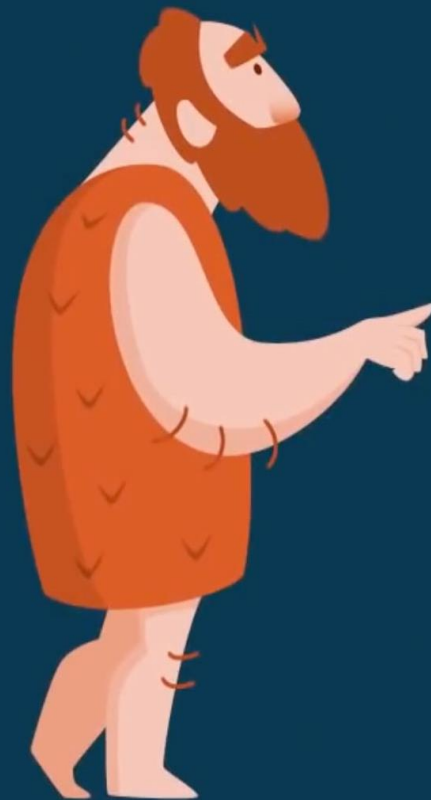




Impact of antimicrobial resistance on poultry gut health and metabolism

Speaker: mzaghari@ut.ac.ir
Available at www.minatoyoor.com



BACTERIA

BEETHOVEN

Outline of presentation

- Antibiotics:

- Mode of action and mechanisms of resistance

- Genetic basis of antimicrobial resistance

- Applied approach (disc diffusion, PCR, . . . , metagenomics)

- How does resistance spread?

- Human:

- GLASS (global antimicrobial resistance surveillance system)

- Antibiotic use, resistance and the link to human nutrition

- Poultry:

- Antibiotic application in poultry production

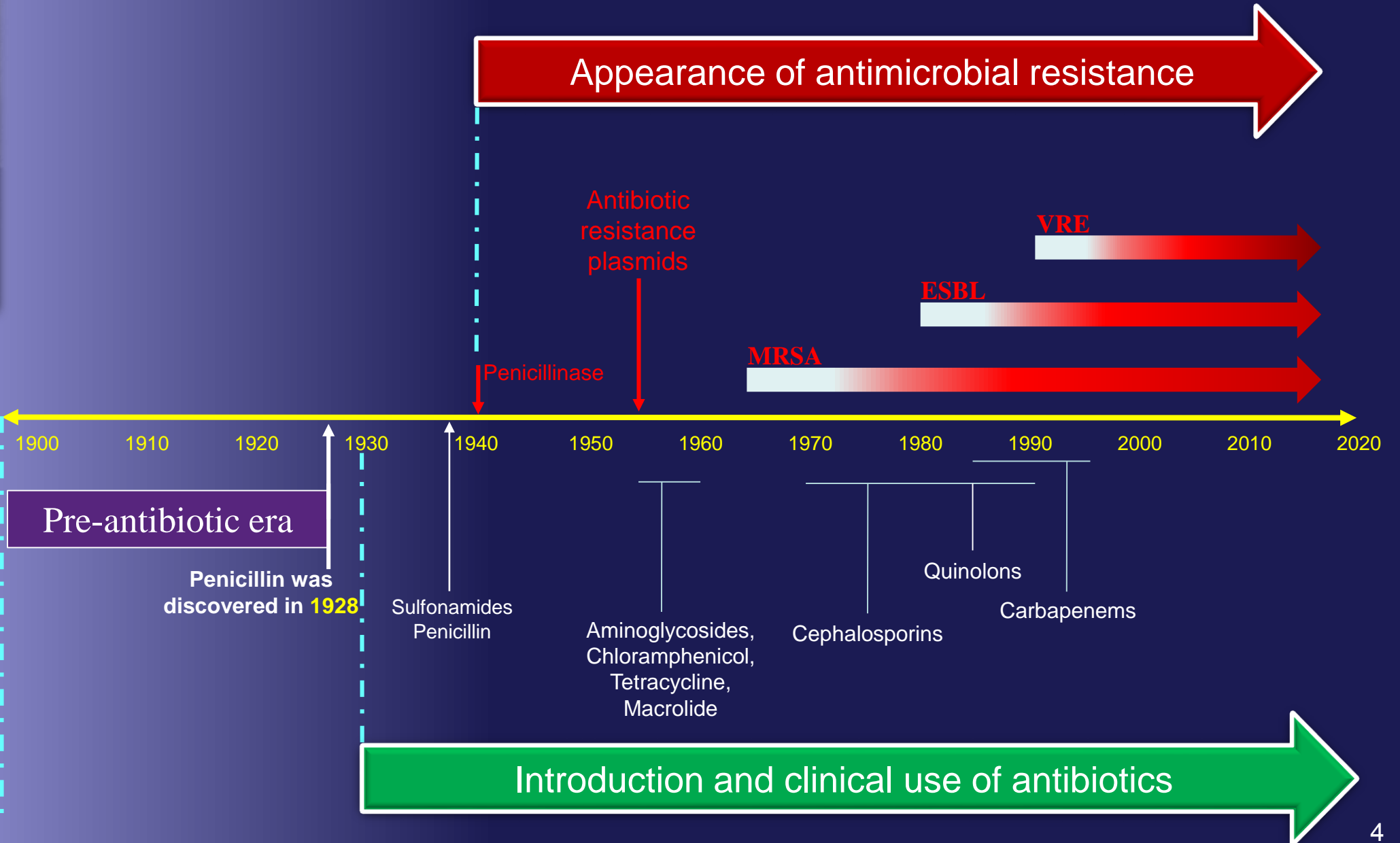
- Antibiotic resistance of some selected organisms in poultry

- Antibiotic use, resistance and the link to poultry nutrition



Paul Ehrlich

Natural dyes
Heavy metal





Antimicrobial agents (mechanism)

