Hygienic-sanitary quality of ready-to-eat salmon sashimi (Salmo Salar)
Qualidade higiênico-sanitária de sashimi de salmão (Salmo salar) pronto para o consumo
Calidad higiénico-sanitaria del sashimi de salmón (Salmo salar) listo para consumir

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Abstract
Fifty sashimi samples from 5 restaurants were characterized for enumeration of Aerobic Psychrotrophic Heterotrophic Bacteria (APHB) and Aerobic Mesophilic Heterotrophic Bacteria (AMHB), Enterobacteriaceae, coliforms at 45°C (thermotolerant coliforms), coagulase-positive Staphylococci, presence of Vibrio parahaemolyticus and Salmonella sp., and determination of hydrogen potential (pH) and total volatile basic nitrogen (TVB-N). The bacterial isolates were evaluated for their resistance profile to the antimicrobial agents Penicillin, Ampicillin, Cefoxitin, Cefotaxime, Amikacin, Gentamicin, Tetracycline, and Trimethoprim-sulfamethoxazole. The results showed that APHB, AMHB and coagulase-positive Staphylococci counts and the determination of pH and TVB-N were in accordance with national and international standards adopted as safe limits for consumption. In contrast, the Enterobacteriaceae and thermotolerant coliforms counts were found to be unsatisfactory hygienic-sanitary quality. The Staphylococcus aureus and V. parahaemolyticus isolates showed resistance to Penicillin, Ampicillin, Cefoxitin, Cefotaxime, Tetracycline, Gentamicin, and Amikacin, while the Salmonella sp. isolate showed no resistance to all the antimicrobial agents studied. The results showed that 48% of the samples were fit for consumption while 52% had unsatisfactory hygienic-sanitary quality for the parameters evaluated.

Keywords: Brazil; Raw fish; Food security; Consumption; Bacterium; Antibiotics.

Resumo
Cinquenta amostras de sashimi de 5 restaurantes foram caracterizadas quanto à contagem de Bactérias Heterotrófiticas Psicotróficas Aeróbicas (APHB) e Bactérias Heterotrófiticas Mesofílicas Aeróbicas (AMHB), Enterobacteriaceae, Coliformes a 45°C (Coliformes Termotolerantes), Estafilococos coagulase positiva e presença de Vibrio parahaemolyticus e Salmonella sp. e determinação do potencial de hidrogênio (pH) e nitrogênio básico volátil total (TVB-N). Foi avaliado o perfil de resistência dos isolados bacterianos aos agentes antimicrobianos penicilina, ampicilina, cefoxitina, cefotaxima, amicacina, gentamicina, tetraciclina e trimetripim-sulfametoxazol. Os resultados mostraram que as contagens de APHB, AMHB e estafilococos coagulase positiva e os valores das determinações de pH e TVB-N correspondem (estão de acordo) com as normas nacionais e internacionais adotadas como limites seguros de consumo. Em contraste, a contagem de Enterobacteriaceae e coliformes termotolerantes e a presença de Salmonella sp. e V. parahaemolyticus não corresponde ao que estabelece (discordam) com essas normas, o que suscita preocupações quanto à qualidade higiênico-sanitária do sashimi. Os isolados de Staphylococcus aureus e V. parahaemolyticus não corresponde ao que estabelece (discordam) com suas normas, o que suscita preocupações quanto à qualidade higiênico-sanitária do sashimi.
parahaemolyticus mostraram resistência à penicilina, ampicilina, cefoxitina, cefotaxima, tetraciclina, gentamicina e amikacina, enquanto Salmonella sp. o isolado não apresentou resistência a todos os antimicrobianos estudados. Os resultados mostraram que 48% das amostras eram adequadas para consumo enquanto 52% apresentavam qualidade higiénico-sanitária insatisfatória para os parâmetros avaliados.

Palavras-chave: Brasil; Peixe cru; Alimento seguro; Consumo; Bactéria; Antibióticos.

Resumen
Cincuenta muestras de sashimi de 5 restaurantes fueron caracterizadas para el recuento de Bacterias Heterotróficas Psicrótróficas Aeróbicas (APHB) y Bacterias Heterotróficas Mesofólicas Aerobias (AMHB), Enterobacteriaceae, Coliformes a 45°C (Coliformes termotolerantes), Estafilococos coagulasa positivos y presencia de Salmonella sp. y Vibrio parahaemolyticus y determinación del potencial de hidrógeno (pH) y nitrógeno básico volátil total (TVB-N). Se evaluó el perfil de resistencia de los aislados bacterianos a los agentes antimicrobianos penicilina, ampicilina, cefoxitina, cefotaxima, amikacina, gentamicina, tetraciclina y trimetripim-sulfametoxazol. Los resultados mostraron que los recuentos de APHB, AMHB y estafilococos coagulasa positivos y los valores de las determinaciones de pH y TVB-N se corresponden (estaban de acuerdo) con los estándares nacionales e internacionales adoptados como límites seguros para el consumo. A contrario, el recuento de Enterobacteriaceae e coliformes termotolerantes y a presencia de Salmonella sp. e V. parahaemolyticus no se corresponde con que establece (estaban en desacuerdo) con esas normas, lo que suscitaba preocupación por la calidad higiénico-sanitaria de los sashimi. Los aislados de Staphylococcus aureus y V. parahaemolyticus mostraron resistencia a penicilina, ampicilina, cefoxitina, cefotaxima, tetraciclina, gentamicina y amikacina, mientras que Salmonella sp. el aislado no mostró resistencia a todos los agentes antimicrobianos estudiados. Los resultados arrojaron que el 48% de las muestras eran aptas para el consumo mientras que el 52% presentaba una calidad higiénico-sanitaria insatisfactoria para los parámetros evaluados.

