Phenotypic and genotypic characterization of Staphylococcus aureus isolated from

pork sausage

Caracterização fenotípica e genotípica de Staphylococcus aureus isolados de linguiça suína

Caracterización fenotípica y genotípica de Staphylococcus aureus aislado de salchicha porcina

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Abstract

The objective of this study was to characterize the genotype and phenotype of *S. aureus* isolates from pork sausages. Fifty samples of pork sausages were collected, counts of coagulase-positive *Staphylococcus* were made and isolates were obtained to identify *S. aureus* species. In the isolates, the presence of genes *sea, seb, sec,* and *sed* was surveyed, the methicillin-resistance was assessed and the production of biofilm in Congo red agar, stainless steel, polyethylene, glass, and pork casing was tested. The capacity of biofilm formation was assessed after the exposure to sublethal stress. Of the samples tested, 12% had counts superior to what is permitted by the legislation. *S. aureus* was isolated in 44% of the samples. Of these, 54% had only the gene *sed* and 32% had genes *sec* and *sed*, 73% were classified as methicillin-resistant *S. aureus* (MRSA). Of the MRSA isolates, 62% had only gene *sed* and 35% had both genes found in this study. Regarding the biofilm formation in Congo red agar, 68% of *S. aureus* isolates were considered as biofilm former isolates responded in a distinct manner. The condition in which the sublethal stress greatly induced the biofilm formation was the cold. Biofilm production was observed only in the stainless steel and pork casing in 71% and 57% of the isolates tested, respectively. Thus, we stress the importance of implementing good manufacturing practices within the industry to control microbial contamination and biofilm formation. **Keywords:** Biofilm; Enterotoxins; Methicillin; Public health; Sublethal stress.

Resumo

O objetivo desse estudo foi caracterizar o genótipo e fenótipo de *S. aureus* isolados de linguiça suína. Cinquenta amostras de linguiça suína foram coletadas, foram realizadas contagens de *Staphyloccus* coagulase positiva e obtidos isolados para identificar a espécie *S. aureus*. Nos isolados, foi pesquisada a presença dos genes *sea, seb, sec* e *sed*, feita a avaliação da resistência a meticilina e testada a produção de biofilme em Ágar Vermelho Congo, aço inoxidável, polietileno, vidro e tripa suína. Também foi verificada a capacidade de alguns isolados formarem biofilme após a exposição a estresses subletais. Das amostras testadas, 12% apresentaram contagens acima do permitido pela legislação. *S. aureus* foi isolado de 44% das amostras. Destes, 54% possuíam apenas o gene *sed* e 32% possuíam os genes *sec* e *sed*, 73% foram classificados como *S. aureus* resistentes à meticilina (MRSA). Dos MRSA 62% possuíam apenas o gene *sed* e 35% possuíam ambos os genes encontrados no estudo. Quanto a formação de biofilme em ágar vermelho congo, 68% dos isolados de S. aureus foram considerados formadores de biofilme. Após serem submetidos

aos estresses subletais, a maioria dos isolados formadores de biofilme continuou formando biofilme e os isolados não formadores de biofilme responderam de maneira distinta. A condição de estresse subletal que mais induziu a formação de biofilme foi o frio. A produção de biofilme foi observada apenas no aço inoxidável e na tripa suína em 71% e 57% dos isolados testados, respectivamente. Assim, ressalta-se a importância da implementação de boas práticas dentro da indústria para controlar a contaminação microbiana e a formação de biofilme.

Palavras-chave: Biofilme; Enterotoxinas; Meticilina; Saúde pública; Estresse subletal.