1

Inhibition of the cell wall synthesis

2

Inhibition of the nucleic acid synthesis

3

Inhibition of the ribosome function

4

Inhibition of the cell membrane function

5

Inhibition of the folate metabolism

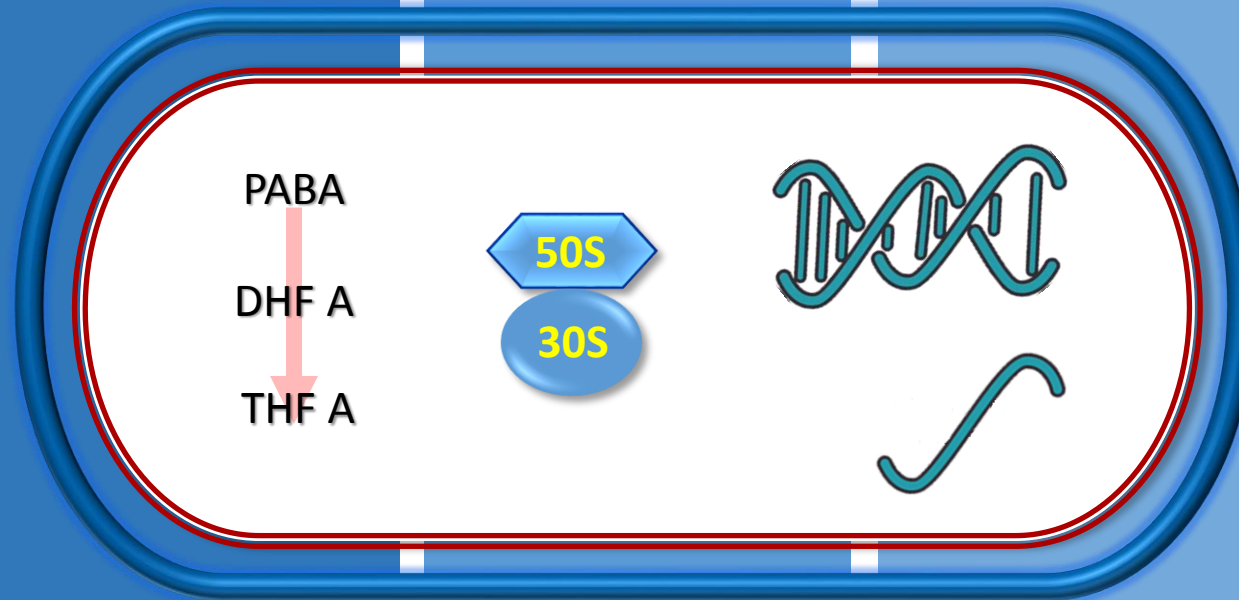
Folic acid inhibitors

Cell membrane

Protein synthesis

DNA, RNA processing

Cell wall synthesis



Folic acid inhibitors

Cell membrane

Protein synthesis

DNA, RNA processing

Cell wall synthesis

PABA
↓
DHF A
↓
THF A

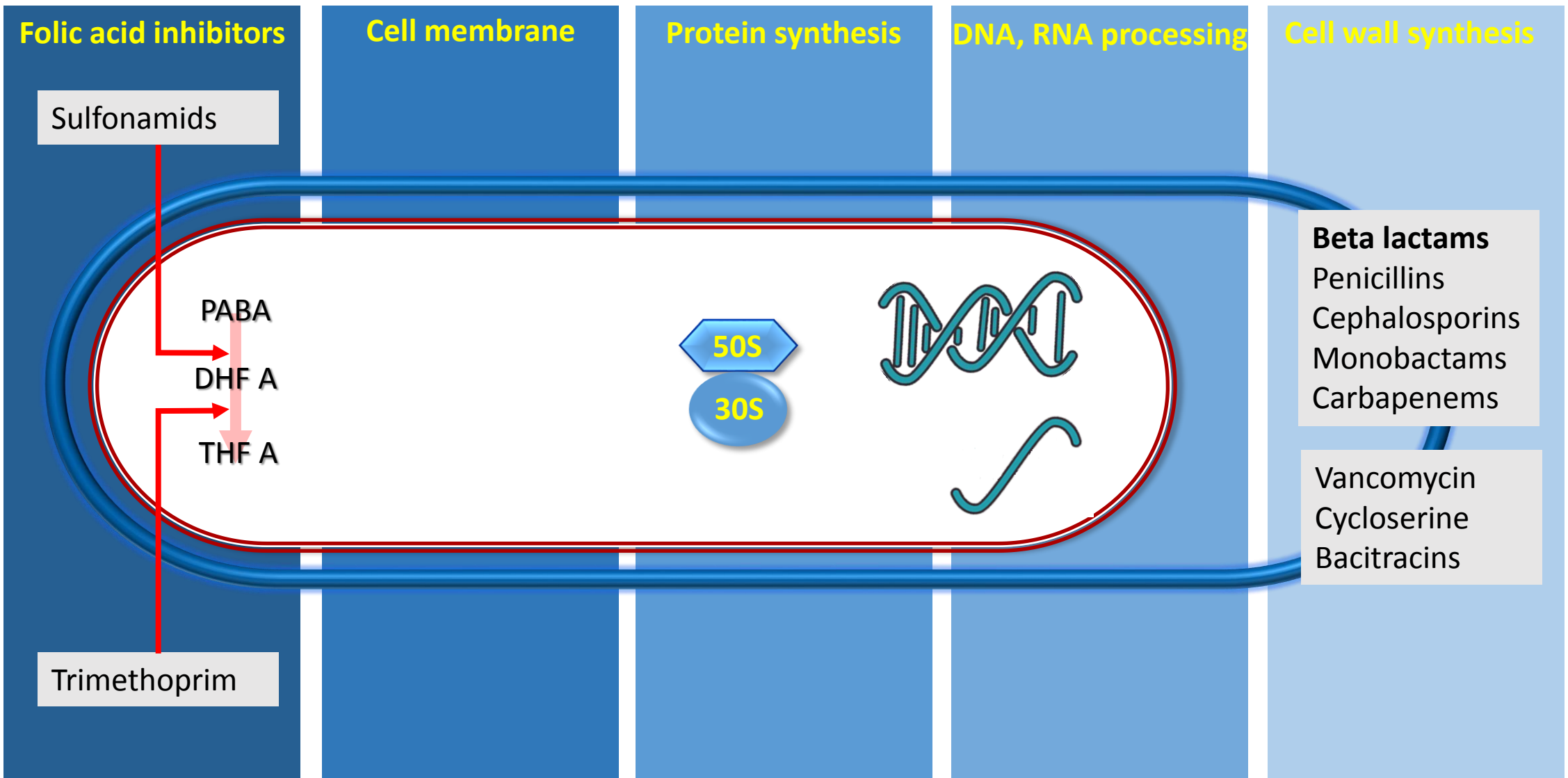
50S
30S

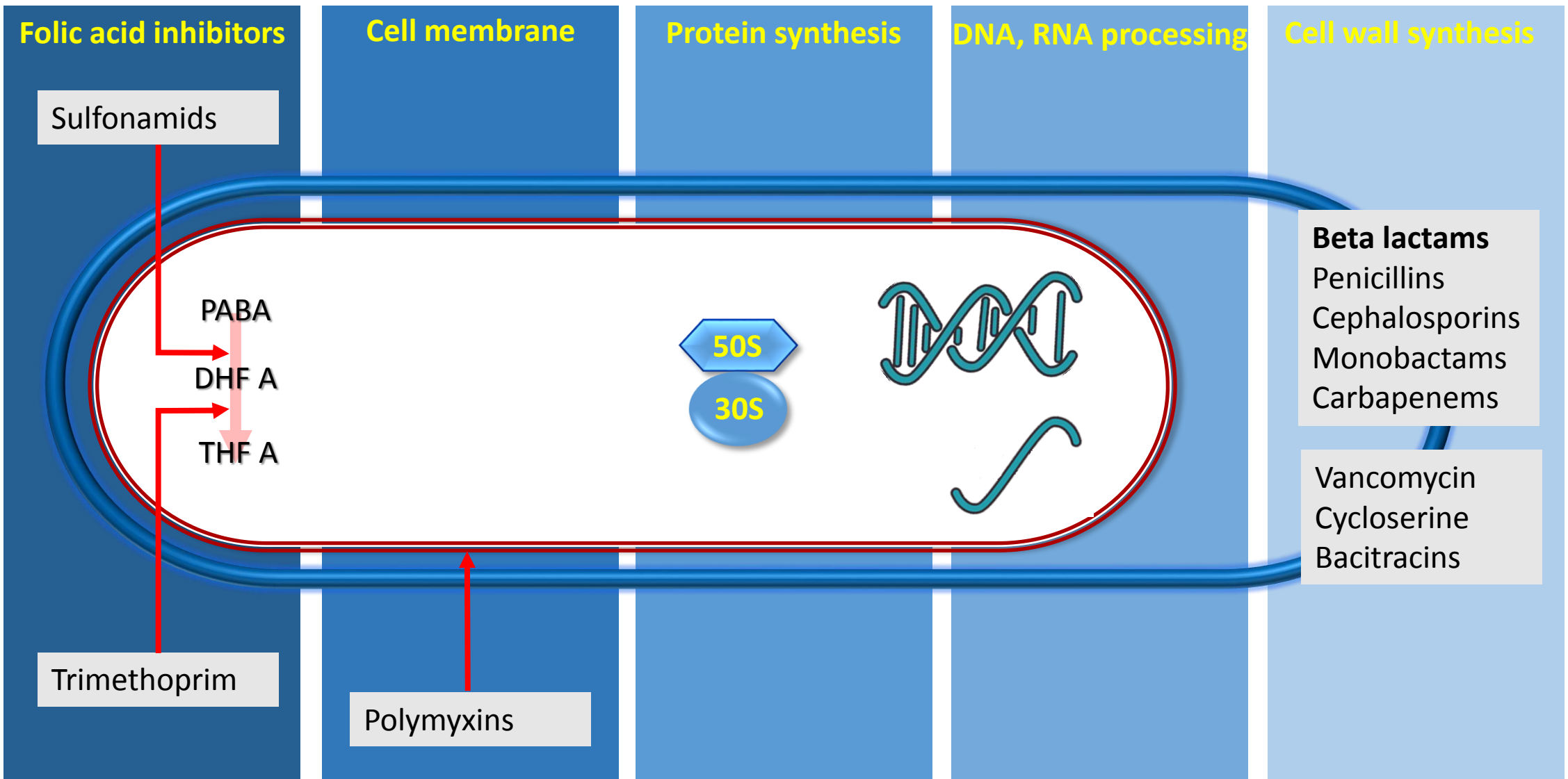


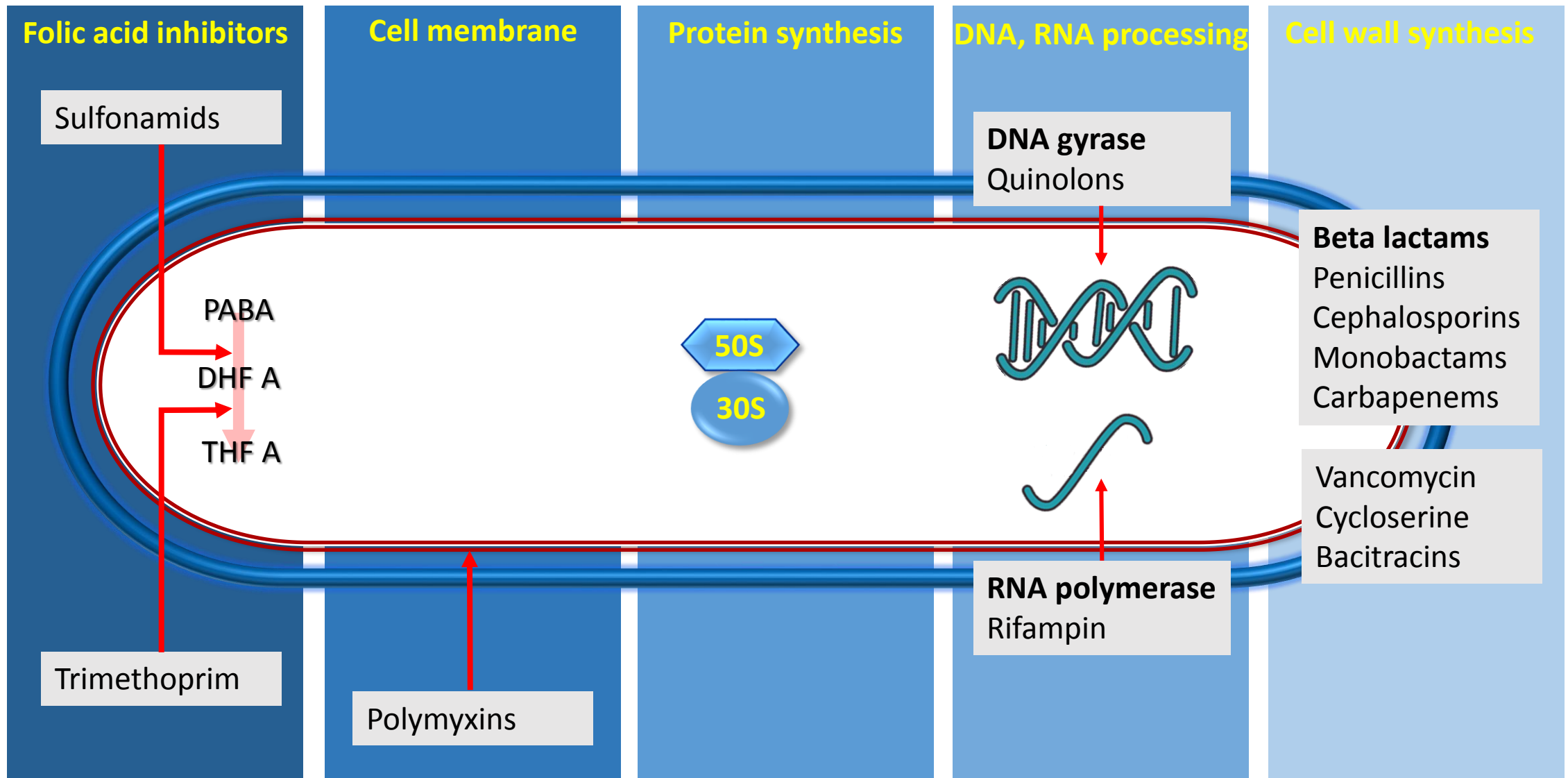
Beta lactams

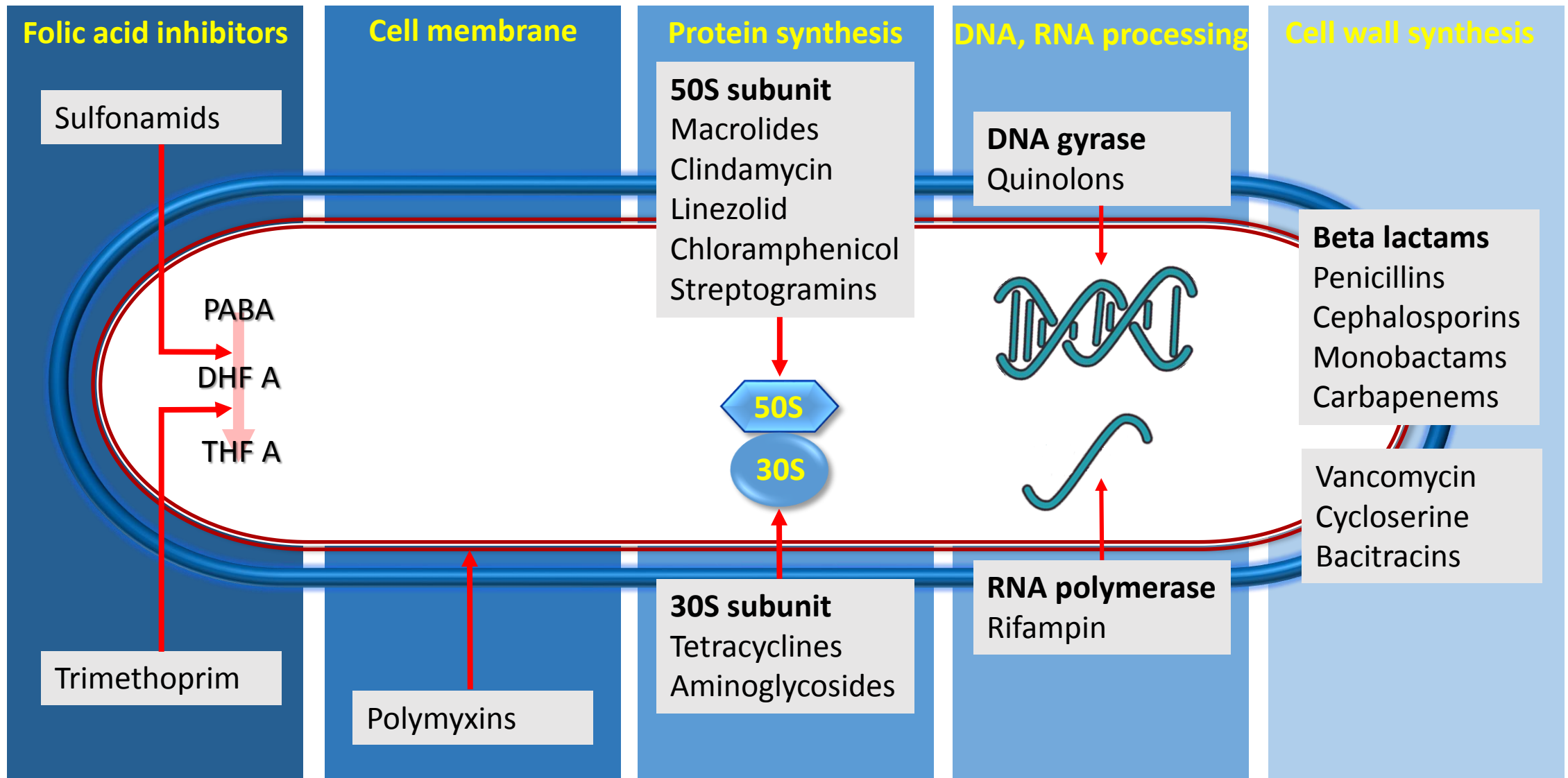
Penicillins
Cephalosporins
Monobactams
Carbapenems

Vancomycin
Cycloserine
Bacitracins

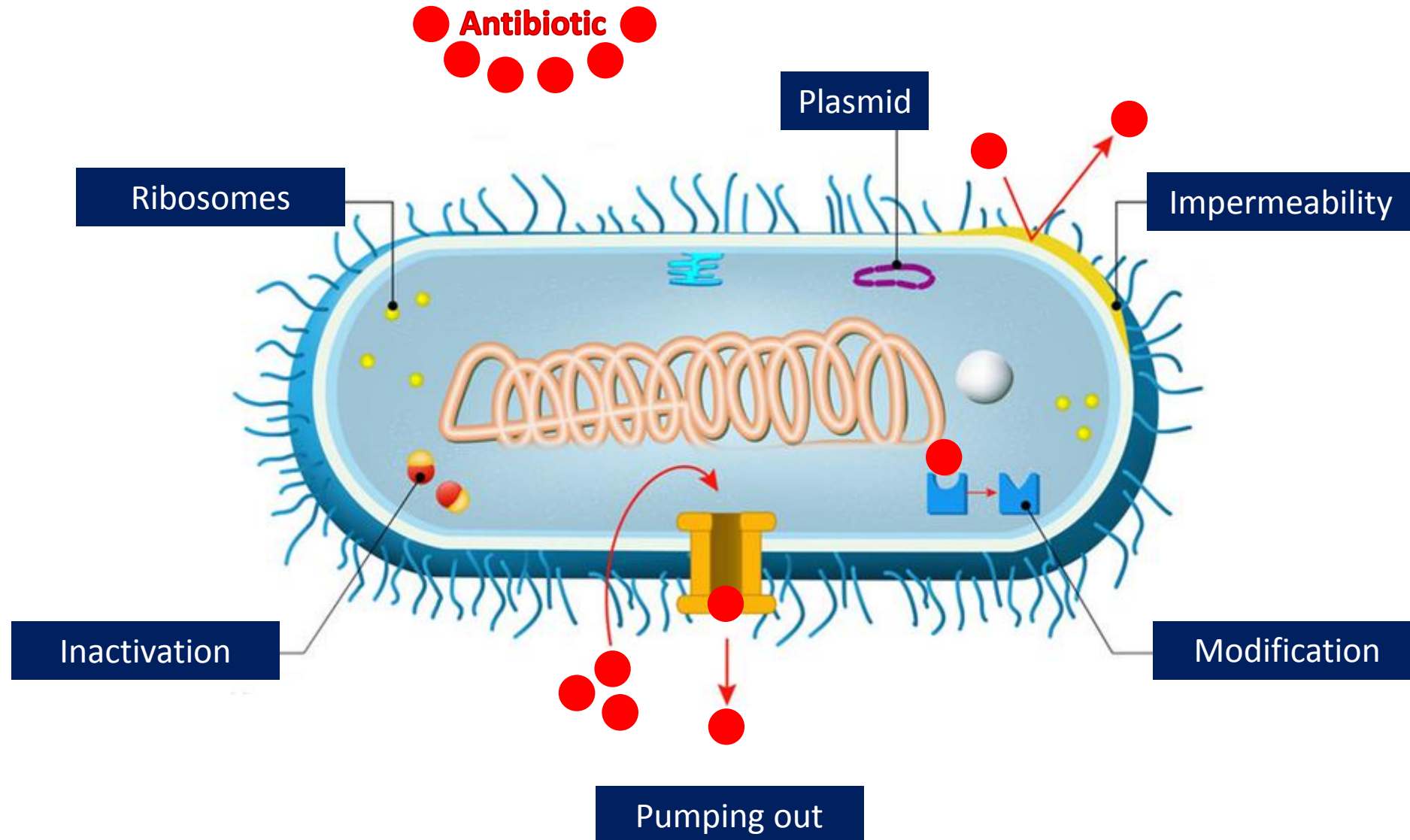








Mechanism of action of **AMR**



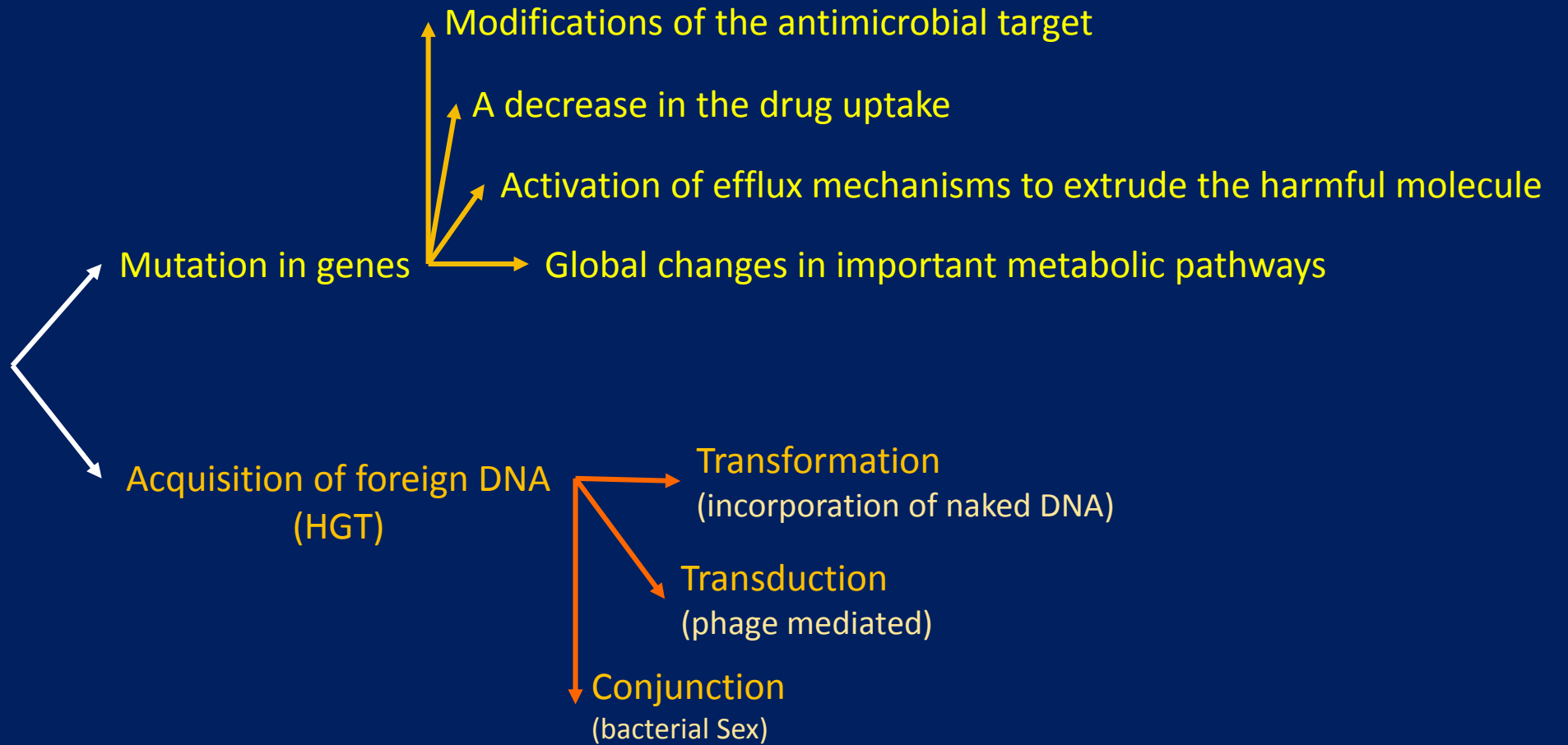
Resistance mechanism of individual antibiotics

Antibiotic class	Resistance type	Resistance mechanism	Common example
Aminoglycoside	Decreased uptake	Changes in outer membrane	<i>P. aeruginosa</i>
	Enzymatic modification	AGE's	Gram-negative bacteria
Beta-lactams	Altered PBP	PBP 2a	Mec A in <i>S. aureus</i> , CONS, <i>S. pneumoniae</i>
	Enzymatic degradation	Penicillinase which are classified as per ambler classification	Gram-negative bacteria
Glycopeptides	Altered target	D-alanyl-alanine is changed to D-alanyl-D-lactate	Vancomycin resistance in <i>E. faecium</i> and <i>E. faecalis</i>
Macrolides	Altered target	Methylation of ribosomal active site with reduced binding	<i>erm</i> -encoded methylases in <i>S. aureus</i> , <i>S. pneumoniae</i> , and <i>S. pyogenes</i>
	Efflux pumps	Mef type pump	<i>S. pneumoniae</i> and <i>S. pyogenes</i>
Oxazolidinones	Altered target	Mutation leading to reduced binding to active site	<i>E. faecium</i> and <i>S. aureus</i>
Quinolones	Altered target	Mutation leading to reduced binding to active site(s)	Mutations in gyr A in enteric Gram-negative bacteria and <i>S. aureus</i>
	Efflux	Membrane transporters	Mutations in gyr A and par C in <i>S. pneumoniae</i> . Nor-A in <i>S. aureus</i>
Tetracyclines	Efflux	New membrane transporters	<i>tet</i> genes encoding efflux proteins in Gram-positive and Gram-negative bacteria
	Altered target	Production of proteins that bind to the ribosome and alter the conformation of the active site	<i>tet</i> (M) and <i>tet</i> (O) in Gram-positive and Gram-negative bacteria species
Chloramphenicol	Antibiotic inactivation	Chloramphenicol acetyl transferase	CAT in <i>S. pneumonia</i>
	Efflux pump	New membrane transporters	<i>cml</i> A gene and <i>flo</i> gene efflux in <i>E. coli</i>
Sulfa drugs	Altered target	Mutation of genes encoding DHPS	<i>E. coli</i> , <i>S. aureus</i> , <i>S. pneumoniae</i>

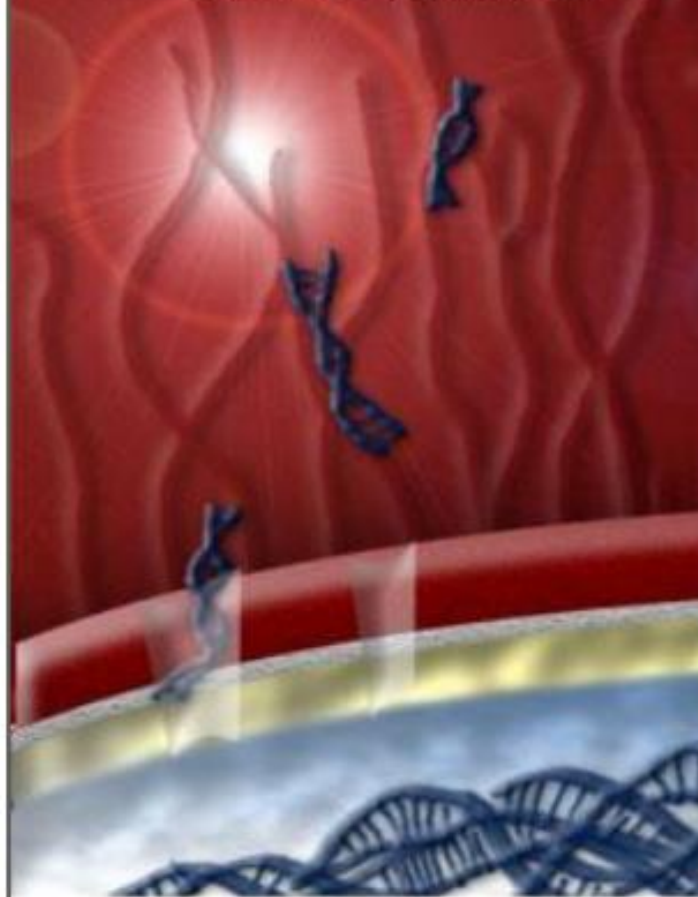
DHPS=Dihydropteroate synthase, *P. aeruginosa*=*Pseudomonas aeruginosa*, *S. aureus*=*Staphylococcus aureus*, *S. pneumoniae*=*Streptococcus pneumoniae*, *E. faecium*=*Enterococcus faecium*, *E. faecalis*=*Enterococcus faecalis*, *S. pyogenes*=*Streptococcus pyogenes*, *E. coli*=*Escherichia coli*, PBP=Penicillin binding protein, AGE's=Aminoglycoside modifying enzymes, CAT=Chloramphenicol acetyl transferases



GENETIC BASIS of AMR



TRANSFORMATION



Transformation involves uptake of short fragments of naked DNA by naturally transformable bacteria.

