

# MLST-Based Analysis and Resistance to Antimicrobial Agents of *Staphylococcus epidermidis* from Cases of Sheep Mastitis in Greece

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**Table S1.** Primers used for detection of genes for inclusion in the MLST database, in 33 *S. epidermidis* isolates recovered from subclinical mastitis in ewes in Greece (MLST: Multi Locus Sequence Typing).

Gene	Primers	Sequences	Product size (bp)
<i>arcC</i> (carbamate kinase)	arcC-F arcC-R	TGTGATGAGCACCGTACCGTTAG TCCAAGTAAACCCATCGGTCTG	465
<i>aroE</i> (shikimate dehydrogenase)	aroE-F aroE-R	CATTGGATTACCTCTTGTCAGC CAAGCGAAATCTGTTGGGG	420
<i>gtr</i> (ABC transporter)	gtr-F gtr-R	CAGCCAATTCTTTATGACTTT GTGATTAAAGGTATTGATTGAAT	438
<i>mutS</i> (DNA mismatch repair protein)	mutS-F3 mutS-R3	GATAATAAGAATAAGGGTTGTGAA GTAATCGTCTCAGTTATCATGTT	412
<i>pyrR</i> (pyrimidine operon regulatory protein)	pyr-F2 pyr-R4	GTTACTAATACTTTGCTGTGTTT GTAGAAATGTAAAGAGACTAAAATGAA	428
<i>tpiA</i> (triosephosphate isomerase)	tpi-F2 tpi-R2	ATCCAATTAGACGCTTAGTAAC TTAATGATGCGGCCACCTACA	424
<i>yqiL</i> (acetyl coenzyme A acetyltransferase)	yqiL-F2 yqiL-R2	CACGCATAGTATTAGCTGAAG CTAATGCCTTCATCTTGAGAAATAA	416

**Table S2.** Primers used and work conditions undertaken for detection of resistance genes in 33 *S. epidermidis* isolates recovered from subclinical mastitis in ewes in Greece.

Gene	Primer sequence	Concentration ( $\mu\text{M}$ )	Product size (bp)	Annealing temperature ( $^{\circ}\text{C}$ )	Reference
<i>ermA</i>	Fw-TCTAAAAGCATGTAAAAGAA Rv-CTTCGATAGTTATTAAATTAGT	0.2	645	55	[1]
<i>ermB</i>	Fw-GAAAAGGATCTAACCAAATA Rv-AGTAACGGTACTTAAATTGTTAC	0.2	639	55	[1]
<i>ermC</i>	Fw-TCAAAACATAATATAGATAAA Rv-GCTAATATTGTTAAATCGTCAAT	0.2	649	52	[1]
<i>ermT</i>	Fw-GGTGTAATTATGTAACCGCCA Rv-ACTTCCTGTAGCTGTGCTTTC	0.2	218	52	[1]
<i>ermY</i>	Fw-ATTACTTCGAAACGTAATATAGAT Rv-ATAGCTATTGAAAAGAGACAAGA	0.2	~ 700	60	[2]
<i>lnuA</i>	Fw-GGTGGCTGGGGGGTAGATGTATTAACGG Rv-GCTTCTTTGAAATACATGGTATTTTCGA	0.2	323	55	[3]
<i>lnuB</i>	Fw-CCTACCTATTGTTGAA Rv-ATAACGTTACTCTCCTATT	0.2	944	50	[4]
<i>lnuC</i>	Fw-GTAGATGCTCTTCTGGAT Rv-TTCTACCGGAAAACAATCC	0.2	246	55	[5]
<i>lsaA</i>	Fw-GATCAGGCAAATATCACGATG Rv-CTTCATGAAATGTGCATCATGC	0.2	~ 1400	60	[6]
<i>mphC</i>	Fw-ATCTCATTGAATGAATCAGGAC Rv-CTACTCTTCCATACCTAACTC	0.2	~ 850	60	[2]
<i>msrA</i>	Fw-GCAAATGGTGTAGGTAAGACAAC Rv- ATCATGTGATGTAAACAAAAT	0.2	399	55	[1]
<i>tetK</i>	Fw-TATTTGGCTTGATTCTTCAT Rv-GCTATACCTGTTCCCTCTGATAA	0.2	1159	59	[7]
<i>tetL</i>	Fw-ATAAATTGTTCGGGTCGGTAAT Rv-AACCAGCCAACTAATGACAATG	0.2	1077	59	[7]
<i>tetM</i>	Fw-GAACTCGAACAAAGAGGAAAGC Rv-ATGGAAGCCCAGAAAGGAT	0.2	740	54	[8]
<i>tetS</i>	Fw-ATCAAGATATTAAGGAC Rv-TTCTCTATGTGGTAATC	0.2	573	55	[9]

<i>tetT</i>	Fw-CAGTGCAGATATAAGGACACGTC Rv-CAAGCCTCTCTACAGCATC	0.2	644	61	[10]
<i>vgaA</i>	Fw-CTTGTCTCCCGAATAC Rv-AGTGGTGGTGAAGTAACACG	0.2	659	55	[11]

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**Table S3.** Husbandry factors applied in sheep flocks, which were evaluated for potential association with resistance to antimicrobial agents of 33 *S. epidermidis* isolates recovered from subclinical mastitis in ewes in Greece.

