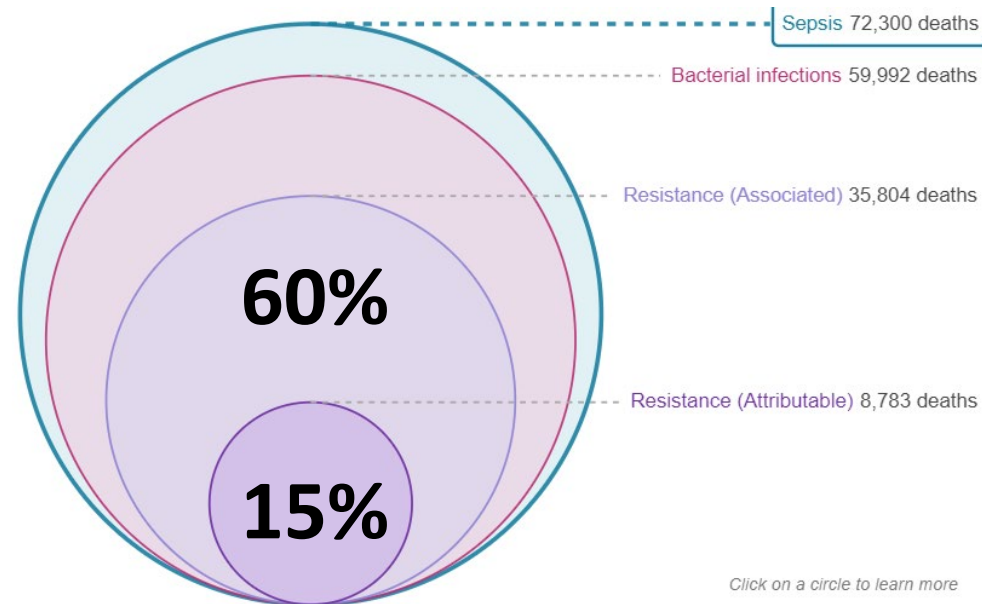
Associated deaths measures people with a drug-resistant infection that contributed to their death. The infection was implicated in their death, but resistance may or may not have been a factor.

Attributable deaths

Most conservative estimate of AMR burden

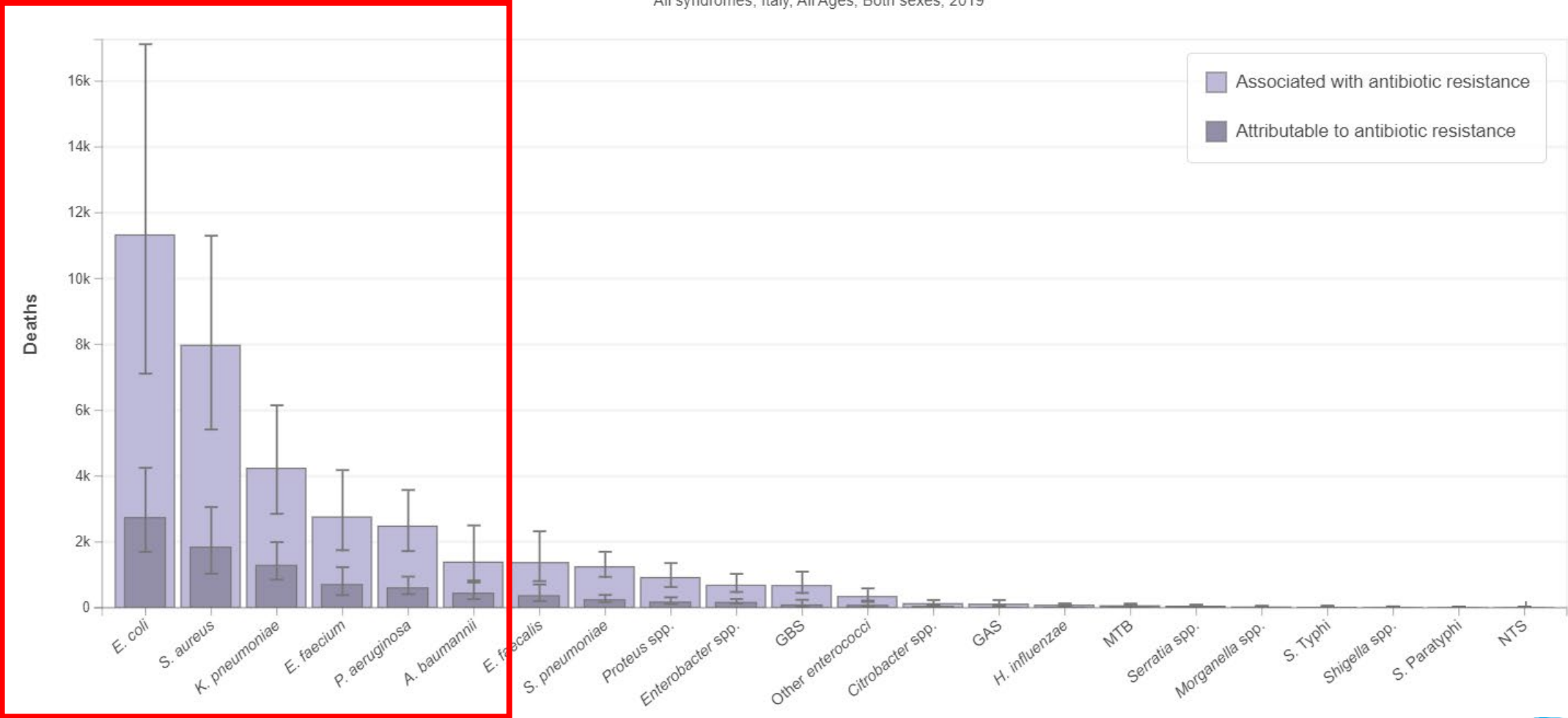
Attributable deaths measures people who would not have died of infection if it was treatable (i.e., if there was no AMR) for whom resistance can be said to have caused their death.

Italia 119 decessi per 100K



Deaths both associated with and attributable to bacterial antimicrobial resistance by pathogen

All syndromes, Italy, All Ages, Both sexes, 2019



Box 6: WHO priority pathogens list for R&D of new antibiotics

On 27th February 2017, the WHO published a catalogue of 12 families of bacteria that should be priorities for R&D efforts, in addition to TB. The list was drawn up in a bid to guide and promote R&D and highlights in particular the threat of gram-negative bacteria that are resistant to multiple antibiotics.

Priority 1: CRITICAL

- *Acinetobacter baumannii*, carbapenem-resistant
- *Pseudomonas aeruginosa*, carbapenem-resistant
- *Enterobacteriaceae*, carbapenem-resistant, ESBL-producing

Priority 2: HIGH

- *Enterococcus faecium*, vancomycin-resistant
- *Staphylococcus aureus*, methicillin-resistant, vancomycin-intermediate and resistant
- *Helicobacter pylori*, clarithromycin-resistant
- *Campylobacter* spp., fluoroquinolone-resistant
- *Salmonellae*, fluoroquinolone-resistant
- *Neisseria gonorrhoeae*, cephalosporin-resistant, fluoroquinolone-resistant

Priority 3: MEDIUM

- *Streptococcus pneumoniae*, penicillin-non-susceptible
- *Haemophilus influenzae*, ampicillin-resistant
- *Shigella* spp., fluoroquinolone-resistant

The objective of the WHO prioritization exercise was to identify previously unrecognised health threats due to increasing antibiotic resistance.

Batteriemie **multidrug resistant (MDR)** - resistenti contemporaneamente a vari antibiotici:

- causano infezioni difficili da trattare; vi sono poche o nessuna opzione di trattamento.
- facilita la diffusione della resistenza agli antibiotici.
- complica gli sforzi per ridurre la resistenza.

Piano Nazionale di Contrasto all'Antibiotico-Resistenza (PNCAR) 2022-2025



Piano Nazionale di Contrasto all'Antibiotico-Resistenza PNCAR 2022-2025

Appendice: funghi, virus e parassiti



SORVEGLIANZA E MONITORAGGIO

- ABR
- ICA
- Uso antibiotici
- Monitoraggio ambientale



PREVENZIONE DELLE INFEZIONI

- ICA
- Malattie infettive e zoonosi



BUON USO ANTIBIOTICI

- Ambito umano
- Ambito veterinario
- Corretta gestione e smaltimento

Governance

Formazione

Informazione, comunicazione e trasparenza

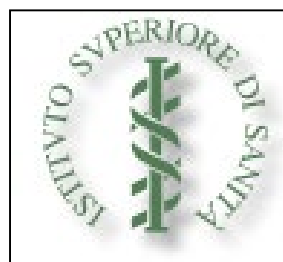
Ricerca, innovazione e bioetica

Cooperazione nazionale e internazionale

La sorveglianza dell'antibiotico resistenza



**European Antimicrobial Resistance
Surveillance Network (EARS-Net)**



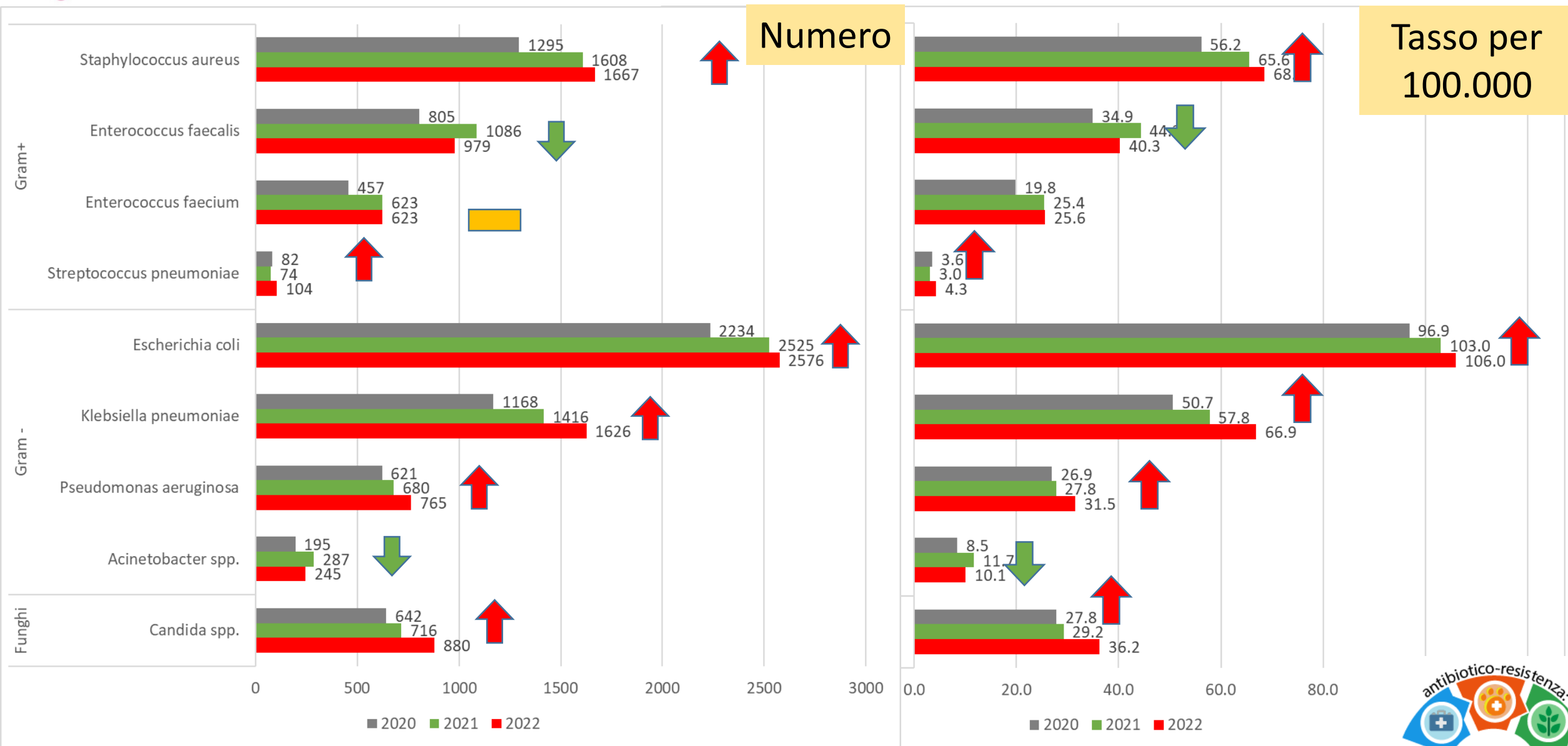
**AR-ISS:
sorveglianza nazionale
dell'Antibiotico-Resistenza**