Palabras clave: Brasil; Pescado crudo; Seguridad alimentaria; Consumo; Bacteria; Antibióticos.

1. Introduction
The general public's view of the modern diet and human health has led to new eating habits in recent years. Fish consumption has increased worldwide, due to its high nutritional value and intrinsic characteristics. However, this food matrix is very susceptible to microbiological contamination, requiring care from capture to consumption, such as hygiène, cold storage, and heat treatment.

Although an adequate cooking process represents food safety, the globalization of Japanese cuisine has led to the consumption of fish in the form of sashimi, which consists of thin slices of raw fish served with sauces, and not subjected to cooking steps. Therefore, it is essential to apply quality control tools, such as the adoption of good manufacturing practices, with temperature control in all stages after the capture and use of ice made with treated water\(^{50}\) for ensuring hygienic-sanitary quality.

Food safety, one of the general food hygiene principles of the Codex Alimentarius, is defined as the assurance that food will not cause harm to the consumer during preparation and consumption (FAO, 1998). Failures in the hygienic-sanitary aspects can contribute to foodborne illness occurrences, with risks to consumers' health (WHO, 2018; Brasil, 2019). In this context, raw fish stand out for being more susceptible to deterioration and microbial contamination.

National and international microbiological guidelines have been adopted to ensure food safety and quality, including those of the National Health Surveillance Agency (ANVISA) (Brasil, 2001), International Commission on Microbiological Specifications for Foods (ICMSF, 1986), Center for Food Safety (CFS, 2014) and Ryser and Schuman (2015), as well as the Regulation of the Industrial and Sanitary Inspection of Products of Animal Origin (RIISPQA), which has established values for the hydrogen potential (pH) and total volatile basic nitrogen of fresh fish (Brasil, 2017).

The emergence of bacteria resistant to antimicrobial agents is another trouble related to fish consumption, which has been a concern of international health agencies and several interrelated institutions, in addition to consumers, once antimicrobial resistance is one of the most serious global public health threats. The resistance of bacteria increases in places with intense transit, such as hospitals, food animal production and aquaculture (Cabello et al., 2016; Cabello e Godfrey, 2016; Venter, Henning e Begg, 2017).
Therefore, this study aimed to investigate the hygienic-sanitary quality of salmon (*Salmo salar*) sashimi ready for consumption in the city of Cuiaba, Mato Grosso, Brazil, through the microbiological, physicochemical, and antimicrobial resistance profile, and the potential risks to consumers’ health.

### 2. Methodology

A survey of all Japanese cuisine restaurants in the city of Cuiaba, Mato Grosso, Brazil (n = 28) was carried out. Five restaurants were randomly selected for sample collection, four of which specialized in Japanese cuisine. The samples consisted of portions of approximately 200 grams of salmon sashimi, and 10 samples were collected in each restaurant (R1, R2, R3, R4, and R5), totaling 50 samples. The samples were ordered and taken from each restaurant at the beginning of the working day. The sashimi portions were packed in the packages offered by the establishment, which were immediately placed in a thermal box with ice packs to maintain the temperature during transport to the Food Microbiology Laboratory of the Federal Institute of Education, Science and Technology of Mato Grosso (IFMT) Cuiaba - Bela Vista Campus, for analysis. The time interval between collecting the samples and starting the analysis did not exceed one hour. This quantitative study used a completely randomized design (Pereira, 2018).

#### 2.1 Microbiological characterization

The microbiological analyses consisted of the enumeration of Aerobic Psychrotrophic Heterotrophic Bacteria (APHB) (APHA 13.61: 2015), Aerobic Mesophilic Heterotrophic Bacteria (AMHB) (APHA 08: 2015), *Enterobacteriaceae* (APHA 9.62: 2015), Coliforms at 45 °C (APHA: 2015 and APHA / AWA / WEF 9221: 2012), coagulase-positive Staphylococci (ISO 6888-1: 1999 Amd 1: 2003 and APHA 39.61: 2015), presence of *Salmonella* sp. (ISO 6579 method) and presence of *Vibrio parahaemolyticus* (APHA 40.62 / 40.63: 2015) (Silva et al., 2017). The microbiological assays were performed in triplicate, with the results expressed in Log CFU/g.