Resumen

El objetivo de este estudio fue caracterizar el genotipo y fenotipo de S. aureus aislado de salchicha porcina. Se recolectaron 50 muestras de salchicha porcina, se realizaron recuentos positivos de Staphyloccus coagulasa positivo y se obtuvieron aislamientos para identificar la especie S. aureus. En los aislamientos se investigó la presencia de los genes sea, seb, sec y sed, se evaluó la resistencia a la meticilina y se probó la producción de biopelículas en Agar Rojo Congo, acero inoxidable, polietileno, vidrio y tripa porcina. También se verificó la capacidad de algunos aislamientos para formar biopelículas después de la exposición a tensiones subletales. De las muestras analizadas, el 12% tenía recuentos superiores a los permitidos por la legislación. Se aisló S. aureus del 44% de las muestras. De estos, el 54% tenía solo el gen sed y el 32% tenía los genes sec y sed, el 73% se clasificó como S. aureus resistente a la meticilina (MRSA). Del MRSA, el 62% tenía solo el gen sed y el 35% tenía ambos genes encontrados en el estudio. En cuanto a la formación de biopelículas en agar rojo Congo, el 68% de los aislados de S. aureus se consideraron formadores de biopelículas. Después de someterse a tensiones subletales, la mayoría de los aislados formadores de biopelículas continuaron formando biopelículas y los aislados que no eran formadores biopelículas respondieron de manera diferente. La condición de estrés subletal que más indujo la formación de biopelículas fue el frío. La producción de biopelículas se observó solo en acero inoxidable y tripa porcina en el 71% y el 57% de los aislamientos probados, respectivamente. Por lo tanto, enfatiza la importancia de implementar buenas prácticas dentro de la industria para controlar la contaminación microbiana y la formación de biopelículas.

Palabras clave: Biopelícula; Enterotoxinas; Meticilina; Salud pública; Estrés subletal.

1. Introduction

Pork is the most consumed meat worldwide (USDA, 2019). Among the industrialized products made from pork, pork sausage stands out for being generated through a simple processing and for having a reasonable price. Sausage is a meat product obtained from ground meat of different domestic animals, with condiments, added or not with other ingredients, stuffed in natural or artificial casing and submitted to specific technological processes (Brasil, 2017).

During the production of fresh sausage, the meat is grounded, which increases the surface exposure to the contamination and microorganism growth and, since it is a fresh product, it does not undergo thermal treatment that eliminates or reduces this contamination. Furthermore, the manufacturing process requires a series of manipulation stages, increasing the possibility of contamination, which would compromise the hygienic-sanitary quality of the final product in case of failures and nonconformity in the processing (Tutenel, et al., 2003). Among the bacteria that could be present in the final product, *Staphylococcus aureus* stands out. It can be found in the nasal cavities and skin of healthy individuals and also in hogs (Robert & Chambers, 2005, Linhares, et al. 2015, Strube, et al., 2018).

S. aureus is considered the most relevant species within *Staphylococcus* genus and is frequently associated with outbreaks of food poisoning due to the great capacity of producing enterotoxins that, once ingested in sufficient amounts, provoke nausea, vomits, cramps, and diarrhea (Lee, et al., 2018; Germano & Germano, 2015). Staphylococcal enterotoxins (SE) are thermoresistant, therefore, once they are produced in the food, they are not eliminated during thermal processing (Franco & Landgraf, 2008).

Moreover, *S. aureus* stands out for being able to develop resistance to different antimicrobials in a fast and effective manner. In this context, methicillin-resistant *S. aureus* (MRSA) are highly relevant for being resistant to all beta-lactam antibiotics that are commonly used in the treatment of infections caused by this microorganism (De Buyser, et al., 2001, Lowy, 1998). Food contamination caused by MRSA poses a risk of transmission of this microorganism to people that manipulate these foods. In addition, it could occur the methicillin resistance transfer from MRSA to methicillin-sensitive *S. aureus* strains (MSSA). Another concerning factor is that some *S. aureus* strains have the capacity to form biofilms on the surfaces of various

equipment in food industries, such as stainless steel, plastic, and glass, which could turn them into a persistent source of microbial contamination that could lead to food deterioration and threat the safety of the products to the consumers (Giaouris, et al., 2014). Biofilms are communitarian structures of bacteria covered by a polymeric matrix produced by the bacteria and constituted of polysaccharides, proteins, and nucleic acids, named together as exopolysaccharides (EPS), with the capacity of adhesion to biotic or abiotic surfaces (Haaber, et al., 2012).

During food processing, the microorganisms are submitted to stress conditions such as high temperatures, cooling and acid pH that could favor the formation of biofilm, since the association of microorganisms in biofilms constitutes a form of protection to its development, permitting their survival in hostile environments (Donlan & Costerton, 2002).

The objective of the present study was to assess the presence of genes that encode staphylococcal enterotoxins in *S. aureus* isolates from pork sausage and to assess phenotypical traits of the isolates, such as methicillin-resistance and the capacity of biofilm formation in distinct surfaces and under distinct environmental conditions.