TRANSDUCTION



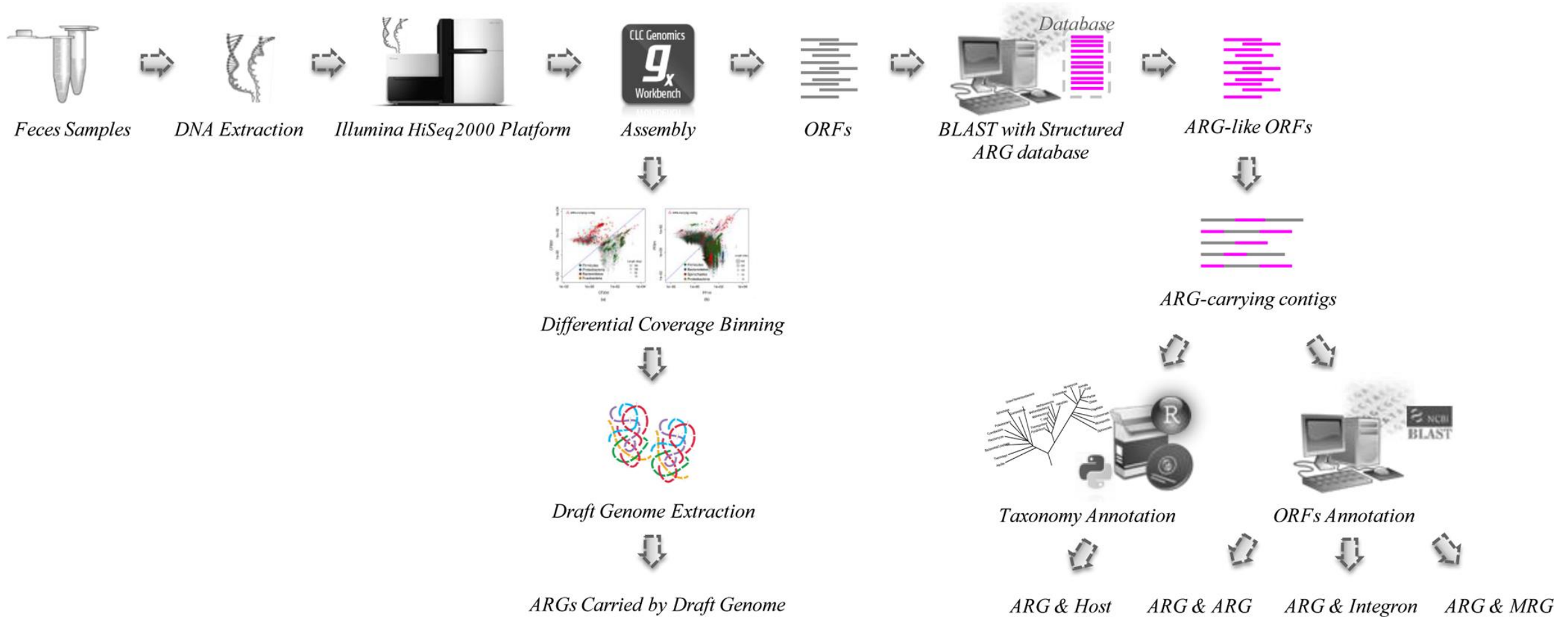
Transduction involves transfer of DNA from one bacterium into another via bacteriophages

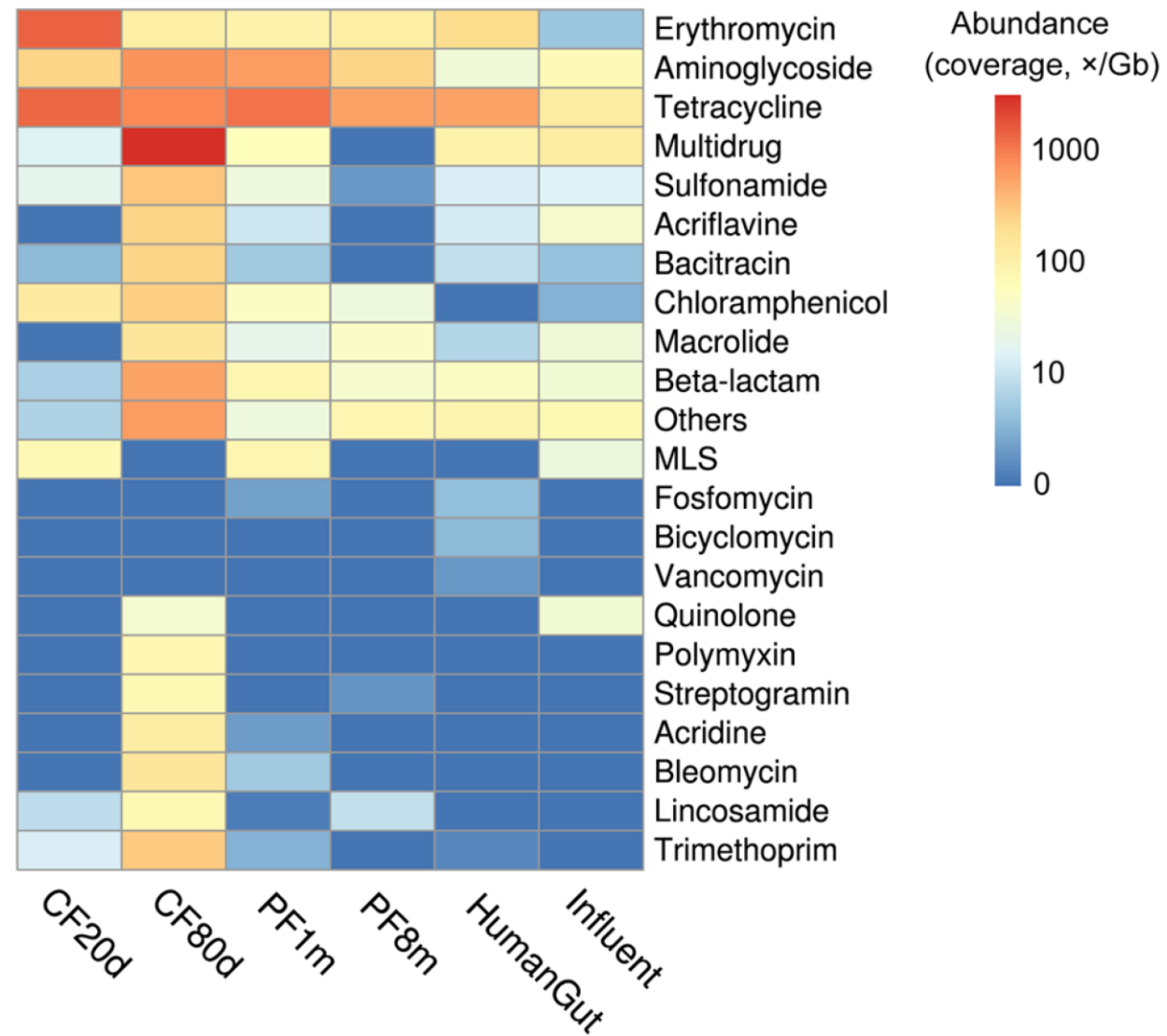
CONJUGATION

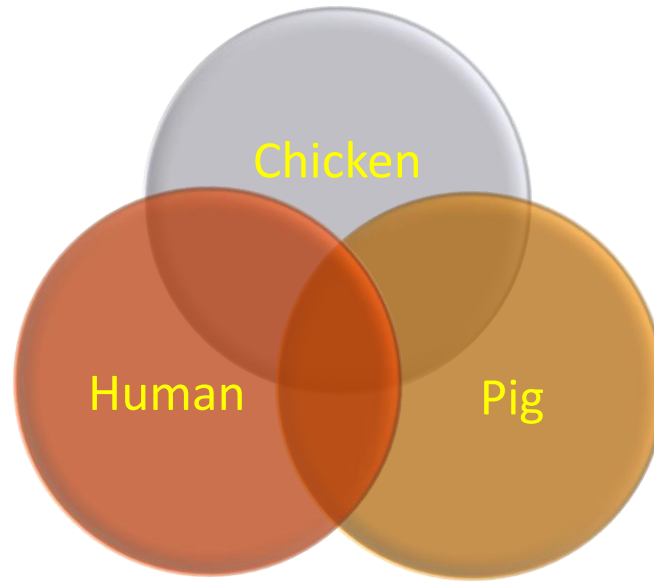


Conjugation involves transfer of DNA material via sexual pilus and requires cell-to-cell contact

Metagenomic Assembly Reveals Hosts of Antibiotic Resistance Genes and the Shared Resistome in Pig, Chicken, and Human Feces

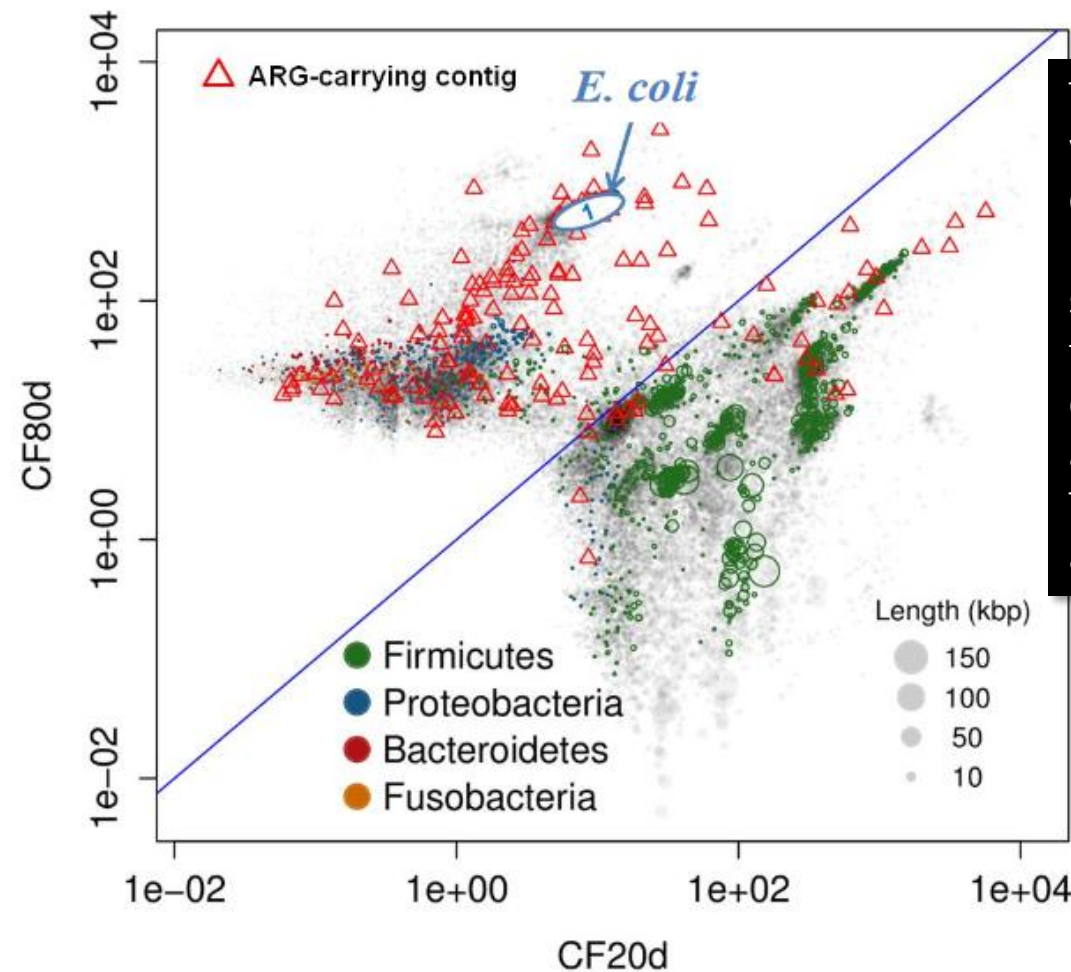






A comparison of metagenomes from feces of chickens, pigs and humans showed a high homology to **tetracycline genes** (tetA) and the presence of gene combinations of individual resistance elements, which encode for resistance to **beta-lactams**, **aminoglycosides**, **macrolides** and multidrug.

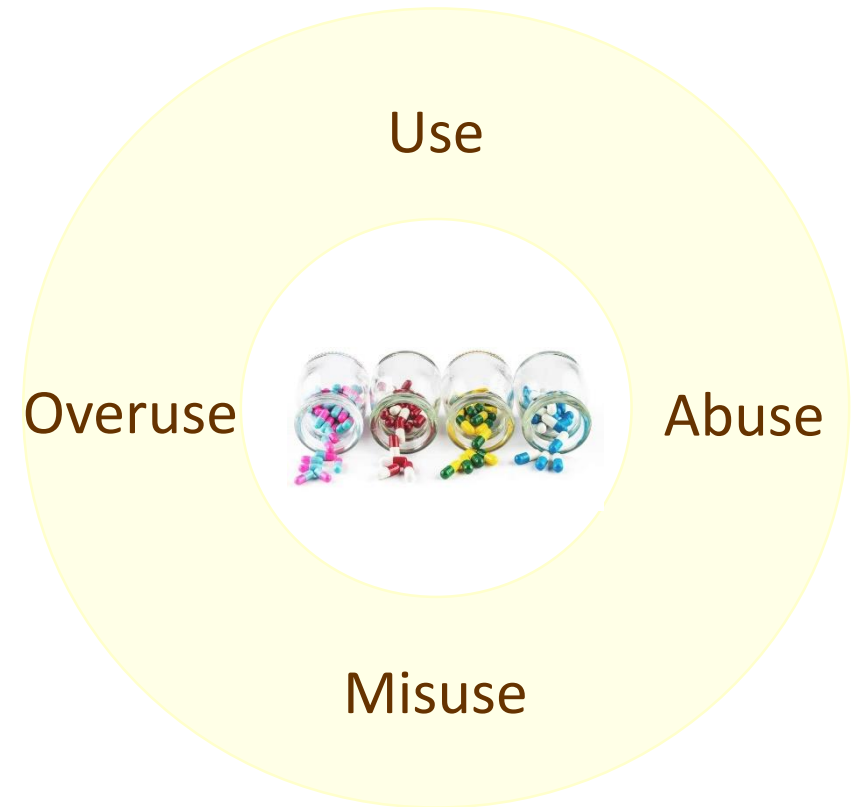
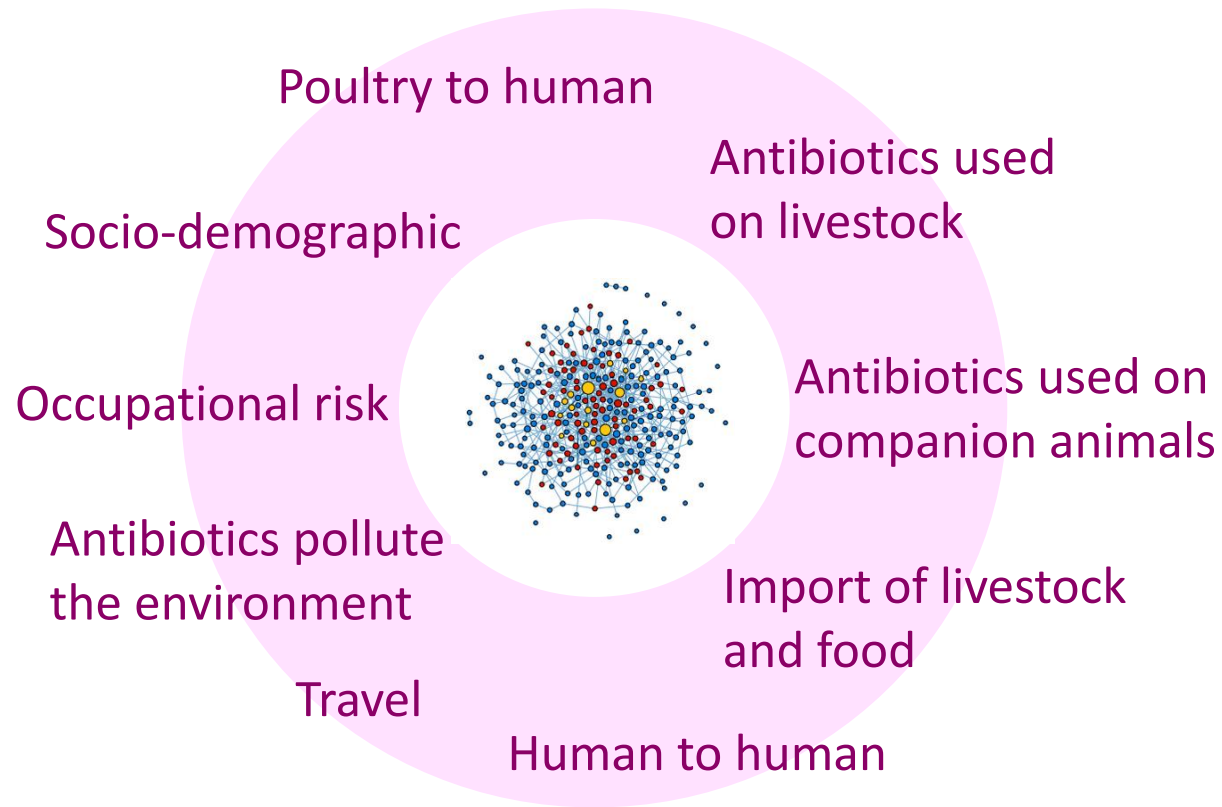
Distribution of ARG-carrying contigs in chicken metagenome data sets via differential coverage-based binning. All nodes represent scaffolds, scaled by the square root of their length, and colored by community taxonomy.

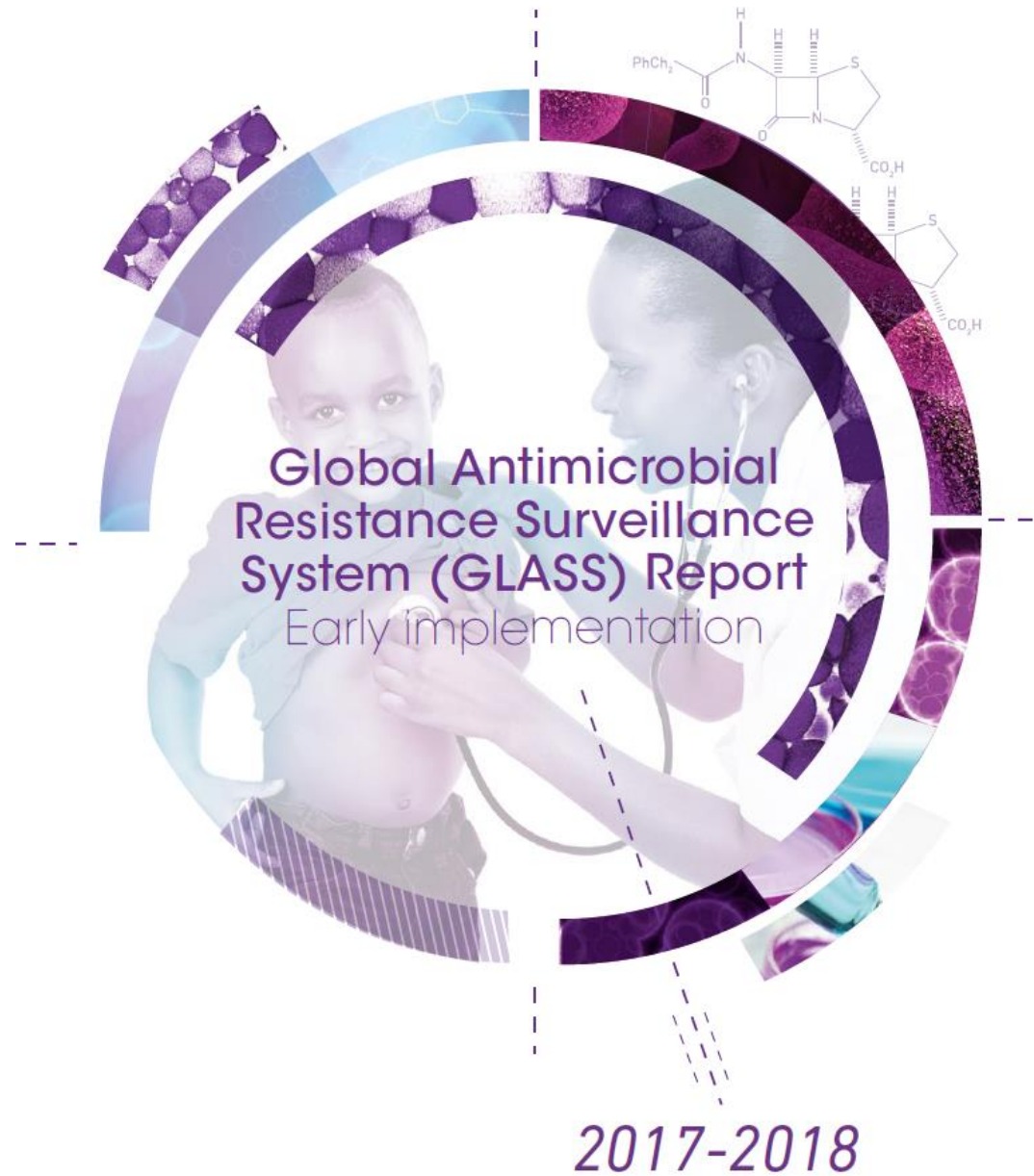


The abundance of this *E. coli* bin was increased 77-fold in CF80d compared to CF20d. Functional analysis of this *E. coli* strain using MEGAN KEGG revealed that the strain might cause human disease and could possess antibiotic and metal resistance. Therefore, this *E. coli* strain is likely a high-risk human pathogen.