**La rete Sorveglianza Microbiologica e
dell'Antibiotico-Resistenza Toscana**



Isolati da sangue non ripetuti, Toscana, 2020-2022

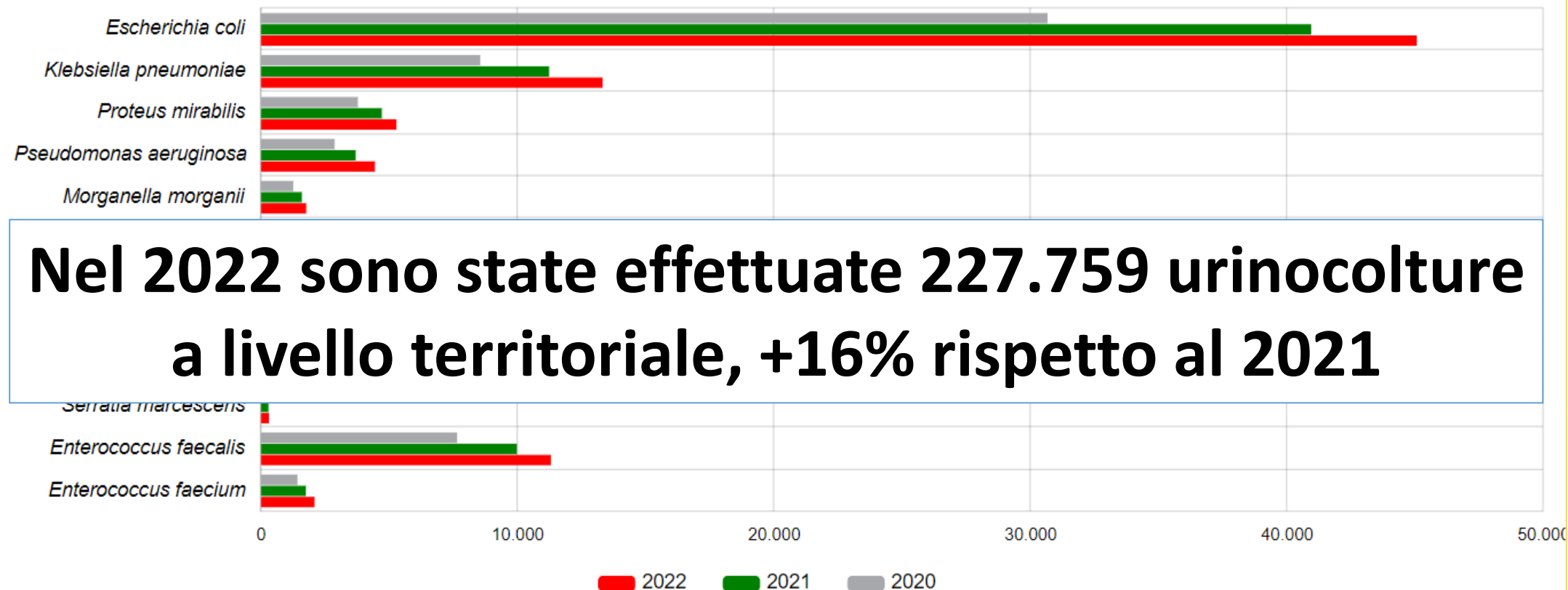


Isolati da urine non ripetuti, Toscana, 2020-2022

Numerosità delle specie analizzate in Toscana

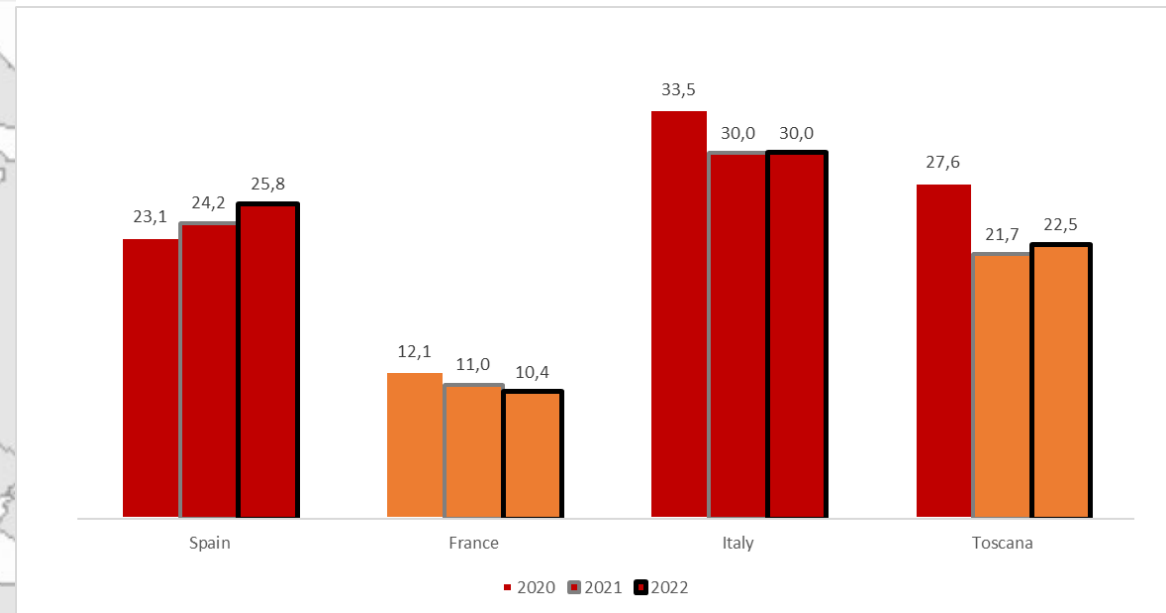
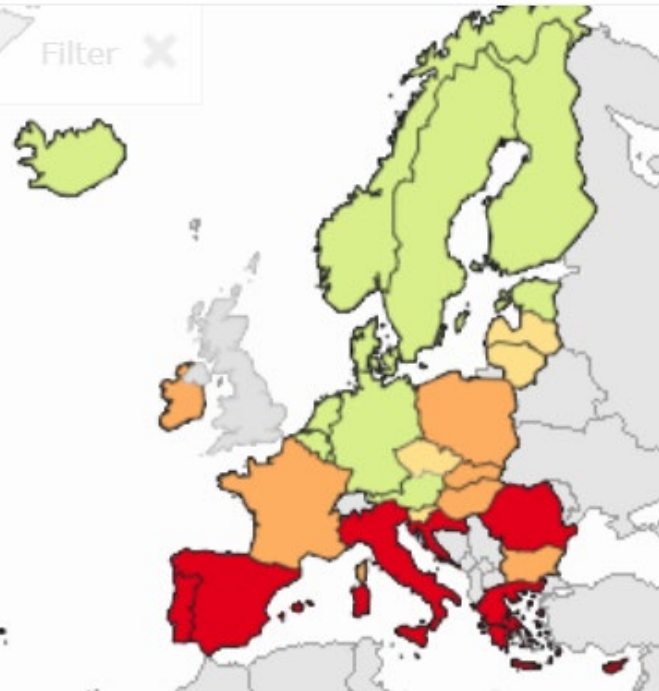
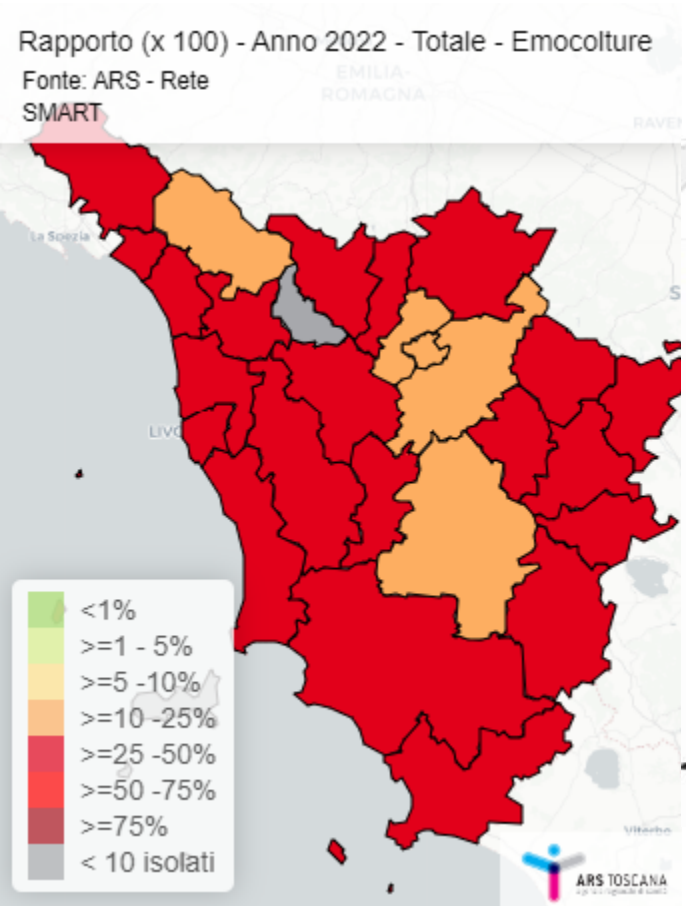
Rapporto (x 100) - Anno 2022 - Totale - Urinocolture

Fonte: ARS - Rete SMART



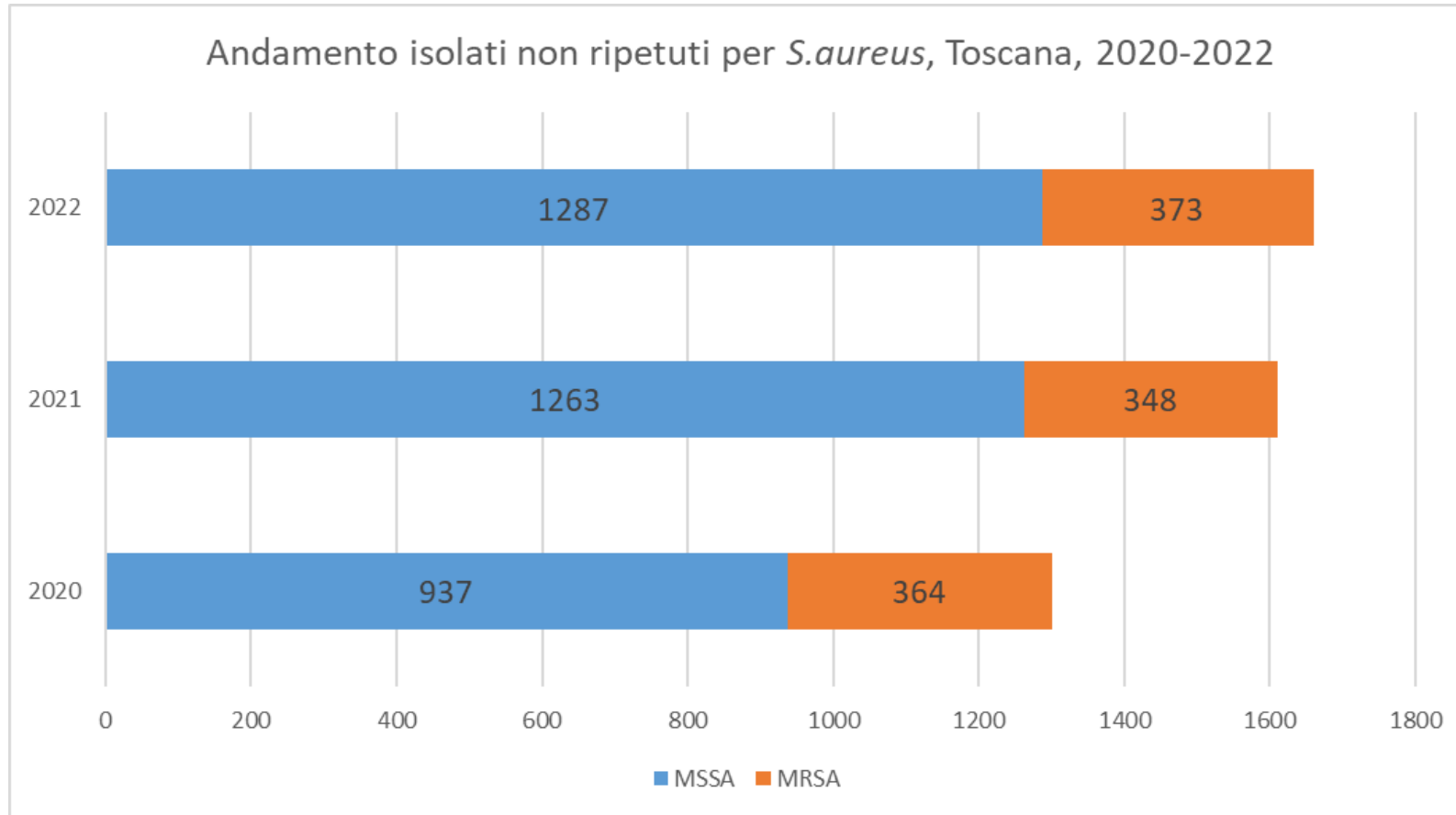
Staphylococcus aureus, meticillino-resistente (MRSA)

(1667 isolati di *S.aureus* da sangue non ripetuti, Toscana, 2022)



Staphylococcus aureus, meticillino-resistente (MRSA)

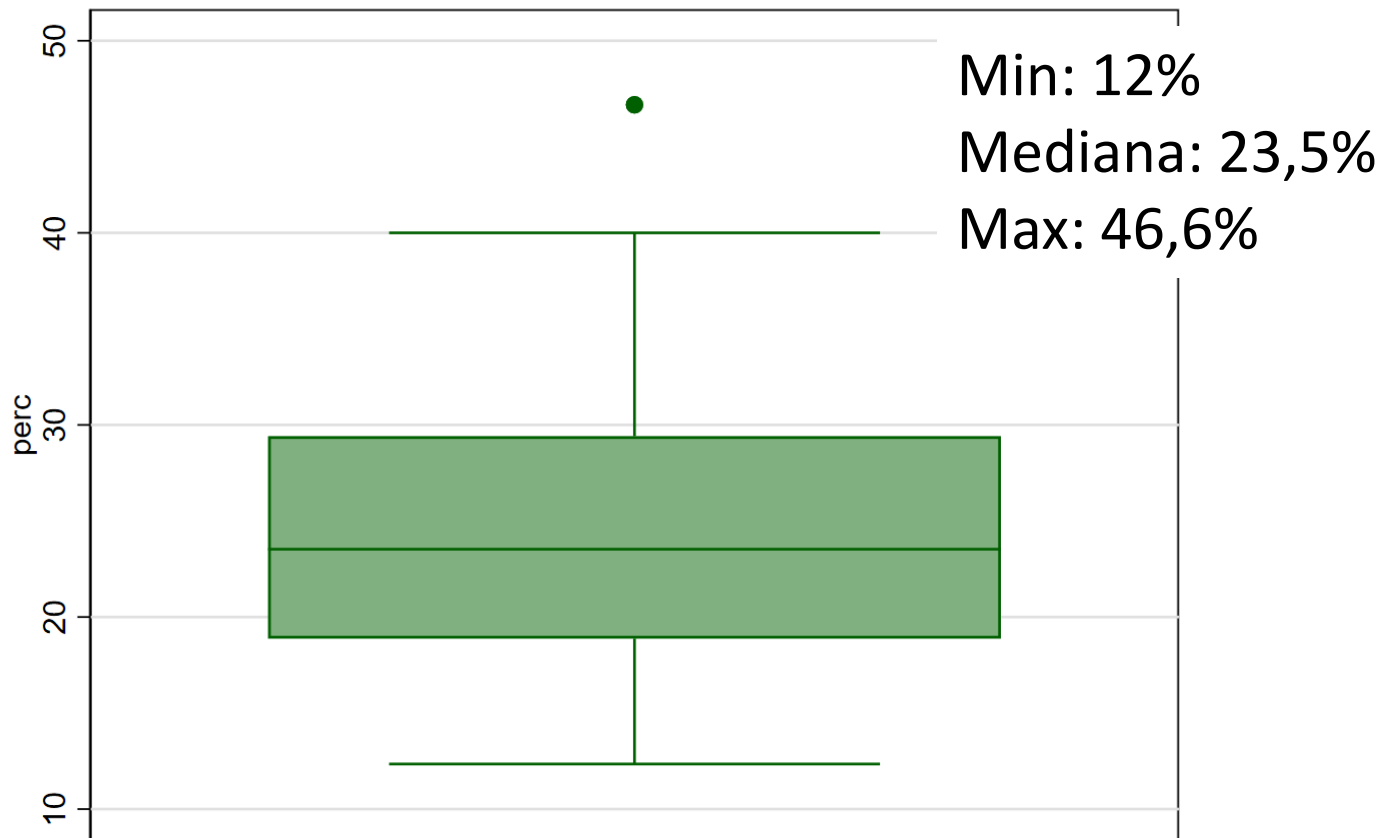
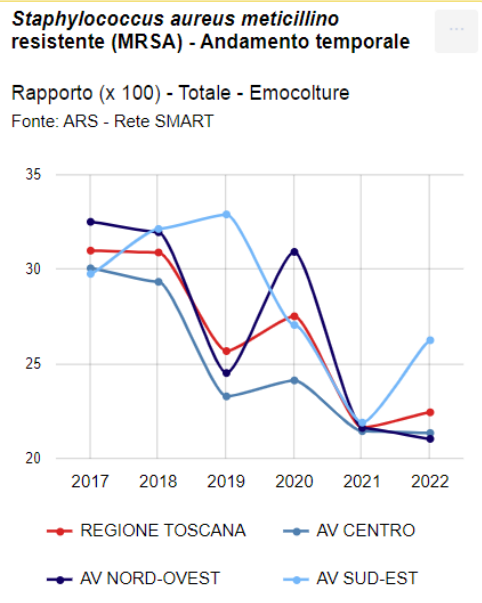
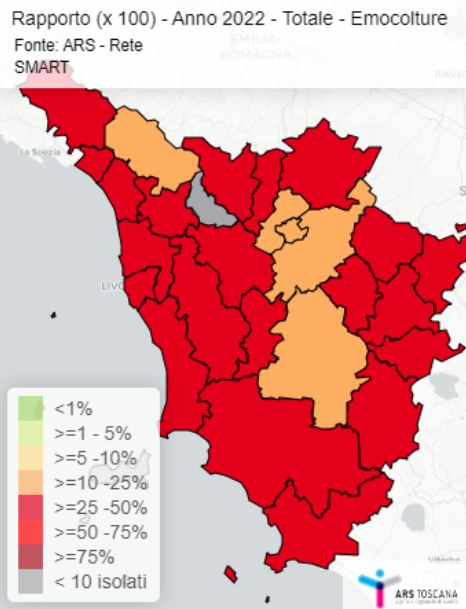
La riduzione «apparente» della prevalenza di MRSA tra il 2020 e il 2021 è principalmente dovuta all'aumento degli isolati MSSA