2. Methodology

During the time of our study, 50 samples of pork sausage were collected from butcher shops and grocery stores in the municipality of Pelotas, Rio Grande do Sul, Brazil, making a total of 19 commercial properties, collecting up to three samples in each local. Samples were stored in isothermal boxes with ice and immediately dispatched to the laboratory to carry out the analysis.

At first, we executed the count of coagulase-positive *Staphylococcus* (CoPS) collected in Baird-Parker agar (Himedia, Mumbai, India) obtained from sausage samples, according to Tallent et al. (2016). Afterwards, one coagulase-positive isolate of each sample was inoculated in Brain and Heart Infusion (BHI; Himedia, India) broth and incubated at 37°C for 24h. To the BHI culture isolates, 20% of glycerol was added to maintain a stock culture at -70°C and to identify *S. aureus* by PCR.

The confirmation of *S. aureus* was performed through PCR assay, by surveying the gene *nuc*. The DNA extraction of coagulase-positive isolates was executed according to Sambrook & Russel (2001). For the PCR assay, we used the protocol as reported by Sasaki et al. (2010), with modifications. In each 25µL of reaction mixture comprised 1µL DNA extracted, 2U of Taq DNA polymerase, 10pmol of each primer, 0.2nM of deoxyribonucleotide triphosphate (dNTP) and buffer reaction. Amplification was made at 95°C for 2min, followed by 30 cycles of 30°C for 30s, 56°C for 30s, and 72°C for 1min, and final elongation at 72°C for 2min. PCR products were stained with Diamond Nucleic Acid Dye (Promega, USA) for visualization in standard gel electrophoresis in a 1.5% agarose gel (Panreac Química SA, Spain).

The isolates that were identified as *S. aureus* by PCR were submitted to another round of PCR assay to identify the genes that encoded the staphylococcal enterotoxins A (gene *sea*), B (gene *seb*), C (gene *sec*), and D (gene *sed*), separately. The protocol used was as described by Cunha *et al.* (2007) with modifications, to survey genes encoding the staphylococcal enterotoxins from A to C. Twenty pmol of each primer, 2.5U of Taq DNA polymerase, 200µM of dNTPs, 20mM of Tris-HCl, pH 8.4, 0.75mM of MgCl2 and 5µL of DNA were added into a microtube. Amplification consisted of one cycle of 94°C for 4min, denaturation at 94°C for 2min, annealing at 55°C for 1.5min and extension from oligonucleotide primers at 72°C for 1.5min. In the third cycle, the annealing temperature was reduced to 51°C, followed by additional 37 cycles at 94°C for 2min, 42.5°C for 1.5min and 72°C for 1.5min. PCR products were visualized as described previously. The survey of gene coding staphylococcal enterotoxin D in the isolates was carried out as described by Andretta (2019).

Disk diffusion test was performed to assess methicillin resistance in Müeller-Hinton agar (Kasvi, Brasil), according to Bauer et al. (1966), in which two disks impregnated with 30µg the antibiotic cefoxitin (CFO). The results were analyzed according to the Clinical and Laboratory Standards Institute (CLSI) (2015), that considers that a microorganism is resistant to

cefoxitin when the halo formed has a diameter inferior than or equal to 21mm and sensitive when the halos had a diameter higher than or equal to 22mm.

Biofilm production by *S. aureus* strains was determined by Congo Red agar (CRA, Êxodo Científica, Brazil) culture, as described by Freeman et al. (1989). The production of rough and black colonies was considered as a positive result of biofilm formation and the production of smooth and red colonies as a negative result.

The effect of sublethal stress over the biofilm formation in CRA was assessed. The isolates were submitted to different types of sublethal stress. The cultures were left overnight in BHI maintained in water-bath at 42°C for 45min to expose cells to heat shock, according to Chang et al. (2004). Cells were also exposed to cold shock and the cultures were left overnight in BHI and maintained at 4°C for 4h (Silva, 2019). The exposure to acid environment was performed according to Wong et al. (1998). Cultures in BHI left overnight had the pH adjusted to 5.0 with HCl 6N and were incubated at 37°C for 30min.

