



HADEA SERVICE CONTRACT 20197409

Provision of EU networking and support for public health reference laboratory functions for antimicrobial resistance in *Salmonella* species and *Campylobacter* species in human samples



FWD AMR.
RefLabCap

EU protocol for harmonised monitoring of antimicrobial resistance in human *Salmonella* and *Campylobacter* isolates.

EUCAST protocols, guidelines, clinical/epidemiological breakpoints, interpretation and website.

EQA-AST 6

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- Has a mandate to gather and analyse data and information on emerging public health threats
- The collection antimicrobial resistance (AMR) data is included as part of the European Surveillance System (TESSy) through several networks:
- EARS-Net (*S. pneumoniae*, *S. aureus*, *E. faecalis*, *E. faecium*, *E. coli*, *Klebsiella pneumoniae*, *P. aeruginosa*, and *Acinetobacter* spp.).
- • HAI-Net collects data on AMR in selected pathogens associated with healthcare-associated infections.
- • ESAC-Net collects data on the consumption of antimicrobial agents in humans.
- • **FWD-Net collects data on AMR in *Salmonella* spp., *Campylobacter* spp. and Shiga toxin/verocytotoxin-producing *Escherichia coli* (STEC/VTEC)**

- Directive 2003/99/EC requires Member States to monitor and report comparable data on AMR in zoonoses and zoonotic agents in food-producing animals and food
- Commission Implementing Decision (EU) 2020/1729 of 17 November 2020 on the monitoring and reporting of antimicrobial resistance in zoonotic and commensal bacteria



Read the report




Publication

The European Union Summary Report on Antimicrobial Resistance in zoonotic and indicator bacteria from humans, animals and food in 2019–2020

Technical report - 29 Mar 2022

Data on antimicrobial resistance (AMR) in zoonotic and indicator bacteria from humans, animals and food are collected annually by the EU Member States (MSs), jointly analysed by the EFSA and the ECDC and reported in a yearly EU Summary Report.

 [The European Union Summary Report on Antimicrobial Resistance in zoonotic and indicator bacteria from humans, animals and food in 2019–2020 - EN - \[PDF-67.39 MB\]](#)

 [Antimicrobial consumption](#) | [Antimicrobial resistance](#) | [Antimicrobial stewardship](#) |




EU HARMONIZED PROTOCOL FOR AMR TESTING OF SALMONELLA AND CAMPYLOBACTER


EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates – June 2016

ecdc.europa.eu/en/publications-data/eu-protocol-harmonised-monitoring-antimicrobial-resistance-human-salmonella-and-0

An official website of the European Union How do you know?

Other sites: ECDC European Antibiotic Awareness Day ESCAIDE - Scientific conference Eurosurveillance journal EVIP - Vaccination portal

 **European Centre for Disease Prevention and Control**
An agency of the European Union


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> EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates – June 2016

Publications & data

EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates – June 2016

Technical guidance
7 Jun 2016
Cite: 



This protocol for harmonised monitoring of antimicrobial resistance in Salmonella and Campylobacter from human isolates was updated from the March 2014 version. While the revised version introduces a number of new antimicrobials and resistance breakpoints, its overall objectives – to increase the quality and comparability of EU/EEA antimicrobial resistance data – remain unchanged.


The Protocol is targeted at the national public health reference laboratories to guide the susceptibility testing needed for EU surveillance and the reporting to ECDC.

Note that annex 1 and 2 were updated in August 2021 and are available below

Download

- EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates, June 2016 - EN - [PDF-928.92 KB]
- EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates - Annexes August 2021 - EN - [PDF-100.96 KB]
- EU protocol for harmonised monitoring of antimicrobial resistance in human Salmonella and Campylobacter isolates, March 2014 - EN - [PDF-1.2 MB]

Antimicrobial resistance | Campylobacteriosis | Europe | Food- and Waterborne Diseases and Zoonoses Programme | Laboratories | Salmonellosis






Page last updated: 3 Sep 2021

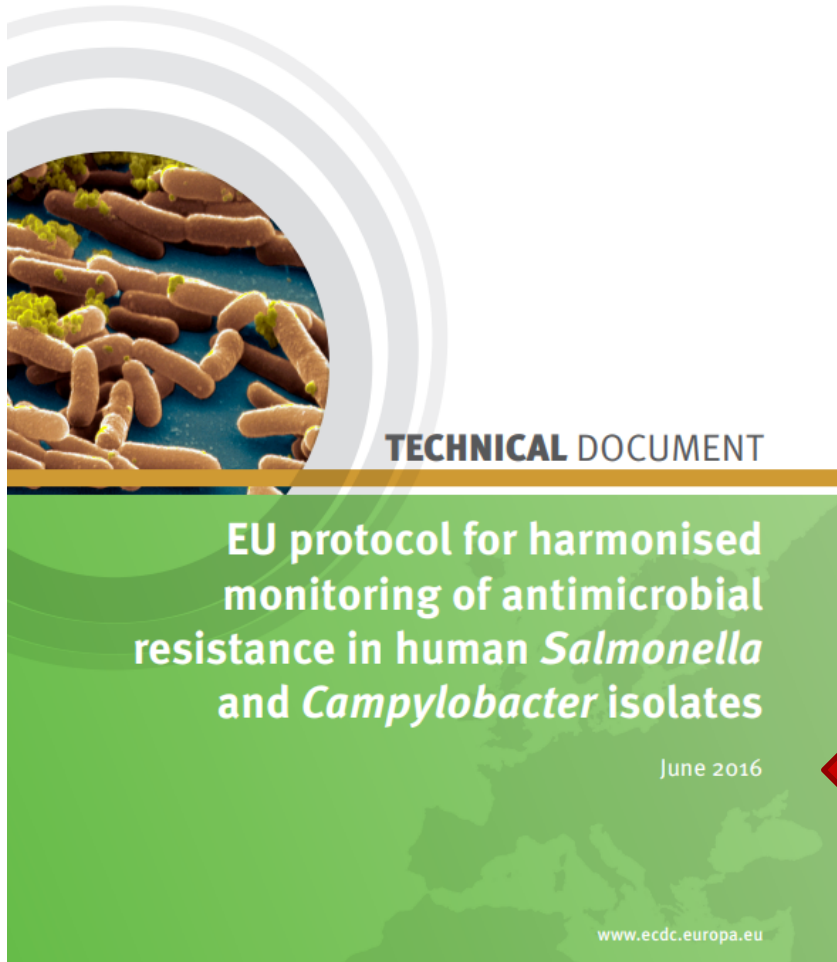
<https://www.ecdc.europa.eu/en/publications-data/eu-protocol-harmonised-monitoring-antimicrobial-resistance-human-salmonella-and-0>



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“The content of this report was developed at three expert workshops arranged by ECDC. The report was sent for consultation to the Food- and Waterborne Diseases and Zoonoses network.”

- a) To monitor, in human clinical isolates, trends in the occurrence of resistance to antimicrobial agents **relevant for treatment of human** *Salmonella* and *Campylobacter* infections, including comparison with food/animal isolates
- b) To monitor, in human clinical isolates, trends in the occurrence of resistance to **other antimicrobial agents** of public and animal health importance, including comparison with food/animal isolates
- c) To monitor, in human clinical isolates, the prevalence of ESBL, plasmid-encoded Ambler class C β lactamases (pAmpC) and carbapenemase phenotypes
- d) To use antimicrobial resistance patterns to characterise human clinical isolates, i.e. as an epidemiological marker, to **support identification of** outbreaks and related cases

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- e) To identify and monitor, in human clinical isolates, genetic determinants of resistance that are important for public health e.g. to aid recognition of epidemic cross-border spread of multi-drug resistant *Salmonella* strains
- f) To monitor, in human clinical isolates, trends in the occurrence of resistance to antimicrobial agents that may be needed for **future therapeutic** use

Data should be reported quantitatively (mm or mg/l)

- No specific requirements for the extent of surveillance/monitoring are defined in the EU harmonized protocol
- One of the tasks for the FVD AMR-RefLabCap project is to propose minimum requirements for national AMR surveillance

ANTIMICROBIALS FOR HUMAN SALMONELLA ISOLATES (1)

Class	Name (abbreviation*)	Surveillance objectives	Comments
First priority			
Aminoglycosides	Gentamicin (GEN)	b, d	
Aminopenicillins	Ampicillin (AMP)	a, b, d	
Amphenicols	Chloramphenicol (CHL)	a, d	
Carbapenems	Meropenem (MEM)	a, b, c, d, e	EUCAST recommend meropenem as it offers the best compromise between sensitivity and specificity in terms of detecting carbapenemase-producers
Cephalosporins	Cefotaxime (CTX)	a, b, c, d, e	May be insensitive for detection of ceftazidimase-type ESBLs
	Ceftazidime (CAZ)	a, b, c, d, e	Added to increase sensitivity of screening for full range of ESBL with diverse substrate specificities
Dihydrofolate reductase inhibitors	Trimethoprim (TMP)	d	Value as an epidemiological marker, e.g. in the resistance pattern ASuT common among <i>S. Typhimurium</i> .
Macrolides	Azithromycin (AZM)	f	May be considered as a last resort drug for invasive salmonellosis.