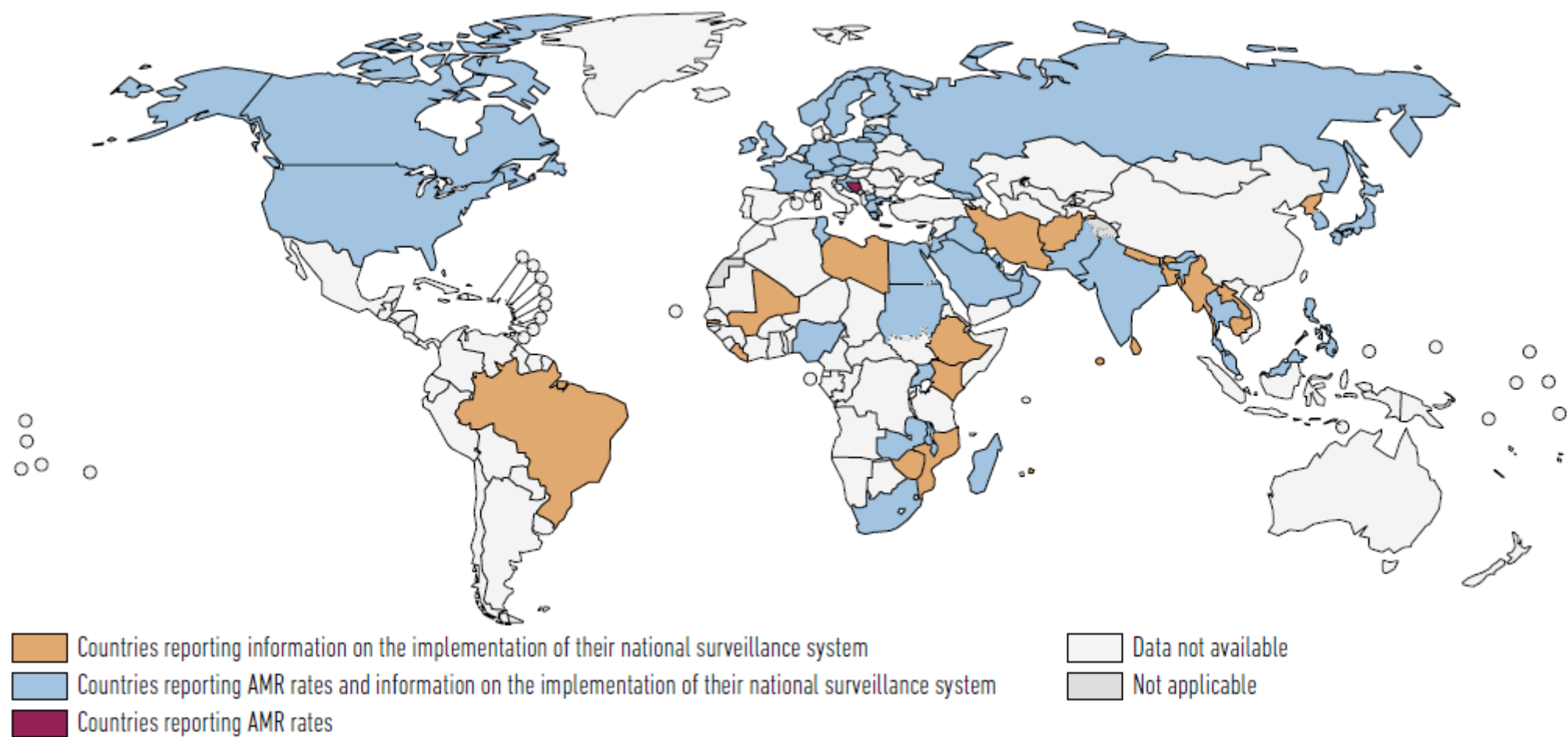


How does resistance spread and develops?







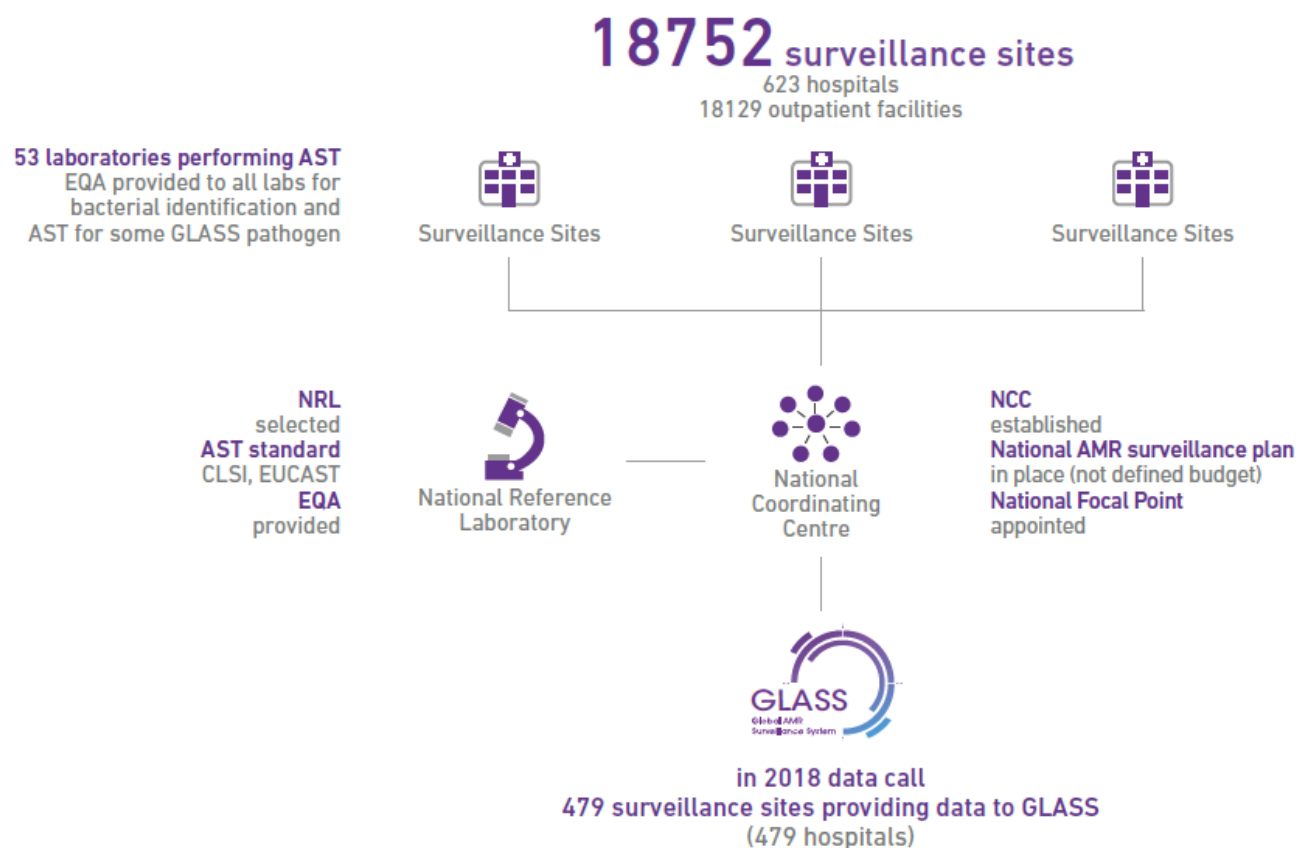


Germany

Population 82.11 million

The national surveillance of AMR is coordinated by the Robert Koch Institute, offering a publicly accessible interactive database for data of the AMR surveillance system (Antibiotika Resistenz Surveillance – ARS). The National action plan on prevention of AMR (DART 2020) was published in 2015. Germany participates in the EARS-NET and has been enrolled in GLASS since September 2016.

Current status of the national AMR surveillance system



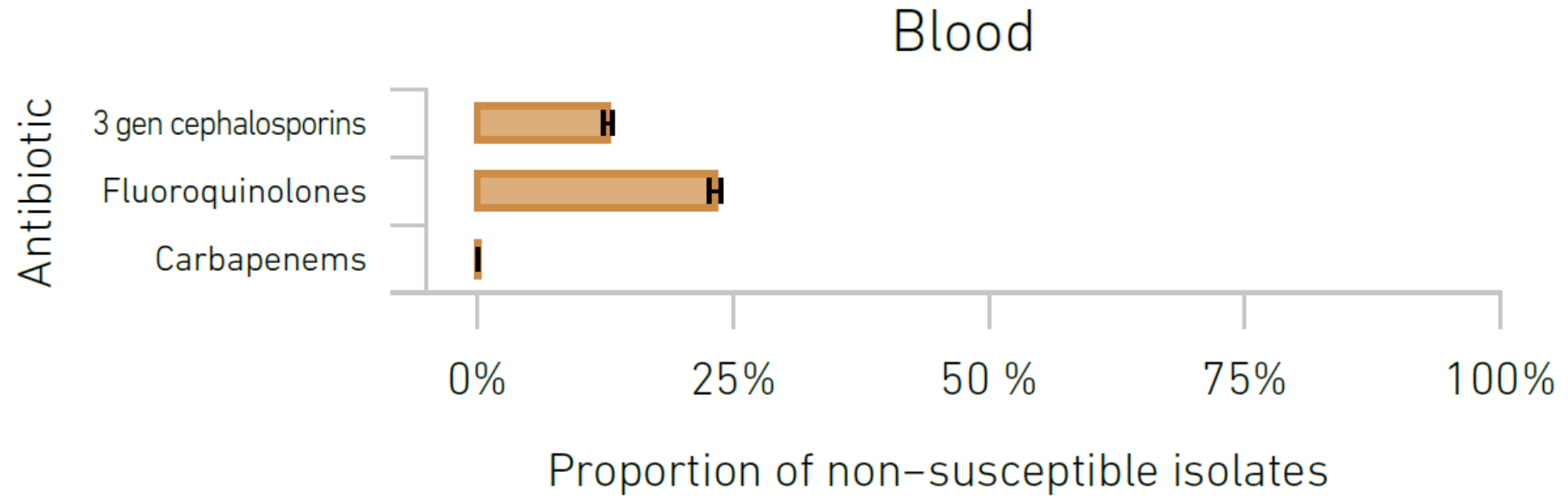
Data submission

Specimen type	Data on number of tested patient	Pathogen	AST results	Age	Gender	Infection origin
Blood	●	<i>Acinetobacter</i> spp.	●	●	●	●
		<i>E. coli</i>	●	●	●	●
		<i>K. pneumoniae</i>	●	●	●	●
		<i>Salmonella</i> spp.	●	●	●	●
		<i>S. aureus</i>	●	●	●	●
		<i>S. pneumoniae</i>	●	●	●	●
Urine	●	<i>E. coli</i>	●	●	●	●
		<i>K. pneumoniae</i>	●	●	●	●
Stool	●	<i>Salmonella</i> spp.	●	●	●	●
		<i>Shigella</i> spp.	●	●	●	●
Genital	●	<i>N. gonorrhoeae</i>	●	●	●	●

● 100% data collected ● 99-70% data collected ● <70% data collected

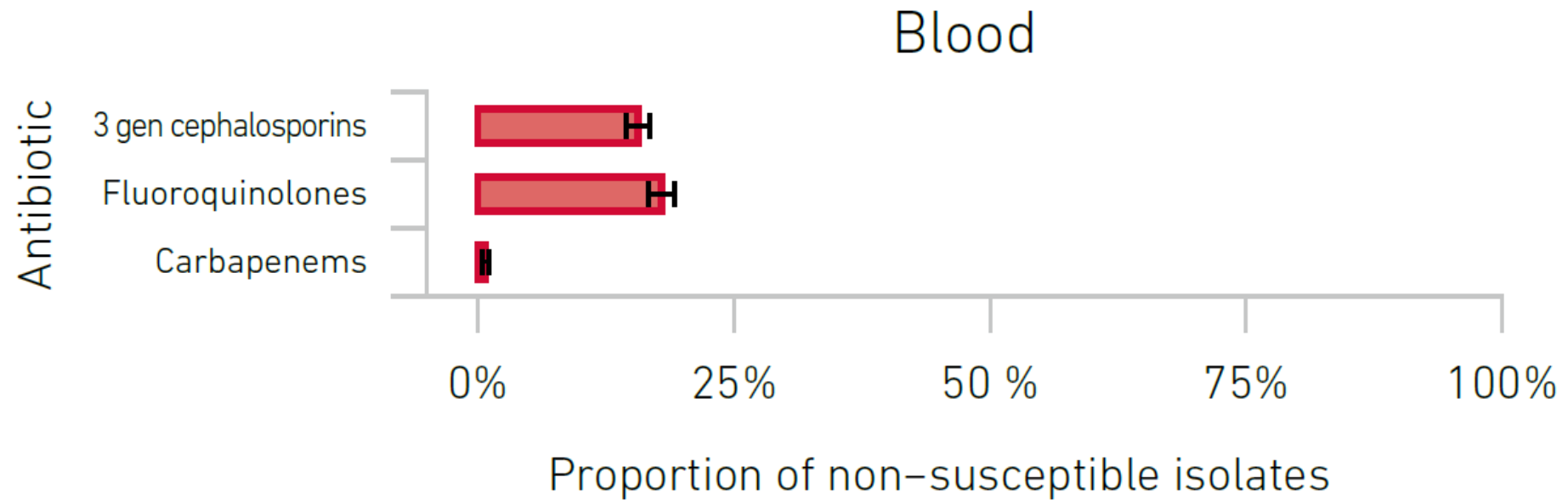
Escherichia coli

■ $\leq 30\%$ unknown AST results □ $> 30\%$ unknown AST results



Klebsiella pneumoniae

■ $\leq 30\%$ unknown AST results □ $> 30\%$ unknown AST results



Iran (Islamic Republic of)

















































Population 81.16 million




Iran has developed its National Action Plan on AMR with promotion and development of AMR surveillance included in the NAP.
Iran has been enrolled in GLASS since May 2016.

Current status of the national AMR surveillance system



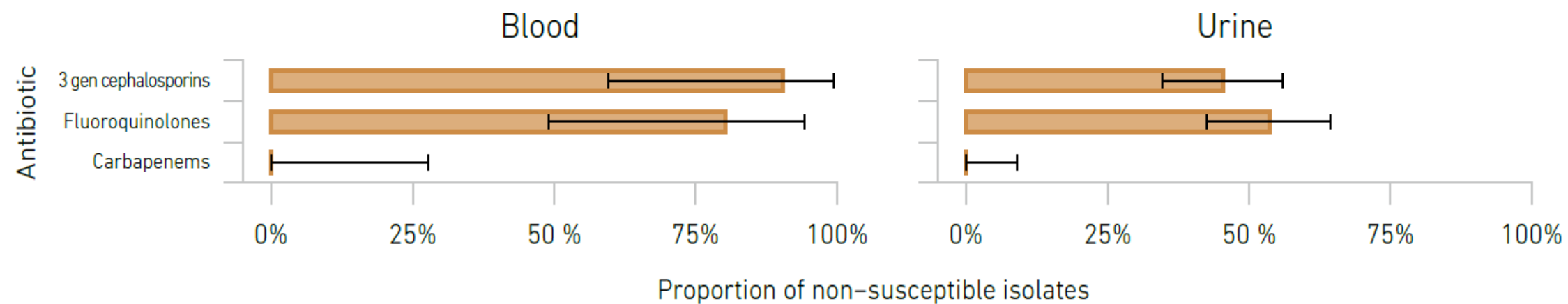
Data submission

Specimen type	Data on number of tested patient	Pathogen	AST results	Age	Gender	Infection origin
Blood		<i>Acinetobacter</i> spp.				
		<i>E. coli</i>				
		<i>K. pneumoniae</i>				
		<i>Salmonella</i> spp.				
		<i>S. aureus</i>				
		<i>S. pneumoniae</i>				
Urine		<i>E. coli</i>				
		<i>K. pneumoniae</i>				
Stool		<i>Salmonella</i> spp.				
		<i>Shigella</i> spp.				
Genital		<i>N. gonorrhoeae</i>				

 100% data collected
  99-70% data collected
  <70% data collected

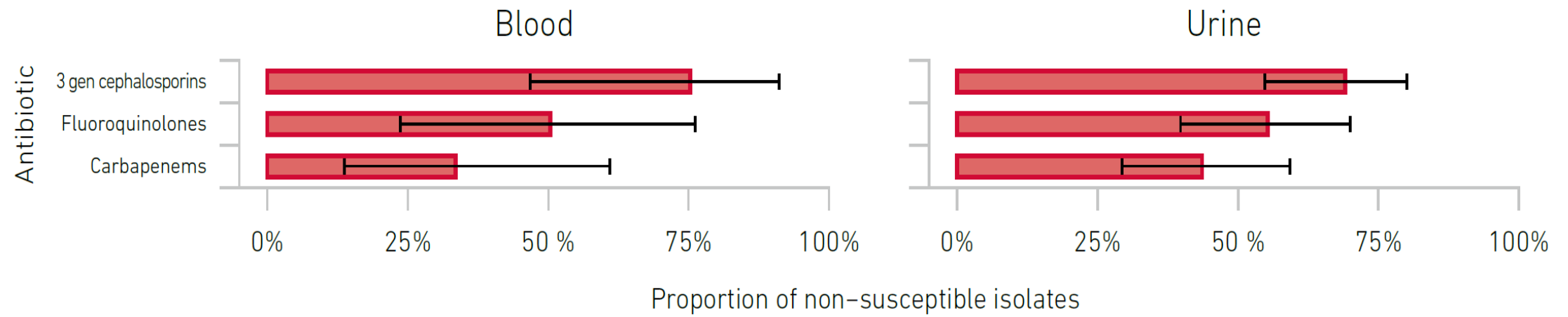
Escherichia coli

■ ≤ 30% unknown AST results □ > 30% unknown AST results



Klebsiella pneumoniae

■ ≤ 30% unknown AST results □ > 30% unknown AST results



Antibiotic use, resistance and the link to nutrition



21st century

Antibiotic resistance as one of the three most important public health

- According to a recent report, antibiotic resistance is estimated to cause around **300 million** premature deaths by 2050.
- with a loss of up to **\$100 trillion** to the global economy.
- This situation is worsened by a paucity of a robust antibiotic pipeline, resulting in the emergence of infections that are almost untreatable and leaving clinicians with **no reliable alternatives** to treat infected patients.