#### 2.2 Physicochemical characterization

The samples were characterized for hydrogen potential (pH) according to ISO 2917 (1999) with direct reading on bench pH and total volatile basic nitrogen (TVB-N) according to Brazil (1981), and the results expressed in mg TVB-N/100 g. All determinations were performed in triplicate.

#### 2.3 Antimicrobial resistance profile

After biochemical confirmation (Silva et al., 2017), the bacterial strains isolated from salmon sashimi were identified and kept refrigerated until analysis. The antimicrobial resistance profile was performed according to the antibiotic disk diffusion method (Bauer et al., 1966).

In *vitro* antimicrobial susceptibility tests were performed according to the recommendations of the Clinical and Laboratory Standards Institute (CLSI, 2010, 2016, 2018). For that, 10 antimicrobial agents were used, as follows: Penicillin 10 µg - PEN; Ampicillin 10 µg – AMP; Gentamicin 10 µg - GEN; Oxacillin 1 µg - OXA; Trimethoprim: sulfamethoxazole (1:5 ratio) 25 µg - TMP-SMX; Tetracycline 30 µg - TET; Amikacin 30 µg - AMI; Cefotaxime 30 µg - CTX; and Cefoxitin 30 µg - CFO.

#### 2.4 Statistical analysis

The results of the physicochemical and microbiological characterization were analyzed by analysis of variance (ANOVA) at a 5% level of significance (p <0.05). The Kolmogorov-Smirnov normality test was applied to all data. The normally distributed data were analyzed through a completely randomized design (CRD) using the Tukey’s test (pH, TVB-N,
APHB, Enterobacteriaceae). The Kruskal-Wallis and the Wilcoxon mean comparison tests were applied to data that presented no normality (AMHB, coliforms at 45 °C, coagulase-positive staphylococci). The microbiological counts (APHB, AMHB, Enterobacteriaceae, coliforms at 45 °C, coagulase-positive staphylococci) were transformed into Log10 by the statistical program. The chi-square test was used for the results of the presence of Salmonella sp. and Vibrio parahaemolyticus, to determine whether the results were homogeneous concerning the five restaurants. Principal Component Analysis (PCA) was also performed to correlate the physicochemical and microbiological data between restaurants, using a correlation matrix. To determine the hygienic-sanitary quality of the samples, the results were compared with legislation standards and international references, determining the number of satisfactory and unsatisfactory samples, in percentage. The statistical program R version 3.6.1 was used (R Core Team, 2019).

3. Results and Discussion

3.1 Hygienic-sanitary quality

APHB, also known as spoilage bacteria, grow in refrigerated products (0 - 7 °C) leading to food deterioration due to the activity of proteolytic and lipolytic enzymes (Lanzarin et al., 2011). Some authors have reported counts above 7, 4.59, and 6.18 Log CFU/g in sashimi (Muscolino et al., 2014, Miguéis et al., 2015, Miguéis et al. 2016). In the present study, as shown in Table 1, R2 presented the lowest APHB count (3.43 Log CFU/g) (p <0.05), ranging from 2.76 to 4.90 Log CFU/g, while R1, R3, and R5 presented intermediate results (p> 0.05), with values ranging from 3.95 to 5.55; 3.96 to 5.53; and 3.64 to 5.90 Log CFU/g, respectively. R4 had the highest APHB count (5.76 Log CFU/g) (p <0.05), ranging from 5.2 to 6.41 Log CFU/g. These results are lower than those found in the literature and satisfactory according to the ICMSF standard (1986), which has established a maximum count of 7 Log, showing little deteriorating activity.

The AMHB counts ranged from 4.69 to 5.52; 3.27 to 4.27; 4.14 to 4.74; 4.37 to 5.32; and 4.81 to 5.40 Log CFU/g for R1, R2, R3, R4, and R5, respectively, and the lowest count (Table 1) was observed for R2 with 3.69 Log CFU/g (p <0.05) while R1 and R5 presented the highest counts with values of 5.06 and 5.14 Log CFU/g, respectively (p> 0.05). These microorganisms are considered indicators of food quality, demonstrating hygienic conditions during the manufacturing process, in addition to contributing to the determination of the shelf life. The present counts were satisfactory, with microbial counts lower than 6 log CFU/g, according to the classification of the Center for Food Safety (2014) and Ryser and Schuman (2015). Several authors have found different mesophilic bacteria counts in sushi and sashimi, with values of 5.5 Log CFU/g; 5.3 Log CFU/g; 5.11 Log CFU/g; and 7.0 Log CFU/g (Liang et al., 2016; Kim et al., 2016; Miguéis et al., 2015; Muscolino et al., 2014).

