

Laboratory Protocol

Isolation of methicillin-resistant *Staphylococcus aureus* (MRSA) from food-producing animals and farm environment

June 2018

Version 1

Written by EURL-AR

HISTORY OF CHANGES				
Version	Sections changed	Description of change		Approval
1	New document	This document replaces previous EURL-AR protocols. The main difference is the exclusion of the second enrichment step with cefoxitin and aztreonam. Editorial changes were also performed.	17 Apr 2018	Valeria Bortolaia, Rene Hendriksen

Background

The most common method for detection of methicillin-resistant *Staphylococcus aureus* (MRSA) in samples from food-producing animals and farm environment (e.g., nasal, skin, and dust swabs) consists of a pre-enrichment step followed by a selective enrichment step before plating on chromogenic MRSA-selective and -indicative agar (1). A recent study at two laboratories has, however, shown that inclusion of the selective enrichment step leads to a higher number of false-negative results when analysing nasal and skin swabs from pigs and environmental swabs from pig stables than the same procedure omitting this stage (2). Studies in poultry (layers and broilers) and cattle did not detect significant differences in the performance of the two methods (3,4). Based on these findings, the EURL-AR modified the method for isolation of MRSA from samples from food-producing animals and farm environment.

References

1. European Food Safety Authority (EFSA). Technical specifications on the harmonised monitoring and reporting of antimicrobial resistance in methicillin-resistant *Staphylococcus aureus* in food-producing animals and food. EFSA J. **2012**; 10(10):2897. Available from: <https://www.efsa.europa.eu/en/efsajournal/pub/2897>
2. Larsen J, Sunde M, Islam MZ, Urdahl AM, Barstad AS, Larsen AR, Grøntvedt CA, Angen Ø. Evaluation of a widely used culture-based method for detection of livestock-associated methicillin-resistant *Staphylococcus aureus* (MRSA), Denmark and Norway, 2014 to 2016. Euro Surveill. **2017**; 22(28):pii=30573. Available from: <http://www.eurosurveillance.org/content/10.2807/1560-7917.ES.2017.22.28.30573>
3. Nemeghaire S, Roelandt S, Argudín MA, Haesebrouck F, Butaye P. Characterization of methicillin-resistant *Staphylococcus aureus* from healthy carrier chickens. Avian Pathol. 2013;42(4):342-6.
4. Nemeghaire S, Argudín MA, Haesebrouck F, Butaye P. Epidemiology and molecular characterization of methicillin-resistant *Staphylococcus aureus* nasal carriage isolates from bovines. BMC Vet Res. 2014;10(1):153.

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Procedure	Theory/comments
1. Sample collection and storage	
<p>1.1 Collect samples for the MRSA monitoring according to the guidelines listed in Table 1 and Table 2 of the EFSA document on “Technical specifications on the harmonised monitoring and reporting of antimicrobial resistance in methicillin-resistant <i>Staphylococcus aureus</i> in food-producing animals and food”. EFSA J. 2012; 10(10):2897. Available from: https://www.efsa.europa.eu/en/efsajournal/pub/2897</p> <p>1.2 It is recommended to store swab samples at 4-8 °C and dust samples at ambient temperature (18-22 °C) to avoid condensation which may occur on refrigeration. Moreover, it is recommended to initiate analysis i) within 3 days of collection for qualitative purposes (i.e. for scoring samples as positive/negative); and ii) on the same day of collection or, if not possible, one day after collection for quantitative purposes.</p>	<p>Always refer to the latest recommendations by EFSA for sample types to be collected for MRSA monitoring.</p> <p>There is limited knowledge on the effect of duration and temperature of storage of different sample types on MRSA viability and counts. Different commercially available transport media ensure viability of bacteria at room temperature for different amount of times. Room temperature, however, varies greatly across seasons and EU countries, which might lead to results that are not comparable across EU. As the purpose of harmonised protocols is to ensure comparability across EU countries, samples should be kept at comparable conditions.</p>
2. Sample analysis	<p>IMPORTANT: the present protocol was validated with nose, skin and dust samples from pigs and pig stables in countries with high and low prevalence of livestock-associated (LA)-MRSA (Euro Surveill. 2017; 22(28):pii=30573). As sensitivity and specificity of detection methods might be affected by sample type, animal species of origin and within-sample MRSA prevalence, laboratories are invited to contact the EURL-AR for sharing their experiences with this and/or other MRSA isolation protocols, with the overall aim to ensure the highest sensitivity and specificity of MRSA detection in samples from food-producing animals, the farm environment and food across EU.</p>