Antimicrobial-resistant, from farm to fork



Antibiotic applications in poultry production

1

Therapeutics

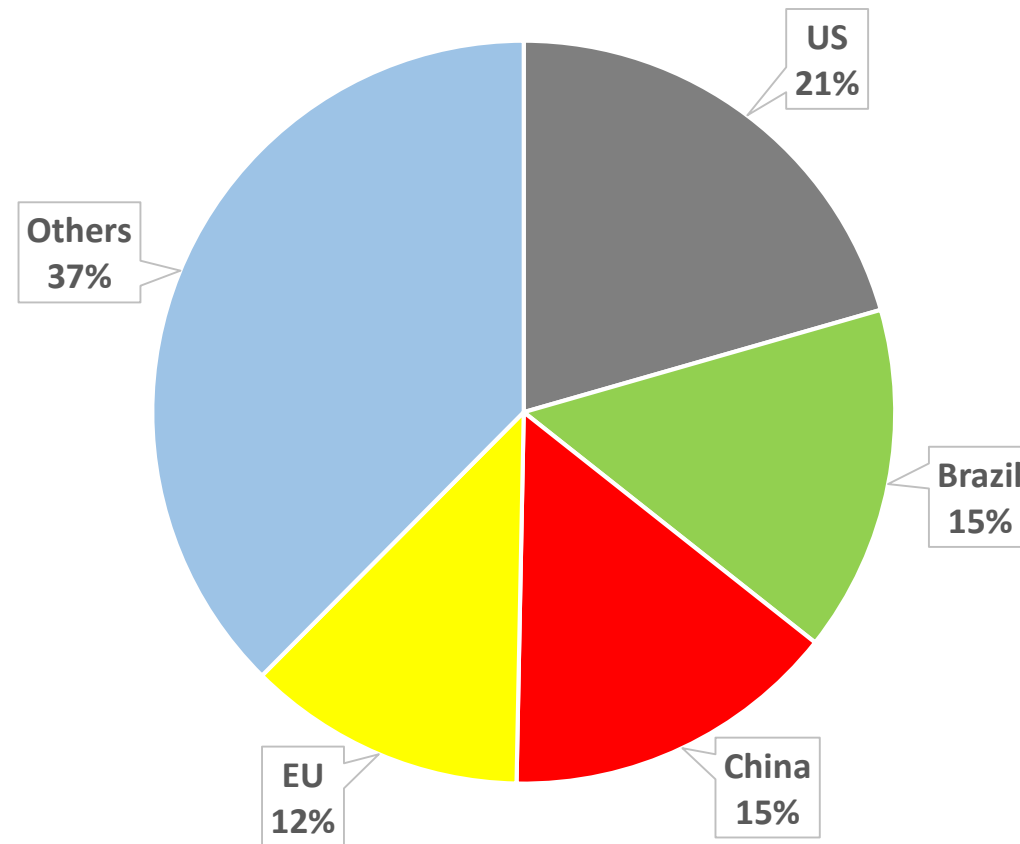
2

Prophylaxis

3

Growth promotion

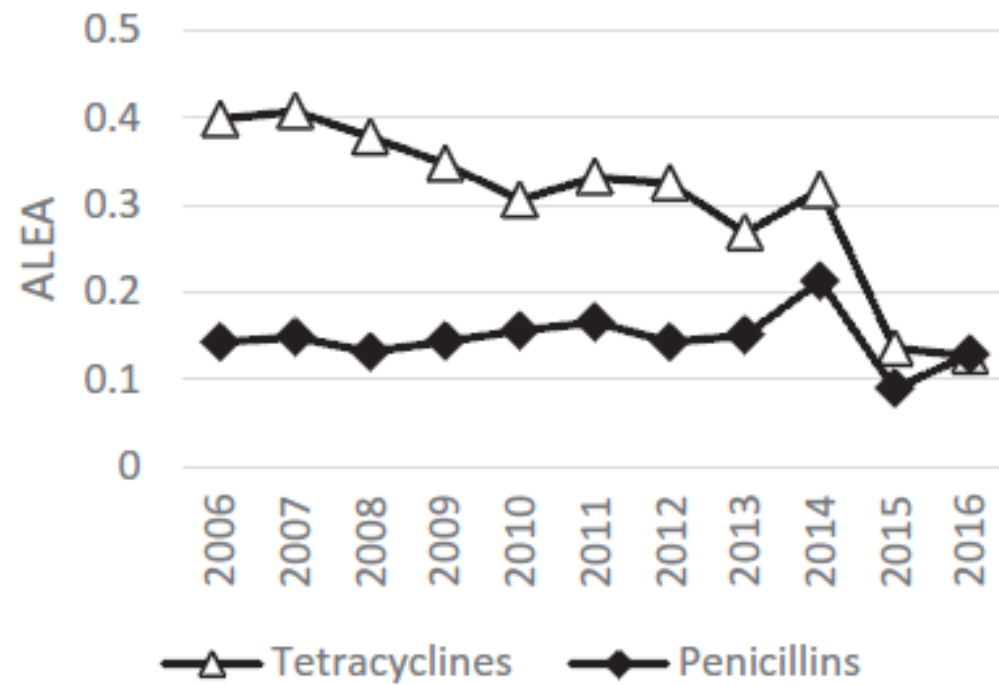
63% of the total worldwide broiler production



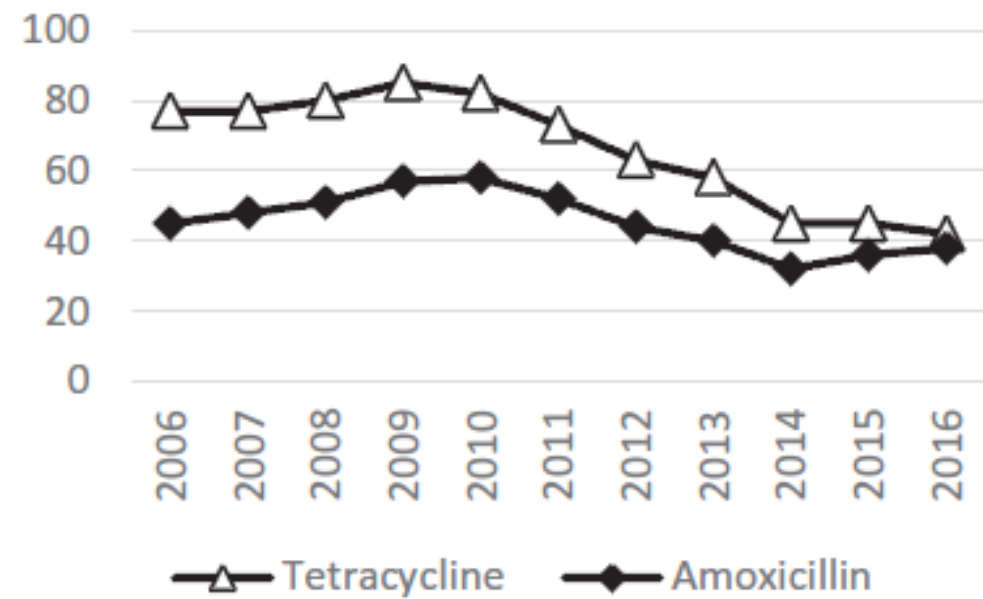
Roth et al. 2019, Poult. Sci.

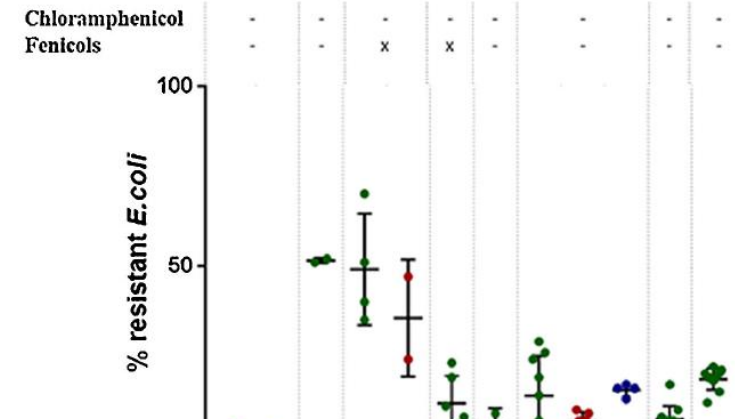
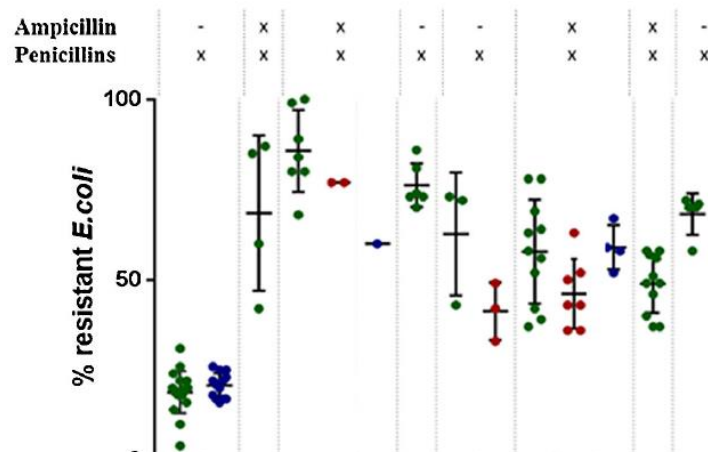
Escherichia coli are commensal bacteria that are ubiquitous in animals and humans. Because of their widespread availability, monitoring of commensal bacteria allows the comparison of the selective pressure effects in all relevant populations and is considered useful as an early alert system, for tracking emerging resistance in livestock and possible spread to animal-derived food (EFSA, 2008). Due to this prevalence, they are widely accepted as indicator bacteria for **AR** in Gram-negative bacteria populations and serve as a model for studying the emergence of AR (Kaesbohrer et al., 2012).

Antibiotic use in broilers

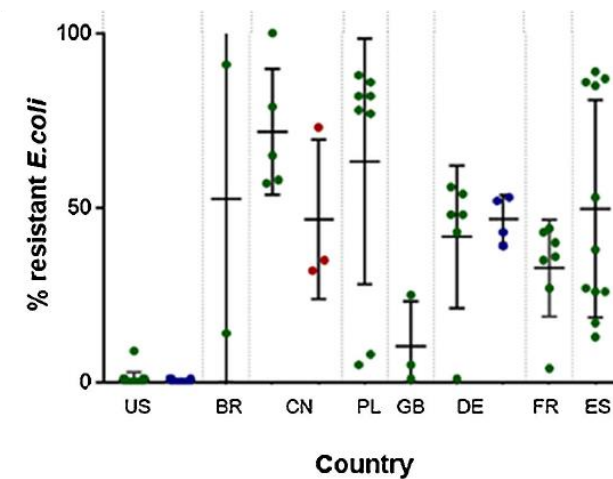
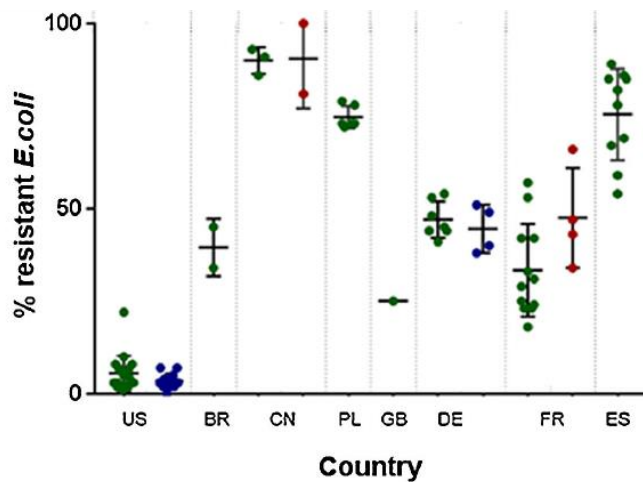


Antibiotic resistance in *E. coli*, %





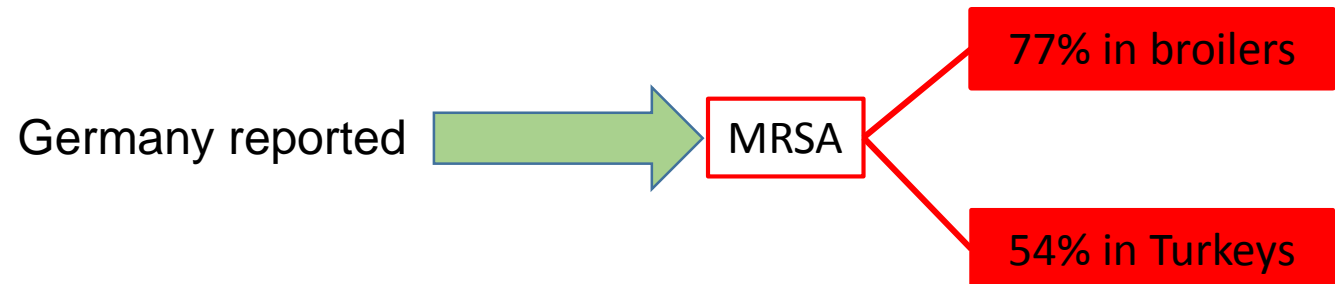
❑ The average resistance rates in *E. coli* to representatives of these antibiotic classes are higher than **40%** in all countries



Staphylococcus species

- Staphylococcosis
- pododermatitis (bumblefoot)
- septicaemia

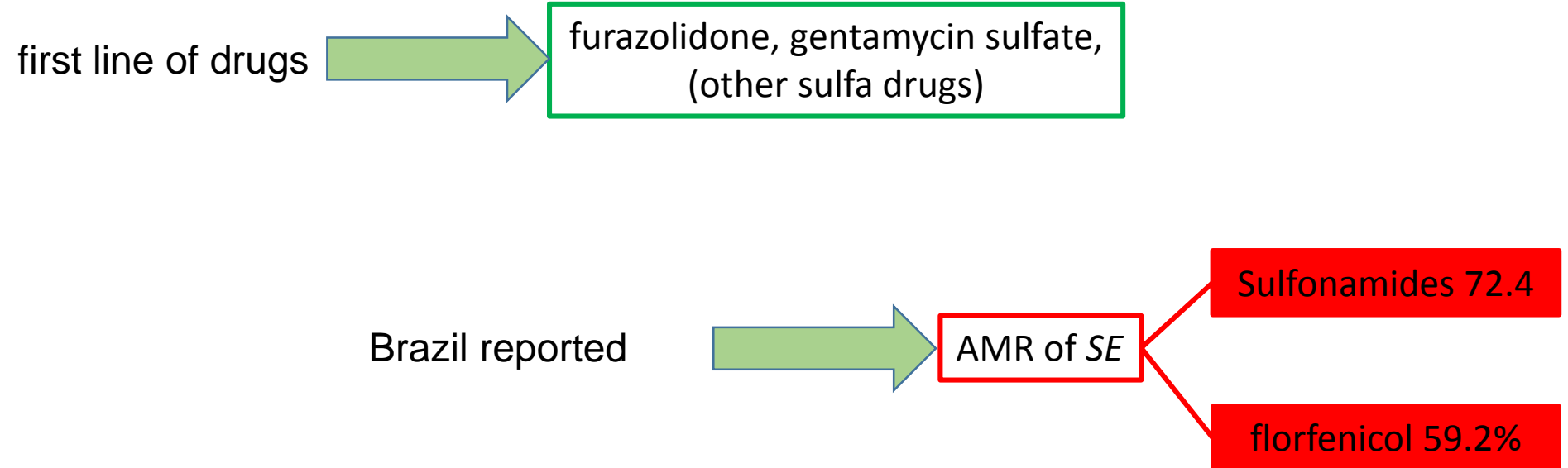
first line of drugs → β-lactams



It is worth noting that most of these organisms showed a high level of resistance to oxacillin and [tetracycline](#), which would be disastrous if these oxacillin-resistant strains are transferred to humans