The enumeration of Enterobacteriaceae (ENT) ranged from 3.26 to 4.06; 0 to 1.49; 1.49 to 3.36; 3.20 to 4.42; and 4.43 to 5.30 Log CFU/g for R1, R2, R3, R4, and R5, respectively, with the lowest and highest counts observed for R2 and R5, with values of 0.43 and 4.76 Log CFU/g, with significant differences between them (p <0.05). The restaurants R1, R3, and R4 presented intermediate counts, with no differences between R1 and R4 (p> 0.05) (Table 1). Bacterial counts <2 Log are considered satisfactory, from 2 to ≤ 4 Log are acceptable, and bacterial counts >4 Log are considered unsatisfactory (CFS, 2014). In this research, R1, R4, and R5 presented high contamination levels, with unsatisfactory results observed in 10%, 40%, and 100% of the samples, respectively, thus the samples were unsuitable for consumption and/or potentially dangerous to human health. Enterobacteriaceae can be indicative of hygienic conditions in manufacturing processes, as they can colonize environments with poor sanitation, and some strains are pathogenic and represent a risk to public health (SILVA et al. 2017). Studies with salmon and tuna sashimi found Enterobacteriaceae counts of 3.39; 3.43; and 3.25 Log CFU/g, respectively (Miguéis, et al., 2015; Miguéis, et al., 2016) corroborating the findings of this study.
Table 1. Microbiological and physicochemical characterization of salmon (*Salmo salar*) sashimi ready for consumption in the city of Cuiabá, Mato Grosso, Brazil.

| Log 10 CFU / g mean ± standard deviation |
|------------------------------------------|
| pH | TVB-N | APHB | AMHB | ENT | COL | STA | SAL | VIB |
|----|-------|------|------|-----|-----|-----|-----|-----|
| R1 | 6.28 ± 0.04 | 17.71 ± 1.78 | 4.82 ± 0.56 | 5.06 ± 0.26 | 3.69 ± 0.30 | 3.49 ± 0.33 | Presence | Presence |
| R2 | 6.24 ± 0.03 | 16.35 ± 1.09 | 3.43 ± 0.66 | 3.69 ± 0.32 | 0.43 ± 0.10 | 0.10 ± 0.10 | Absence | Absence |
| R3 | 6.22 ± 0.02 | 16.61 ± 1.58 | 4.90 ± 0.45 | 4.45 ± 0.19 | 2.74 ± 0.59 | 0.28 ± 0.60 | Absence | Absence |
| R4 | 6.30 ± 0.04 | 17.47 ± 2.40 | 5.76 ± 0.37 | 4.79 ± 0.28 | 3.96 ± 0.36 | 0.13 ± 0.41 | Absence | Presence |
| R5 | 6.28 ± 0.01 | 18.98 ± 1.68 | 5.05 ± 0.62 | 5.14 ± 0.19 | 4.76 ± 0.28 | 2.11 ± 1.30 | Absence | Presence |

ENT: *Enterobacteriaceae*, COL: *coagulase-positive staphylococci*, SAL: *Salmonella*; VIB: *Vibrio parahaemolyticus*. Same letters in the same column indicate no difference at the 5% significance level (p <0.05). Tukey’s test was used for pH, TVB-N, APHB, and ENT. Kruskal-Wallis test was used for AMHB, COL, and STA. The chi-square test was used for SAL and VIB.

As can be seen in Table 1, the lowest thermotolerant coliforms (COL) counts were observed for R2, R3, and R4, with values of 0.10; 0.28; and 0 Log CFU/g, respectively (p > 0.05) (ranging from 0 to 1, 0 to 1.51, and 0 Log CFU/g, respectively). The highest count was found in R1 with 3.49 Log CFU/g (2.94 to 4.03 Log CFU/g), followed by R5 with 2.11 Log CFU/g (0 to 3.76 Log CFU/g) (p < 0.05). All samples (100%) from the restaurant R1 and half (50%) of the samples from R5 had counts higher than those established by the Brazilian legislation (2 Log CFU/g) (Brasil, 2001), corroborating the results of *Enterobacteriaceae* and showing concern with hygienic-sanitary quality since this group of microorganisms may contain potentially pathogenic and FBD-causing strains. In this study, although *Escherichia coli* was not detected, other bacteria from the coliform group were detected, such as *Enterobacter* sp., *Klebsiella* sp., and *Pantoea* sp.