<p>2.1 Cover the samples in Mueller-Hinton broth containing 6.5% sodium chloride (NaCl) and incubate at 35-37°C for 16-24 h.</p> <p>2.2 Spread a 10-µl loopful of the broth on <i>Brilliance</i> MRSA 2 agar (Oxoid) and incubate at 35-37°C for 16-24 h.</p> <p>2.3 Subculture presumptive MRSA colonies on blood agar and incubate at 35-37°C for 22-24 h.</p>	<p>This step selects for staphylococci and other salt-tolerant bacteria. For this validation protocol, the following volumes of Mueller-Hinton broth containing 6.5% sodium chloride (NaCl) were used:</p> <ul style="list-style-type: none"> • 10 mL for pools of five cotton swabs • 500 mL for pools of two to three cloth swabs • 300 mL for individual cloth swabs <p>The effect of using different volumes is unknown.</p> <p>Presumptive MRSA appear as denim blue colonies after overnight incubation.</p> <p>MRSA colonies on blood agar are greyish or yellowish and usually surrounded by a zone of haemolysis. The catalase test can be used to distinguish staphylococci from enterococci, which sometimes produce similar colony morphology on <i>Brilliance</i> MRSA 2 agar.</p> <p>See appendix 2 for pictures of MRSA colony morphology on different media.</p>
<p>3. Identification of MRSA</p>	
<p>3.1 Confirm presumptive MRSA colonies by PCR, according to the protocol described in the document entitled “Protocol for PCR Amplification of <i>mecA</i>, <i>mecC</i>, <i>spa</i> and <i>pvl</i>” on the EURL-AR website.</p>	<p>MRSA isolates are positive for <i>spa</i> and either <i>mecA</i> or <i>mecC</i>, whereas the presence of <i>pvl</i> is variable. MRSA-confirmed isolates can be further <i>spa</i>-typed in order to determine the corresponding Clonal Complex (CC). Isolates for which no CC can be inferred from the <i>spa</i>-type should be further typed by MLST-typing. These methods are described in the documents entitled “Protocol for <i>spa</i>-typing” and “MLST typing” on the EURL-AR website https://www.eurl-ar.eu/CustomerData/Files/Folders/21-protocols/289_7-protocols-for-spa-typing.pdf and https://www.eurl-ar.eu/CustomerData/Files/Folders/21-protocols/284_8-protocols-for-mlst-typing.pdf .</p>

APPENDIX 1

Composition and preparation of culture media and reagents

The media and reagents are available from several companies including Oxoid, BD, Merck and Difco. The composition of the dehydrated media given below is an example and may vary among the different manufacturers. Also, the media should be prepared according to the manufacturer's description if that differs from the description given here.

Mueller Hinton Broth with 6.5% NaCl

<i>Formula</i>	g/L
Dehydrated beef infusion	300
Casein hydrolysate	17.5
Starch	1.5
Sodium Chloride	65
pH 7.3 +/- 0.1	

Oxoid *Brilliance*[™] MRSA 2 Agar

Prepare according to the manufacturer's description.

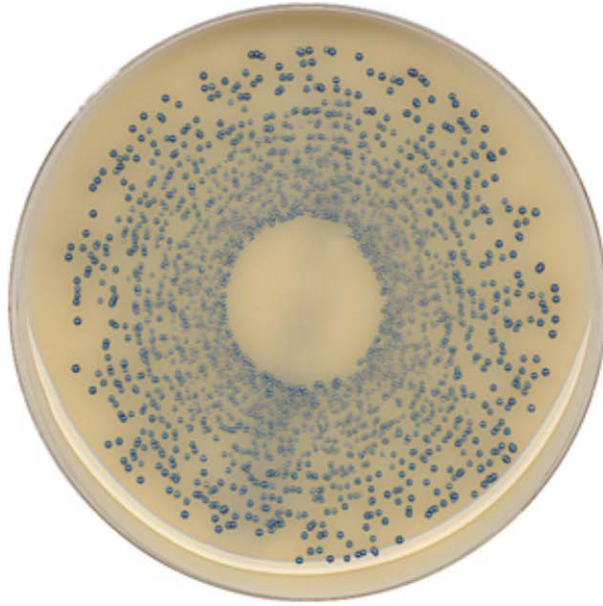
Blood Agar

<i>Formula</i>	g/L
"Lab-Lemco" powder	10.0
Neutralised peptone	10.0
Sodium chloride	5.0
Agar	15.0
pH 7.3 +/- 0.2	

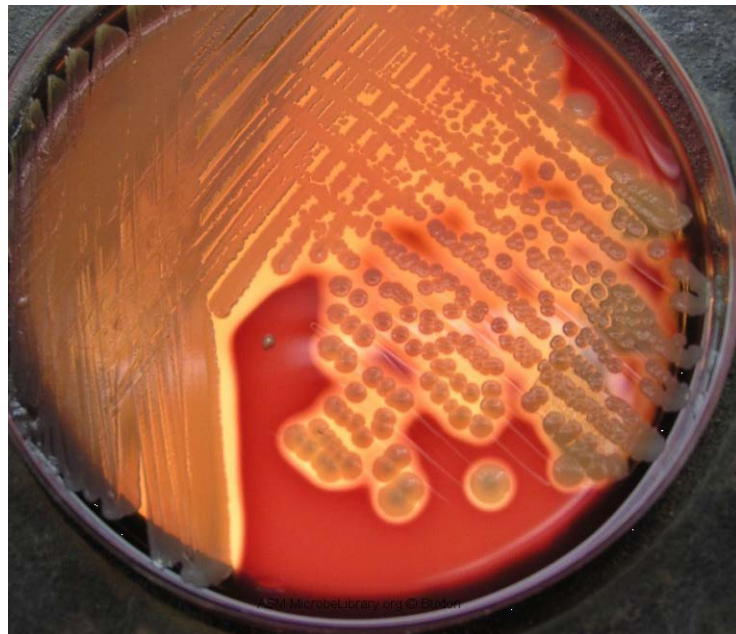
After cooling to 50°C, add 7% of defibrinated sheep or horse blood.

APPENDIX 2

Pictures showing typical colony morphology of MRSA on Oxoid *Brilliance*[™] MRSA 2 Agar and Blood Agar



Methicillin-resistant *Staphylococcus aureus* on Oxoid *Brilliance*[™] MRSA 2 Agar



Staphylococcus aureus on Blood agar

APPENDIX 3

Flow Diagram

Isolation of MRSA from samples from food-producing animals and their environment