The isolates that were considered as biofilm formers in CRA, classified as MRSA, and/or that had at least one gene that coded any staphylococcal enterotoxin, among the surveyed, were tested for the capacity of biofilm formation in different surfaces, as described by Milan et al. (2015), with modifications. Coupons with flat surfaces of 4cm² of high density polyethylene, stainless steel and sterilized glass in autoclave were used. The coupons were deposited on Petri dishes (150 X 20mm) containing 100mL of Tryptone soya broth (TSB, Acumedia, USA) and 2mL of overnight cultures of each isolate standardized in spectrophotometer at 600nm for 0.5 of optical density. Every 24h of incubation at 37oC, coupons were gently rinsed twice with phosphate buffer saline (PBS, 0.1 M, pH 7.0) to remove cells that could not adhere and then returned to the Petri dishes with 100mL of TSB, but without inoculum, and re-incubated. After three repetitions, the coupon surfaces were rubbed with sterile swabs that were stored in test-tubes containing 10mL of PBS. Then, serial dilutions were performed to count microorganisms that grew in Baird-Parker agar. One of the isolates in CRA non-biofilm former was used as negative control.

The isolates that were considered as biofilm-former in CRA and classified as MRSA and/or had at least one gene coding staphylococcal enterotoxin were tested for their ability to form biofilm in pork casing. Strips of dehydrated pork casing, sterilized by plunging in alcohol for 30min and dried in drying oven at 60°C were used. The rest of the assay was conducted as described aforementioned. Count was performed by rubbing the swab on a 4cm² area, delimited by a sterile sampling template.

The assays of biofilm formation in different surfaces were performed in triplicates and the means of *S. aureus* counts were assessed by analysis of variance and compared by Tukey's test.

3. Results and Discussion

Of the 50 samples of fresh pork sausage analyzed, 48% (24/50) exhibited CoPS, whereas 12% (6/50) had counts of CoPS above the standard as established by the Resolution of the Collegiate Board (RDC) n. 12/01 (BRASIL, 2001), that admits CoPS counts up to 5.0×10^3 CFU/g for sausages. The maximum count found in this study was 6.4×10^4 CFU/g. A result inferior than what we found in our study was reported by Valiatti et al. (2016), that assessed fresh sausage samples from grocery stores at the municipality of Ji-Paraná, Rondônia, and found that 6.6% (2/30) exhibited CoPS count above the limit determined by the Brazilian legislation. However, a higher prevalence was reported by Santa et al. (2012), that assessed 50 samples of pork sausage and mixed-meat sausage collected from industries in Southern Brazil and, of these, 15 (30%) presented CoPS, with 11 (22%) samples with counts above the limit established by the legislation. On the other hand, Botelho (2017) reports that of the nine samples of pork sausages collected in an abattoir in Viçosa, Minas Gerais, all of them presented CoPS, which could be attributed to extensive manipulation during the prepare of the product; however, none of them had counts higher than the limit stablished. The high counts of CoPS found in our study could be attributed to: (1) inappropriate

hygienic-sanitary conditions during the manipulation to prepare the sausages, (2) previous contamination of feedstock used in the process, and/or (3) storage under inadequate temperature.

S. aureus was isolated from 44% (22/50) of the samples. In 92% (22/24) of the samples in which CoPS growth was observed in the plates, we identified *S. aureus* and this microorganism was present in all samples where counts were superior to the permitted by the legislation.

Since *S. aureus* is a natural inhabitant of human skin and upper respiratory tract, the presence of this microorganism in the sausage samples surveyed could be attributed to the non-adoption of hygienic-sanitary measures by manipulators. Moreover, some studies report the presence of *S. aureus* in the skin, tonsils, and rectum of pigs (Linhares, et al., 2015, Strube, et al., 2018), indicating that these animals may host this bacterium and, if proper care are not adopted during the slaughter and the meat processing, this microorganism can contaminate the meat and other carcasses and equipment and utensils due to cross contamination. This could also be a possible cause of the presence of *S. aureus* in the samples analyzed.

Of the *S. aureus* isolates, 73% (16/22) were resistant to cefoxitin, being classified as MRSA, which means that 32% (16/50) of sausage samples was contaminated with microorganisms that were resistant to this antimicrobial. Few studies have reported the presence of MRSA in pork sausages. In a study conducted by Botelho (2017), methicillin resistance was observed in 61.5% (8/13) of *S. aureus* isolates of seven pork sausage samples, which apparently (the author does not elucidate whether all isolates, even those obtained within the same samples, were distinct strains) was similar in our study. On the other hand, an inferior result was reported by Thapaliva et al. (2017) that found a MRSA prevalence of 3.4% (10/293) in pork sausages within 293 samples analyzed in this product in the United States. Regardless the prevalence, the presence of MRSA in food is considered as threat to human health.