ANTIMICROBIALS FOR HUMAN SALMONELLA ISOLATES (2)



Class	Name (abbreviation*)	Surveillance objectives	Comments
First priority			
Polymyxins	Colistin (COL)	b	<p>Last-resort drug in human medicine and extensively used in animal medicine. Plasmid-mediated resistance detected in <i>E. coli</i> and <i>Salmonella</i> in Europe in 2015. Its chemical properties however cause unreliable results with dilution and render it impossible to test with disk diffusion. Please follow the dilution method agreed between CLSI and EUCAST [10].</p> <p>Note: Any laboratory that wants to report an isolate as resistant to colistin must get the result confirmed at a reference laboratory that is up to date with the latest method developments for testing of colistin.</p>
Quinolones	Ciprofloxacin (CIP)/pefloxacin (PEF)	a, b, c, d, e	<p>Preferably test ciprofloxacin with broad MIC range. For disk diffusion, EUCAST recommend screening with pefloxacin [11] since ciprofloxacin is poor at detecting low-level fluoroquinolone resistance in <i>Salmonella</i> spp. with this method and nalidixic acid is often not detecting plasmid-mediated fluoroquinolone resistance [12]. Only for isolates having the <i>aac(6)-Ib-cr</i> gene, pefloxacin does not work well.</p>
Sulphonamides	Sulfamethoxazole (SMX)	d	<p>Value as an epidemiological marker, e.g. in the resistance pattern ASuT common among <i>S. Typhimurium</i>. No ECOFF available however due to methodological problems and little harmonisation between disk manufacturers.</p>
Tetracyclines	Tetracycline (TCY)	b, d	Used both in veterinary and human medicine.
	Tigecycline (TGC)	f	

OPTIONAL ANTIMICROBIALS FOR HUMAN SALMONELLA ISOLATES



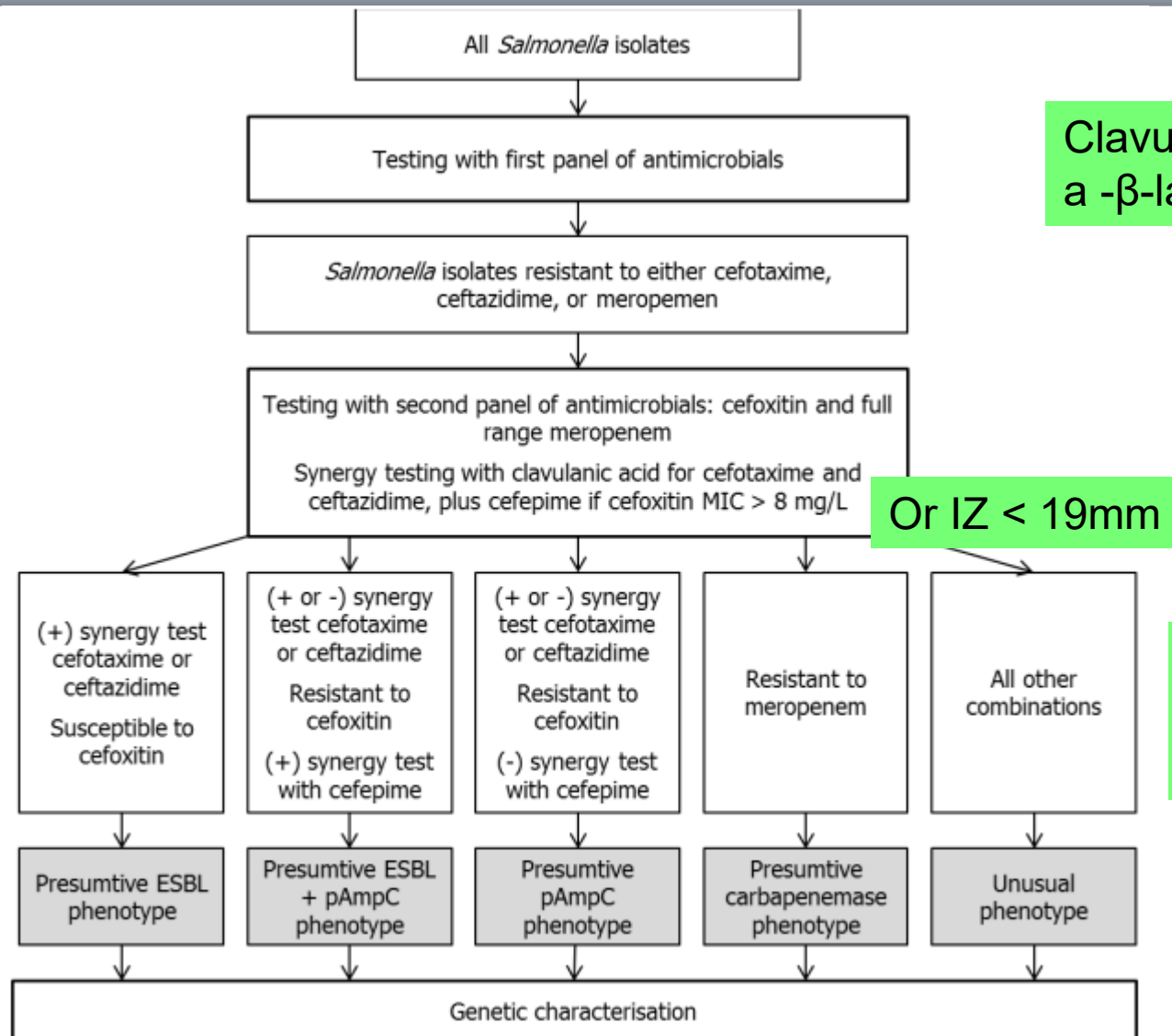
Optional			
Aminopenicillins	Amoxicillin (AMX)		Alternative for testing and reporting if AMP not tested.
Carbapenems	Ertapenem (ETP)		Many human laboratories test for ertapenem so should be possible to report.
Cephalosporins	Ceftriaxone (CRO)	a, b, c, d, e	Alternative for cefotaxime with disk diffusion method as has similar spectrum of activity.
Combination drugs	Trimethoprim + sulfamethoxazole (co-trimoxazole) (SXT)		No need to test if the substances are tested separately.
Quinolones	Nalidixic acid (NAL)		For laboratories using disk diffusion, nalidixic acid (NAL) can be tested in addition to pefloxacin for easier identification of QRDR mutations (<i>gyr</i> and <i>par</i>) since such mutations may result in clinical treatment failure (Le Hello, Institut Pasteur Paris, personal communication, Sep 2015).