Luogo di isolamento	MSSA	MRSA
H	65.8%	65.1%
PS	26.2%	24.7%
RSA/Hospice	0.8%	1.6%
Altro, non noto	7.2%	8.6%

Staphylococcus aureus, meticillino-resistente (MRSA)

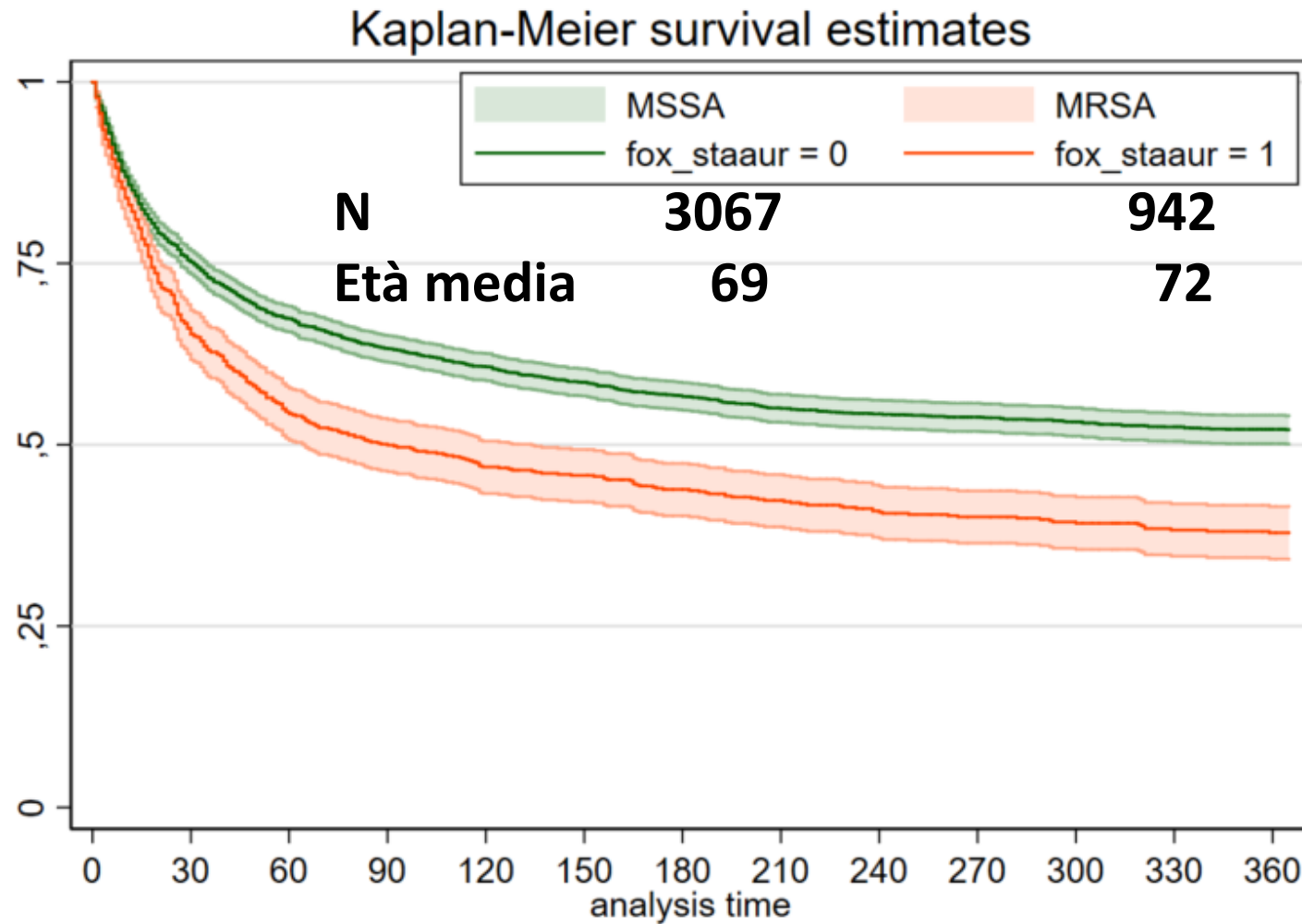
Prevalenza MRSA per ospedale, Toscana, 2022 (30 H con oltre 10 isolati)




Median Odds Ratio (MOR) 1,18

Passando da un ospedale a minor prevalenza di MRSA ad uno a maggior prevalenza, il rischio di MRSA aumenta del 18%

Staphylococcus aureus, meticillino-resistente (MRSA) – sopravvivenza a 1 anno Toscana 2020-2022



MICROBE
Measuring Infectious Causes and Resistance Outcomes for Burden Estimation

 The drug-bug combination with the greatest **attributable** fatal burden was

Methicillin-resistant *S. aureus*

[Explore more pathogen-drug combinations](#)

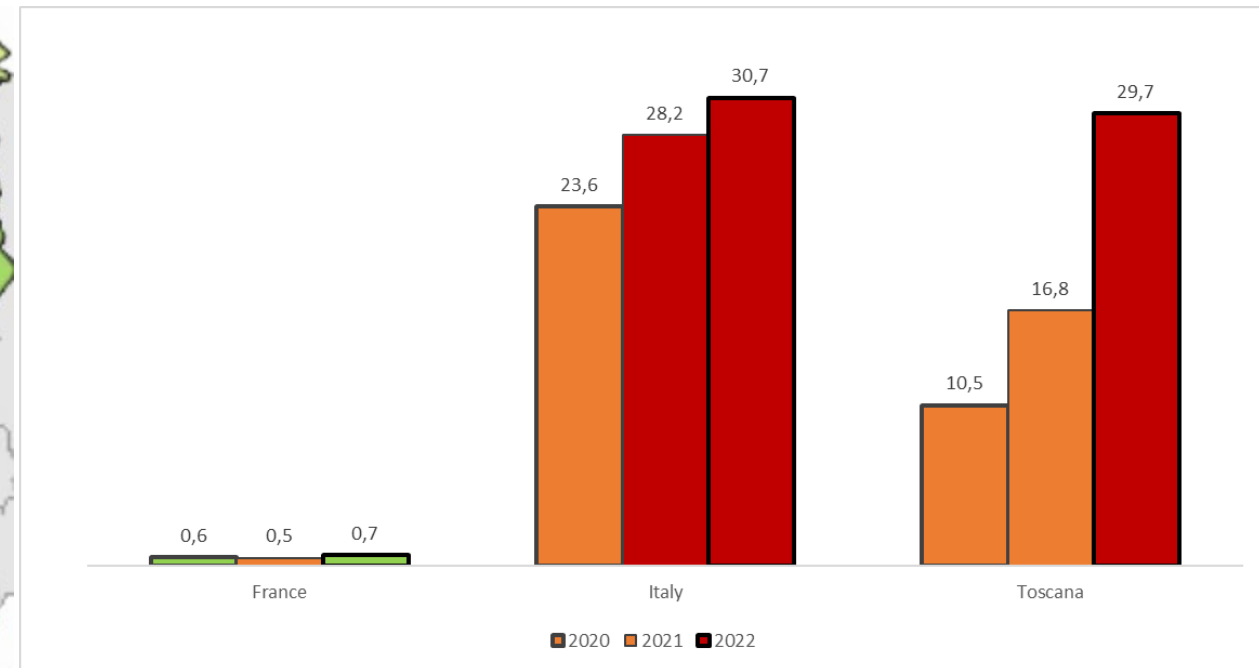
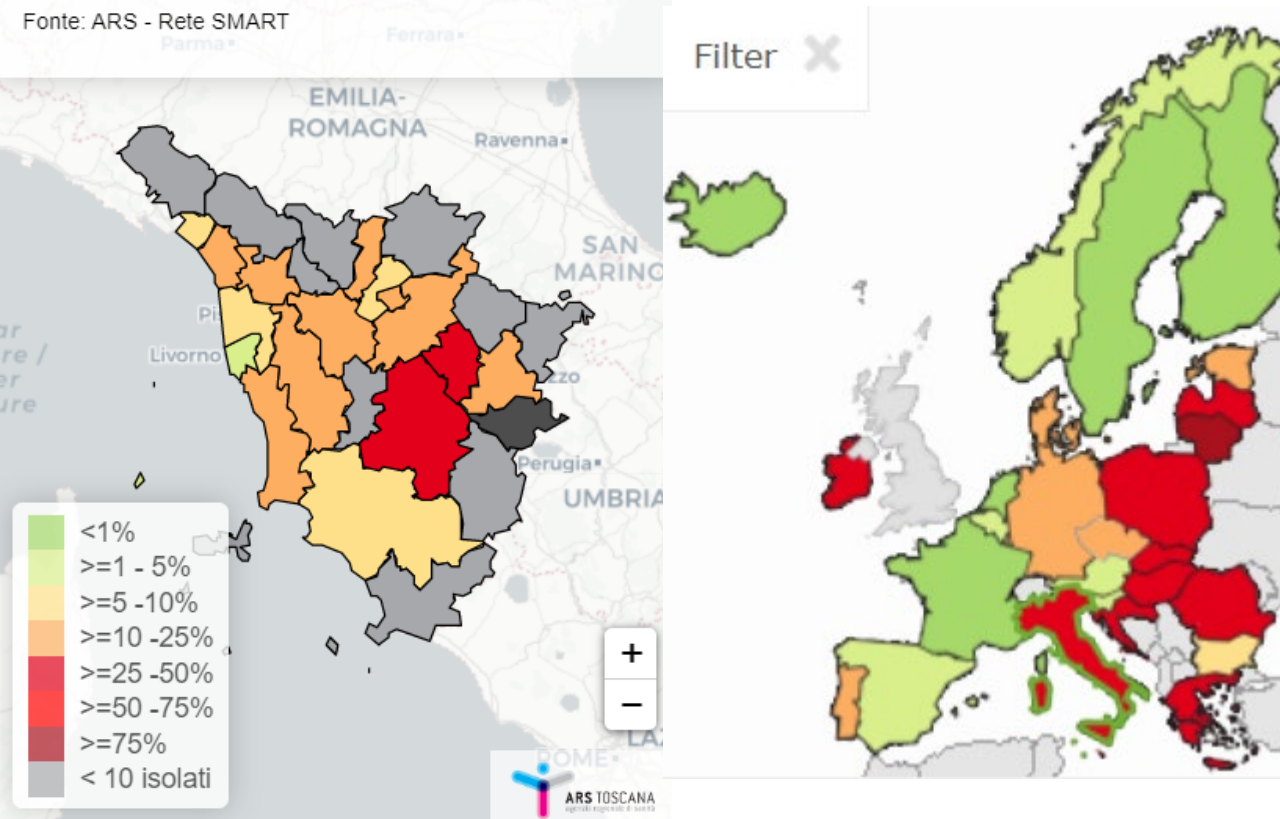
Enterococcus faecium, resistente alla vancomicina (VRE)

(623 isolati di *E.faecium* da sangue non ripetuti, 2022)

Enterococcus faecium resistente alla vancomicina - Mappa per zona distretto

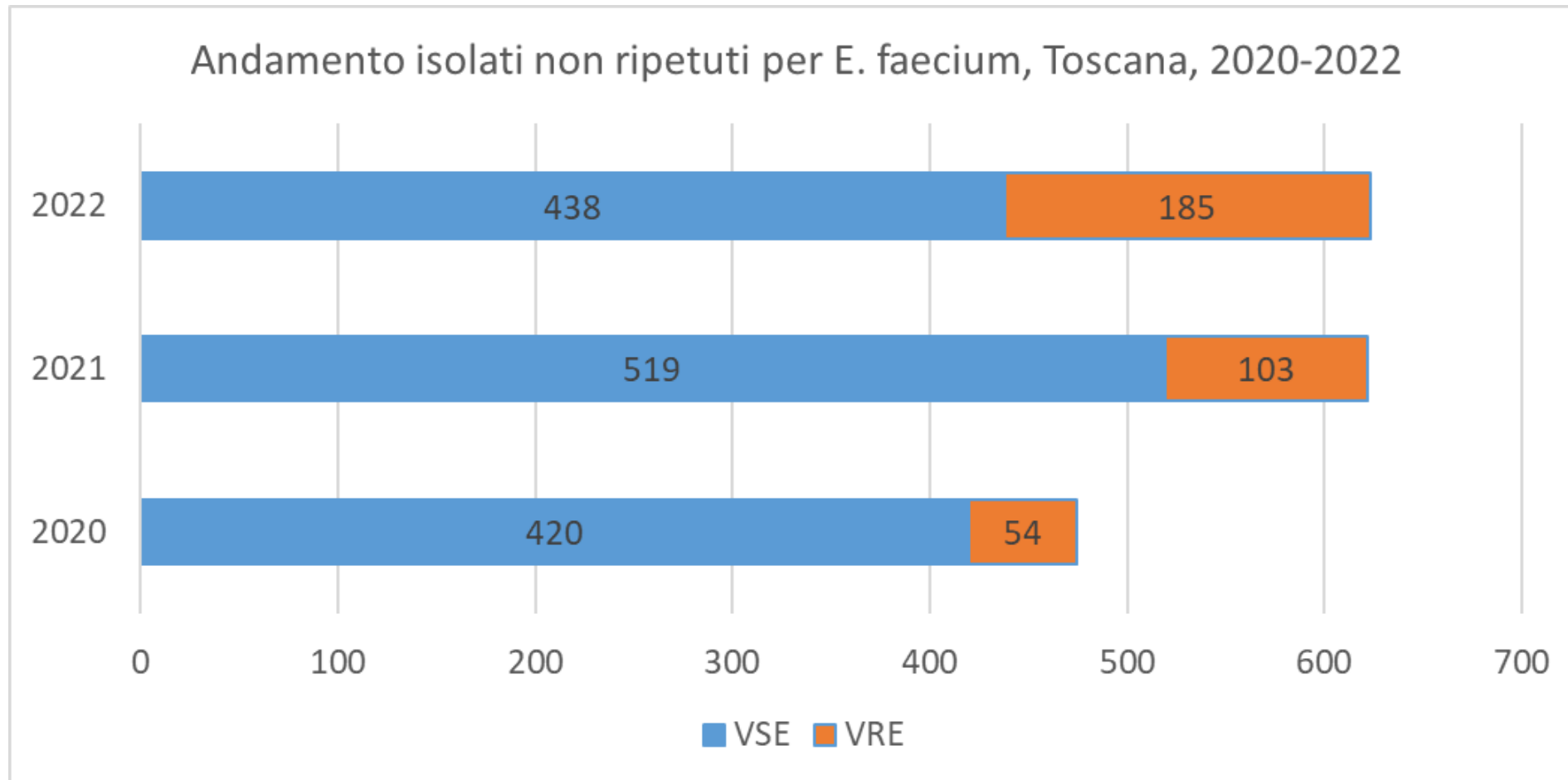
Rapporto (x 100) - Anno 2022 - Totale - Emocolture