Regarding the enumeration of positive coagulase staphylococci (STA), all samples were classified as satisfactory, with counts lower than 3.69 Log CFU/g (Brasil, 2001), with the lowest values observed for R1, R2, R3, and R4, with 0; 0.13; 0.38; 0.13 Log CFU/g, respectively (p > 0.05) (Table 1) (counts ranging from 0; 0 to 1.30; 0 to 1.43; and 0 to 1.30 Log CFU/g for R1, R2, R3, and R4, respectively). In contrast, the highest count was presented by the highest count, 1.38 Log CFU/g, (p <0.05) (Table 1), ranging from 0 to 2.58 Log CFU/g. Although the samples presented satisfactory counts, this group of microorganisms stands out as is enterotoxins producers and the contamination may be due to the lack of hygiene during handling or processing since this bacterium is natural in the animal and human microbiota. From 2009 to 2018, this agent was considered one of the main bacteria responsible for foodborne outbreaks in Brazil, being the third etiological agent with 9.5% of cases (Brasil, 2019). In the present study, *Staphylococcus aureus*, a species with great potential for toxin production, was found in the restaurants R2, R3, R4, and R5. Staphylococci are popularly identified in sashimi, with recent studies in Italy (Muscolino et al., 2014), Portugal (Miguéis et al., 2015; Miguéis et al., 2016) and Malaysia (Puah et al., 2017). They are also found in oysters, shrimp, sushi, and a variety of fresh marine fish in several countries such as Turkey (Onmaz et al., 2015), Switzerland (Boss et al., 2016), Iran (Arfatahery et al., 2016), and Denmark (Li et al., 2019).

The presence of *Salmonella* sp. in food is a worrying factor, as it is the main cause of food-borne illnesses with tens of millions of cases per year worldwide (WHO, 2018). In Brazil, from 2000 to 2017, 12,503 outbreaks were reported by etiologic agents, with 236,403 sick people registered with the Ministry of Health, and *Salmonella* sp. was the most common bacterial cause, with 30% of cases (Brasil, 2018). As can be seen in Table 1, the presence of *Salmonella* sp. was detected in a sample from the restaurant R1, which is in disagreement with the legislation (Brasil, 2001) and considered inappropriate for human consumption.
consumption, becoming an alert factor with potential pathogenic risk. Prevention requires care in all stages of the food chain, from production to consumption (WHO, 2018). Puah et al. (2017) found *Salmonella enterica* in 21.66% (13/60) of retail sashimi samples in Valley Klang, Malaysia, which was also observed by other authors in fresh fish in Nigeria (Beshiru et al., 2019), Jordan (Obdait e Salman, 2017) and Turkey (Onmaz et al., 2015).

*Vibrio parahaemolyticus* is a marine bacterium responsible for illness of gastroenteritis after consumption of raw, inadequately cooked, or cross-contaminated seafood (Sakazaki, 2003). The present study found the presence of *Vibrio parahaemolyticus* in 24% of the samples studied, consisting of twelve samples (6 samples in R1, 5 samples in R4, and 1 sample in R5), which is of concern, as this bacterium is related to cases of gastroenteritis evolving to septicemia in immunocompromised individuals, even leading to death. The presence of this agent in the samples of this study may be due to the origin of the raw material, suggesting that it comes from contaminated water and different producers. Other *Vibrio* sp. species were also verified in this study, such as *V. mimicus*, *V. metchnikovii*, which serves as a warning for the occurrence of unknown pathogenic strains that can be risk factors associated with outbreaks (FDA, 2012). Some authors have reported *Vibrio* sp. in sashimi, with counts ≥ 2 and ≥ 3 Log CFU/g (Muscolino et al., 2015; Kim et al., 2016). *V. parahaemolyticus* has also been found in several fresh seafood worldwide, such as in Ecuador (Sperling et al., 2015), China (Xu et al., 2016), Vietnam (Vu et al., 2016), Poland (Lopatek et al., 2018), Brazil (Silva et al., 2018), and Korea (Kang et al., 2019, Ryu et al., 2019).

Some compounds are produced from bacterial metabolism and fish decomposition, which can be indicative of fish quality, such as total volatile basic nitrogen (TVB-N) and hydrogen potential (pH) (Gonçalves, 2011). For TVB-N, the results ranged from 14.56 to 20.22; 14.57 to 17.71; 14.70 to 19.18; 13.88 to 20.62; and 15.79 to 21.87 mg of TVB-N/100g for R1, R2, R3, R4, and R5, respectively. The restaurants R2 and R3 had the lowest mean value, with 16.35 and 16.61 mg TVB-N/100g, respectively, when compared to R5, which presented 18.98 mg TVB-N/100g. The restaurants R1 and R4 presented TVB-N levels similar to those observed for the restaurants R2, R3, and R5 (p> 0.05). The determination of volatile basic nitrogen (TVB-N) are used to characterize the freshness of fish since the formation of nitrogen compounds results from the enzymatic and bacterial deterioration, forming products from the decomposition of the amino acids such as dimethylamine, trimethylamine, ethylamine, monomethylamine, ammonia, putrescine, cadaverine, and spermidine (Ogawa e Maia, 1999; Howgate, 2010; Nellet e Toldrá, 2010; Gonçalves, 2011). Therefore, the TVB-N levels can be indicative of fish conservation, especially regarding the stage of deterioration.

However, some species can present higher levels of these compounds, even though they are not under the decomposition process, such as the *Salmo salar* species for example, which has a higher level up to 35 mg N/100g, according to the Annex I of the European Communities (1995). For the Brazilian legislation, fish, in general, must have total volatile bases levels lower than 30 mg of N/100g of muscle tissue (Brasil, 2017). All samples analyzed in this study met this parameter.