EUCAST CLINICAL BREAKPOINTS AND EPIDEMIOLOGICAL CUT-OFF VALUES FOR THE PRIORITY LIST OF ANTIMICROBIALS TO BE TESTED FOR SALMONELLA ENTERICA AS OF 31 AUGUST 2021



Antimicrobial	Criteria based on MIC dilution (mg/L)			Recommended concentration range ¹ (mg/L) (number of wells)	Criteria based on disk diffusion (mm)			Disk load (µg)
	S≤	R>	NWT>		S≥	R<	NWT<	
First priority								
Ampicillin (AMP)	8.0	8.0	4.0	1-32 (6)	14	14	18	10
Azithromycin (AZM)	ND	ND	16	2-64 (6)	ND	ND	12	15
Cefotaxime (CTX)	1.0	2.0 (1.0) ²	0.5	0.25-4 (5), 0.25-64 (9) ³	20	17 (21) ²	20	5
Ceftazidime (CAZ)	1.0 ²	4.0 (1.0) ²	2.0	0.25-8 (6), 0.25-128 (10) ³	22 ³	19	20	10
Chloramphenicol (CHL)	8.0	8.0	16.0	8-64 (4)	17	17	19	30
Ciprofloxacin (CIP)	0.06	0.06	0.064	0.015-8 (10)	NA	NA	NA	NA
Colistin (COL)	2.0	2.0	NA	1-16 (5)	NA	NA	NA	NA
Gentamicin (GEN)	2.0	2.0	2.0	0.5-16 (6)	17	17	17	10
Meropenem (MEM)	2.0	8.0	0.06 (0.125) ²	0.03-16 (10)	22	16	27 (28) ²	10
Pefloxacin	NA	NA	NA	NA	24	24	24	5
Sulfamethoxazole (SMX)	ND	ND	ND	8-512 (7)	ND	ND	ND	100
Tetracycline (TCY)	ND	ND	8.0	2-32 (5)	ND	ND	17	30
Tigecycline (TGC)	ND	ND	ND	0.25-8 (6)	ND	ND	16	15
Trimethoprim (TMP)	4.0	4.0	2.0	0.25-16 (7)	15	15	23	5
Second level testing ESBL-producers								
Cefepime (FEP)	1.0	4.0	ND		27	24	ND	30
Cefoxitin (FOX)	ND	ND	8.0 ²	0.5-64 (8)	19	19 ²	21	30
Optional								
Amoxicillin (AMX)	8.0	8.0	4.0		ND	ND	ND	10
Ceftriaxone (CRO)	1.0	2.0 (1.0) ²	0.25		25	22 (23) ²	ND	30
Ertapenem (ETP)	0.5	0.5	ND (0.125) ²	0.015-2 (8)	25	25 ³	ND	10
Nalidixic acid (NAL)	ND	ND	8.0	4-64 (5)	ND	ND	16	30
Trimethoprim-sulfamethoxazole (SXT)	2.0	4.0	ND		14	11	22	1.25-23.75

SCHEMATIC VIEW OF THE PROPOSED PHENOTYPIC TESTING FOR DETECTION AND CONFIRMATION OF ESBL-, ACQUIRED AMPC, AND CARBAPENEMASE-PRODUCING *SALMONELLA* SPP.



Clavulanic acids works as a β -lactamase inhibitor

Or IZ < 19mm

Positive synergy test:
IZ > 5 mm
MIC ratio >8

Mechanisms of antibiotics

- **Bacteriostatic**

Stops growth of the infectious agent but does not kill it
The immune system has to kill the bug

- **Bactericidal**

Actively kills the infectious agent (some only growing bacteria)

Bacteriostatic antibiotic classes

- **Tetracyclines**
- **Aminoglycosides** (Gentamicin, Apramycin, Neomycin, Spectinomycin, Streptomycin)
- **Sulphonamides** (Sulphamethoxazole)
- **Macrolides** (Erythromycin)
- **Amphenicols** (Chlorphenicol, Florphenicol)
- **Trimethoprim**

Bactericidal antibiotics classes

- Beta-lactams
- **Penicillins** (ampicillin, methicillin)
 - **Cephalosporins** (Cefotaxime, Ceftazidime, Ceftiofur)
 - **Monobactams** (Aztreonam)
 - **Carbapenems** (Imipenem, Meropenem, Ertapenem)
 - **Quinolones** (Nalidixan)
 - **Fluoroquinolones** (Ciprofloxacin)
 - **Polymoxins** (Colistin)

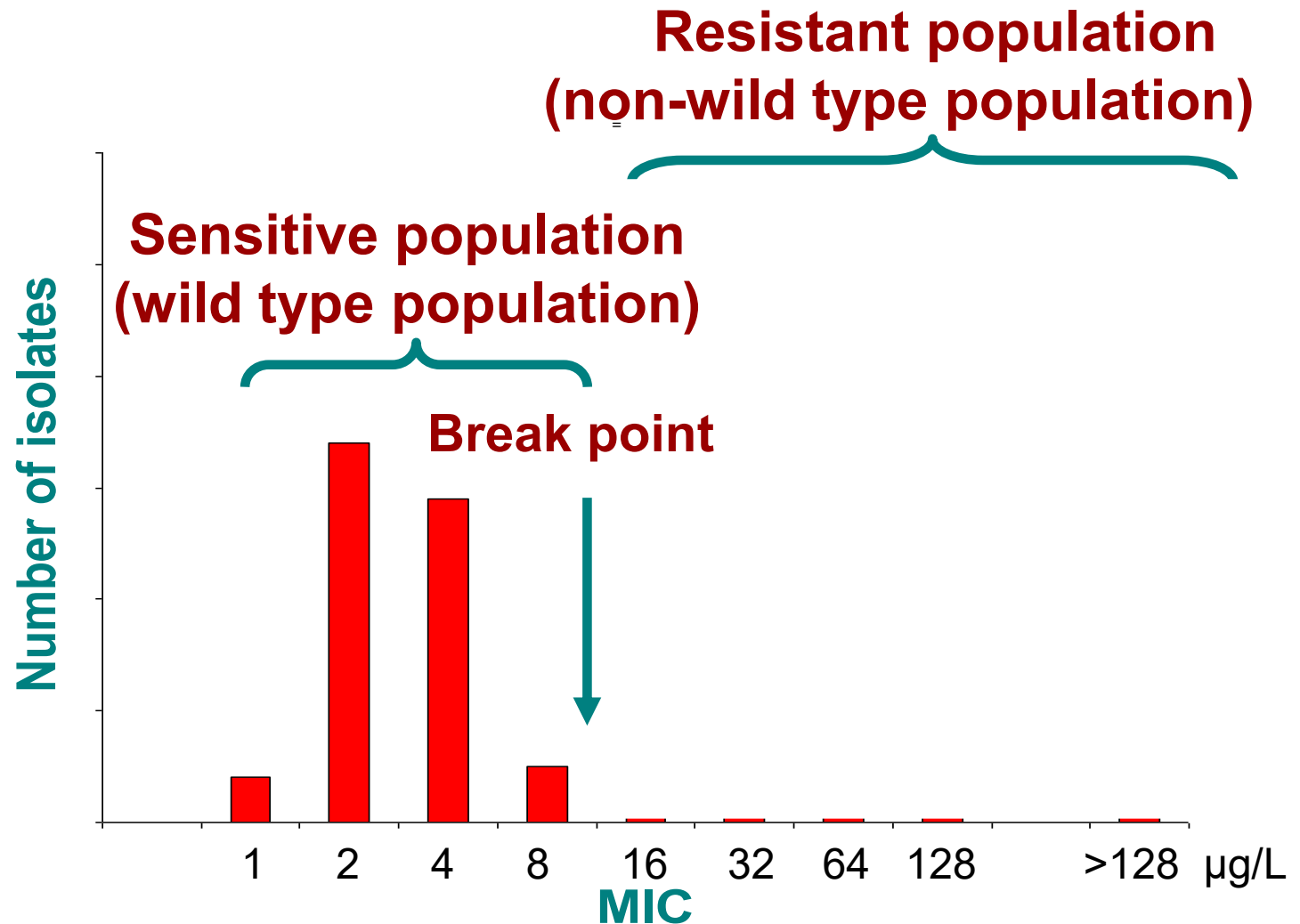
What is antimicrobial resistance I?

The ability of a microorganism to survive at a given concentration of an antimicrobial agent at which the wild type population of the microorganism would be killed

This is called the
“epidemiological/microbiological breakpoint”.

EUCAST* defines epidemiological breakpoints – ECOFFs

Population distribution



MIC > Breakpoint → Resistant ($R > 8$ or $R \geq 16$)

What is antimicrobial resistance II?