Salmonella species

- salmonellosis

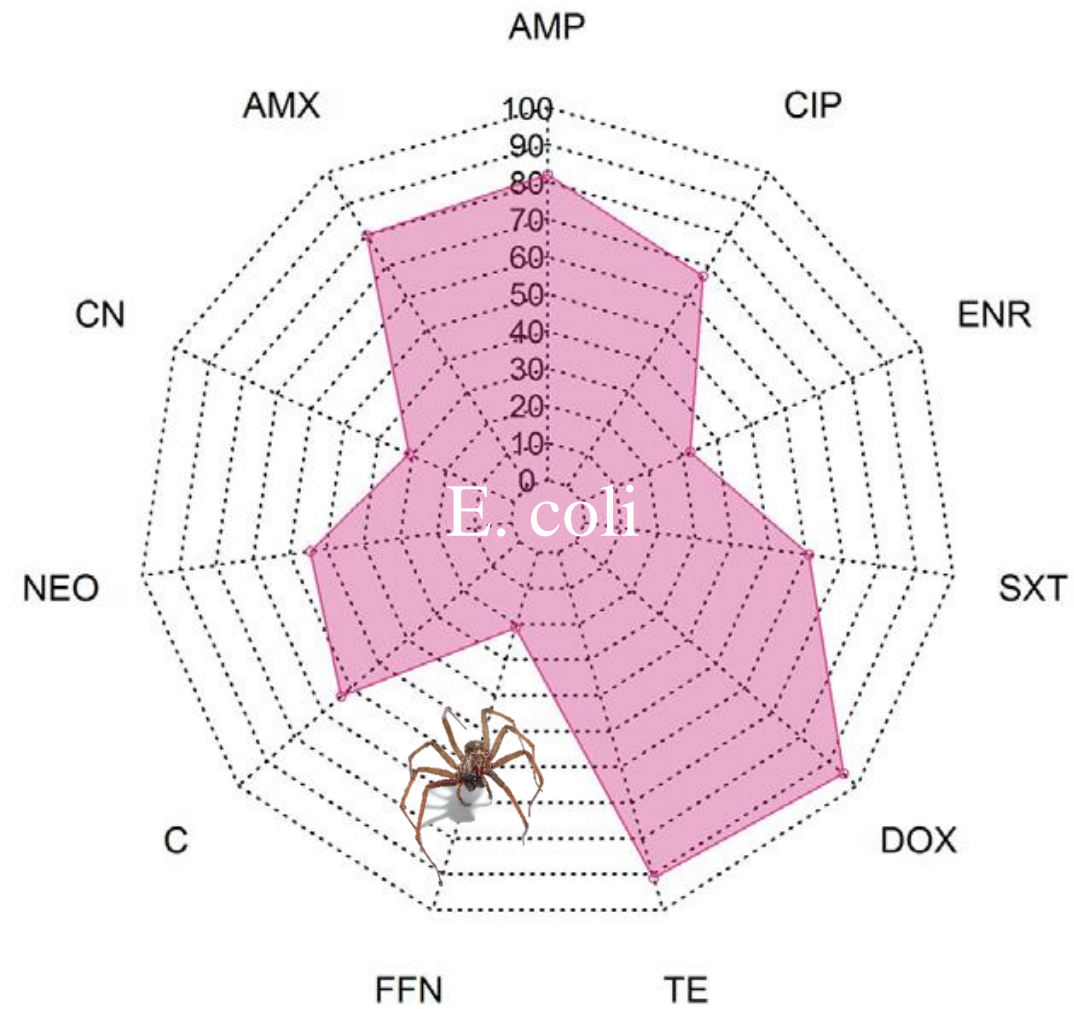


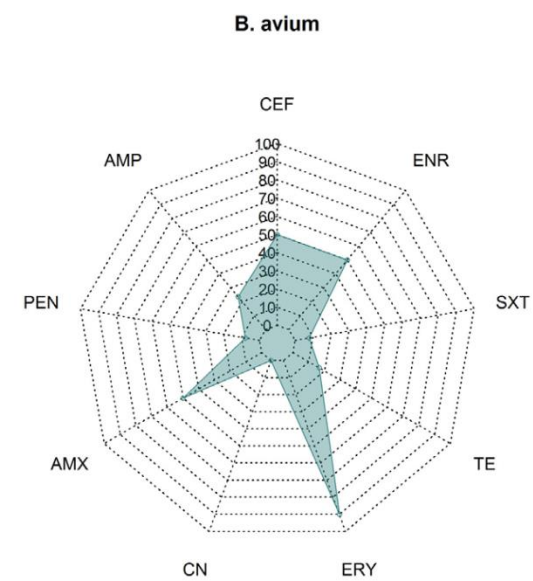
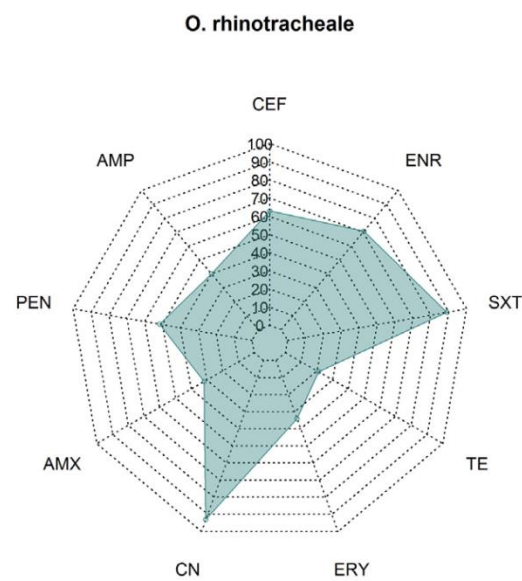
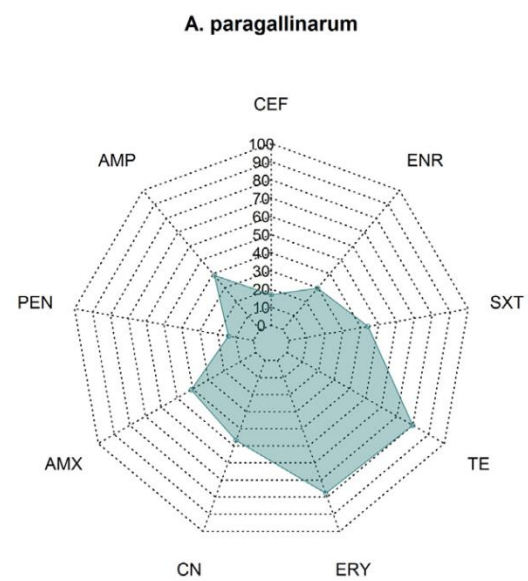
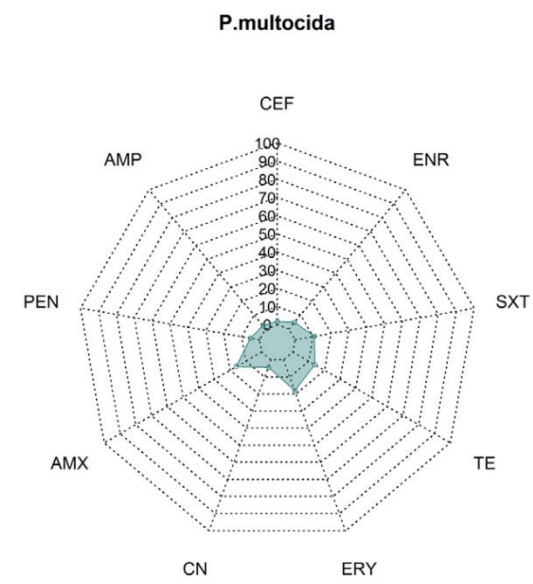
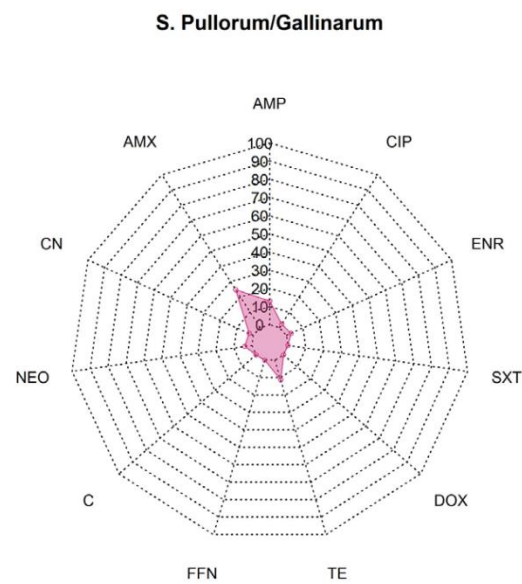
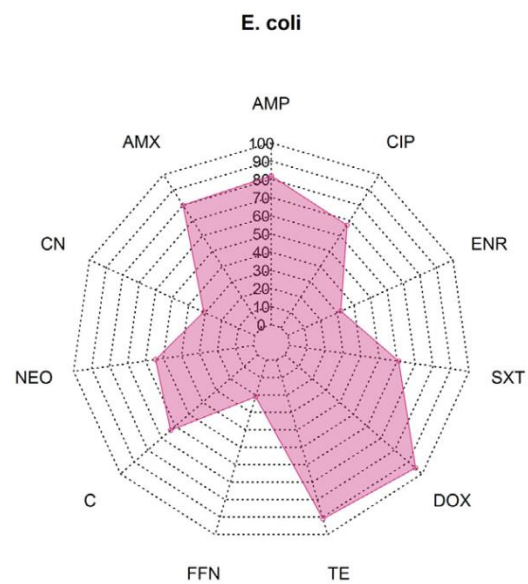
70 Published Articles

2000 - 2016

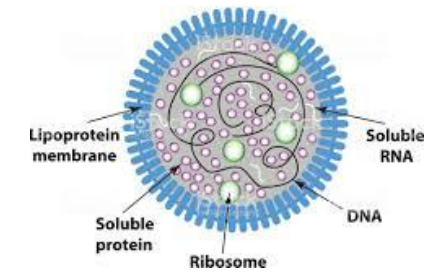
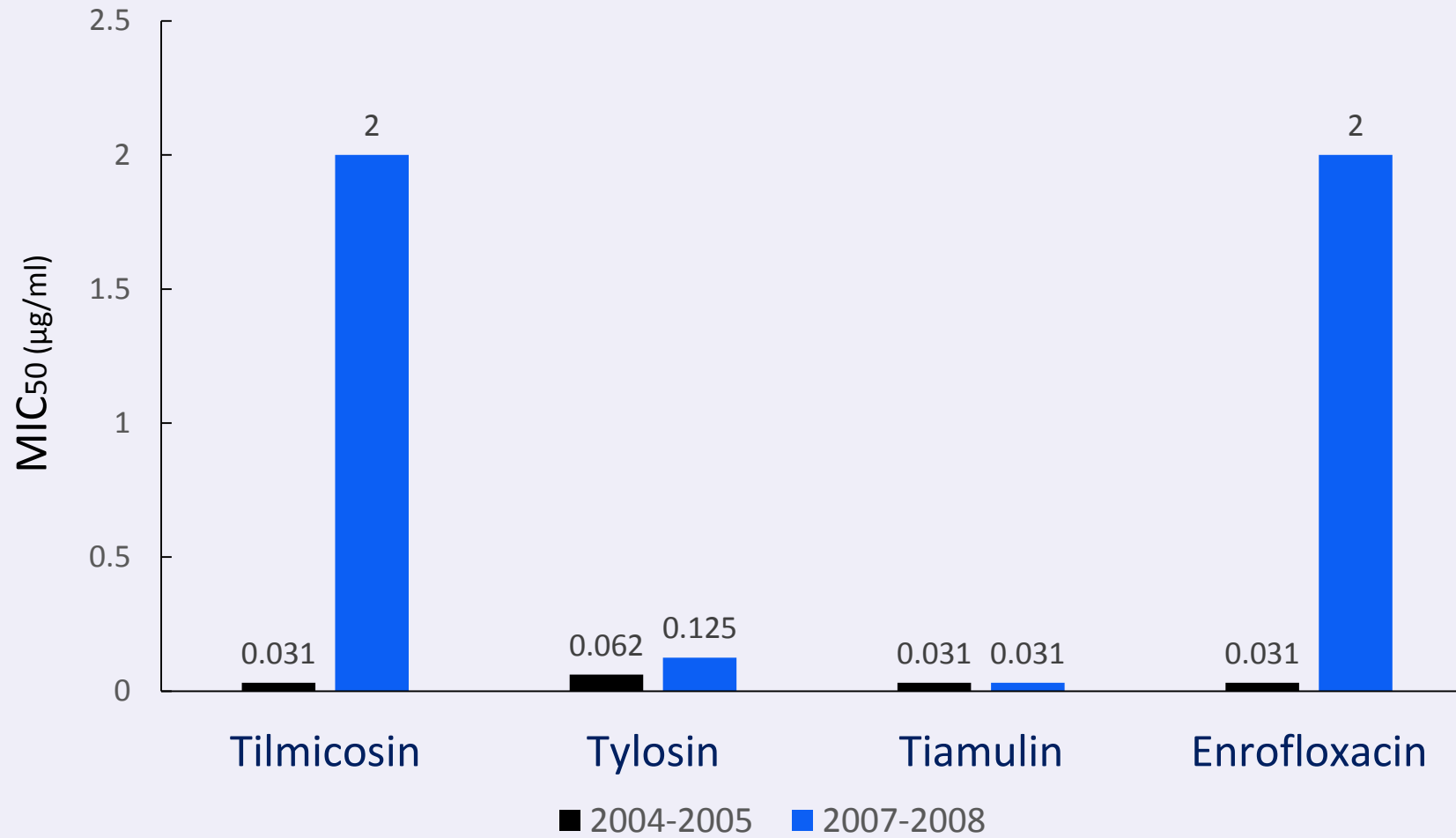
China, US, Canada, Jordan, Iran, Egypt, Brazil, Spain, Thailand, Korea, India, Zimbabwe, Hungary, Indonesia, Taiwan, Mexico, Uganda, Peru, Ecuador, Panama, Belgium, Netherlands, Germany, Sweden, Denmark and Norway

"E. coli," "S. pullorum," "S. gallinarum," "Pasteurella multocida," "Avibacterium paragallinarum," "Haemophilus para-gallinarum," "Mannheimia haemolytica," "Gallibacterium anatis," "Ornithobacterium rhinotracheale," "Mycoplasma," "Chlamydia psittaci," "Bordetella avium," "Riemerella anatipestifer," "Pseudomonas aeruginosa," "Mycobacterium avium," "Clostridium perfringens," and "Erysipelothrix rhusiopathiae."



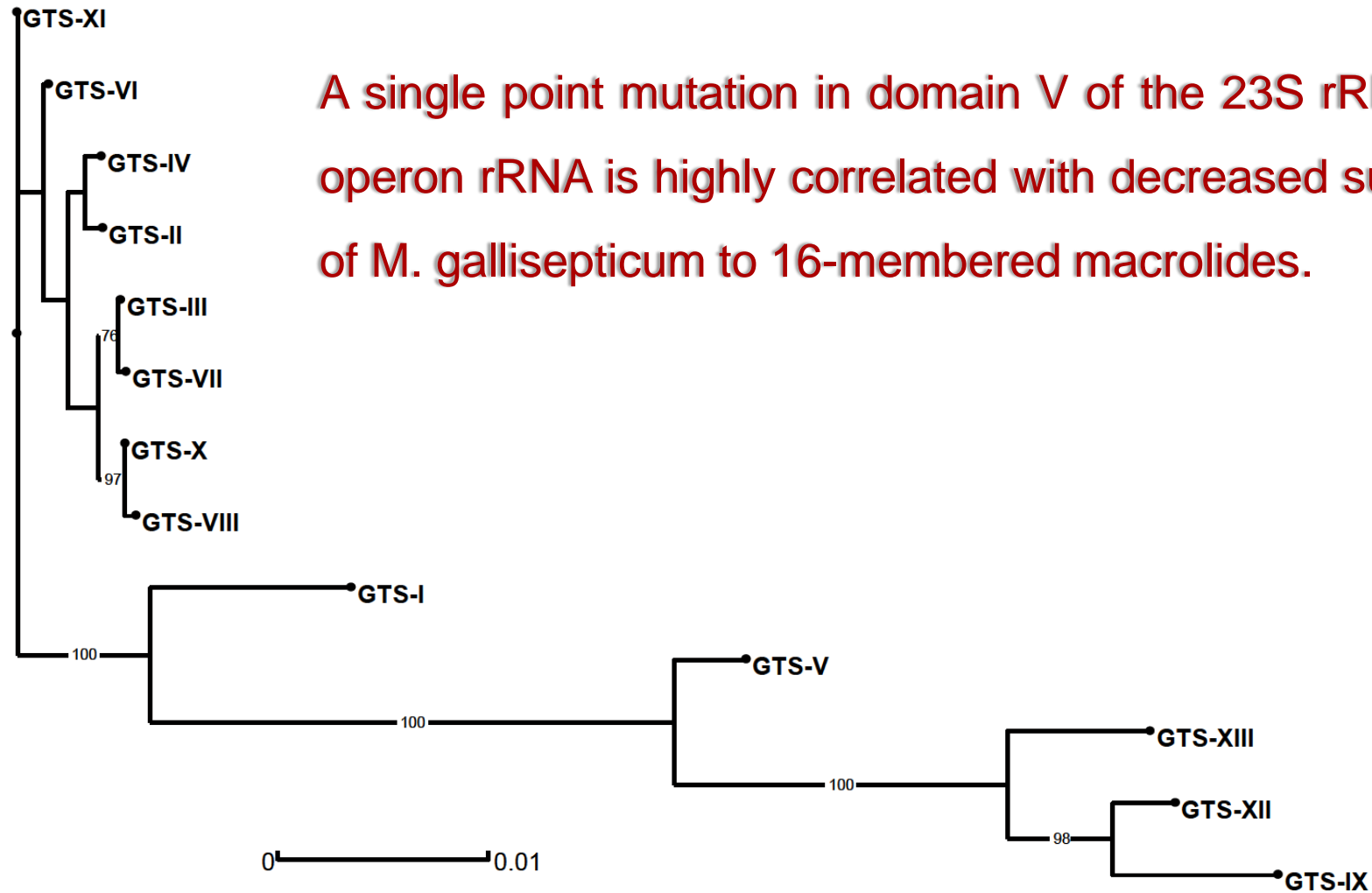


Change in antimicrobial susceptibility of *Mycoplasma gallisepticum* field isolates



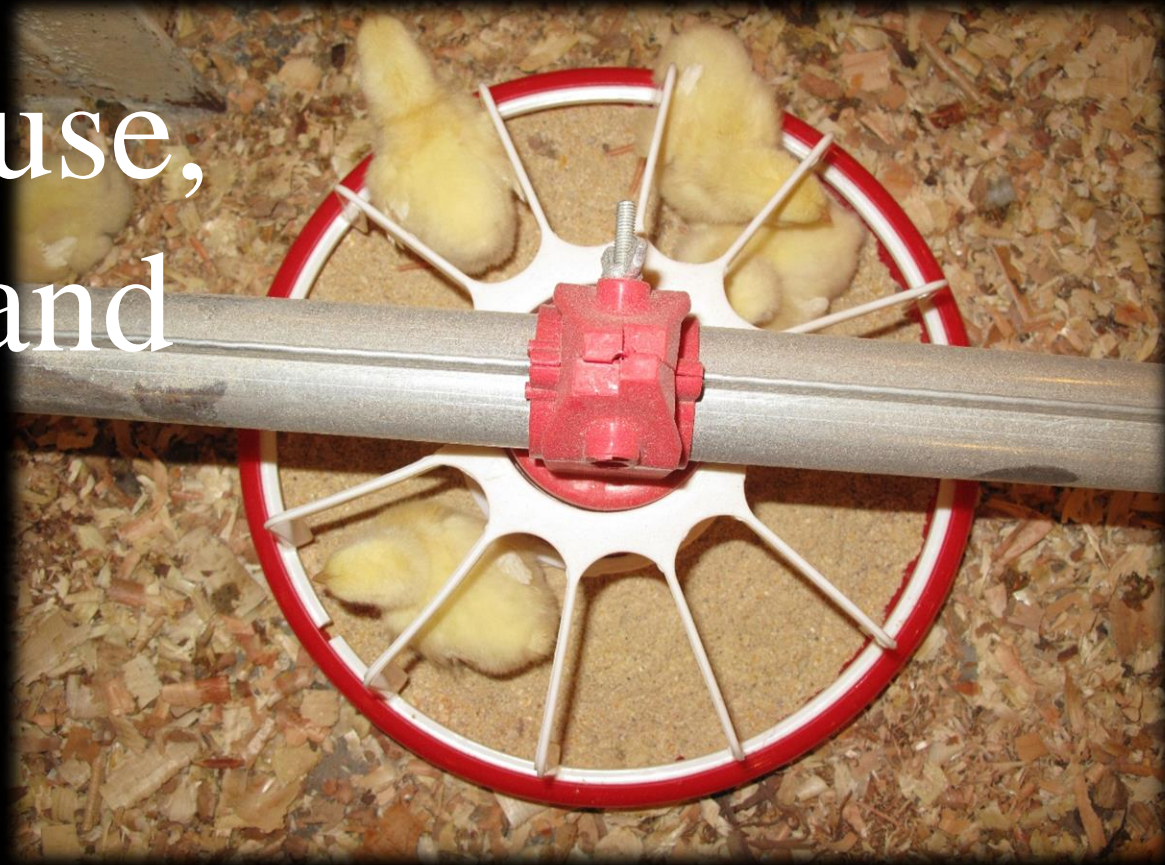
Characterization of in vivo-acquired resistance to macrolides of *Mycoplasma gallisepticum* strains isolated from poultry

Antimicrobial agent	Number of isolates with MIC (μg/mL) of												MIC ₅₀	MIC ₉₀	% Resistance
	≤0.0032	0.0063	0.0125	0.025	0.05	0.1	0.25	0.63	1.25	2.5	5	≥10			
Tylosin	1	11	10	2	1			3	8	13	1		0.05	2.5	50
Tilmicosin	16	6		2		1			3		7	15	0.1	≥10	50
Enrofloxacin*				3	10	9	5		2	11	8	2	0.25	5	46