Fonte: ARS - Rete SMART



Enterococcus faecium, resistente alla vancomicina (VRE)

Il numero di *E. faecium* isolati in Toscana nel 2022 non è aumentato, cresce il numero di VRE



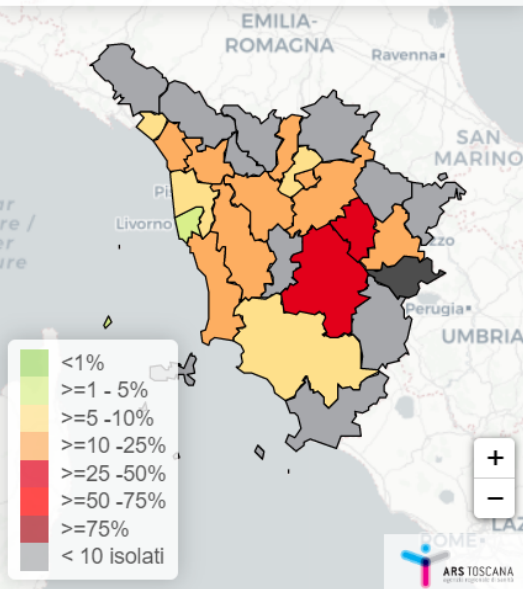
Enterococcus faecium, resistente alla vancomicina

(VRE)

Enterococcus faecium resistente alla vancomicina - Mappa per zona distretto

Rapporto (x 100) - Anno 2022 - Totale - Emocolture

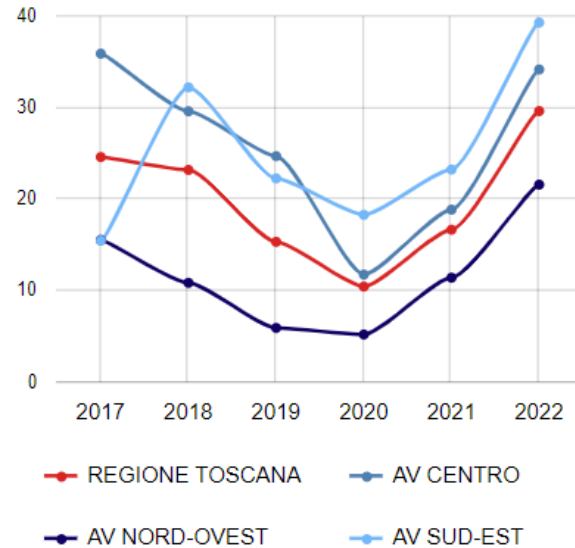
Fonte: ARS - Rete SMART



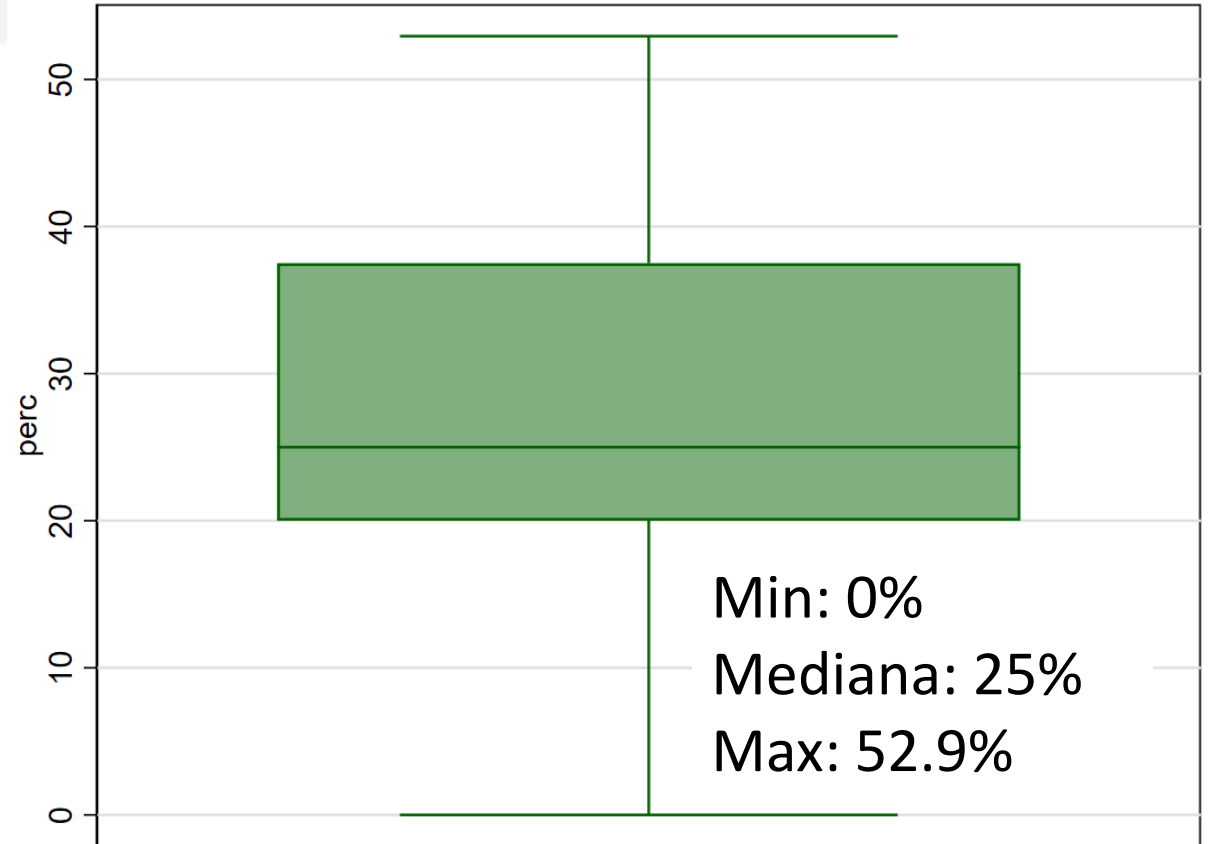
Enterococcus faecium resistente alla vancomicina - Andamento temporale

Rapporto (x 100) - Totale - Emocolture

Fonte: ARS - Rete SMART



Prevalenza VRE per ospedale, Toscana, 2022 (15 H con 10 + isolati)



Median Odds Ratio (MOR) 1,31

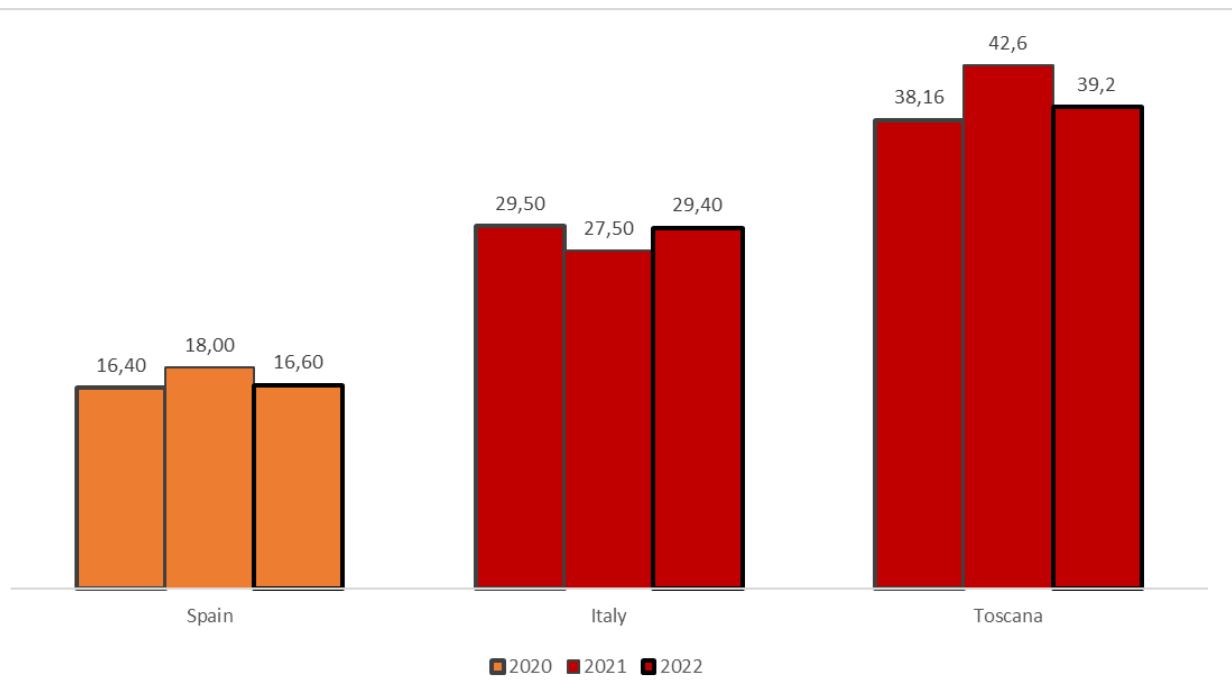
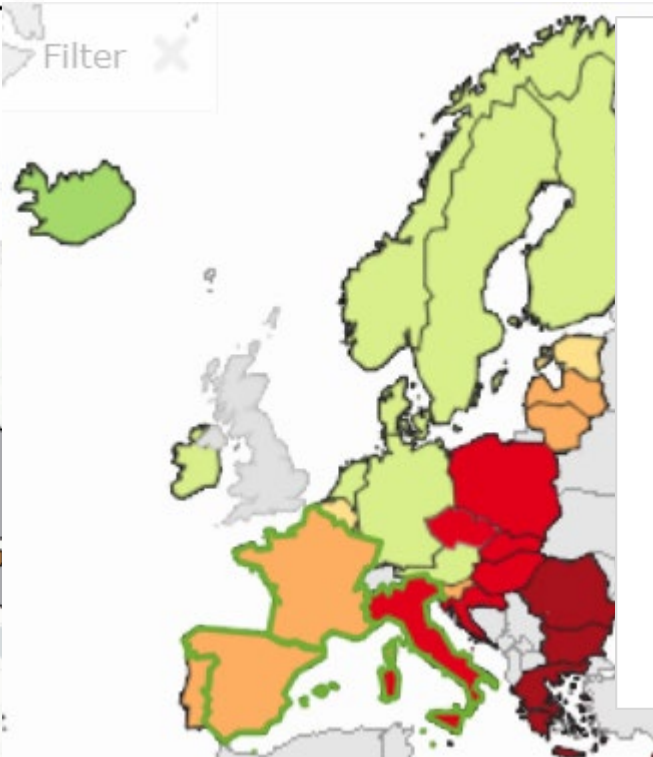
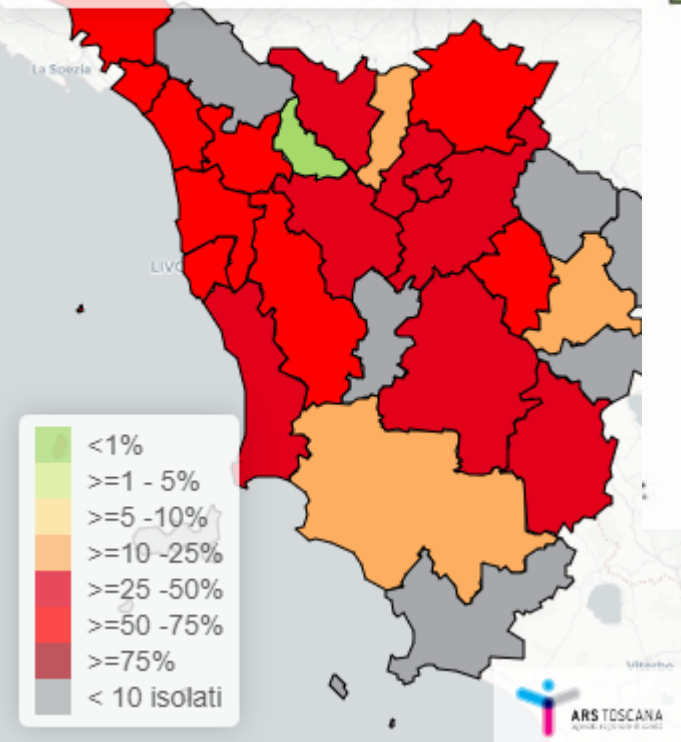
Passando da un ospedale a minor prevalenza di VRE ad uno a maggior prevalenza, il rischio di VRE aumenta del 31%

Klebsiella pneumoniae, multiresistente (cefalosporina di III generazione, fluorochinoloni e aminoglicosidi)

Klebsiella pneumoniae multiresistente (MDR) -
Mappa per zona distretto

Rapporto (x 100) - Anno 2022 - Totale -
Emocolture

Fonte: ARS - Rete SMART



(1625 isolati di *K.pneumoniae* da sangue non ripetuti, 2022)