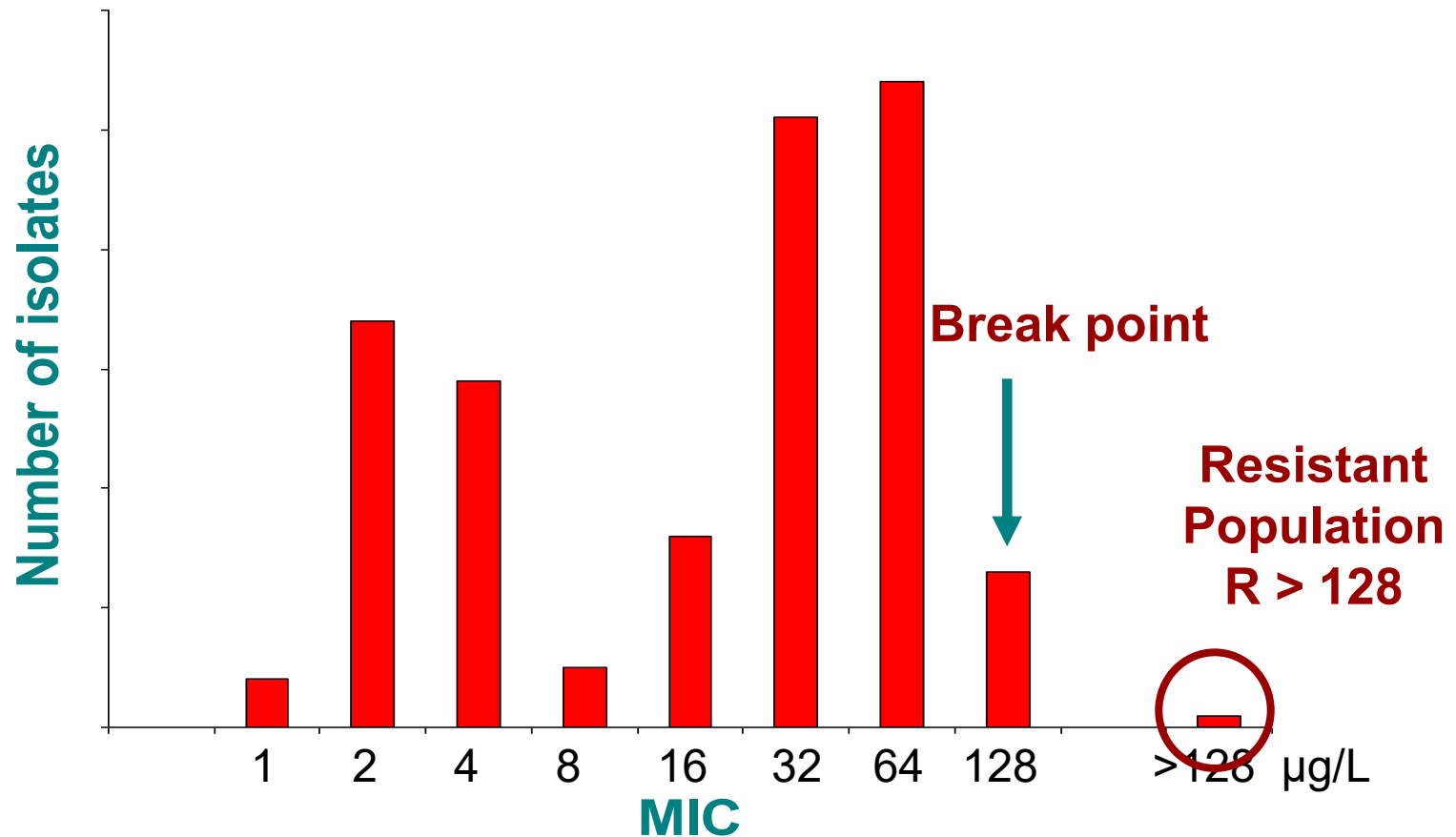
The ability of a microorganism to survive treatment with a clinical concentration of an antimicrobial agent in the body.

This is called the
“Clinical breakpoint”.

EUCAST and CLSI* is defining the clinical breakpoints.

Population distribution

Drug concentration in infection site: 128 $\mu\text{g/L}$



MIC > Breakpoint \rightarrow Resistant ($R > 128$)



mic.eucast.org/search/?search%5Bmethod%5D=mic&search%5Bantibiotic%5D=-1&search%5Bspecies%5D=431&search%5Bdisk_content%5D=-1&search%5Blimit%5D=50

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MIC EUCAST

Login

Antimicrobial wild type distributions of microorganisms

Mic distributions include collated data from multiple sources, geographical areas and time periods and can never be used to infer rates of resistance

Search database

Method

☒ MIC ☐ Disk diffusion

Antimicrobial

Species

Antimicrobial ...

Salmonella enterica

Elements per page 50

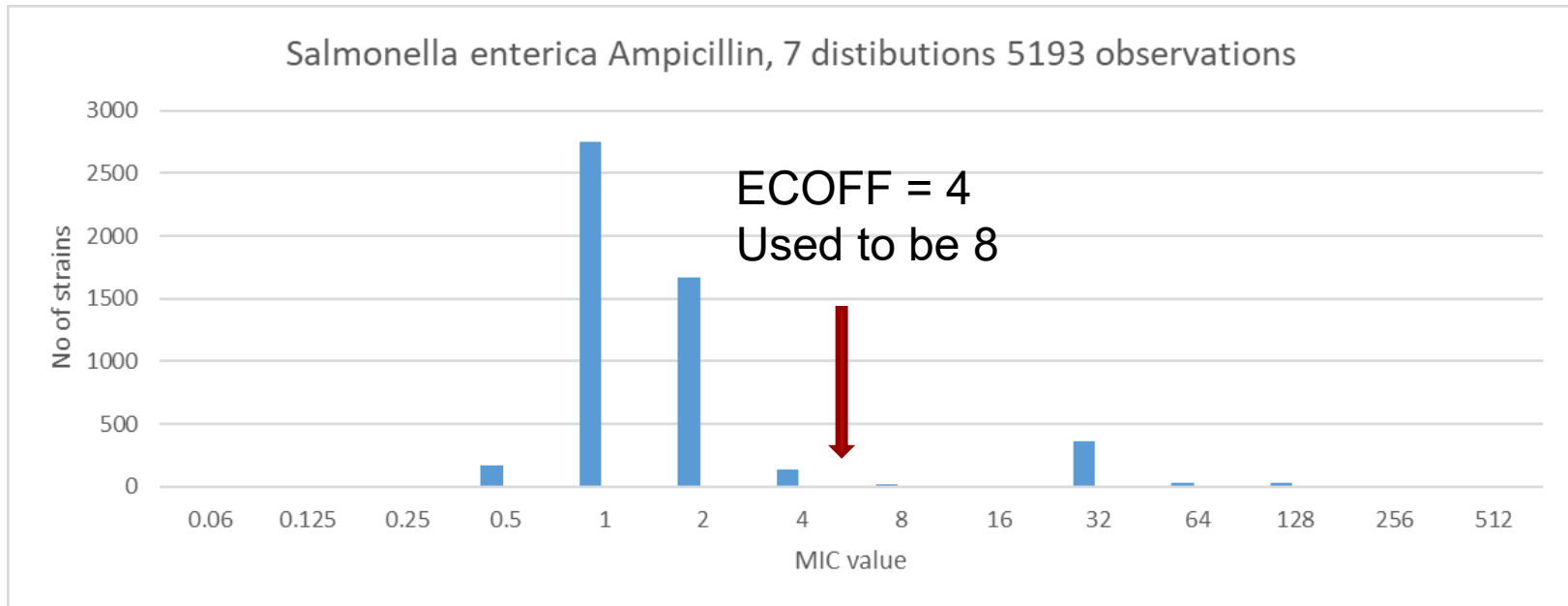
MIC distributions for Salmonella enterica, 2022-05-15

Species: Salmonella enterica (Method: MIC)