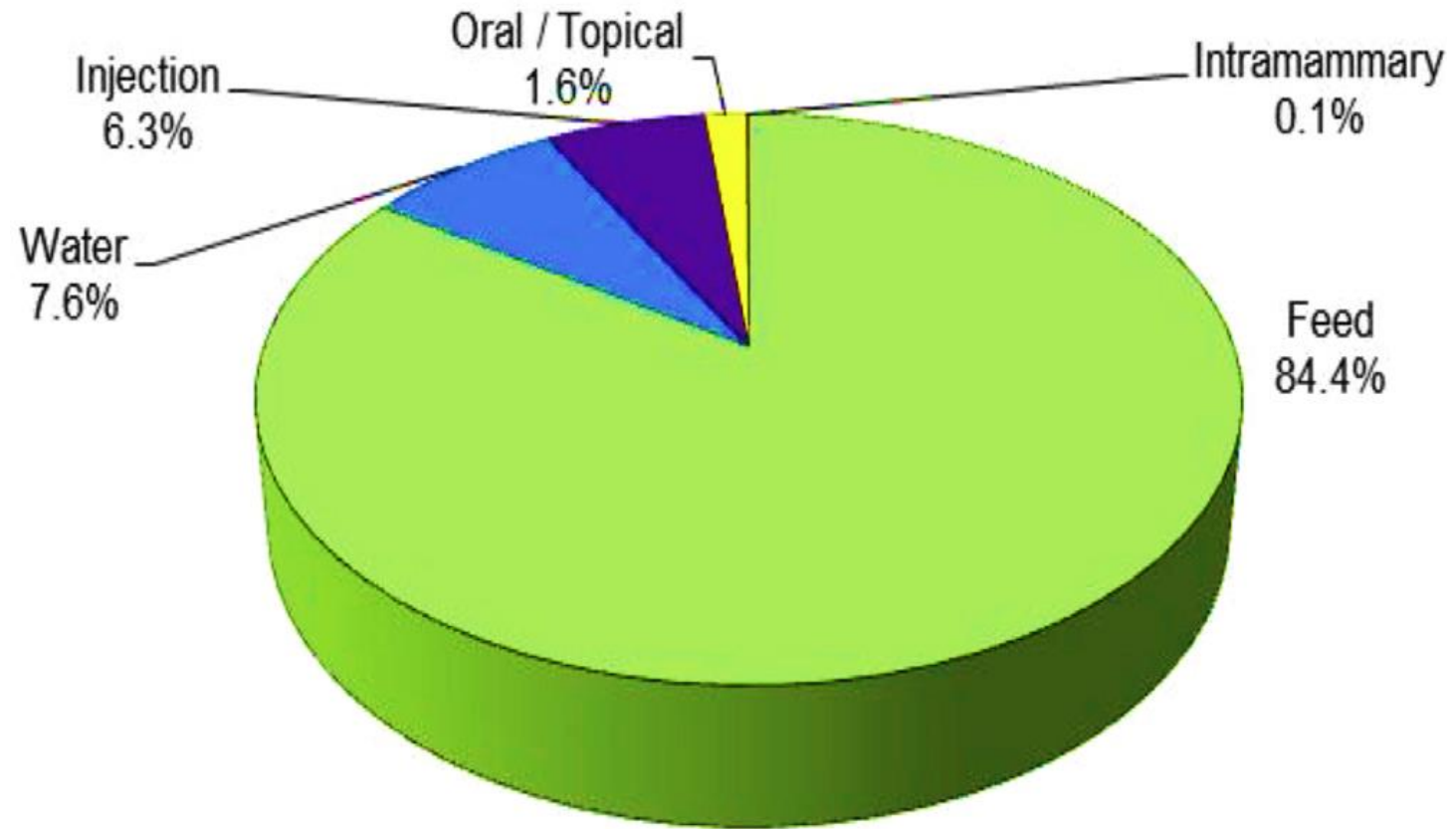


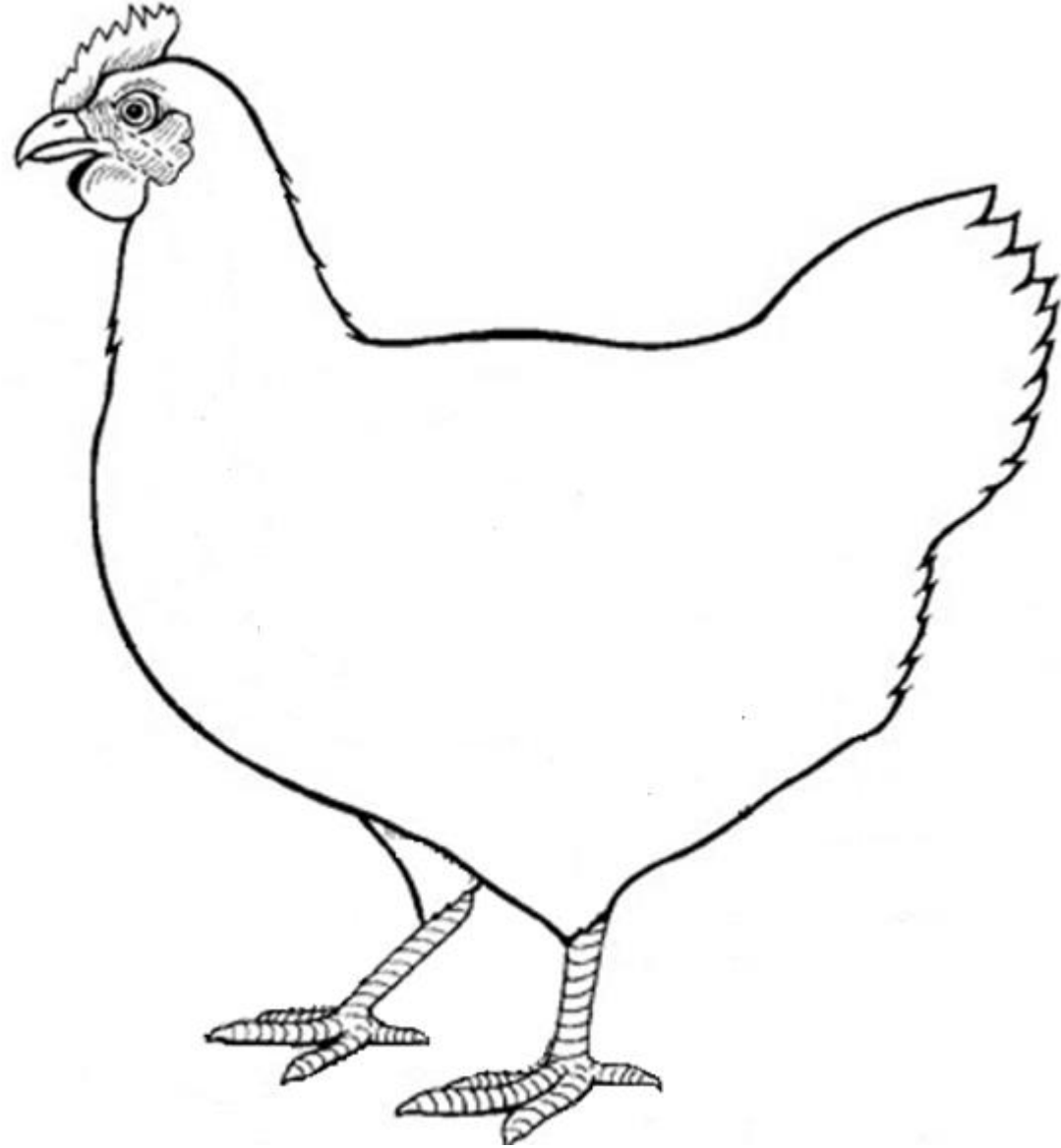
A single point mutation in domain V of the 23S rRNA gene of operon rRNA is highly correlated with decreased susceptibility of *M. gallisepticum* to 16-membered macrolides.

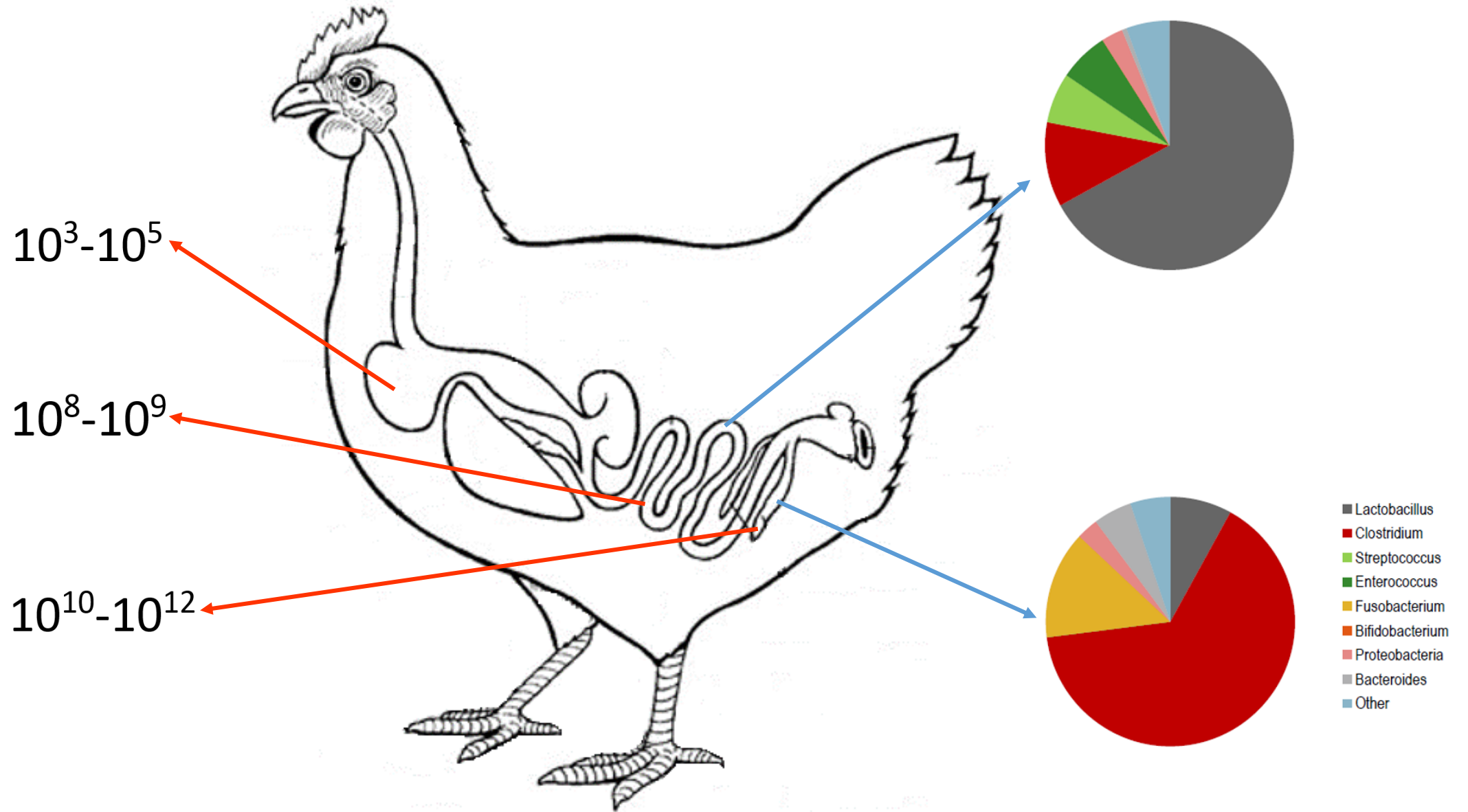
Antibiotic use, resistance and the link to nutrition

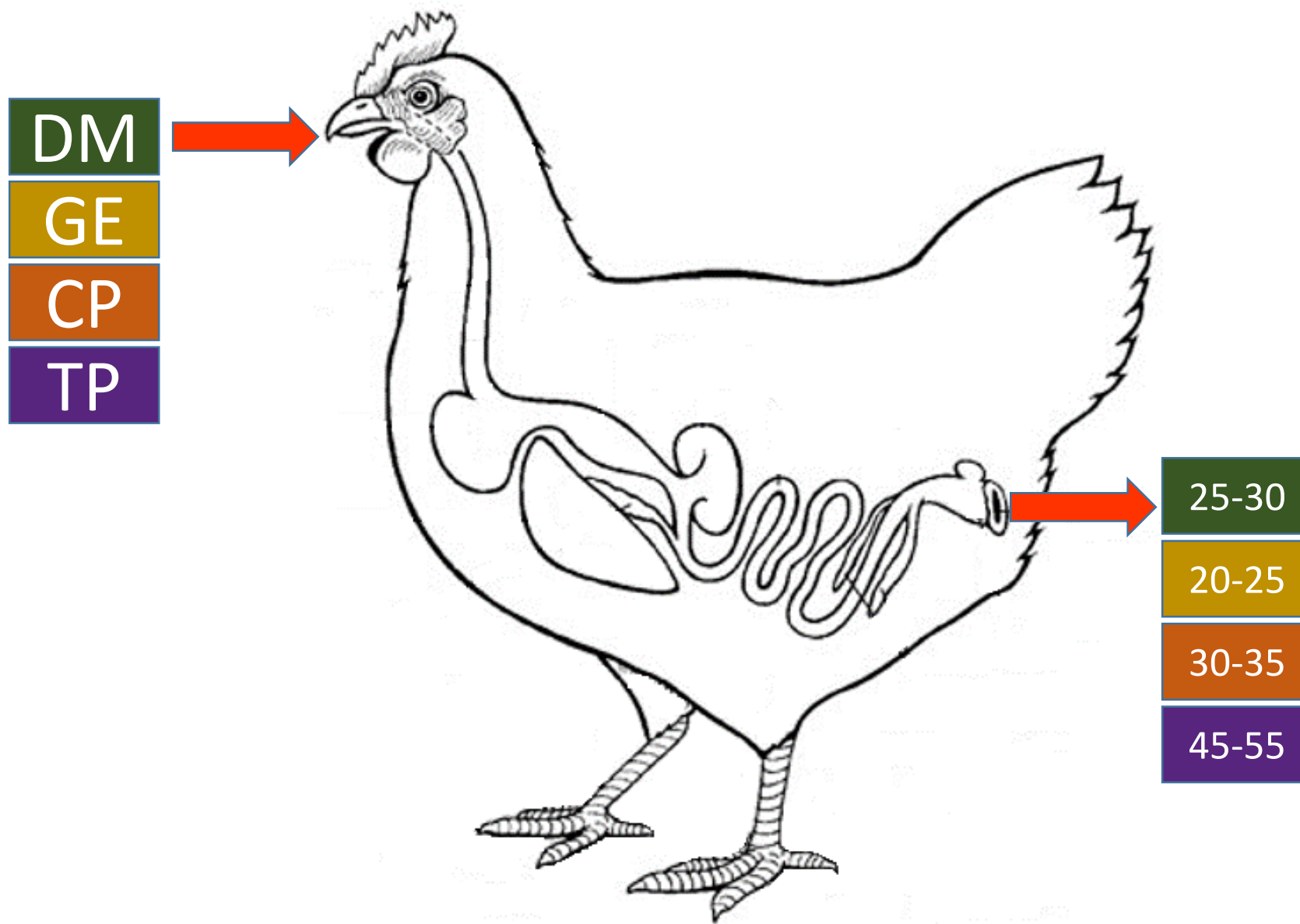


Quantity of antimicrobials (% of total weight in kg) distributed for veterinary use by route of administration in Canada (CSCRA, 2016).

