



### Annex F – WGS data reported complementary to MIC values

#### Annex to:

EFSA (European Food Safety Authority) and ECDC (European Centre for Disease Prevention and Control), 2024. The European Union Summary Report on Antimicrobial Resistance in zoonotic and indicator bacteria from humans, animals and food in 2021/2022.

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### F.1 WGS data reported complementary to MIC values in 2021 and 2022

In 2021 and 2022, some reporting countries **reported both MIC and WGS data** for isolates showing a presumptive ESBL/AmpC/CP phenotype after susceptibility testing using panel 1. The tables presented in this Annex gives an overview of the ESBL-/AmpC-/CP-encoding genes reported for these isolates, as it was indicated by the reporting countries that the MIC values should be used for the analysis in the report.

In 2021, both MIC- and WGS data were reported for indicator commensal *Escherichia coli* collected in the routine monitoring (NL), for presumptive ESBL-/AmpC-/CP-producing *E. coli* collected in the specific monitoring for ESBL-/AmpC-/C-producing *E. coli* (Belgium, Iceland, Netherlands, Norway and Sweden), and for presumptive CP-producing *E. coli* collected in the specific monitoring for CP-producing *E. coli* (Spain). None of the countries reported both MIC- and WGS data for *Salmonella* spp. collected in the routine monitoring.

In 2022, MIC- and WGS data were reported for indicator commensal *E. coli* collected in the routine monitoring (Ireland and Norway), and for presumptive ESBL-/AmpC-/CP-producing *E. coli* collected in the specific monitoring of ESBL-/AmpC-/CP-producing *E. coli* (Austria and Ireland). None of the countries reported both MIC- and WGS data for presumptive CP-producers collected in the specific monitoring for CP-producing *E. coli* or for *Salmonella* spp. collected in the routine monitoring.







# F.2 WGS data from *Escherichia coli* isolates collected within the routine monitoring and subjected to supplementary testing in 2021

**Table 1**: WGS data in indicator *Escherichia coli* isolates from food-producing animals and meat thereof per country, 2021.

Country	Matrix	Sampling stage	ESBL family	ESBL gene		AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
Netherlands	Cattle meat	ВСР	CTX-M	CTX-M-15	-	-	-	-	161	1
Netherlands	Cattle	Slaughterhouse	CTX-M	CTX-M-1	-	-	-	-	306	1
Sweden	Fattening pigs	Slaughterhouse	-	-	AmpC mutation or insertion	C-42T	-	-	173	1

<sup>\*</sup>Cattle are cattle under one year of age, ESBL: extended-spectrum beta-lactamase, AmpC: AmpC beta-lactamase, CP: carbapenemase





### F.3 WGS data from *Escherichia coli* isolates collected within the specific monitoring of ESBL-/AmpC-/CP-producers and subjected to supplementary testing in 2021

**Table 2**: WGS data in presumptive ESBL/AmpC/CP-producing *Escherichia coli* isolates from food producing animals and meat thereof per country, 2021.

Country	Matrix	Sampling stage	ESBL family	ESBL gene	AmpC family	AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
	Dia mont	Retail	CTX-M	CTX-M-55	-	-	-	_	300	1
	Pig meat	Retail	TEM	TEM-52	-	-	-	-	300	1
	0 111			CTX-M-1	-	-	-	-	300	1
	Cattle meat	Retail	CTX-M	CTX-M-32	-	-	-	-	300	1
	incat			CTX-M-65	-	-	-	-	300	1
	Cattle	Slaughterhouse		CTX-M-1	-	-	-	-	301	4
			CTX-M	CTX-M-15	-	-	-	-	301	2
Belgium				CTX-M-32	-	-	-	-	301	4
Deigiain			-	-	AmpC mutation or insertion	AmpC mutation or insertion	-	-	300	1
				CTX-M-1	-	-	-	-	300	1
	Fattening	Slaughterhouse	CTX-M	CTX-M-14	-	-	-	-	300	2
	pigs	Sidagileariidase	CIX-M	CTX-M-32	-	-	-	-	300	1
				CTX-M-55	-	-	-	-	300	1
			SHV	SHV-12	-	-	-	-	300	1
			TEM	TEM-52	-	-	-	-	300	1
	Eattoning		-	-		CMY-2	-	-	152	1
Iceland	Fattening pigs	Slaughterhouse	-	-	CMY	CMY-22	-	-	152	1
	P193		-	-		CMY-61	-	-	152	1
Netherlands	Pig meat	Retail	CTX-M	CTX-M-1	-	-	-	-	320	3