	0.002	0.004	0.008	0.016	0.03	0.06	0.125	0.25	0.5	1	2	4	8	16	32	64	128	256	512	Distributions	Observations	(T)ECOFF	Confidence interval
Amikacin	0	0	0	0	0	0	0	1	2358	9523	3498	376	26	0	1	0	1	0	0	3	15784	(4)	1 - 16
Amoxicillin	0	0	0	0	0	0	0	6	2418	5791	207	6	4	2	14	1148	553	0	0	6	10149	4	1 - 2
Ampicillin	0	0	0	0	0	0	0	12	172	2752	1673	137	16	11	360	32	28	0	0	7	5193	4	1 - 4
Apramycin	0	0	0	0	0	0	0	0	0	2	2	14	97	48	3	0	0	0	0	1	166	-	
Azithromycin	0	0	0	0	0	0	2	2	14	116	3672	13721	2577	176	19	0	0	0	0	7	20299	16	4 - 16
Aztreonam	0	0	0	0	12	37	18	3	1	0	0	0	0	0	0	0	0	0	0	2	71	-	
Cefalothin	0	0	0	0	0	0	0	0	3	4	1954	2554	603	162	57	13	10	0	0	4	5360	(16)	
Cefazolin	0	0	0	0	0	0	0	0	4	88	87	38	2	0	0	0	0	0	0	5	219	4	1 - 4
Cefepime	0	0	0	2	38	234	63	33	18	3	2	5	2	0	0	0	0	0	0	4	400	(0.25)	0.06 - 0.25
Cefixime	0	0	0	0	0	0	10	15	1	2	0	0	0	0	0	0	0	0	0	1	27	-	

[Antimicrobial wild type distributions](#)

SALMONELLA ENTERICA AMP MIC DISTRIBUTION



Data from EUCAST
2022-05-12

EUCAST CLINICAL BREAKPOINTS: NEW DEFINITIONS OF S, I AND R FROM 2019

- S - Susceptible, standard dosing regimen: A microorganism is categorised as "Susceptible, standard dosing regimen", when there is a high likelihood of therapeutic success using a standard dosing regimen of the agent.
- I - Susceptible, increased exposure*: A microorganism is categorised as "Susceptible, Increased exposure*" when there is a high likelihood of therapeutic success because exposure to the agent is increased by adjusting the dosing regimen or by its concentration at the site of infection.
- R - Resistant: A microorganism is categorised as "Resistant" when there is a high likelihood of therapeutic failure even when there is increased exposure.
- ATU: The Area of Technical Uncertainty

EUCAST CLINICAL BREAKPOINTS AND EPIDEMIOLOGICAL CUTOFF VALUES FOR THE PRIORITY LIST OF ANTIMICROBIALS TO BE TESTED FOR SALMONELLA ENTERICA AS OF 31 AUGUST 2021

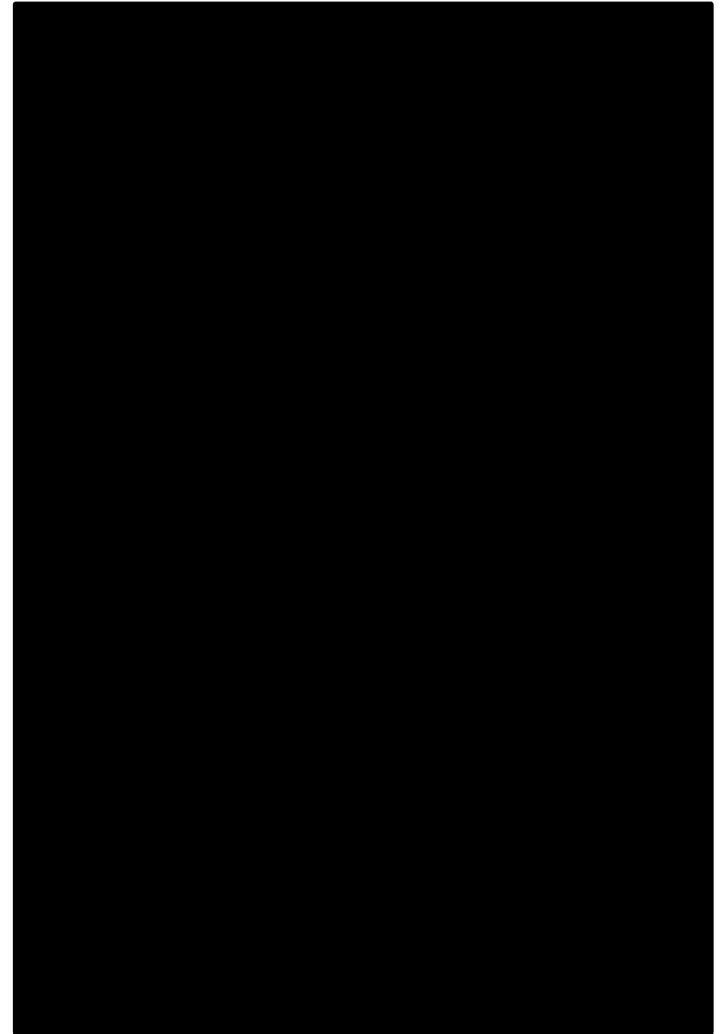
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HOW DO WE MEASURE ANTIMICROBIAL SUSCEPTIBILITY *IN VITRO*?

Phenotypic methods

- Agar diffusion method
 - Disks (tablet) mm
 - Gradient strips quantitative

- Dilution methods (quantitative)
 - Liquid media
 - MicroBrothDilution
 - Solid media



- Dilution methods - minimum inhibitory concentration (MIC) is determined (mg/L) is considered the **gold standard** for AST by EUCAST

ISO 20776-1:2019

- Disk diffusion – inhibition zones in mm - according to EUCAST guidelines v10 (1 January 2022)

- Gradient strips (MIC) – according to EUCAST and producer – should be validated
- Other methods, e.g. Trek sensititre, Vitek should be validated

Validation protocol:

ISO 20776-2:2021

- Website [EUCAST: EUCAST](#)
- Disk diffusion methodology [EUCAST: Disk diffusion methodology](#)
- Broth microdilution reading guide [EUCAST: MIC determination](#)
- QC tables [EUCASTQuality: Control](#)
- Breakpoint table
 - [EUCAST: Clinical breakpoints and dosing of antibiotics](#)
 - [V. 12 v 12.0 Breakpoint Tables.xlsx \(live.com\)](#)
- ECOFFS [EUCAST: MIC and zone distributions and ECOFFs](#)
- Warnings [EUCAST: Warnings!](#)
- Instruction videos [Instruction videos](#)

Aims:

- support the implementation of the harmonized EU AST protocol for *Salmonella* and *Campylobacter*
- assess the quality of the AST data obtained using MIC and/or DD methods in NPHRLs across Europe
- evaluation of serotyping of *Salmonella*

Objectives:

- identify common laboratory problem(s)
- assess the overall comparability of routinely collected AST results from European NPHRLs

EQA6-AST for *Salmonella*

- Participants - Laboratories in the FWD-Net
- Laboratories were asked to follow the harmonised EU AST protocol whenever possible
 - Eight strains for AST testing
 - Five mandatory antimicrobials:
Ampicillin, Cefotaxime, Meropenem, Cipro/Pefloxacin, Tetracycline
 - Possible to report ESBL-, acquired AmpC-, and carbapenemase status of the test strains – both pheno- and genotypes
 - Possible to report serotyping results

- Represented commonly reported human strains in the EU/EEA
- Were stable during the testing period in the organising laboratory
- Expected MIC and DD results were established by the EQA provider following the harmonized EU AST protocol
- DD results established using disks from Oxoid by EQA provider
- MIC values established using the micro-broth dilution based MIC system from TREK diagnostic systems© from Thermo Scientific by EQA provider

- Test results were compared to the expected results
 - *Salmonella*: MIC results within +/- one dilution difference and DD results within +/- 3 mm difference were evaluated as correct
- MIC results that were not in the relevant concentration range for comparison with expected results were not evaluated (ND)
- Qualitative results interpreted using EUCAST ECOFF and clinical breakpoints

Salmonella

25 EU/EEA countries

- 16 reported disk diffusion results
- 17 reported MIC results - broth dilution or gradient strip