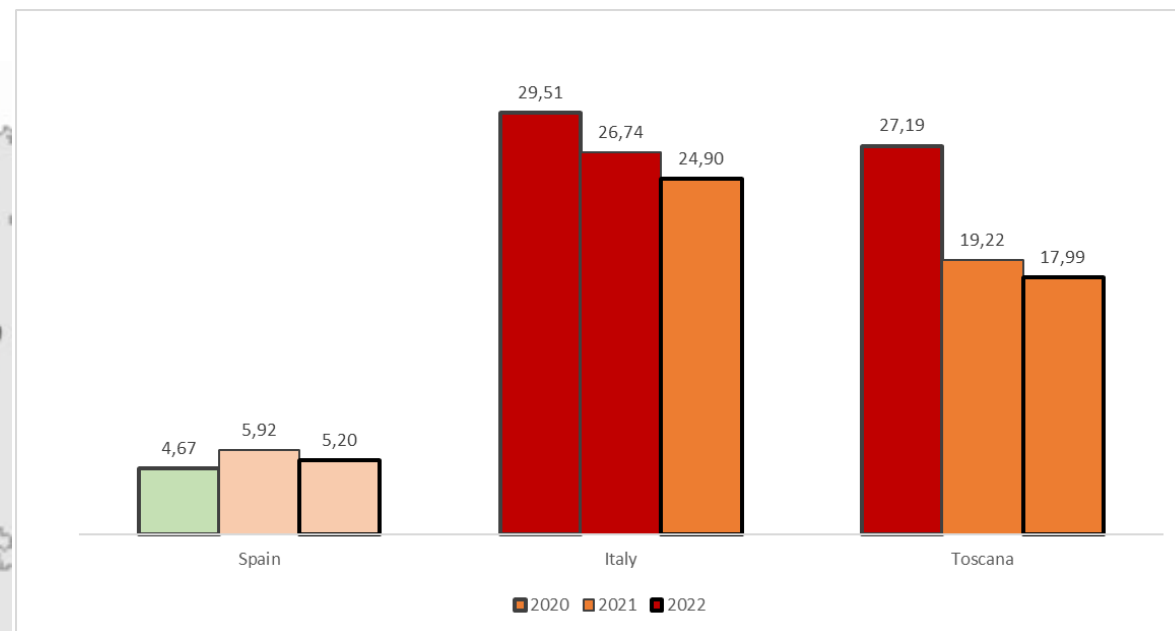
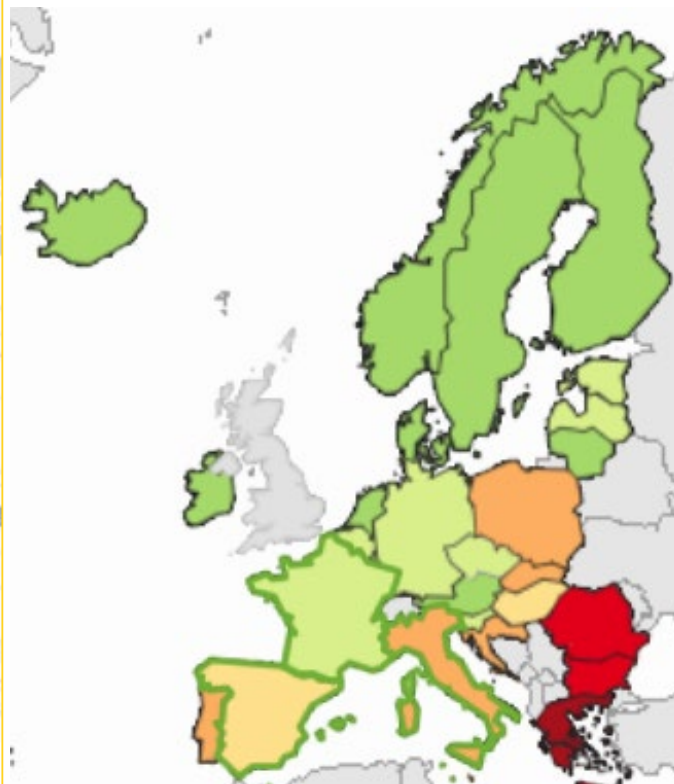
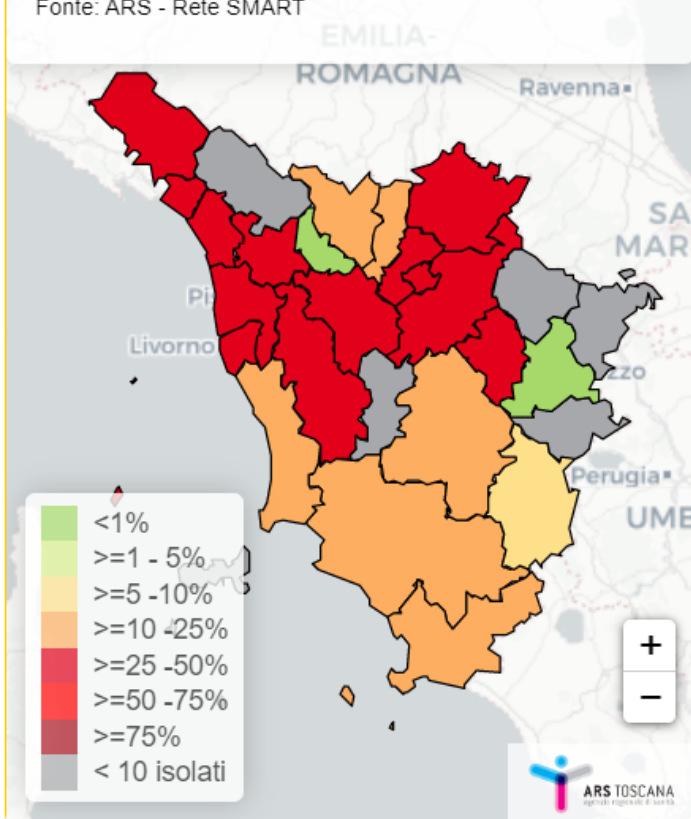
Klebsiella pneumoniae, resistente ai carbapenemi

(1625 isolati di *K.pneumoniae* da sangue non ripetuti, 2022)

Klebsiella pneumoniae resistente ai carbapenemi
- Mappa per zona distretto

Rapporto (x 100) - Anno 2022 - Totale - Emocolture

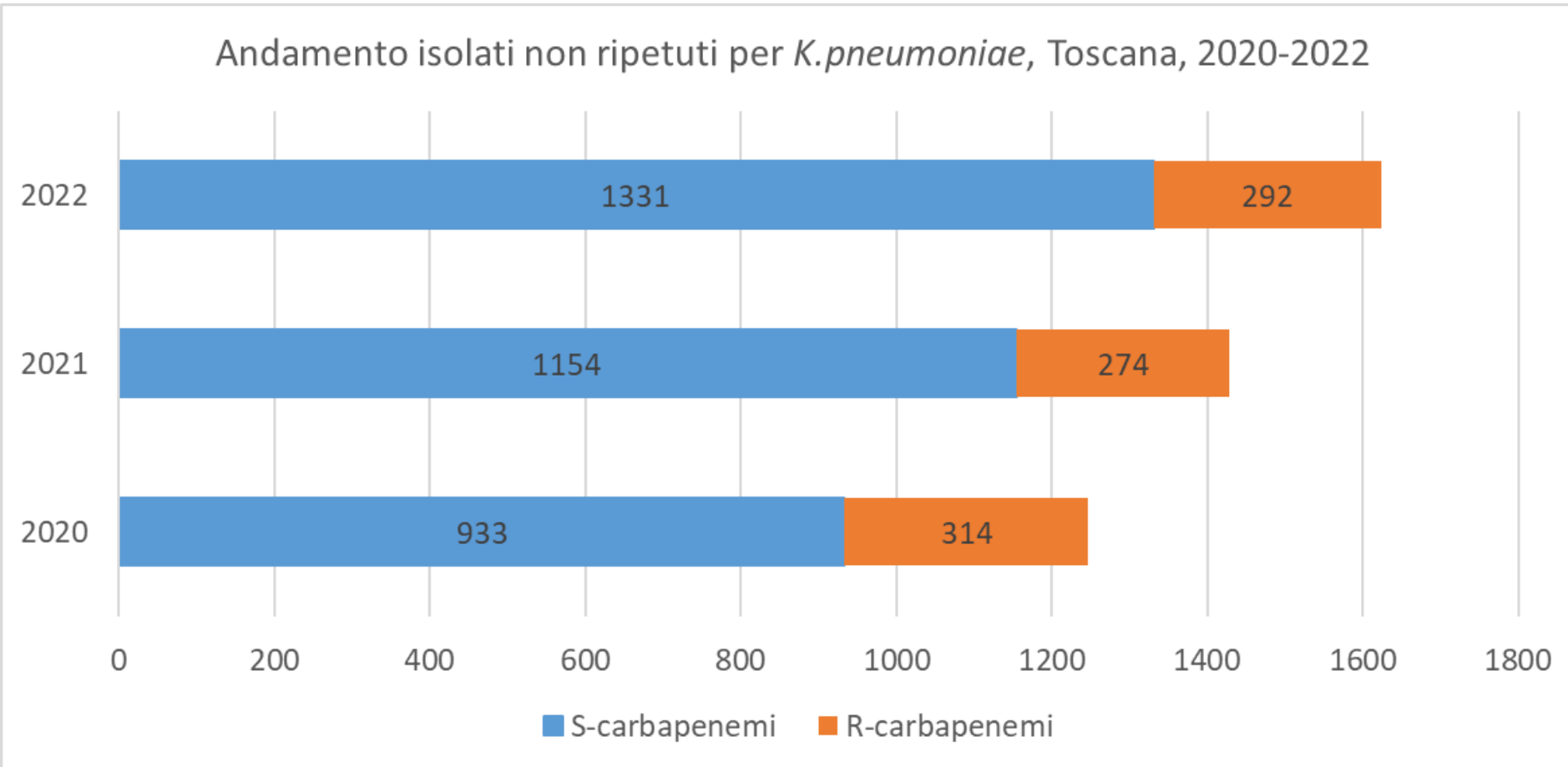
Fonte: ARS - Rete SMART



Klebsiella pneumoniae, resistente ai carbapenemi

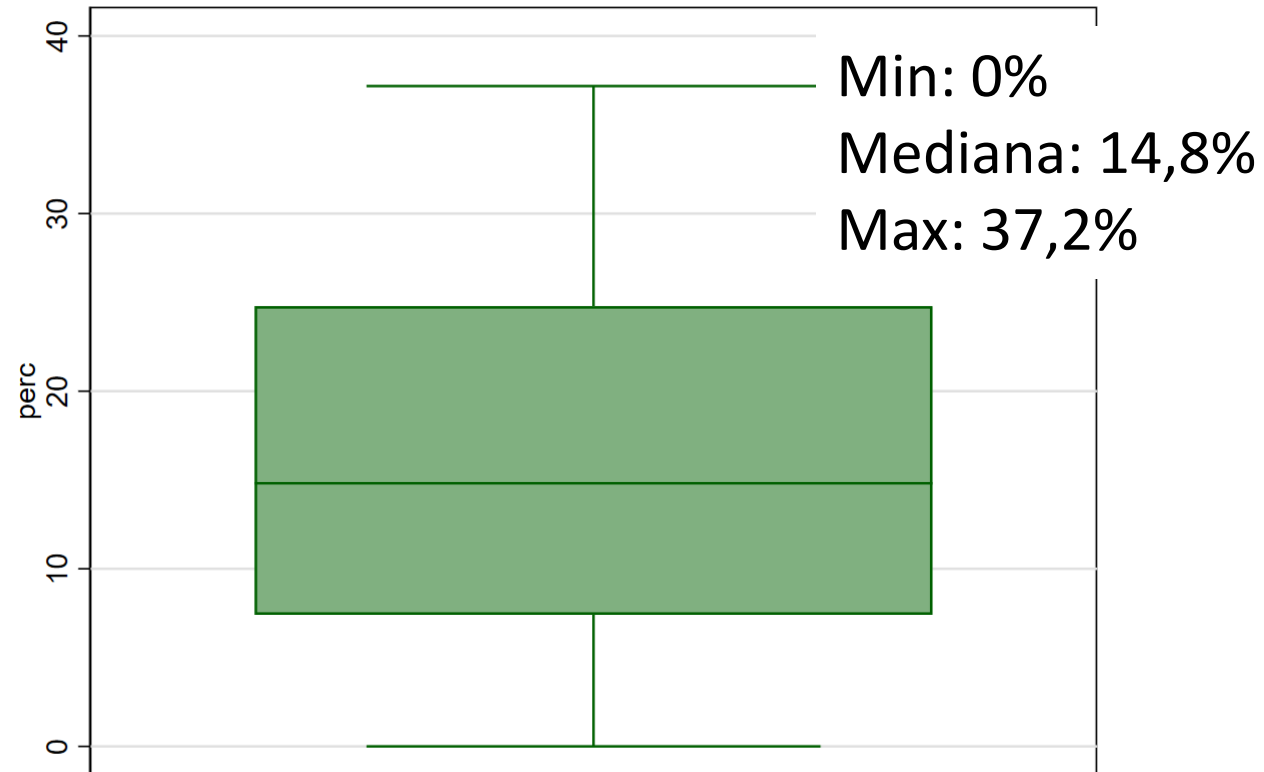
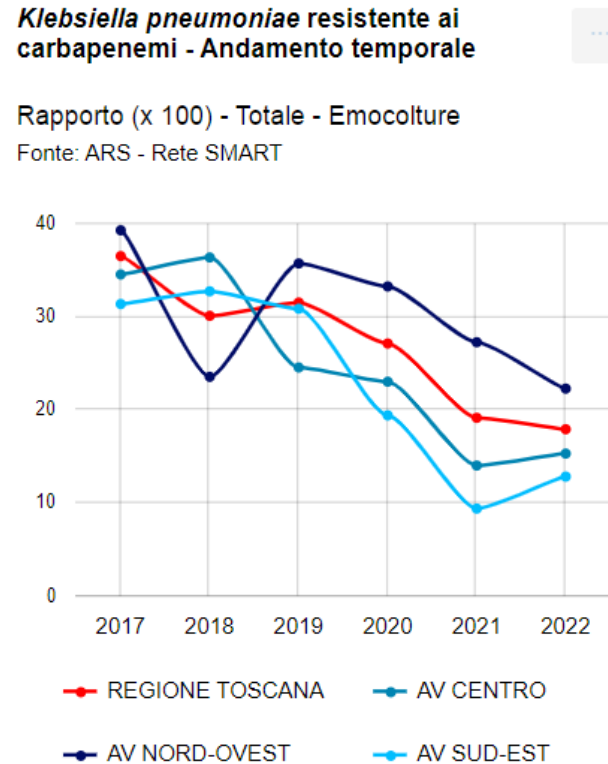
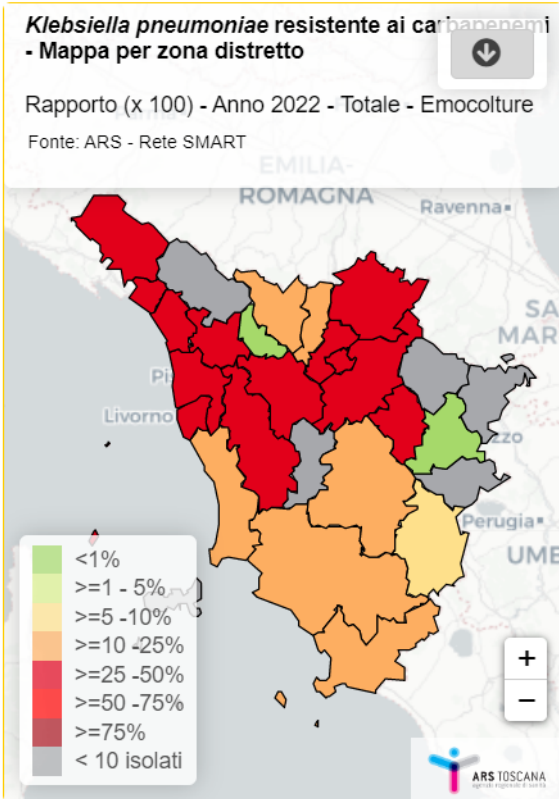
Il numero di *K. pneumoniae* isolate in Toscana nel 2022 è aumentato, in particolare i S-carbapenemi

Andamento isolati non ripetuti per *K.pneumoniae*, Toscana, 2020-2022



Klebsiella pneumoniae, resistente ai carbapenemi

Prevalenza *K. pneumoniae* per ospedale, Toscana 2022 (25 H con 10 + isolati)