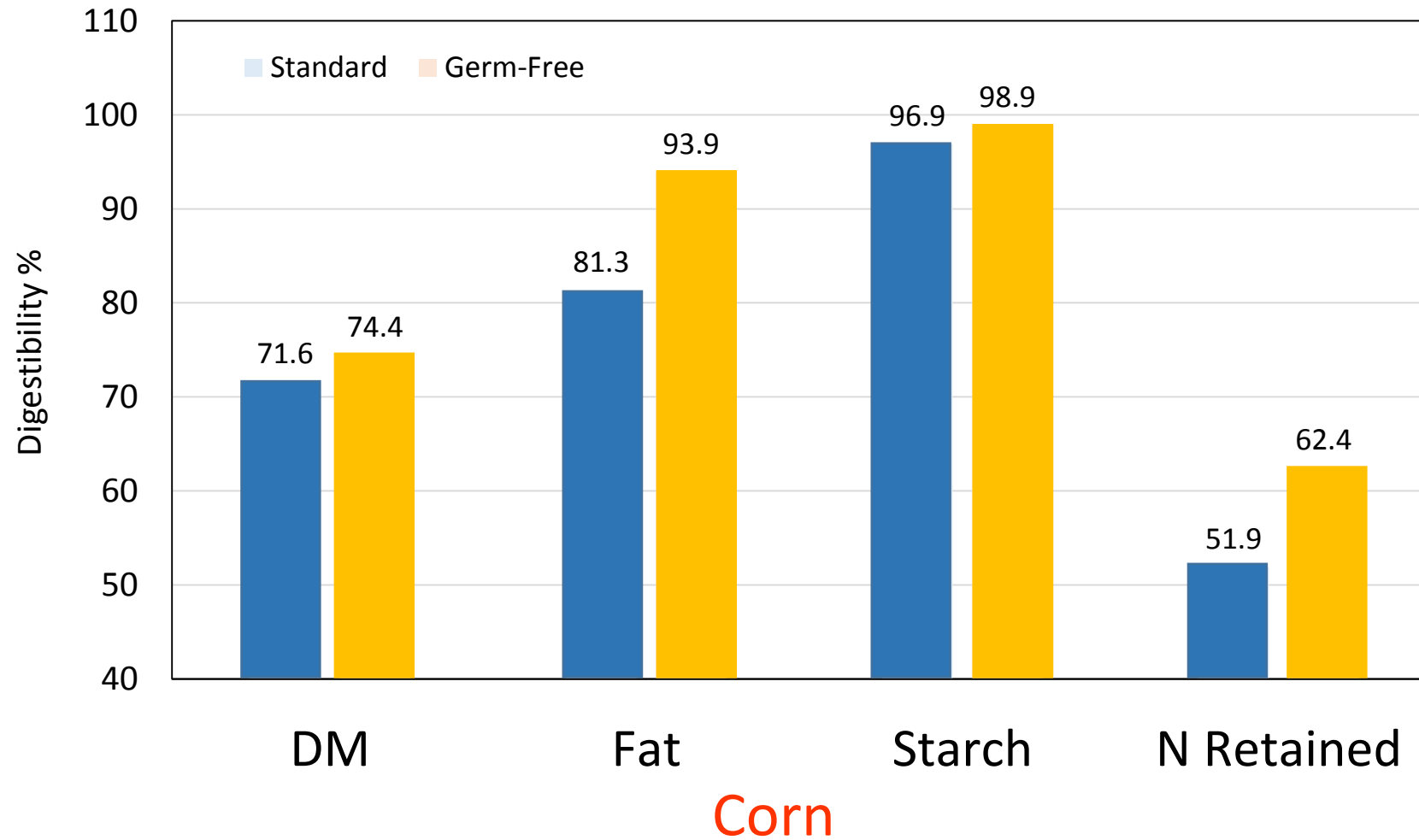




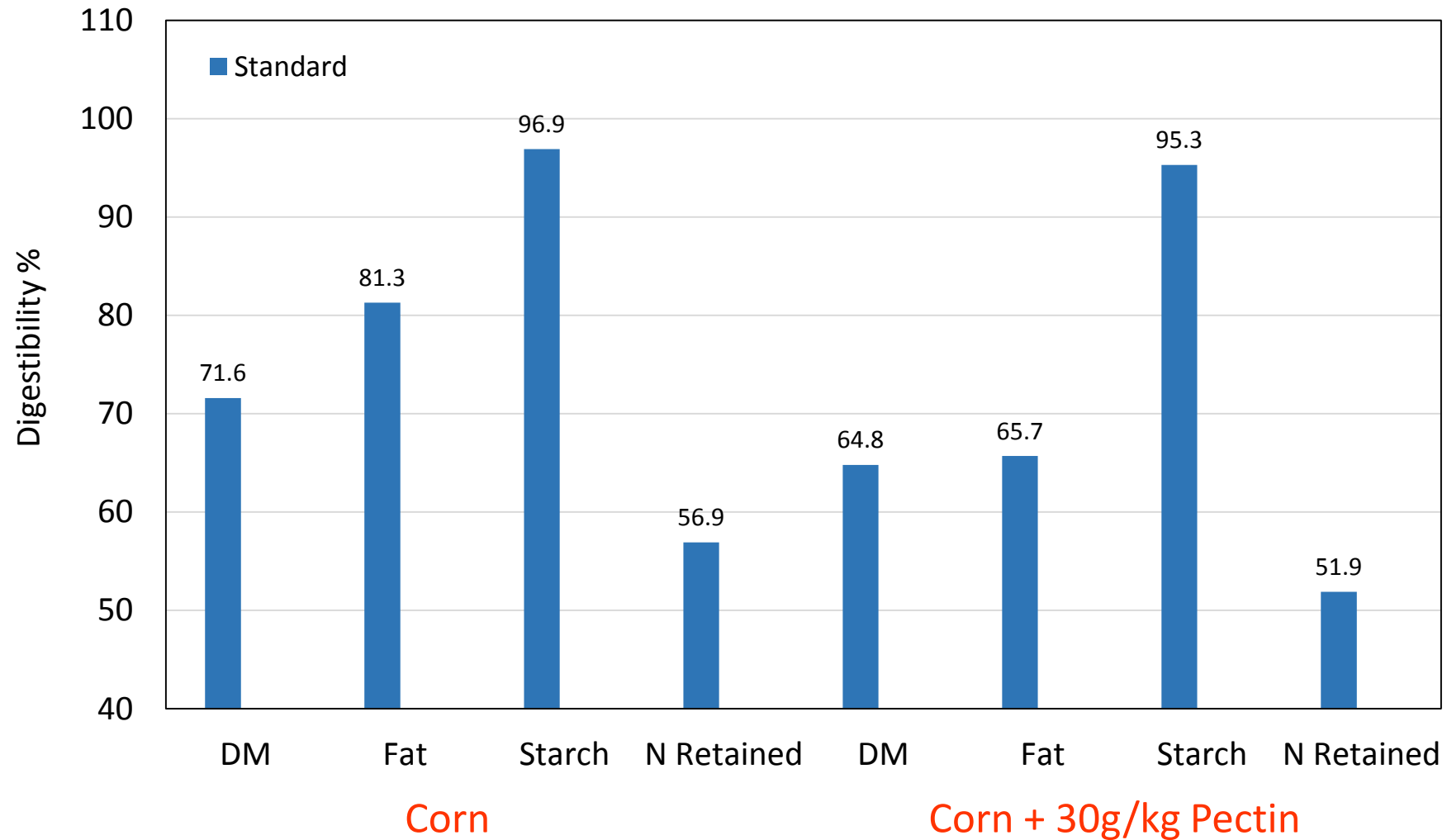




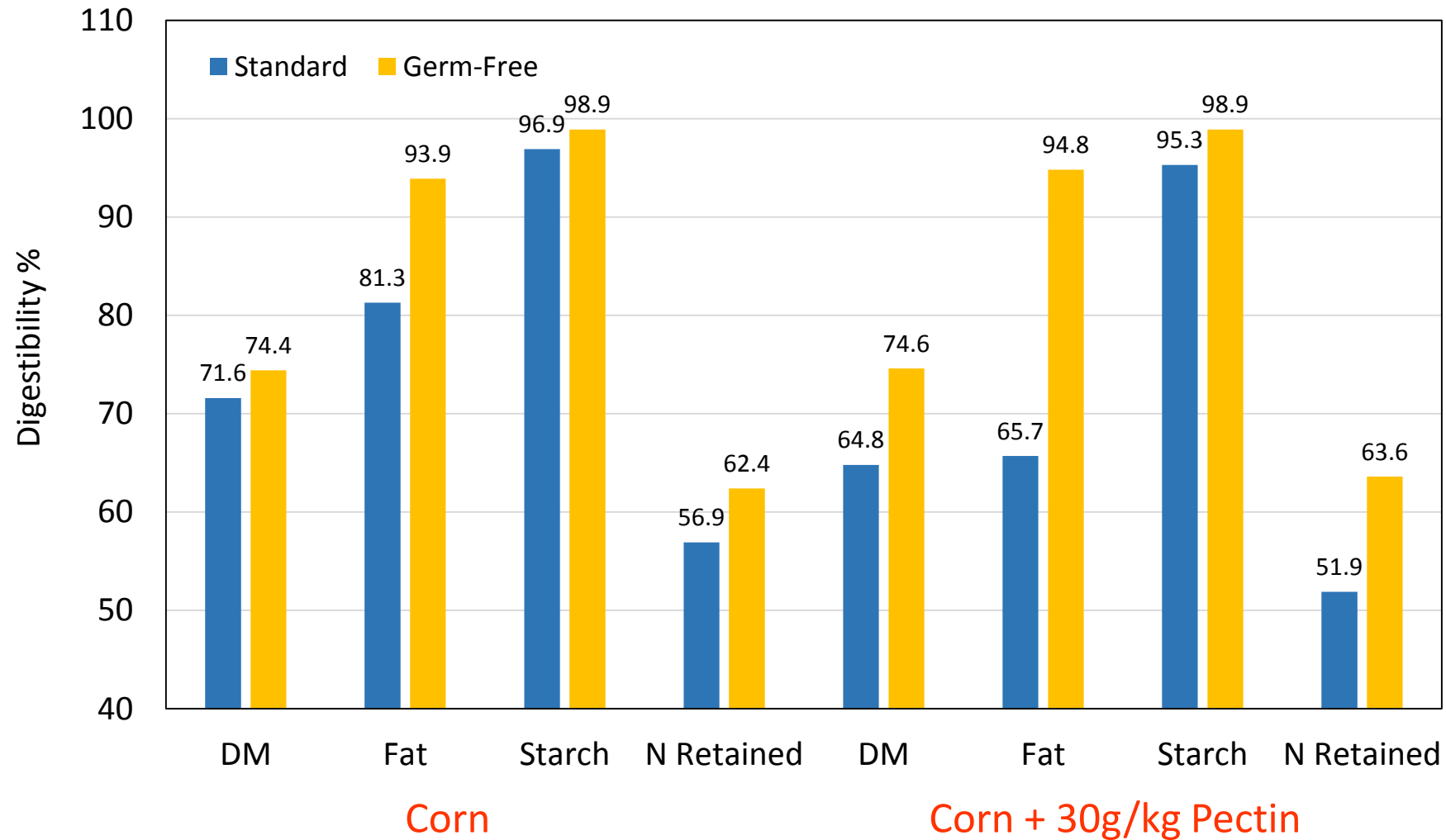
Feed Interacts with Bacteria



Feed Interacts with Bacteria



Feed Interacts with Bacteria



Microbial competition for dietary protein in chicken small intestine

Amino acid requirements of some small intestine bacteria.

	<i>Lactobacillus</i> spp. ^a	<i>Clostridium perfringens</i> ^b	<i>Escherichia coli</i>
Alanine	±	—	—
Arginine	+	+	—
Aspartic acid	+	—	—
Cysteine	+	—	—
Glutamic acid	+	+	—
Glycine	±	—	—
Histidine	±	+	—
Iso-leucine	+	+	—
Leucine	+	+	—
Lysine	±	—	—
Methionine	+	+	—
Phenylalanine	+	+	—
Proline	±	—	—
Serine	±	—	—
Threonine	+	+	—
Tryptophan	+	+	—
Tyrosine	+	+	—

+growth of the tested strains was dependent on the amino acid

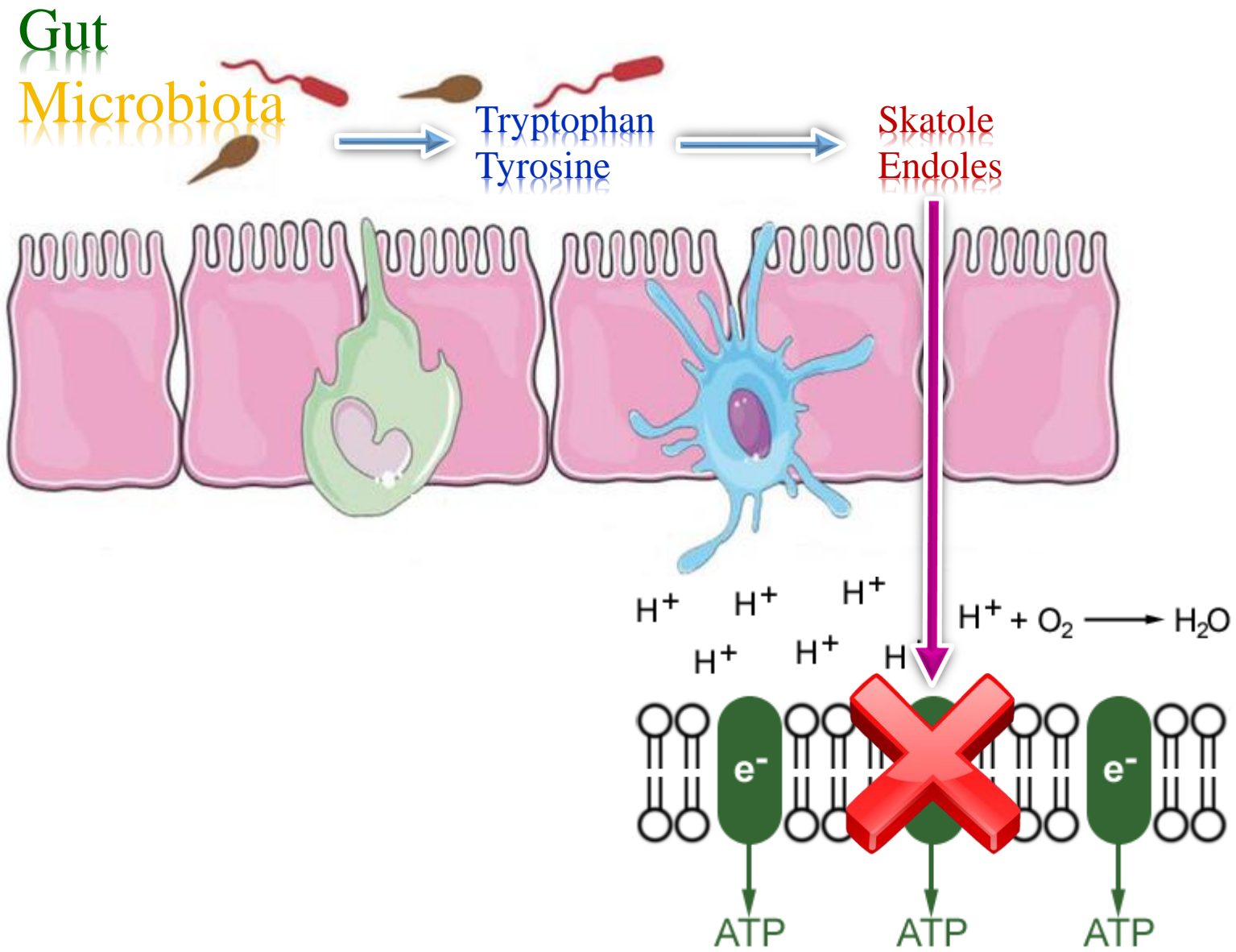
±growth of some of the tested strains was dependent on the amino acid

—growth of the tested strains was not dependent on the amino acid

^a Morishita et al., 1981.

^b Sebald and Costilow, 1975.





Microbial Reconstitution Reverses Maternal Diet-Induced Social and Synaptic Deficits in Offspring



Shelly A. Buffington,^{1,2} Gonzalo Viana Di Prisco,^{1,2} Thomas A. Auchtung,^{3,4} Nadim J. Ajami,^{3,4} Joseph F. Petrosino,^{3,4} and Mauro Costa-Mattioli^{1,2,*}

¹Department of Neuroscience

²Memory and Brain Research Center

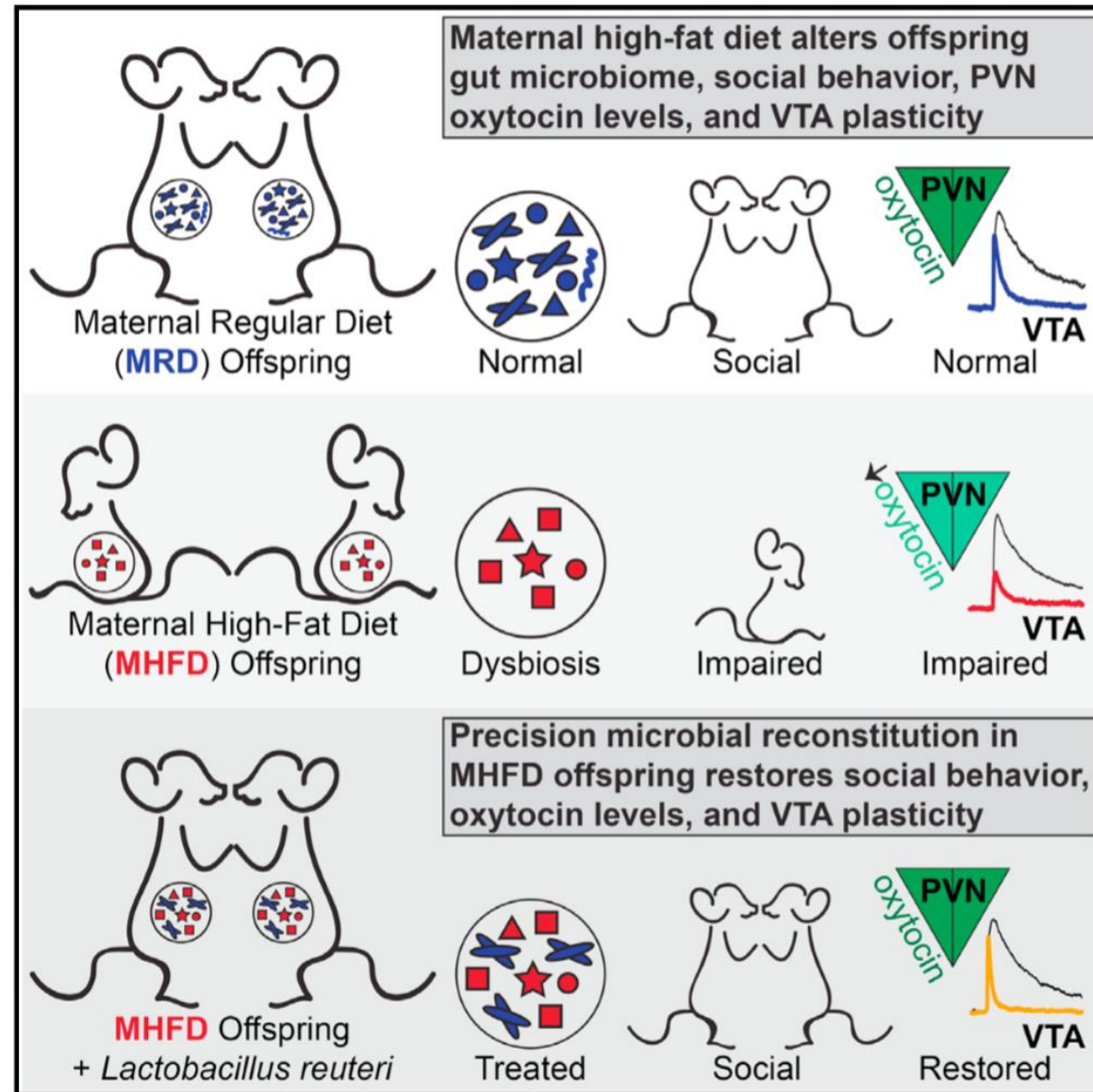
³Alkek Center for Metagenomics and Microbiome Research

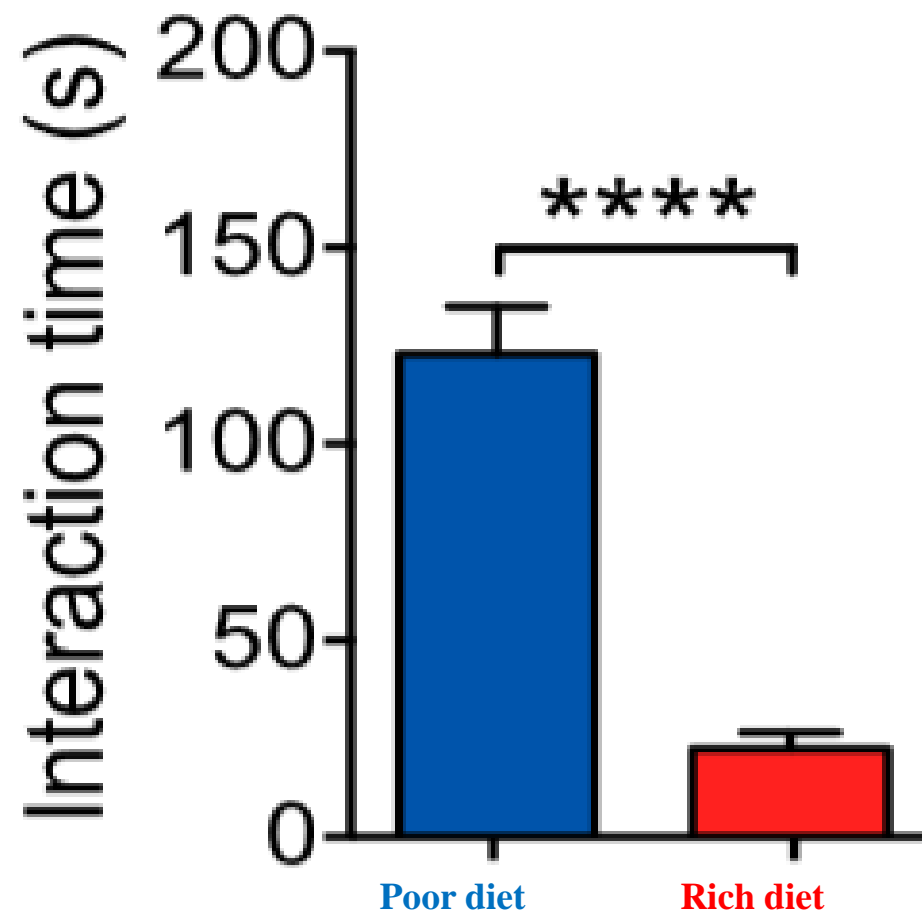
⁴Department of Molecular Virology and Microbiology
Baylor College of Medicine, Houston, TX 77030, USA

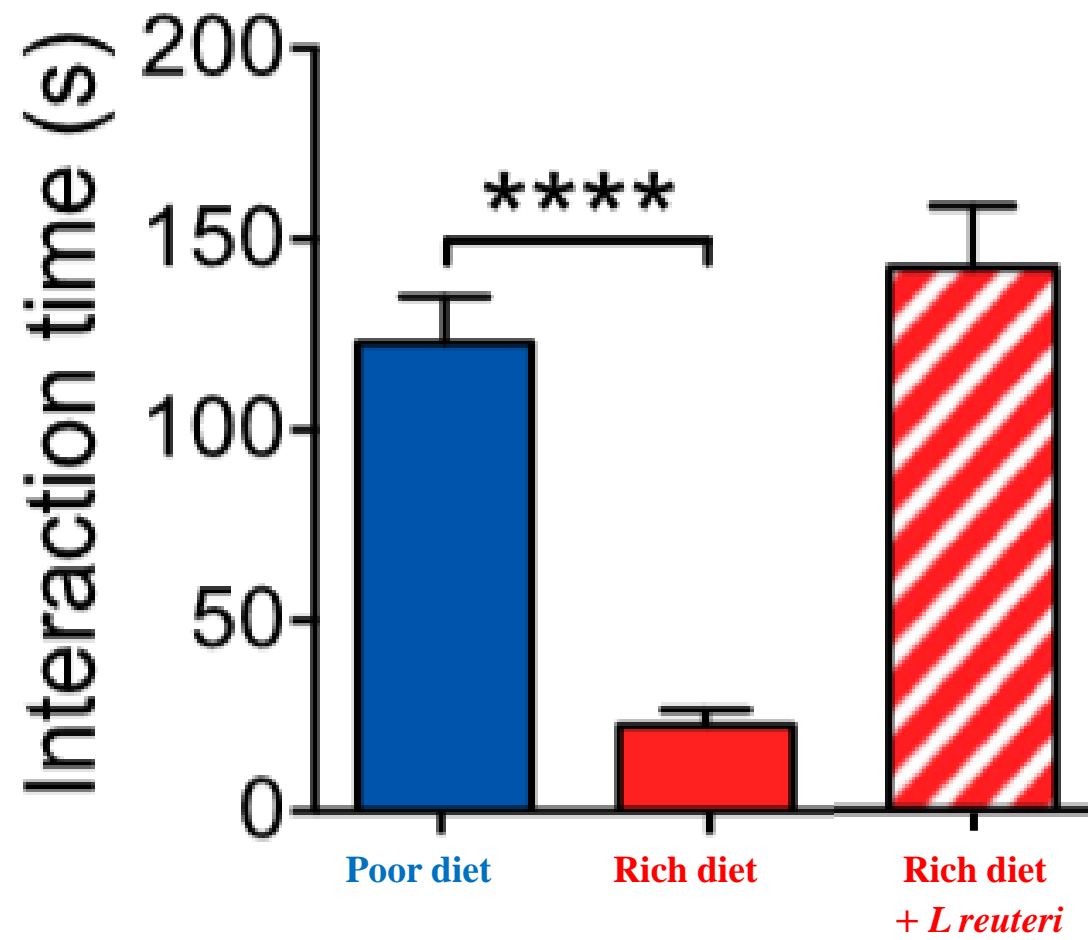
*Correspondence: costamat@bcm.edu

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Graphical Abstract







Antibiotics

-  Reduction in the stability of the microbiota
-  Increases Firmicutes/Bacteroidetes
-  Fewer functions associated with carbohydrate transport and metabolism
-  Smaller arsenal of families involved in the degradation of starch, cellulose, and hemicellulose.

- Lan et al., 2005;
- Danzeisen et al., 2011;
- Allen and Stanton, 2014;
- Mancabelli et al., 2016 ;
- Clavijo et al., 2018.



Some probiotics strains carry antibiotic resistance genes

Ivan & Joyce, 2016