Salmonella test strains EQA6 AST

Strain	Serotype	Microbiological resistance profile (NWT)*	Genotype, selected resistance genes
EQA_AST.S20.0001	Chester	CHL, CIP, COL, PEF, SMX, TCY, TMP	
EQA_AST.S20.0002	Dublin	AMP, AZM, COL, SMX, TCY	
EQA_AST.S20.0003	Stanley	AMP, AZM, CHL, CIP, GEN, PEF, SMX, TMP	
EQA_AST.S20.0004	Infantis	AMP, CEP, CAZ, CTX, FOX, CHL, CIP, GEN, PEF, NAL, SMX, TCY, TMP	<i>blaCTX_M_65</i>
EQA_AST.S20.0005	Rissen	AMP, CEP, CTX, CAZ, CHL, CIP, GEN, NAL, PEF, SMX, TEM, TCY	<i>blaCTX_M_55</i>
EQA_AST.S20.0006	Typhimurium	AMP, CEP, CTX, CHL, CIP, GEN, PEF, SMP, TCY, TMP	<i>mcr_9, blaCTX_M_9</i>
EQA_AST.S20.0007	Enteritidis	AMP, CHL, CIP, PEF, NAL, TCY	
EQA_AST.S20.0008	Heidelberg	AMP, AZM, CEP, CTX, CAZ, CHL, CIP, PEF, SMP, TCY, TMP	<i>blaCTX_M_123</i>

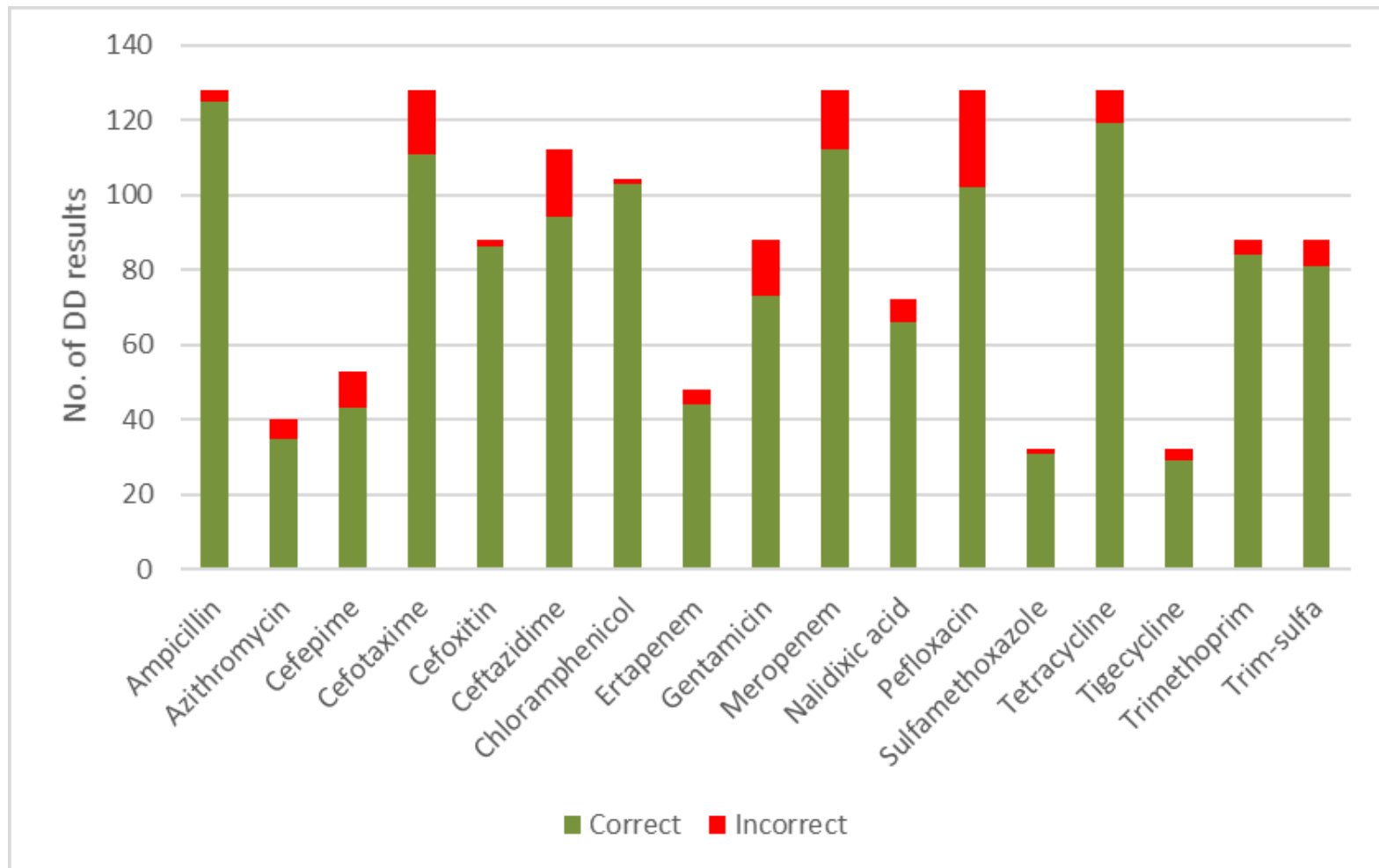
* AMP: ampicillin, AZM: Azithromycin, CEP: cefipime, CAZ: Ceftazidime, CHL: chloramphenicol, CIP: ciprofloxacin, COL: colistin, CTX: cefotaxime, FOX: ceftiofur, PEF: pefloxacin, MEM: meropenem, NAL: nalidixic acid, TCY: tetracycline, TMP: trimethoprim

EQA6-AST SALMONELLA – OVERALL RESULTS

DD and MIC results evaluated against expected quantitative and expected qualitative results using ECOFF's and clinical breakpoints

Results by DD assay	All antimicrobials	Mandatory	Optional
Expected value	1338/1485 (90%)	569/640 (89%)	769/845 (91%)
ECOFF	1204/1264 (95%)	616/640 (96%)	588/624 (94%)
Clinical breakpoints	1138/1181 (96%)	503/512 (98%)	635/669 (95%)
NA (No breakpoints)	221/304	0/128	221/176
Excluded	48	0	48
total	1533		
Results by MIC determination	All antimicrobials	Mandatory	Optional
Expected value	1240/1329 (93%)	433/458 (95%)	807/871 (93%)
ECOFF	1353/1440 (94%)	498/511 (97%)	855/929 (92%)
Clinical breakpoints	911/973 (94%)	399/407 (98%)	512/566 (90%)
NA (No breakpoints)	0/467	0/104	0/363
ND	111	53	58
Excluded	29	24	6
Total	1469		