An additional fact to the possibility of the manipulators being MRSA carriers and act as sources of contamination in the abattoir flowchart is that pigs are considered as MRSA reservoirs (Oppliger, et al., 2012) and, once they are slaughtered, the contamination of carcasses and other products could occur, including cross-contamination (Kluytmans, 2010).

As for the presence of genes that code staphylococcal enterotoxins (SE), 54% (12/22) had only gene *sed*, that codes the enterotoxin D and 32% (7/22) had genes *sec* and *sed*. All isolates that tested positive to gene *sec* were also positive to gene *sed*. Moreover, all samples that had CoPS counts higher than the limit permitted by the Brazilian legislation carried *S. aureus* positive to the gene that coded the staphylococcal enterotoxin D. The presence of genes *sea* and *seb* was not observed.

Of the 16 MRSA isolates, 87% (14/16) had genes that coded enterotoxins, wherein 62% (10/16) had only gene *sed* and 25% (4/16) had both genes found in our study. Of the 12 *S. aureus* isolates that had only gene *sed*, 83% (10/12) were MRSA and of the seven isolates that had genes *sec* and *sed*, 57% (4/7) were MRSA.

Studies that involve the presence of genes that code SE in sausages report the presence of distinct genes or gene combinations in the isolates studied. El-Maghraby et al. (2018) reported the presence of *S. aureus* isolated from sausages that could potentially produce *sea*, *seb*, *sec* and *sed*. Shylaja et al. (2018) and Sankomkai et al. (2020) noted the presence of *S. aureus* isolates from mixed-meat sausages and fermented pork sausages, respectively, with genes *sea* and *seb* and without genes *sec* and *sed*, differently from our results. The staphylococcal enterotoxins C and D were already detected in several cases of staphylococcal food poisoning that involved different foods (Carmo, et al., 2002; Goulart, et al., 2016; Denayer, et al., 2017; Schmid, et al., 2009), evidencing the importance that these enterotoxins have in public health.

The presence of genes that code SE does not necessarily indicate that the microorganism will produce the toxin in the food. However, if the enterotoxigenic strains find favorable conditions to produce enterotoxins, these toxinis will be produced in the food offering risks to public health. In our study, we found a high prevalence of enterotoxigenic *S. aureus* isolated from pork sausage, which indicates that this product could represent a vector of staphylococcal intoxication.

In our study, 87% of MRSA strains were potentially enterotoxigenic. A similar percentage was obtained by Savariraj

et al. (2019) in which 94.3% (66/70) of MRSA isolates from pork sausages had at least one gene coding an enterotoxin, with genes *sea*, *seb*, *sec* and *sed* being reported in the studied samples, differently of what we found in our study in which we observed only the genes *sec* and *sed*. The assessment of enterotoxigenic genes in MRSA isolates have been studied (Song, et al., 2016; Marques, 2017; Carfora, et al., 2015); however, the isolates generally proceed from milk and dairy products, and are rare in meat products, especially in pork sausages.

The high prevalence of MRSA and the presence of enterotoxigenic *S. aureus* observed in this study serve as a warning sign to public health on the necessity of adoption of control measures in food production chain to reduce the contamination of products with these microorganisms, thus decreasing the risk of transmission to people.

Of the 22 *S. aureus* isolates obtained, 68% (15/22) were considered as biofilm formers by CRA test. Of these, 53% (8/15) were MRSA and potentially enterotoxigenic (they had genes *sed* or *sec+sed*), 40% (6/15) were MRSA and had only gene *sed*, 13% (2/15) were not MRSA and simultaneously had genes *sec* and *sed* and 7% (1/15) were not MRSA and none of the genes surveyed were present.

Other studies assessed the biofilm formation by *S. aureus* isolated from food, except for pork sausage, which demonstrates that, although pork is highly consumed, more studies concerning this product are necessary. In a study conducted by Zhang et al. (2018), 130 *S. aureus* isolates from pig farms, abattoirs and ready-to-eat pork products were assessed and all of them were capable of forming biofilm, which is a higher prevalence than the prevalence found herein. Chen et al. (2020) also reported a higher prevalence, 72% (70/97) of *S. aureus* isolates of different foods, including fresh meat, were able to form biofilm.