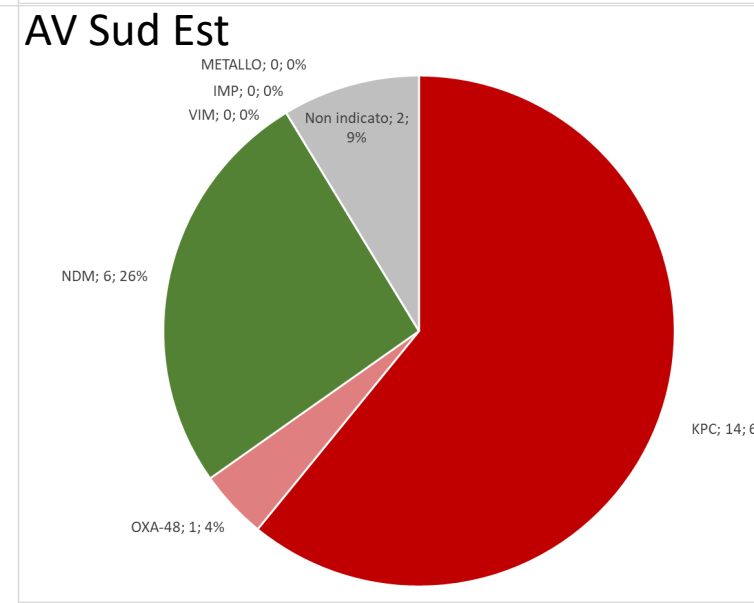
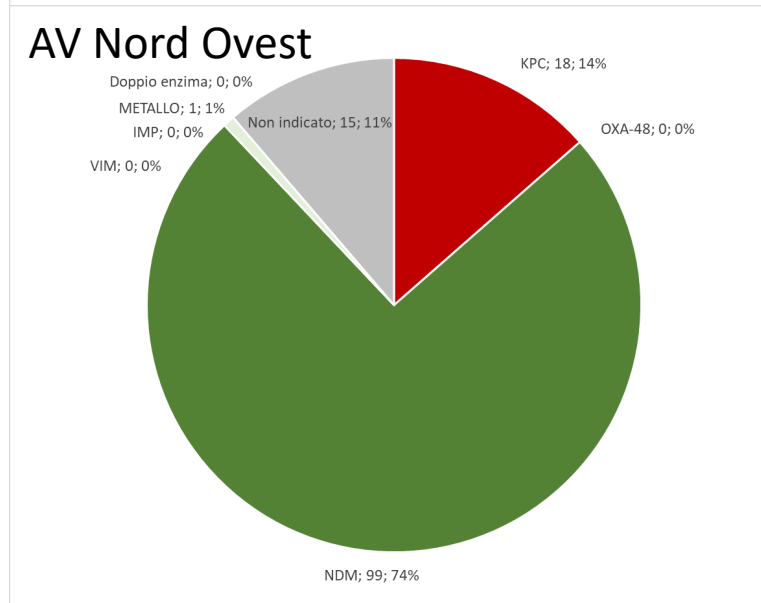
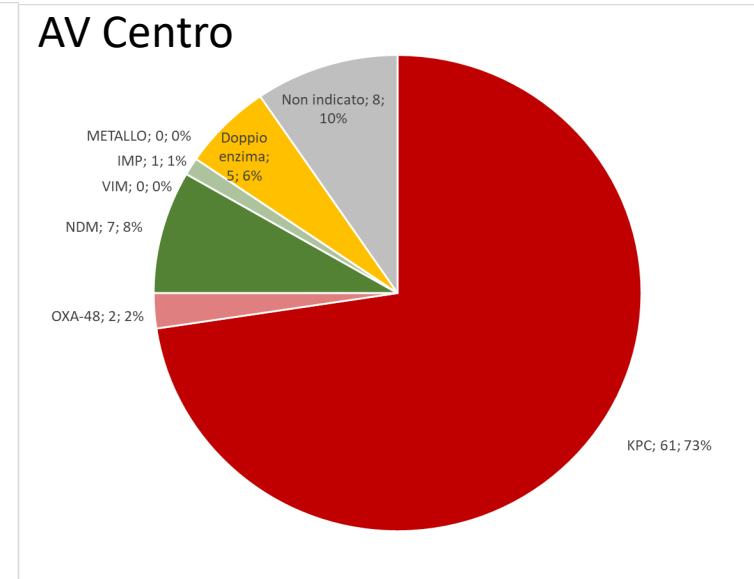
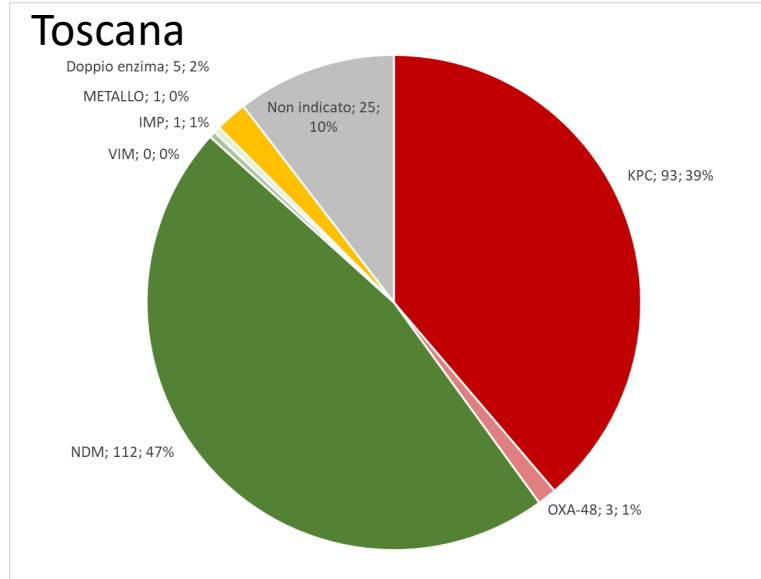


Median Odds Ratio (MOR) 2,59

Passando da un ospedale a minor prevalenza di *K. pneumoniae* resistenti a carbapenemi ad uno a maggior prevalenza, il rischio è più che doppio

Klebsiella pneumoniae, resistente ai carbapenemi

Enzimi responsabili (fonte: Sorveglianza CRE)

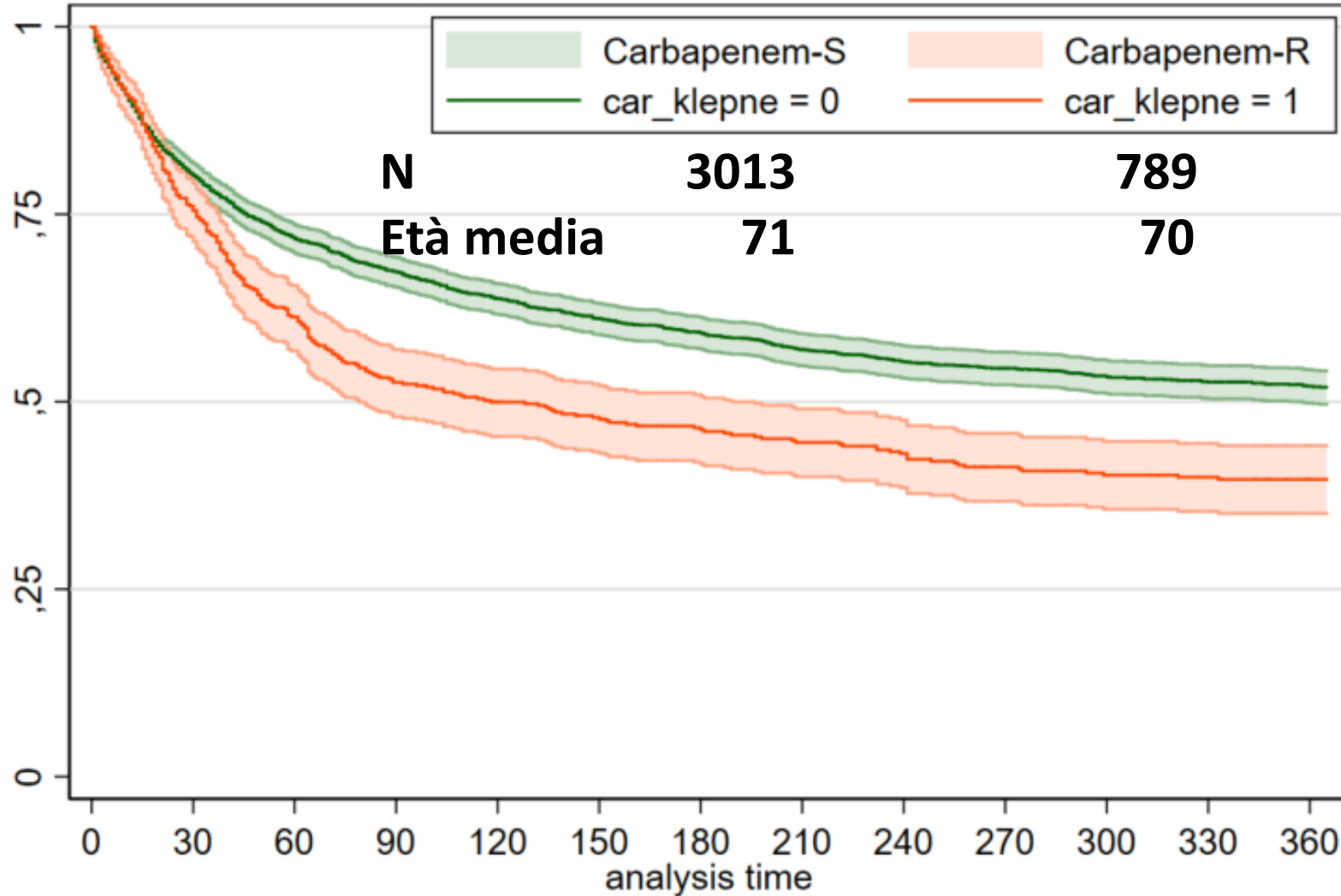


- KPC
- OXA-48
- NDM
- VIM
- IMP
- METALLO
- Doppio enzima
- Non indicato

Klebsiella pneumoniae, resistente ai carbapenemi

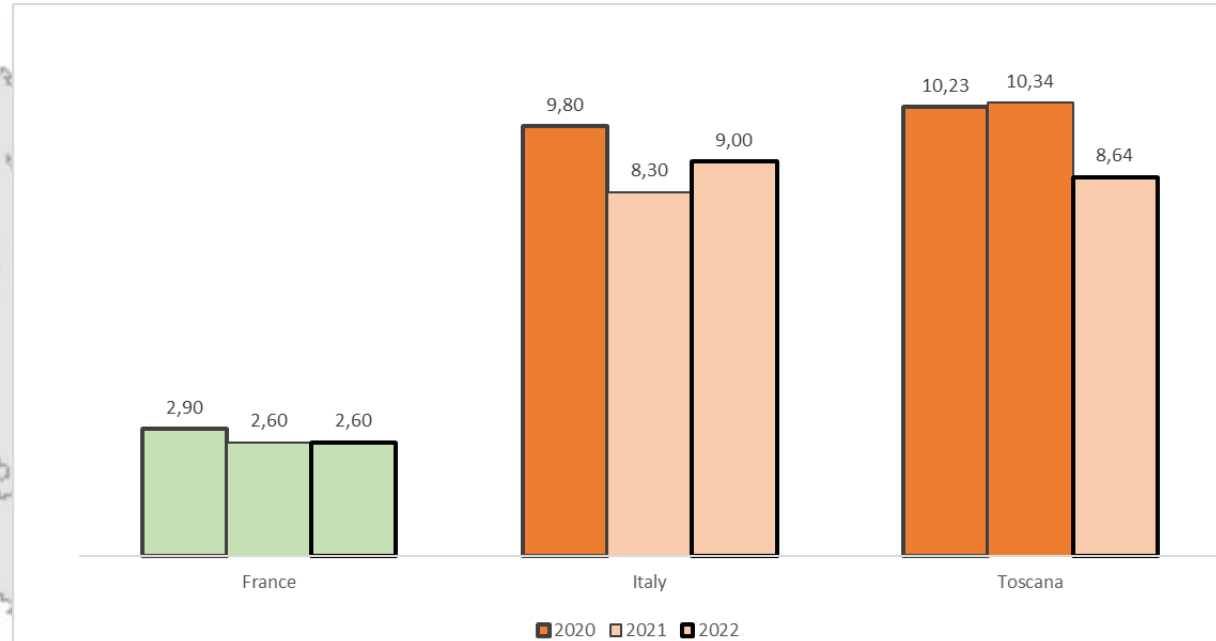
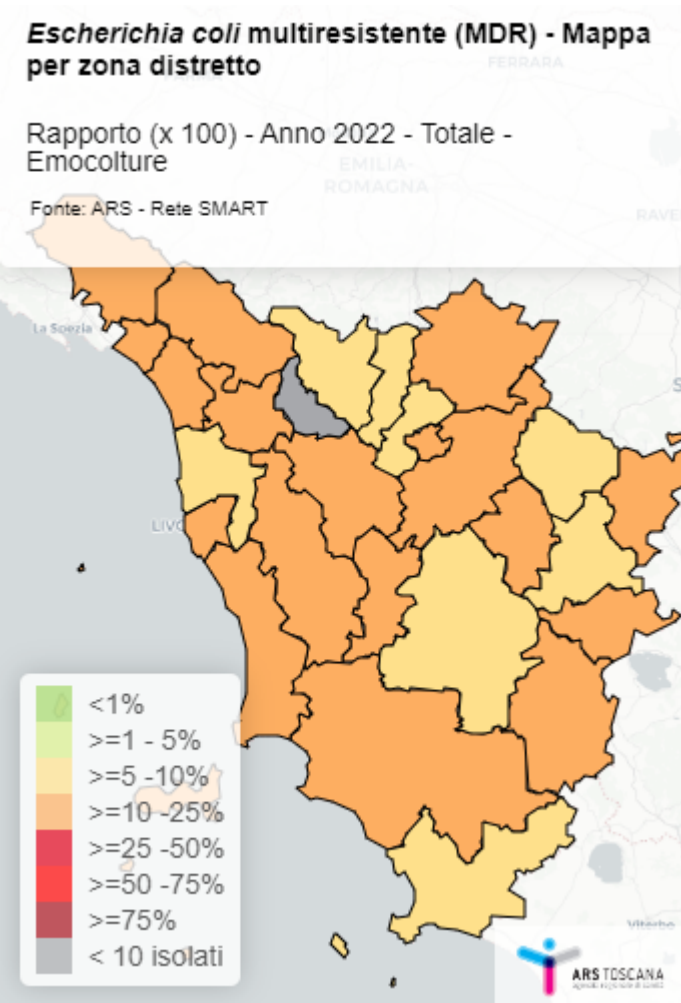
– sopravvivenza a 1 anno Toscana 2020-2022