The pH of the samples ranged from 6.20 to 6.36; 6.19 to 6.27; 6.18 to 6.25; 6.22 to 6.35; and 6.25 to 6.30, for R1, R2, R3, R4, and R5, respectively, with the lowest pH found in the restaurants R3 and R4 (6.22 and 6.24, respectively), and the highest pH in the restaurants R1, R2, and R5 (6.28, 6.30, and 6.28, respectively) as shown in Table 1 (p> 0.05). As reported by Ogawa and Maia (1999), the pH in salmon can range from 6.1 to 6.3. According to the Brazilian legislation, the pH of fish should be lower than 7.0 to be considered fresh and acceptable for consumption (BRASIL, 2017). All samples of this study were within the limit established by law. From the physicochemical point of view, the pH value is directly related to the quality and conservation of fish, and the bacterial load is consistent with the variation of pH. Changes in the concentration of hydrogen ions, due to the hydrolytic, oxidative, or fermentative decomposition of the muscle can lead to higher pH values, thus leading to an increase in the bacterial activity. Fish with a pH below 5.6 has a longer shelf life when compared to those with pH from 6.2 and 6.6 due to the greater availability of bacterial changes (Ogawa e Maia, 1999; Gonçalves, 2011).
The hygienic-sanitary quality of the salmon sashimi samples was determined by comparing the microbiological and physicochemical results of each restaurant with national and international standards. Regarding the samples from the restaurant R1, 10% were unsatisfactory for Enterobacteriaceae, 100% for thermotolerant coliforms, 1% exhibited the presence of Salmonella sp. and 6% Vibrio parahaemolyticus, while in R4, 40% were unsatisfactory for Enterobacteriaceae and 5% were contaminated with V. parahaemolyticus. The samples from the restaurant R5 were 100% unsatisfactory for Enterobacteriaceae, 50% for thermotolerant coliforms and 1% exhibited the presence of V. parahaemolyticus. Finally, the samples from the restaurants R2 and R3 were 100% satisfactory (Table 2).

Table 2. Hygienic-sanitary quality of salmon (Salmo salar) sashimi ready for consumption in the city of Cuiaba, Mato Grosso, Brazil

|     | Satisfactory | Unsatisfactory |
|-----|--------------|----------------|
| R1  | 0%           | 100%           |
| R2  | 100%         | 0%             |
| R3  | 100%         | 0%             |
| R4  | 40%          | 60%            |
| R5  | 0%           | 100%           |

Source: Authors (2021).

Principal Component Analysis (PCA) was used to synthesize the quantitative results considered as variables, aimed to understand the correlation between variables (Figure 1). The principal components 1 and 2, called Dimension 1 (Dim1) and Dimension 2 (Dim 2), are the main explanations of this statistical tool. The two principal dimensions were responsible for 78.1% of the total variation on the contamination rate of sashimi, and Dim 1 explained 55.2% and Dim 2 explained 22.9%.

PCA (Figure 1) showed that both R2 and R3 were clustered together, once both dimensions Dim 1 and Dim 2 decrease together. The restaurants R1 and R4 were located in another group, with an increase in Dim 2 with an increase in Dim 1. The restaurant R5 was located alone, not forming a cluster with the others, and it was more distant from the origin, showing greater discrepancy; while Dim 1 was positive and increasing, dimension Dim 2 was negative and decreasing.
For a better interpretation of the graph, the angle between variables was measured (APHB, AMHB, pH, TVB-N, etc). Dim 1 showed a strong positive correlation between the variables TVB-N and AMHB, thus it is possible to state that AMHB increased with an increase in TVB-N, as they formed an acute angle and the vectors had very close sizes. Concerning Dim 2, the variables thermotolerant coliforms, pH, and APHB presented quite strong correlations, as they form acute angles to each other, that is, the increase in pH was affected by the increase in thermotolerant coliforms, and the APHB growth was the most responsible for increasing the pH. There was a strong correlation between R1 and R4 for the variable APHB, with fast growth in R1. A strong correlation between the variables *S. aureus* and *Enterobacteriaceae* was observed, with a strong correlation with R5. In contrast, R2 and R3 were located in opposite directions to all variables, indicating a strong negative correlation, that is, there was no interaction between the variables of these restaurants, thus they can be classified as the best quality restaurants according to the PCA, confirming the results of microbiological and physicochemical characterization.