Most of the biofilm forming isolates were classified as MRSA, 67% (10/15). Khan et al. (2011) also observed that MRSA isolates tested were more likely to form biofilm than Methicillin-sensitive *Staphylococcus aureus* (MSSA). However, other studies reported that there were no significant differences in biofilm formation between MRSA and MSSA isolates (Reiter, et al., 2011, Smith, et al., 2008).

After being submitted to sublethal stresses, *S. aureus* strains responded in distinct manners, as observed in Table 1. Most of the biofilm former isolates, 80% (12/15), continued to form a biofilm even after the application of stress. One of the biofilm former isolates did not form the biofilm in acid medium and another one, after heat stress and acid environment. Only one of the biofilm form isolates did not show this capacity after being exposed to the sublethal stresses. With respect to the non-biofilm former isolates, after undergoing the stress, two remained unchanged. However, 57% (4/7) of non-biofilm former isolates formed biofilms after undergoing all stress challenges and one non-biofilm former isolate formed biofilm after cold stress.

| Isolates | Heat | Cold | Acid |
|---------------------|------|------|------|
| | | | |
| Biofilm formers | | | |
| 1 | + | + | + |
| 2 | - | + | - |
| 3 | + | + | - |
| 4 | + | + | + |
| 5 | + | + | + |
| 6 | + | + | + |
| 7 | + | + | + |
| 8 | + | + | + |
| 9 | + | + | + |
| 10 | + | + | + |
| 11 | + | + | + |
| 12 | + | + | + |
| 13 | + | + | + |
| 14 | + | + | + |
| 15 | - | - | - |
| Non-biofilm formers | | | |
| 16 | - | + | - |
| 17 | + | + | + |
| 18 | + | + | + |
| 19 | - | - | - |
| 20 | - | - | - |
| 21 | + | + | + |
| 22 | + | + | + |

Table 1. Capacity of biofilm formation by S. aureus isolates after being submitted to different types of sublethal stress.

(+) biofilm formation; (-) non biofilm formation. Source: Authors.

The fact that most of the non-biofilm former isolates formed biofilm after undergoing at least one of the stress condition indicates that the stress condition induce the biofilm formation by *S. aureus*. According to Jefferson (2004), the reason behind the biofilm formation is a defense strategy against stress condition.

The sublethal stress condition that most induced biofilm formation by the isolates used in this study was the cold. This study is of significant importance for the food industry, since the temperature of 4oC is the recommended for storing pork sausages up to consumption time and, moreover, before the stuffing process there is a critical step which is a resting period of 24 hours at 4oC to avoid microorganism multiplication in the product (Ordóñez, et al., 2005). Likewise, sausages are generally stored under refrigeration in retail business, restaurants and households, where sometime they are manipulated in a careless manner, what could induce the capacity of *S. aureus* in forming biofilms when present in the food.

In our study, as previously reported, two out of the seven non-biofilm former isolates (29%) remained as non-biofilm formers, a percentage that is similar as reported by Silva (2019) who tested six *S. aureus* isolates, three isolates were biofilm formers and the other three were non-biofilm formers, and were submitted to cold shock, heat shock and acid environment. This author reported that all biofilm former isolates remained biofilm formers after stresses and the non-biofilm former isolates, just as in our study, reacted in a distinct manner when facing stressor stimuli, despite one of the isolates (1/3, 33%)

that did not form biofilm under none of the stressors tested.

Of the non-biofilm former strains studied herein, 57% (4/7) began forming biofilm after being submitted to heat stress. Thus, heat stress could be considered as a factor that influences biofilm formation. However, Rode et al. (2007) observed that biofilm formation by *S. aureus* after being exposed to heat was low.

The results obtained in the present study, like other studies that assessed the effect of sublethal stress on biofilm formation performed with different microorganisms (Rosa, et al., 2017; Galvão, et al., 2012; Lianou & Koutsoumanis, 2012), demonstrated that each microorganism, just as each strain of the same microorganism, has its own peculiarities when facing hostile environments, and may or may not form a biofilm.

Of the 15 biofilm former isolates in CRA, seven were selected to assess biofilm formation in stainless steel, glass, polyethylene and pork casing, with two of them being MRSA with genes *sec* and *sed*, three MRSA with only gene *sed*, one non-enterotoxigenic MRSA and one MSSA with genes *sec* and *sed*.