Salmonella: 1485 quantitative DD results - antimicrobials



1338/1485 = 90% correct DD results

Salmonella: 1485 quantitative results DD – laboratory

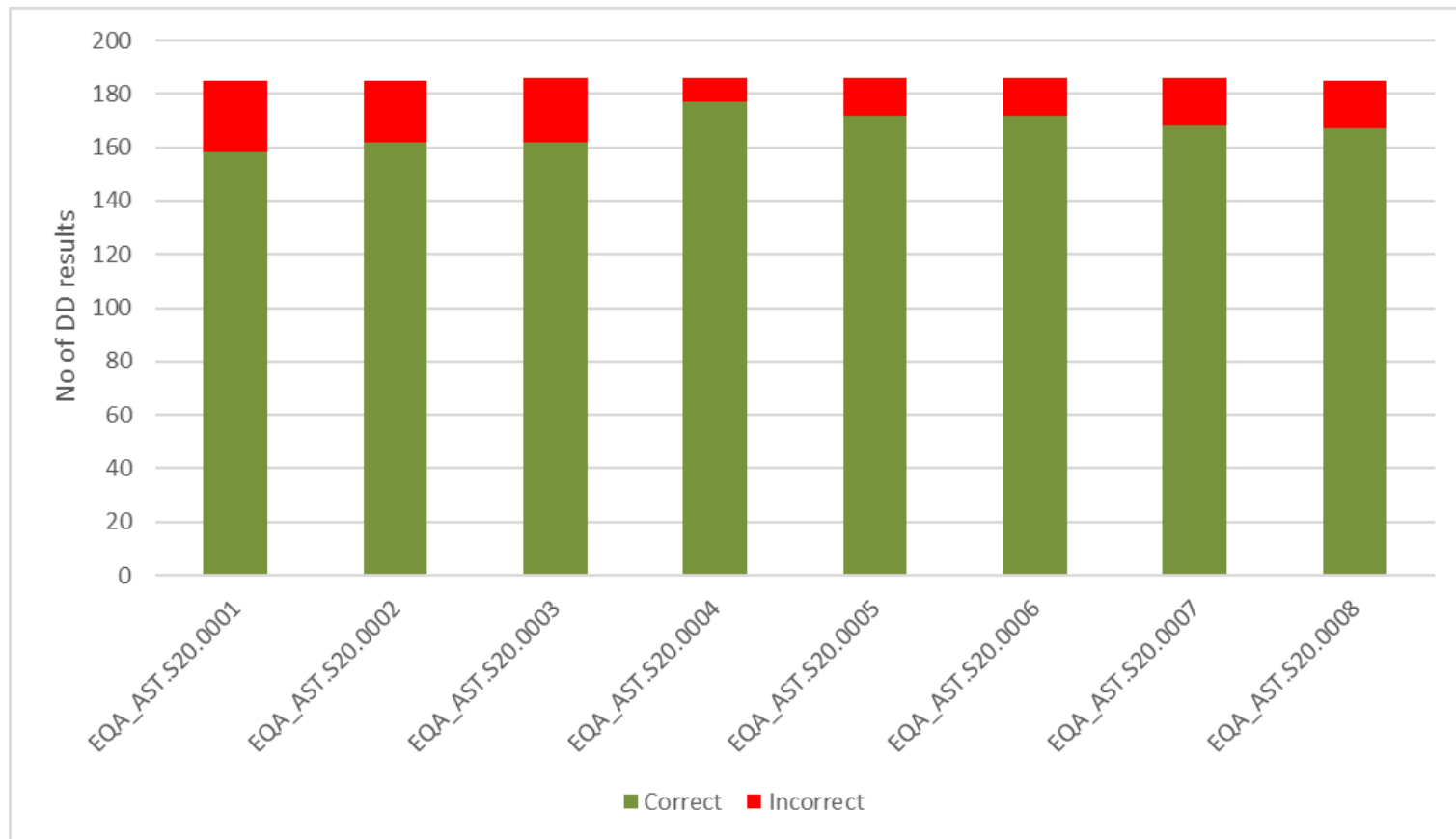
All antimicrobials



1338/1485 = 90% correct DD results

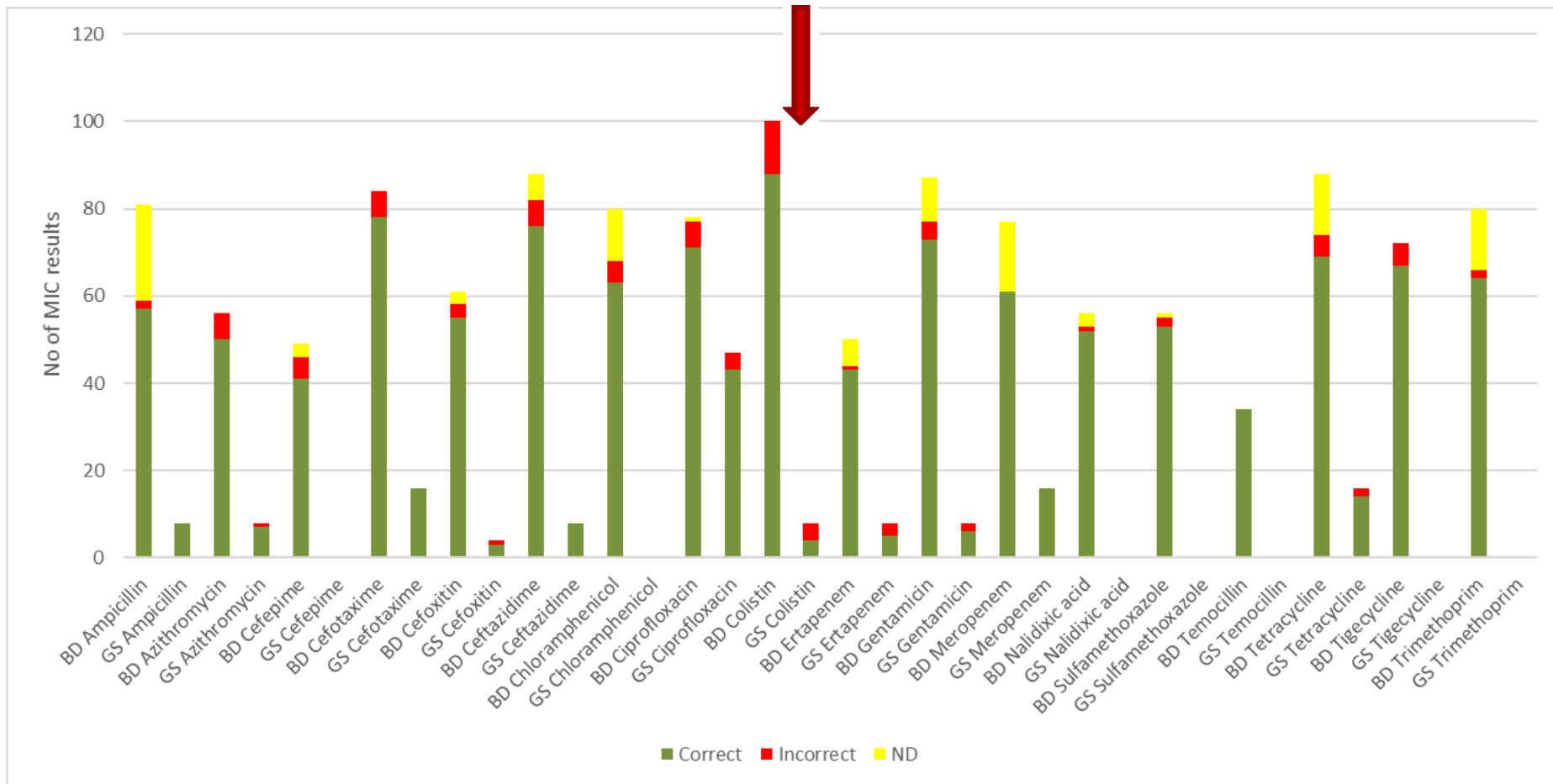
Salmonella: 1485 quantitative DD results for test strains