### 3.2 Antimicrobial resistance profile

Antimicrobial susceptibility tests were applied according to the classification of the Clinical and Laboratory Standards Institute (CLSI, 2010, 2016 e 2018), for drugs of first, second, and third choice for the treatment of pathogenic bacteria isolated in this study. The antimicrobial resistance profile is shown in Table 3, for the isolates *Vibrio parahaemolyticus* (n = 12) *Staphylococcus aureus* (n = 12) and *Salmonella* sp. (n = 1).
Table 3. Results of the antibiogram of *Vibrio parahaemolyticus*, *Staphylococcus aureus* and *Salmonella* sp. from salmon (*Salmo salar*) sashimi ready for consumption in the city of Cuiaba, Mato Grosso, Brazil.

|                          | Resistant N. (%) | Intermediate N. (%) | Sensitive N. (%) |
|--------------------------|------------------|---------------------|------------------|
| **V. parahaemolyticus**  |                  |                     |                  |
| β-lactams                |                  |                     |                  |
| AMP                     | 11/12 (91.6)     | 1/12 (8.3)          | 0/12             |
| CTX                     | 7/12 (58.3)      | 1/12 (8.3)          | 4/12 (33.3)      |
| CFO                     | 9/12 (75.0)      | 1/12 (8.3)          | 2/12 (16.6)      |
| Aminoglycoside          |                  |                     |                  |
| GEN                     | 6/12 (50.0)      | 4/12 (33.3)         | 2/12 (16.6)      |
| AMI                     | 8/12 (66.6)      | 1/12 (8.3)          | (3/12 (25.0)     |
| Tetracycline            |                  |                     |                  |
| TET                     | 11/12 (91.6)     | 1/12 (8.3)          | 0/12             |
| **Salmonella sp.**      |                  |                     |                  |
| β-lactams               |                  |                     |                  |
| AMP                     | 0                | 0                   | 1 (100%)         |
| CTX                     | 0                | 1 (100%)            | 0                |
| CFO                     | 0                | 0                   | 1 (100%)         |
| Aminoglycoside          |                  |                     |                  |
| GEN                     | 0                | 0                   | 1 (100%)         |
| AMI                     | 0                | 1 (100%)            | 0                |
| Tetracycline            |                  |                     |                  |
| TET                     | 0                | 1 (100%)            | 0                |
| **S. aureus**           |                  |                     |                  |
| β-lactams               |                  |                     |                  |
| PEN                     | 12/12 (100.0)    | 0/12                | 0/12             |
| OXA<sup>a</sup>         | 7/12 (58.3)      | 3/12 (25.0)         | 2/12 (16.6)      |
| AMP<sup>b</sup>         | 12/12 (100.0)    | 0/12                | 0/12             |
| CTX<sup>a</sup>         | 3/12 (25.0)      | 9/12 (75.0)         | 0/12             |
| CFO                     | 10/12 (83.3)     | 2/12 (16.6)         | 0/12             |
| Aminoglycoside          |                  |                     |                  |
| GEN                     | 7/12 (58.3)      | 3/12 (25.0)         | 2/12 (16.6)      |
| AMI<sup>b</sup>         | 10/12 (83.3)     | 2/12 (16.6)         | 0/12             |
| Tetracycline            |                  |                     |                  |
| TET                     | 7/12 (58.3)      | 5/12 (41.6)         | 0/12             |
| **Salmonella sp.**      |                  |                     |                  |
| β-lactams               |                  |                     |                  |
| AMP                     | 0                | 0                   | 1 (100%)         |
| CTX                     | 0                | 1 (100%)            | 0                |
| CFO                     | 0                | 0                   | 1 (100%)         |
| Aminoglycoside          |                  |                     |                  |
| GEN                     | 0                | 0                   | 1 (100%)         |
| AMI                     | 0                | 1 (100%)            | 0                |
| Tetracycline            |                  |                     |                  |
| TET                     | 0                | 1 (100%)            | 0                |
| **Sulphonamide**        |                  |                     |                  |
| TMP-SMX                 | 0/12             | 0/12                | 12/12 (100.0)    |

<sup>a</sup>: according to M100-S21 CLSI (2011); <sup>b</sup>: according to M100-S26 CLSI (2016). Source: Authors (2021)

Twelve *S. aureus* isolates were identified, as follows: R2 (n = 1), R3 (n = 3), R4 (n = 1), and R5 (n = 7). The isolates showed 100% resistance to Penicillin and Ampicillin, 83.3% resistance to Cefoxitin and Amikacin, 58.3% resistance to Oxacillin, Gentamicin, and Tetracycline. The lowest resistance was observed for Cefotaxime, with 25%, while Trimethoprim-
sulfamethoxazole was the most sensitive antimicrobial agent for 100% of isolates (Table 3). Given the results, the antibiotic Trimethoprim-sulfamethoxazole can be an effective alternative for the treatment of *S. aureus*, as the bacteria was sensitive to this antibiotic, probably due to the synergistic combination between Sulfamethoxazole and Trimethoprim, which may have impaired the appearance of resistant strains, with a positive classification for first-choice antibiotics (CLSI, 2018). *S. aureus* has been widely studied for antibiotic resistance, and high percentages have been found for multiple drugs. Some authors have reported lower and similar results in fresh marine fish in Iran (Arfatahery et al., 2016) when compared to this study, for Penicillin (91.9% resistance), Ampicillin (89.1% resistance), and Tetracycline (46% resistance), while other authors reported 60% resistance to Tetracycline in Malaysia (Puah, Chua e Tan, 2016). Other studies found low resistance levels to other antimicrobial agents, such as sushi in Denmark (Li et al. 2019), sashimi made with various fish species in Portugal (Moura et al., 2017), imported fish in Switzerland (Boss et al., 2016), and anchovies, trout, and sea bream in Turkey (Onmaz et al., 2015). Moura et al. (2017) reported a higher resistance profile for sashimi made with tuna and sea bass (44.6% and 43.1%), and a lower resistance for salmon (36.9%) and other fish species (35.4 %).