The analysis of variance of counts in each surface demonstrated that there was no effect of repetitions. According to Tukey's test, there was biofilm formation in stainless steel and pork casing in 71% (5/7) and 57% (4/7) of the isolates, respectively. The production of biofilm in glass and polyethylene was not observed (Table 2).

| Isolates | Stainless steel (SD) | Glass (SD) | Polyethilene (SD) | Pork Casing (SD) |
|------------------|----------------------|---------------|-------------------|------------------|
| MRSA+sec+sed | | | | |
| А | 6.2 (0.8) [+] | 5.9 (1.3) [-] | 5.4 (1.15) [-] | 4.3 (0.6) [-] |
| В | 5.5 (0.3) [-] | 5.1 (1.1) [-] | 6.4 (0.60) [-] | 6.0 (0.6) [-] |
| MRSA+sed | | | | |
| С | 5.9 (0.4) [+] | 6.1 (0.4) [-] | 4.6 (0.49) [-] | 6.9 (0.7) [+] |
| D | 6.5 (0.1) [+] | 6.9 (0.4) [-] | 6.5 (1.75) [-] | 6.9 (0.0) [+] |
| Е | 5.8 (1.5) [+] | 5.9 (0.9) [-] | 5.5 (0.24) [-] | 6.7 (0.8) [+] |
| MRSA | | | | |
| F | 5.6 (0.4) [-] | 5.5 (0.4) [-] | 4.8 (0.22) [-] | 7.1 (0.3) [+] |
| MSSA+sec+sed | | | | |
| G | 7.0 (0.5) [+] | 6.8 (0.7) [-] | 5.8 (1.17) [-] | 4.9 (0.3) [-] |
| Negative Control | 3.7 (1.0) | 5.0 (1.1) | 3.9 (0.9) | 4.9 (0.7 |

Table 2. S. aureus counts in different surfaces $(\log/4 \text{ cm}^2)$ and assessment of biofilm formation.

SD = standard deviation; [+] biofilm formation; [-] non biofilm formation. Source: Authors.

None of the isolates in which we tested other abiotic surfaces other than stainless steel formed biofilm. Biofilm formation in the surface of this material was also observed by other authors that studied *S. aureus* isolated from different foods (Di Ciccio, et al, 2014; Lee, et al., 2015; Friedriczewski, et al., 2018). Differently from our results, some studies observed the formation of biofilm by *S. aureus* in glass and/or polyethylene (Lee et al. 2015, Friedriczewski, et al., 2018). The difference in the capacity of forming biofilm in glass could be related to the strains and/or food traits from which they were isolated. The ability of *S. aureus* isolates tested to form biofilm and attach to stainless steel is of great concern for food industry, since this material is commonly present in equipment and utensils within the industries and, once there is biofilm formation, it could turn

into a constant source of contamination to food that they contact with.

This is the first study that reported the capacity of *S. aureus* of producing biofilm in pork casing, which is a relevant finding. Once the biofilm is present in this material, other food, equipment, and surfaces could be contaminated when they contact, in addition to represent a direct risk for people that consume them. According to Iñiguez-Moreno et al. (2018), the multiplication of microorganisms in food surfaces is one of the main causes of deterioration and loss in processed and fresh products, and is also one of the main causes of foodborne illnesses and damage to equipment in food industry. Furthermore, since all biofilm formers that we isolated in pork casing were MRSA positive, they represent a huge risk and become very relevant to public health.

The presence of genes *sec* and *sed*, the methicillin resistance or the combination of both factors in the isolates studied are pathogenic mechanisms that, in addition to the capacity of forming a biofilm, turn the occurrence of *S. aureus* in sausages a bigger issue when it comes to food safety to consumers.

4. Conclusion

Methicillin-resistant *S. aureus* and the potential staphylococcal producers of enterotoxins C and D can be isolated from fresh pork sausages and were capable of forming biofilm, including in surfaces such as stainless steel and pork casing. Some *S. aureus* strains non-biofilm formers, when submitted to determined stress conditions, such as refrigeration, develop the capacity of forming biofilm.

These results highlight the importance of the implementation of good manufacturing practices within the industry to control microbial contamination and biofilm formation.

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