Kaplan-Meier survival estimates



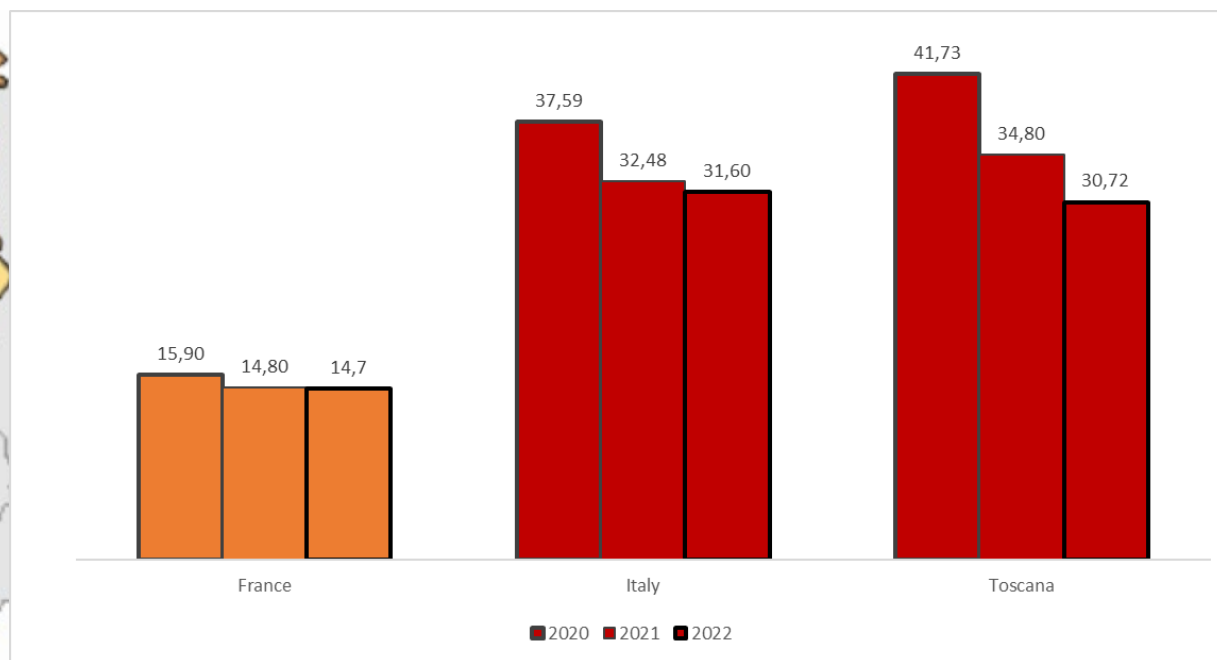
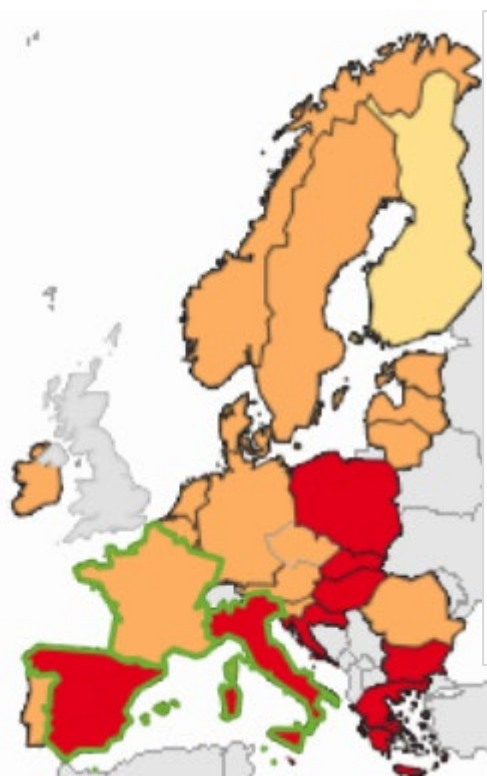
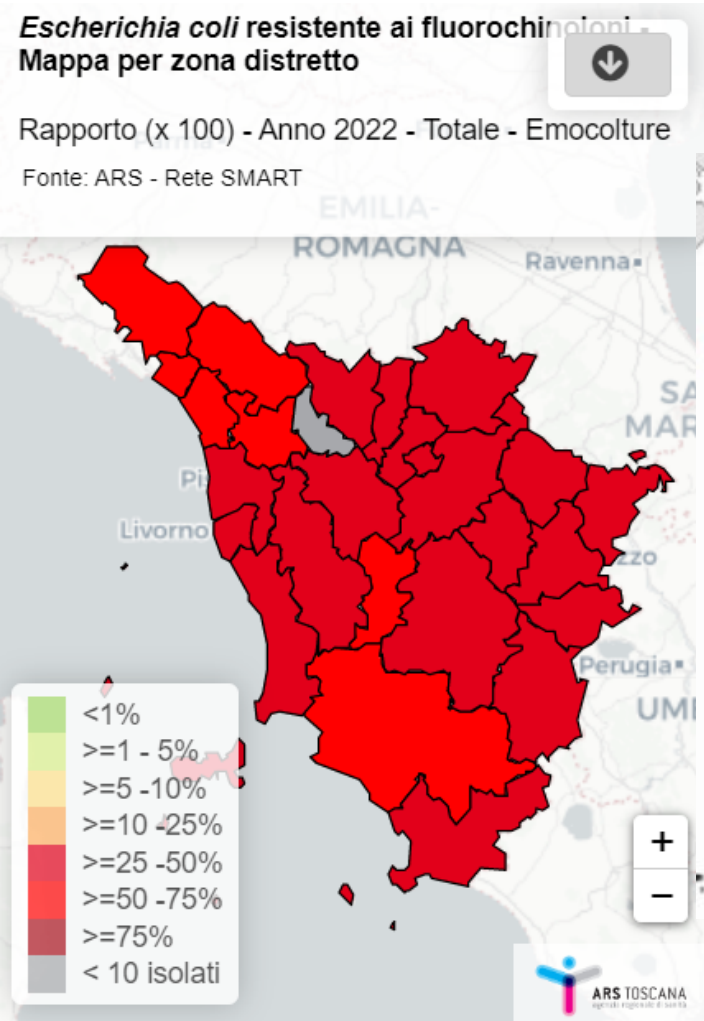
Escherichia coli, resistente ai multiresistente (cefalosporina di III generazione, fluorochinoloni e aminoglicosidi)

(2575 isolati di *E.coli* da sangue non ripetuti, 2022)



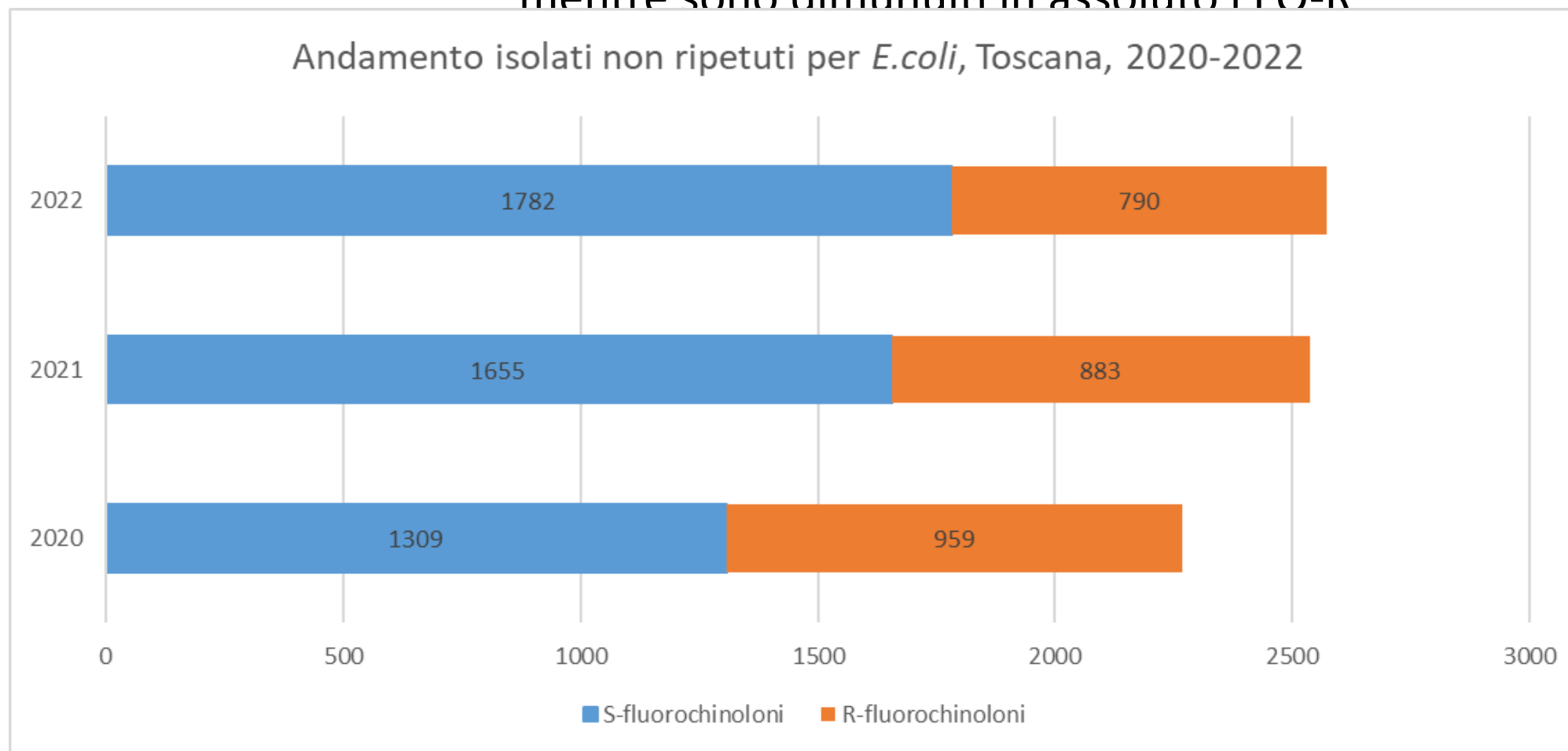
Escherichia coli, resistente ai fluorichinoloni

(2575 isolati di *E.coli* da sangue non ripetuti, 2022)



Escherichia coli, resistente ai fluorichinoloni

Il numero di *E.coli* isolati in Toscana nel 2022 è aumentato, in particolare i S-fluorochinoloni, mentre sono diminuiti in assoluto i FO-R

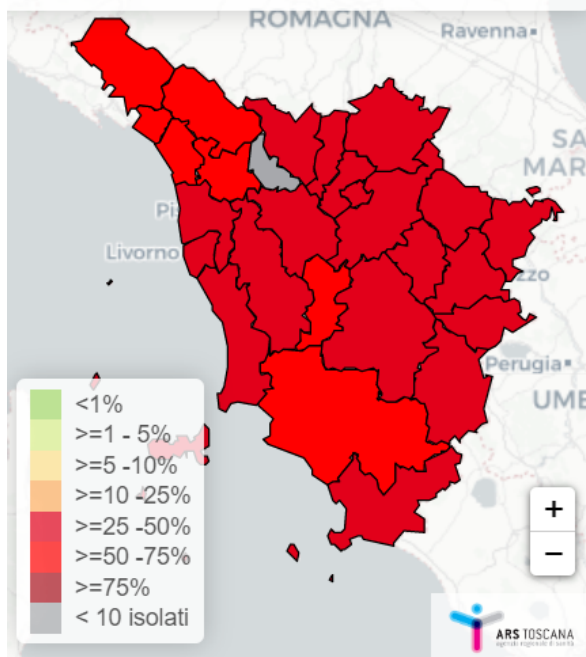


Escherichia coli, resistente ai fluorichinoloni

Escherichia coli resistente ai fluorochinoloni -
Mappa per zona distretto

Rapporto (x 100) - Anno 2022 - Totale - Emocolture

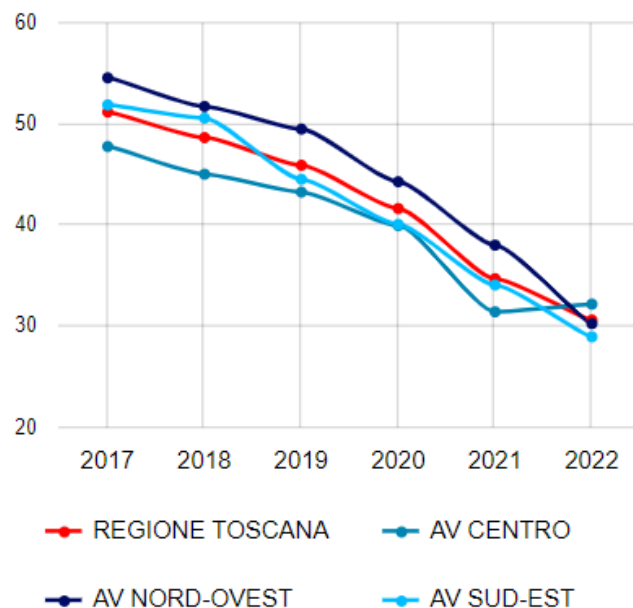
Fonte: ARS - Rete SMART



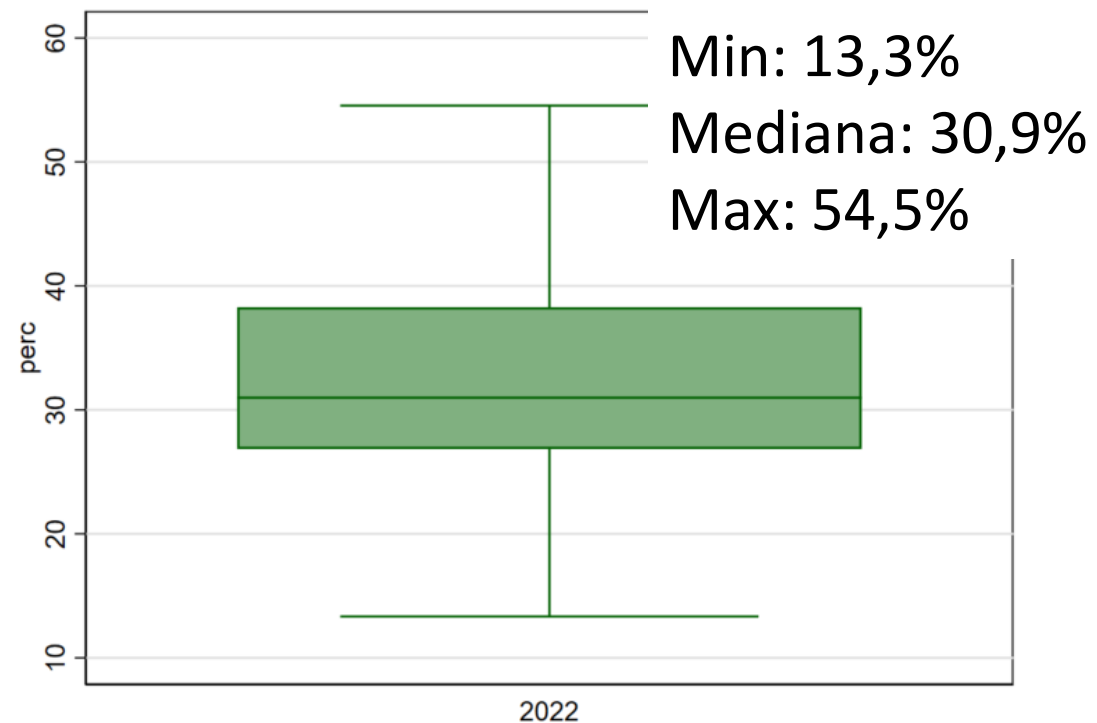
Escherichia coli resistente ai
fluorochinoloni - Andamento temporale

Rapporto (x 100) - Totale - Emocolture

Fonte: ARS - Rete SMART



Prevalenza *E.coli* fluorochinoloni R per ospedale, Toscana,
2020-2022 (32 H con 10 + isolati)