#### Annex F- EUSR on AMR in zoonotic and indicator bacteria from humans, animals and food2021/2022

Country	Matrix	Sampling stage	ESBL family		AmpC family	AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
				CTX-M-55	-	-	-	-	320	1
	Cattle	Retail	CTX-M	CTX-M-1	-	-	-	-	583	3
	meat	Recair	0.7	CTX-M-14	-	-	_	-	583	1
				CTX-M-15	-	-	_	-	583	5
				CTX-M-32	-	-	-	-	583	1
				CTX-M-55	-	-	-	-	583	1
Netherlands			SHV	SHV-12	-	-	-	-	583	1
Netrierianus			-	-	AmpC mutation or insertion	C-42T	-	-	306	3
	Cattle	Slaughterhouse		CTX-M-1	-	-	-	-	306	29
			SE CTX-M	CTX-M-3	-	-	-	-	306	1
				CTX-M-8	-	-	-	-	306	1
				CTX-M-14	-	-	-	-	306	5
				CTX-M-15	-	-	-	-	306	45
				CTX-M-32	-	-	-	-	306	8
				CTX-M-55	-	-	-	-	306	3
				CTX-M-65	-	-	-	-	306	2
			TEM	TEM-52	-	-	-	-	306	6
			-	-	AMPC phenotype/geno type	C-42T	-	-	300	19
			-	-	CMY	CMY-2	-	-	300	2
	Fattening	Slaughterhouse		CTX-M-1	-	-	-	-	300	17
	pigs	Ciaagiicaiiicas	CTV M	CTX-M-9	-	-	-	-	300	1
			CTX-M	CTX-M-15	-	-	-	-	300	5
				CTX-M-55	-	-	-	-	300	1
			TEM	TEM-52	-	-	-	-	300	2
Norway	Pig meat	Retail	-	-	CMY	CMY-2	-	-	311	1
Norway	Cattle	Slaughterhouse	-	-	AmpC mutation or insertion	C-42T	-	-	295	3





#### Annex F- EUSR on AMR in zoonotic and indicator bacteria from humans, animals and food2021/2022

Country	Matrix	Sampling stage	ESBL family	ESBL gene	AmpC family	AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
	Fattening pigs	Slaughterhouse	-	-	AmpC mutation or insertion	C-42T	-	-	321	44
		Sidugificationse	CTX-M	CTX-M-15	-	-	-	-	321	2
			CTX-M	CTX-M-55	-	-	-	-	321	1
	Cattle meat	Retail	CTX-M	CTX-M-3	-	-	-	-	303	1
	Cattle	Slaughterhouse	CTX-M	CTX-M-15	-	-	_	-	20	2
Sweden	Fattening	Slaughterhouse	-	-	AmpC mutation or insertion	C-42T	-	-	300	22
	pigs	Sidugitterilouse	CTV M	CTX-M-1	-	-	-	-	300	1
			CTX-M	CTX-M-15	-	-	-	-	300	2

<sup>\*</sup>Cattle are cattle under one year of age, ESBL: extended-spectrum beta-lactamase, AmpC: AmpC beta-lactamase, CP: carbapenemase





# F.4 WGS data from *Escherichia coli* isolates collected within the specific monitoring of CP-producers and subjected to supplementary testing in 2021

**Table** 3: WGS data in presumptive CP-producing *Escherichia coli* isolates from food producing animals and meat thereof per country, 2021.

Country	Matrix	Sampling stage	l	ESBL gene	AmpC family	AmpC gene /mutation	<u>-</u>	CP gene	Tot units tested	No. isolates having the gene
Spain	Fattening pigs	Slaughterhouse	-	-	-	-	OXA	OXA-48	423	2

ESBL: extended-spectrum beta-lactamase, AmpC: AmpC beta-lactamase, CP: carbapenemase





# F.5 WGS data from *Escherichia coli* isolates collected within the routine monitoring and subjected to supplementary testing in 2022

Table 4: WGS data in indicator Escherichia coli isolates from food-producing animals and meat thereof per country, 2022.

Country	Matrix	Sampling stage		ESBL gene	AmpC family	AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
Ireland	Broiler meat	ВСР	CTX-M	CTX-M-55	CMY	CMY-2	-	-	7	1
Norway	Fattening turkey	Slaughterhouse	ı	-	AmpC mutation or insertion	C-42T	-	1	110	1

ESBL: extended-spectrum beta-lactamase, AmpC: AmpC beta-lactamase, CP: carbapenemase, BCP: border control post





### F.6 WGS data from *Escherichia coli* isolates collected within the specific monitoring of ESBL-/AmpC-/CP-producers and subjected to supplementary testing in 2022

**Table 5:** WGS data in presumptive ESBL/AmpC/CP-producing *Escherichia coli* isolates from food producing animals and meat thereof per country, 2022.

Country	Matrix	Sampling stage	ESBL family	ESBL gene	AmpC family	AmpC gene /mutation	CP family	CP gene	Tot units tested	No. isolates having the gene
				CTX-M-1	-	-	-	-	181	1
			CTX-M	CTX-M-15	-	_	-	-	181	2
	Turkey meat	Retail	CTX-M	CTX-M-27	-	-	-	-	181	2
				CTX-M-32	-	-	-	-	181	1
Austria			TEM	TEM-126	-	-	-	-	181	1
Austria			CTX-M	CTX-M-55	-	-	-	-	338	1_
	Broiler meat	Retail	SHV	SHV-12	-	-	-	-	338	2
	mode		3110	SHV-2a	-	-	-	-	338	1
		Claughterhouse	TEM	TEM-52B	-	-	-	-	301	3
	Broilers	Slaughterhouse	-	-	-	_	VIM	VIM-1	301	2
Ireland	Broiler meat	ВСР	CTX-M	CTX-M-8	-	-	-	-	7	2
	Broilers	Slaughterhouse	CTX-M	CTX-M-1	-	-	-	-	300	1

ESBL: extended-spectrum beta-lactamase, AmpC: AmpC beta-lactamase, CP: carbapenemase, BCP: border control post