All antimicrobials



1338/1485 = 90% correct DD results

Salmonella: 1440 quantitative MIC results – antimicrobials and methods

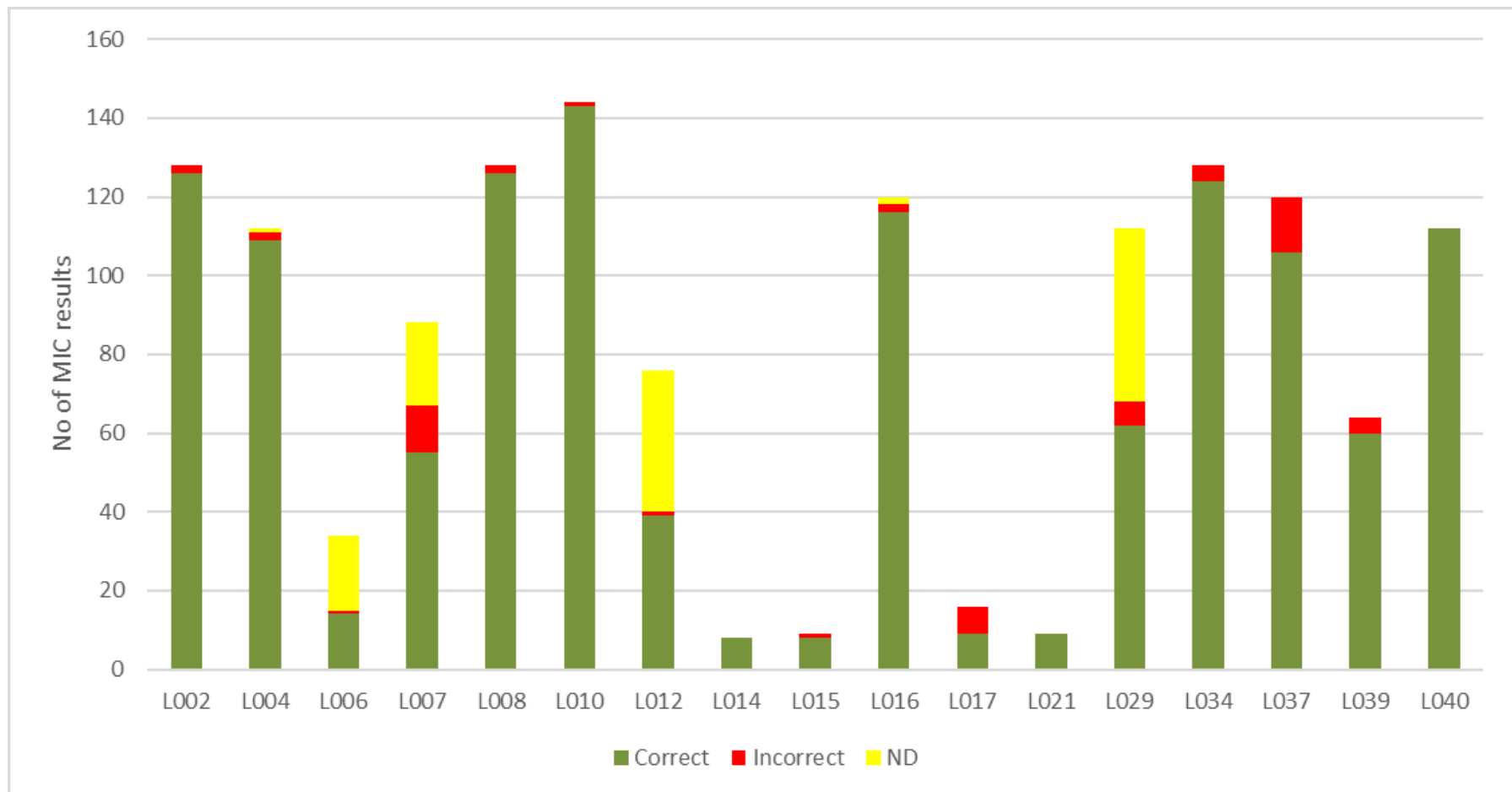


BD: Broth dilution methods
GS: Gradient strip methods

Overall 93% of evaluated MIC results correct
Most ND-results: correct ECOFF interpretation

Salmonella: 1440 quantitative MIC results – Laboratories

All antimicrobials

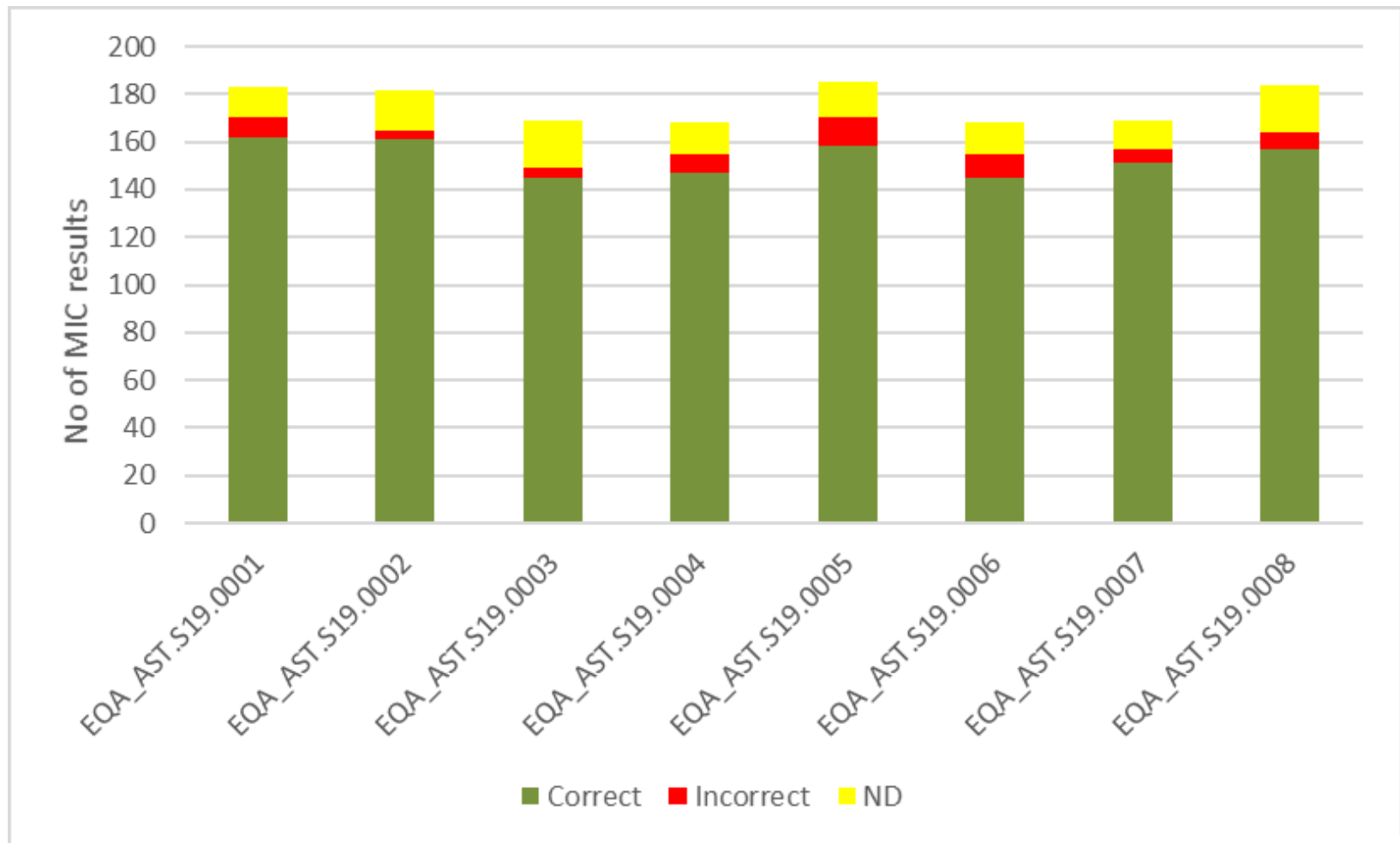


BD: Broth dilution methods
GS: Gradient strip methods

Overall 93% of evaluated MIC results correct
Most ND-results: correct ECOFF interpretation

Salmonella: 1440 quantitative MIC results – by strains

All antimicrobials



BD: Broth dilution methods
GS: Gradient strip methods

Overall 93% of evaluated MIC results correct
Most ND-results: correct ECOFF interpretation

Phenotypic prediction of ESBL-, acquired AmpC and carbapenemase-production

Strain	Expected phenotype	Number of laboratories reporting phenotype	AmpC	ESBL	Carbapenemase	Carbapenemase, AmpC	ESBL, Carbapenemase	ESBL, AmpC	ESBL, AmpC, Carbapenemase
EQA_AST.S20.0001		2		2					
EQA_AST.S20.0002									
EQA_AST.S20.0003		1		1					
EQA_AST.S20.0004	ESBL	20		17				3	
EQA_AST.S20.0005	ESBL	20		19				1	
EQA_AST.S20.0006	ESBL	19		18				1	
EQA_AST.S20.0007		1		1					
EQA_AST.S20.0008	ESBL	19		18				1	
Total		82							

25 laboratories participated in the EQA

A few of the phenotypes could not entirely be justified from the reported data

SALMONELLA SEROTYPING RESULTS

Strain	Serotype	Reported serotype		Incorrectly reported serotype
		Correct	Incorrect	
EQA_AST.S20.0001	Chester	19	1	Chartres
EQA_AST.S20.0002	Dublin	20		
EQA_AST.S20.0003	Stanley	19	1	Typhimurium
EQA_AST.S20.0004	Infantis	20		
EQA_AST.S20.0005	Rissen	19	1	Montevideo
EQA_AST.S20.0006	Typhimurium	19	1	Paratyphi B
EQA_AST.S20.0007	Enteritidis	20		
EQA_AST.S20.0008	Heidelberg	20		
Total		156	4	

Derived from WGS or based on slide agglutination

- Good correspondence between expected and reported results
 - Some laboratories deviated from the recommended testing range (MIC) and disk concentrations specified in the harmonized EU protocol.
 - Some laboratories had issues with the results for the control strain ATCC 25922
 - Colistin (MIC) results could be improved in some laboratories
 - **Results indicate that it is possible to compare phenotypic DD and MIC AST *Salmonella* results from NPHRLs across Europe**
-
- **On this course you are going to work with the EQA7-AST strains**



Thank you for your attention !!