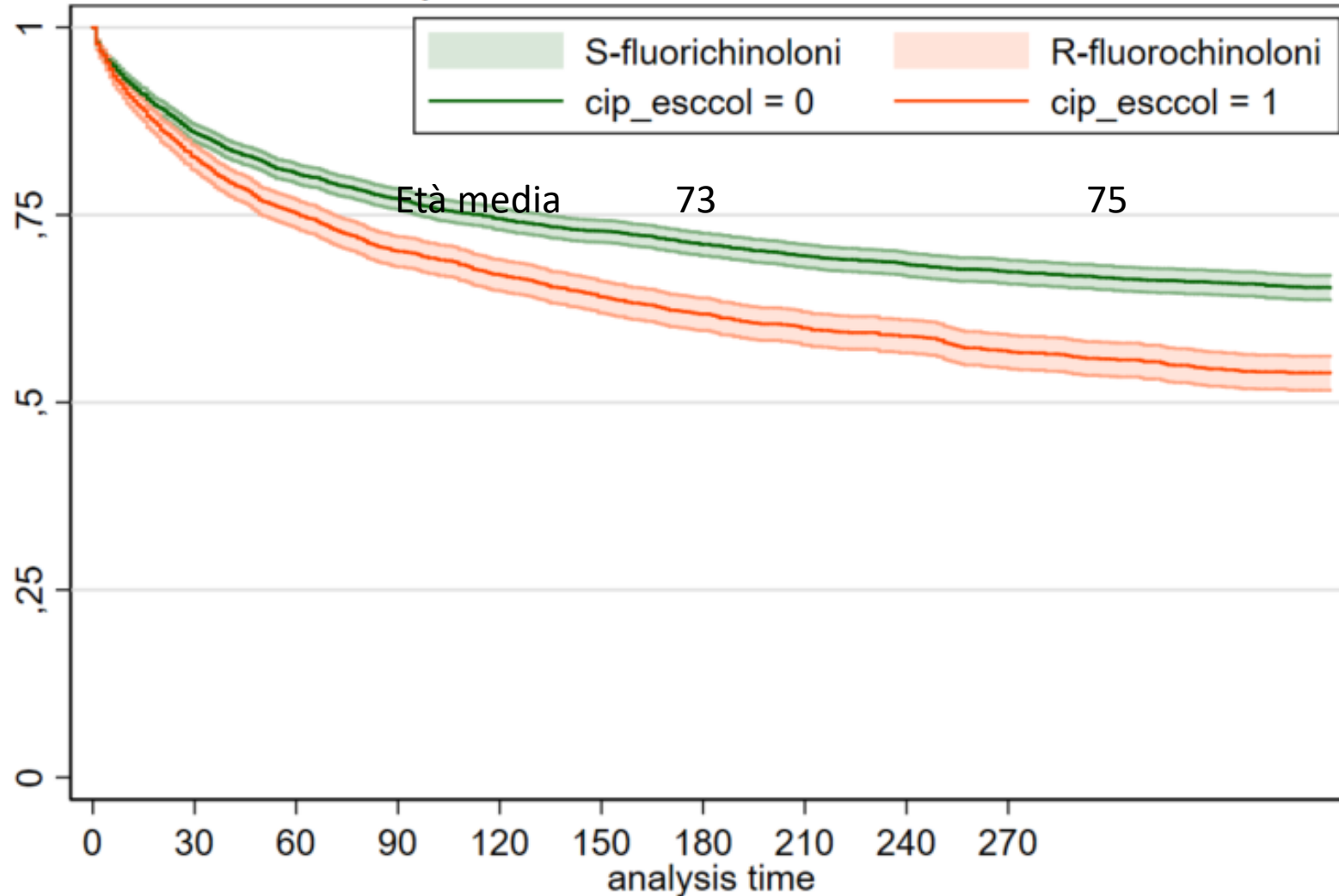
Median Odds Ratio (MOR) 1,12 (non sign)

Passando da un ospedale a minor prevalenza di *E.coli* resistenti a fluorochinoloni ad uno a maggior prevalenza, il rischio cresce in modo non significativo

Escherichia coli, resistente ai fluorichinoloni

– sopravvivenza a 1 anno Toscana 2020-2022

Kaplan-Meier survival estimates



grazie per l'attenzione

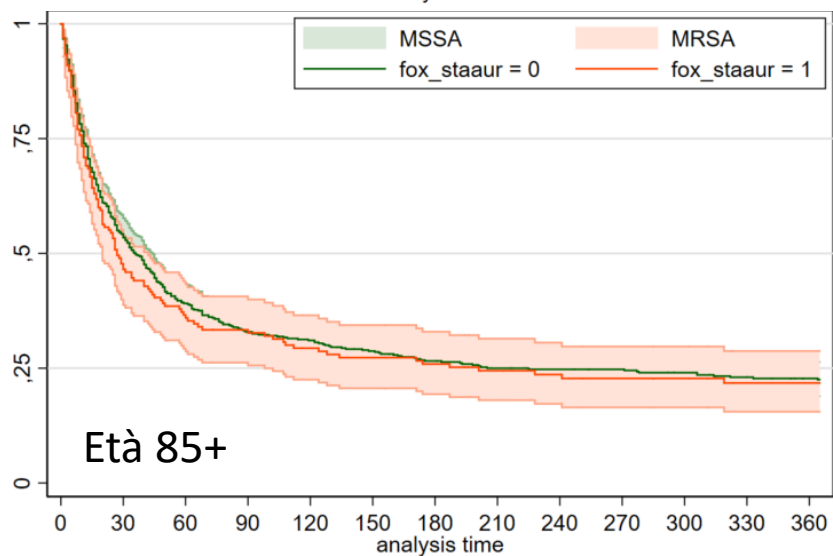
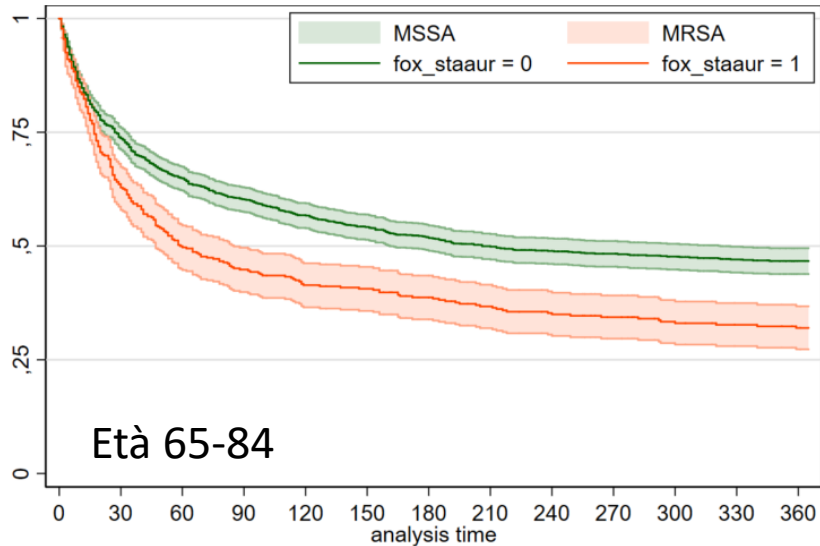
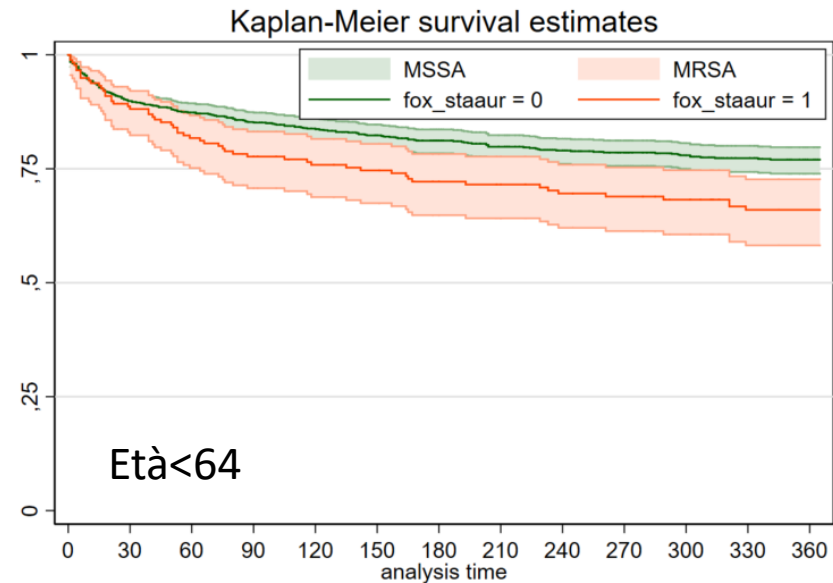
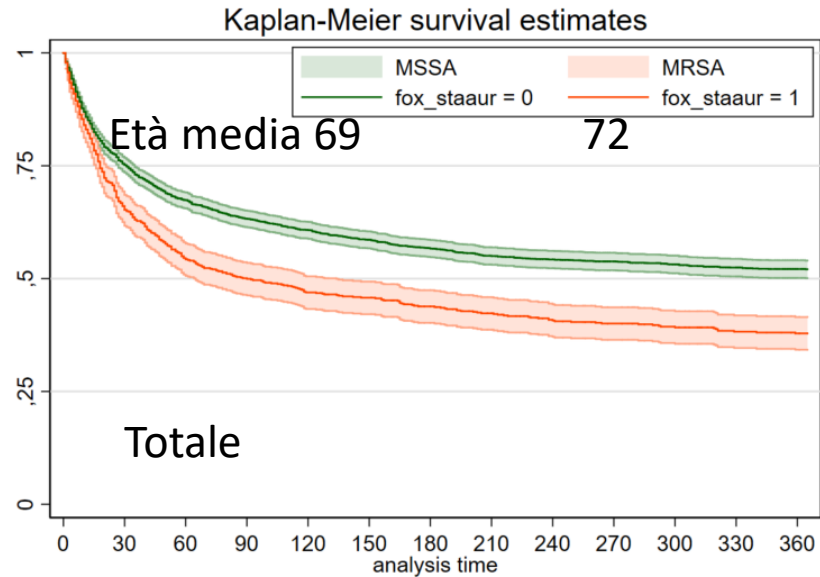
Profili di resistenza, Toscana, 2017-2022

Profili di Resistenza	2022	2021	2020	2019	2018	2017
	Toscana					
Acinetobacter spp. - carbapenemi	65.7%	72.0%	65.8%	63.9%	76.9%	78.0%
E. coli - multiresistente	8.6%	10,3%	10.2%	13.2%	15.6%	17.2%
E. coli - fluorchinoloni	30.7%	34,8%	41.7%	45.8%	48.0%	51.0%
E. coli – cefalosporine III gen.	21.0%	21.5%	28.3%	32.8%	37.0%	39.0%
K. pneumoniae - multiresistente	39.4%	42.7%	38.2%	40.1%	33.0%	38.9%
K. pneumoniae - fluoroquinoloni	57.0%	57.1%	58.1%	63.1%	61.9%	65.4%
K. pneumoniae – cefalosporine III gen.	61.4%	58.9%	58.7%	64.2%	61.5%	63.9%
K. pneumoniae - carbapenemi	18.0%	19.2%	27.2%	33.0%	29.1%	35.5%
P. aeruginosa - multiresistente	9.4%	10.0%	7.3%	8.6%	12.5%	17.5%
P. aeruginosa - carbapenemi	14.0%	13.1%	6.0%	5.7%	8.7%	14.7%
S. aureus - MRSA	22.5%	21.6%	27.5%	26.0%	30.8%	30.8%
E. faecium - Vancomicina	29.7%	16.7%	10.5%	15.7%	23.1%	24.5%

Andamento
17-22



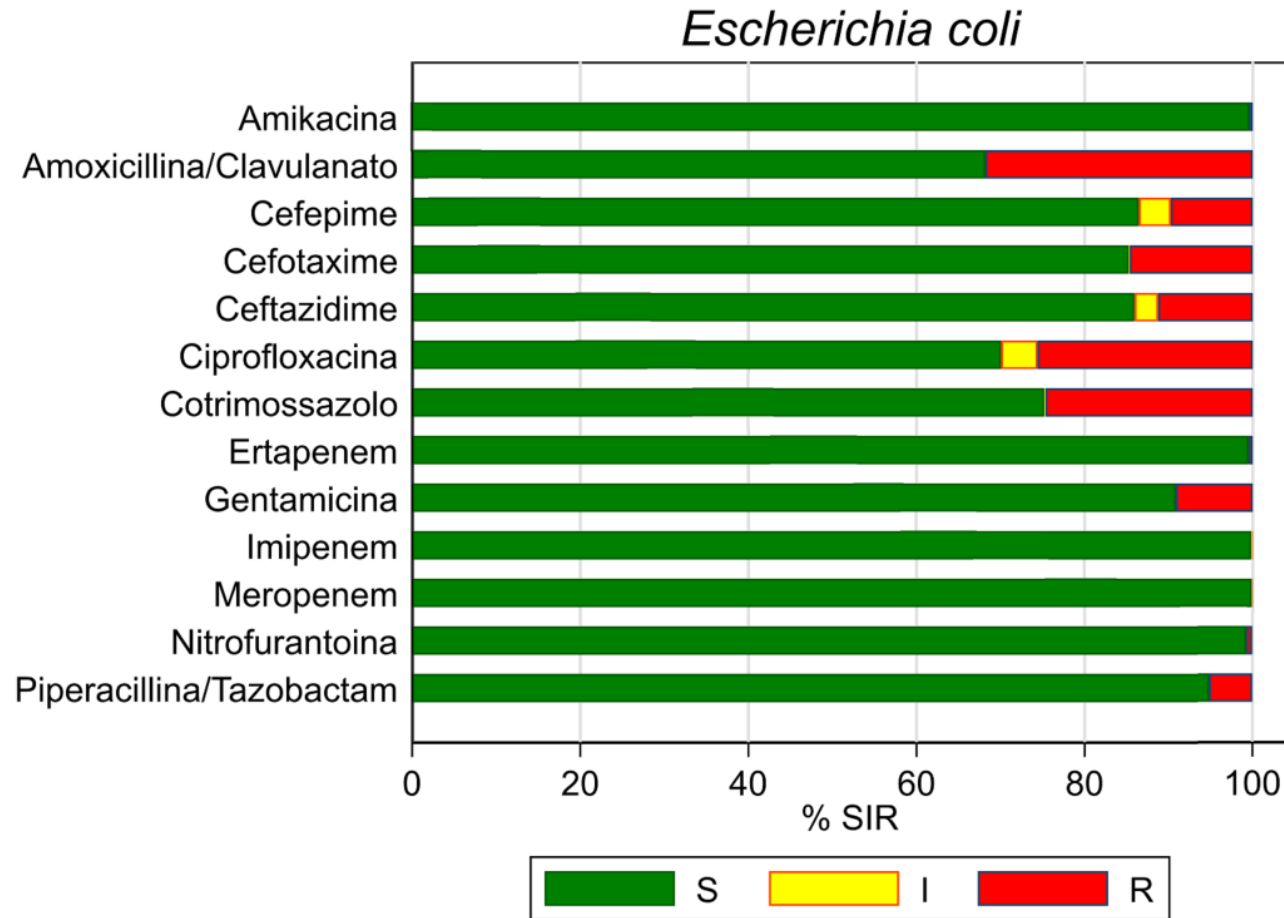
Staphylococcus aureus meticillino-resistente (MRSA), 2020-2022



Classe di età	MSSA	MRSA
<64	36%	27%
65-84	46%	52%
85+	18%	21%

Escherichia coli, resistente ai fluorichinoloni

Urinocolture



R-ciprofloxacina:

2021: 29,4% (44.840)

2022: 25,6% (44.913)