

1 **Table S1.** Primers used in this study for amplification of the virulence factors of pathotypes of *Escherichia coli*

Virulence factor	Primers	Pathotype	T <sub>a</sub> (°C) <sup>a</sup>	Amplicon (bp)	Oligonucleotide sequence (5'--> 3')	Reference
β-D-GUD	<i>uidA</i> -F	All	55	168	TGTGCCRGGMAGTTTTAAC	this study
	<i>uidA</i> -R				TGATTATTKACCCACACTTTG	
Shiga toxin 1	<i>stx1</i> -F	EHEC	55	107	GCAAAGAMGTATGTWGATTTCG	this study
	<i>stx1</i> -R				GWGCCACTATCAATCATCAG	
Shiga toxin 2	<i>stx2</i> -F	EHEC	55	82	AATGCAAATCAGTCGTCAC	this study
	<i>stx2</i> -R				TGCATCTCTGGTCATTGTAT	
Shiga toxin 2 (variant)	<i>stx2</i> -F var	EHEC	55	82	GATGCAGATRRKCGYCAT	this study
	<i>stx2</i> -R var				TGATGCGCGGGTCATGGAAC	
Intimin	<i>eae</i> -F	EHEC/EPEC	52	92	GCTATAACRTCTTCATTGATC	this study
	<i>eae</i> -R				RCTACTTTTTRAAATAGTCTCG	
Enterohaemolysin	<i>ehxA</i> -F	EHEC/EPEC	55	86	GCACCACAACCTTGAYAAACT	this study
	<i>ehxA</i> -R				CCAGATTATTACCTACATTYTCAG	
ST	<i>est</i> -F	ETEC	54	69/72	TGAAAGCATGAATRGTAGCAA	this study
	<i>est</i> -R				TTAATAACATSSAGCACAGG	
ST variant	<i>est</i> -F var	ETEC	54	69/73	TCAGAAAATATGAAYAACACATT	this study
	<i>est</i> -R var				TAATAGCACCCGGTACAAG	

LT	<i>elt</i> -F	ETEC	54	142	GGYAAAAGAGAAATGGTTAT	this study
	<i>elt</i> -R				TCTCGGTCAGATATGYGATTC	
Bundle- forming pilus ( <i>bfpA</i> )	<i>bfpA</i> -F	EPEC	53	109	CMGGTGTGATGTTTTACTAC	this study
	<i>bfpA</i> -R				TGCCCAATATACARACCAT	
Invasion plasmid ( <i>spa24</i> )	<i>pInv</i> -F	EIEC	53	159	CCAATCACAATATCAGTACCA	this study
	<i>pInv</i> -R				AAAGAGCCTTATTACCCATAT	
Enteroaggregative regulator	<i>aggR</i> -F	EAEC	56	94	TTTATCGCAATCAGATTAARC	this study
	<i>aggR</i> -R				GGACAACRCAAGCATCTAC	
<i>rbf</i> <sub>O157</sub>	<i>rbf</i> <sub>O157</sub> -F	EHEC	55	125	CAAAAGGAAACTATATTCAGAAGT	(18)
	<i>rbf</i> <sub>O157</sub> -R				CGATATACCTAACGCTAACAA	
<i>fliC</i> <sub>H7</sub>	<i>fliC</i> <sub>H7</sub> -F	EHEC	55	91	CGACAGGTCTTTATGATCTGA	(18)
	<i>fliC</i> <sub>H7</sub> -R				ACTGTGACTTTATCGCCATT	
<i>wzx</i> <sub>O104</sub>	<i>wzx</i> <sub>O104</sub> -F	EAEC	55	99	GCGCAAAGAATTTCAACTT	(19)
	<i>wzx</i> <sub>O104</sub> -R				TGTAAAATCCTTTAAACTATACG	
<i>fliC</i> <sub>H4</sub>	<i>fliC</i> <sub>H4</sub> -F	EAEC	55	192	CTGGGGGTAAACAAGTCAA	(19)
	<i>fliC</i> <sub>H4</sub> -R				CCAGTGCTTTTAACGGATC	

2 <sup>a</sup> T<sub>a</sub>, annealing temperature.

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4 **Table S2.** Antimicrobial resistance rates found in *Escherichia coli* isolates depending on the host species.

Antimicrobial	Concentration range (mg/L)	Cut-off <sup>a</sup>	Pigs		Broiler		Cattle			
			Non-O157:H7 <sup>b</sup> (n=278) <sup>c</sup>		Non-O157:H7 (n=196)		Non-O157:H7 (n=256)		O157:H7 (n= 50)	
			NR	%R	NR	%R	NR	%R	NR	%R
Ampicillin	0.5-32	8	192	69.1	139	70.9	40	15.6	17	34.0
Cefotaxime	0.06-4	0.25	3	1.1	51	26.0	0	0.0	0	0.0
Ceftazidime	0.25-16	0.5	4	1.4	45	23.0	0	0.0	0	0.0
Nalidixic Acid	4-64	16	51	18.3	164	83.7	8	3.1	17	34.0
Ciprofloxacin	0.008-8	0.06	81	29.1	169	86.2	9	3.5	17	34.0
Streptomycin	2-128	16	211	75.9	133	67.9	91	35.5	32	64.0
Kanamycin	4-128	8	45	16.2	24	12.2	7	2.7	13	26.0
Gentamicin	0.25-32	2	20	7.2	20	10.2	8	3.1	17	34.0
Chloramphenicol	2-64	16	76	27.3	37	18.9	25	9.8	18	36.0
Florphenicol	2-64	16	5	1.8	1	0.5	15	5.9	1	2.0

Tetracycline	1-64	8	251	90.3	147	75.0	125	48.8	32	64.0
Sulphonamide	8-1024	64	195	70.1	105	53.6	90	35.2	32	64.0
Trimethoprim	0.5-32	2	198	71.2	93	47.4	45	17.6	32	64.0
Colistin	2-4	2	0	0.0	0	0.0	0	0.0	0	0.0

5 Abbreviations: NR, number of resistant isolates; %R, percentage of resistant isolates; NA, Not applicable.

6 <sup>a</sup> EUCAST cut-off values ([http://www.eucast.org/mic\\_distributions/](http://www.eucast.org/mic_distributions/))

7 <sup>b</sup> Non-O157:H7: *E. coli* belonging to other serotypes different than O157:H7.

8 <sup>c</sup> For colistin in pigs only 156 isolates were tested.