Management system applied in the flock (description according to EFSA classification) [1]
Stage of the lactation period at sampling (month)
Milking technique applied in the flock (machine milking, hand-milking)
Application of post-milking teat dipping (yes, no)
Intramammary administration of antimicrobial agents at the end of the lactation period (yes, no)
Vaccination against staphylococcal mastitis (yes, no)

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**Table S4.** Identity and details in the MLST database of 33 *S. epidermidis* isolates recovered from subclinical mastitis in ewes in Greece (MLST: Multi Locus Sequence Typing).

Isolate ID	Alleles of relevant genes							ST
	<i>arcC</i>	<i>aroE</i>	<i>gtr</i>	<i>mutS</i>	<i>pyrR</i>	<i>tpiA</i>	<i>yqiL</i>	
1134	2	1	1	17	20	1	1	709
1135	8	51	56	37	9	16	1	710
1136	1	1	57	6	2	16	7	711
1143	65	59	5	5	8	48	11	700
1316	12	25	9	8	6	5	8	315
1317	1	1	2	2	3	1	3	200
1318	2	1	2	2	2	1	1	142
1319	2	1	2	2	2	1	1	142
1320	2	1	2	2	2	1	1	142
1321	1	1	2	6	2	1	1	152
1323	1	1	2	6	2	1	1	152
1325	1	1	2	6	2	1	1	152
1327	1	1	2	1	2	1	8	454 <sup>1</sup>
1328	2	1	1	2	20	1	1	100
1329	2	1	1	2	20	1	1	100
1330	2	1	1	2	20	1	1	100
1331	2	1	1	1	2	1	1	59
1332	1	1	2	6	2	16	7	570
1333	2	1	2	2	2	2	1	142
1366	8	2	1	1	2	1	1	677
1367	2	1	2	2	2	1	1	142
1368	2	1	1	2	20	1	1	100
1369	2	1	1	17	20	1	1	709
1370	8	51	56	37	9	16	1	710
1371	2	1	1	2	20	1	1	100
1372	2	1	1	2	20	1	1	100
1373	2	1	1	2	20	1	1	100
1374	2	1	6	2	2	2	1	153
1375	1	1	2	6	2	1	1	152
1376	1	1	2	6	2	1	1	152
1377	65	59	5	5	8	48	11	700
1378	2	1	2	2	2	1	7	678
1379	2	1	2	2	2	1	7	678

<sup>1</sup> Red lettering indicates missing alleles (completed by means of an imputational method) and missing STs (thereafter completed as the most probable ST).

**Table S5.** Profiles of resistance to antimicrobial agents of 33 *S. epidermidis* isolates recovered from subclinical mastitis in ewes in Greece.

Isolate identity in the MLST database <sup>1</sup>	Antimicrobial agents to which resistance was recorded <sup>2</sup>
1134	P, AMP
1135	TE, FA
1136	none
1143	E, TE
1316	none
1317	P, AMP, TE
1318	P, AMP, FA
1319	P, AMP, TE
1320	P, AMP, TE
1321	P, AMP, TE
1323	P, AMP, TE
1325	none
1327	none
1328	none
1329	none
1330	none
1331	E, CC, TE, FOX
1332	none
1333	none
1366	P, AMP, TE, FA
1367	P, AMP, TE
1368	P, AMP, TE
1369	P, AMP
1370	TE, FA
1371	TE
1372	none
1373	TE
1374	none
1375	none
1376	none
1377	E, TE
1378	TE
1379	none

<sup>1</sup> MLST: Multi Locus Sequence Typing, <sup>2</sup> AMP: ampicillin, CC: clindamycin, E: erythromycin, FA: fucidic acid, FOX: cefoxitin, P: penicillin, TE: tetracycline.

**Table S6.** Frequency of *S. epidermidis* isolates from cattle farms and of isolates from sheep flocks, machine-or hand-milked, in Greece, according to STs in the MLST database (MLST: Multi Locus Sequence Typing, ST: sequence type).

Isolates from cattle farms <sup>1</sup>		Isolates from sheep flocks with machine-milking		Isolates from sheep flocks with hand-milking	
ST (n=36)	no. of isolates (n=50)	ST (n=8)	no. of isolates (n=12)	ST (n=10)	no. of isolates (n=21)
5	2	59	1	100	5
7	1	100	2	142	2
48	2	142	3	152	3
54	1	152	2	153	1
57	1	200	1	454	1
59	5	315	1	677	1
91	6	570	1	678	2
92	1	711	1	700	2
93	1			709	2
94	2			710	2
98	1				
99	1				
100	1				
101	1				
102	1				
103	1				
104	1				
105	1				
106	1				
107	1				
108	1				
109	2				
110	1				
111	2				
112	1				
114	1				
329	1				
441	1				
443	1				
444	1				
452	1				
453	1				
454	1				
473	1				
474	1				
639	1				

1. Details obtained from the MLST-database.

**Table S7.** STs with *S. epidermidis* isolates from sheep flocks, machine-or hand-milked, in Greece, with simultaneous inclusion of isolates from human sources in the MLST database (MLST: Multi Locus Sequence Typing, ST: sequence type).

STs with isolates from sheep flocks with machine-milking (n=8)	STs with isolates from sheep flocks with hand-milking (n=10)	Inclusion of isolates from human sources in the ST
59		yes
100	100	no
142	142	yes
152	152	yes
	153	yes
200		yes
315	454	yes
570	677	no
	678	no
	700	no
	709	no
	710	no
711		no



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