The *Salmonella* sp. strains (R1, n = 1) isolated in this study showed no resistance to all the antimicrobial agents tested. It presented intermediate resistance to Cefotaxime, Amikacin, and Tetracycline, and was sensitive to Ampicillin, Cefoxitin, Gentamicin, and Trimethoprim-sulfamethoxazole, as shown in Table 3. Recent studies in various countries have reported high resistance profile of fresh fish to these antimicrobial agents, such as fresh shrimp (Penicillin 100%, Ampicillin 83.86%, and Tetracycline 45.45%) in Nigeria (BESHIRU et al., 2019); fresh prawn (Ampicillin 21.9%, Tetracycline 25%, and Trimethoprim-sulfamethoxazole 15%) in Vietnam (Nguyen et al., 2016); fresh marine fish (Ampicillin 44.8%) in Jordan (Obdait e Salman, 2017); and anchovies and fresh trout (TMP-SMX and GEN 20%) in Turkey (Onmaz et al., 2015). Knowledge about the sensitivity of *Salmonella* isolates to the antimicrobial agents used in this study is important for public health, as it allows the use of these agents in therapies during outbreaks since *Salmonella* outbreaks have been reported as the first causes of ATD in Brazil and in the world (Brasil, 2019; WHO, 2018).

*Vibrio parahaemolyticus* isolates were found in twelve samples, as follows: R1 (n = 6), R4 (n = 5), and R5 (n = 1). The present results indicated resistance to Ampicillin and Tetracycline (91.6%), Cefoxitin (75%), Amikacin (66.6%), Cefotaxime (58.3%), and Sulfazotrim (25%), with a great sensitivity observed for Sulfazotrim (75%), as shown in Table 3. Although the *V. parahaemolyticus* isolates were resistant to all the antimicrobials used, better results were observed for Trimethoprim-sulfamethoxazole, with sensitivity for the isolates from two restaurants (R4 and R5). According to CLSI (2010), Halophilic *Vibrio* sp. is normally resistant to sulfonamides, penicillins, and other cellelaphosphorins. In this study, resistance to aminoglycosides (Amikacin, Gentamicin), Tetracycline, and semi-synthetic penicillin (Ampicillin) was also observed.

The microorganism *V. parahaemolyticus* is frequently associated with diseases transmitted by seafood, being of concern in fish, mollusk, shrimp and consequently to human health, due to its multi antimicrobial resistance. This microorganism has shown higher resistance rates to Ampicillin (91%), demonstrating the ineffective action of this antibiotic, not being an alternative to combat this species, as also reported by some authors in Korea (Ryu et al., 2019; Kang et al., 2018), Brasil (Silva et al., 2018), Poland (Lopatek et al., 2018), Vietnam (Vu et al., 2016), China (Xu et al., 2016) e no Ecuador (Sperling et al., 2015). Studies have shown that Tetracycline, which exhibited 91% resistance in this study, presented great sensitivity for these bacteria in Korea, Brazil, Poland, and China. Cefotaxime, defined as a first-choice drug by CLSI (2010), showed a significant resistance (58.3%), indicating low *in vitro* activity. Some authors have also reported a strong sensitivity to Trimethoprim-sulfamethoxazole, as observed in the present study (Millanao et al., 2018; Cabello e Godfrey, 2016).

In general, several factors are correlated with increased resistance of antimicrobials, such as overuse in aquaculture (XU et al., 2016) and genetically modified organisms in the aquatic environment. Although some antimicrobial agents behave
similarly for different marine fish and seafood species, such comparison is still difficult, due to the different origin, the collection procedures, and the methodologies used, as observed by Lopatek et al. (2018).

4. Conclusion

It was concluded that 48% of sashimi samples were suitable for human consumption, while 52% presented unsatisfactory hygienic-sanitary quality for the parameters evaluated, which can be a risk to the consumers' health, such as the development of a Foodborne Disease (FBD). The restaurants R2 and R3 presented better hygienic-sanitary conditions when compared to R1, R4, and R5.

The antimicrobial resistance profile revealed that both the S. aureus and V. parahaemolyticus isolates showed sensitivity in vitro only to Trimethoprim-sulfamethoxazole, which is a challenge related to the guidelines to assist with treatment choices in the case of foodborne illness.

Based on this study, it is suggested that similar work be carried out in several cities in the country for a broader survey of the quality of this food matrix, as well as scientific studies related to handling and raw materials.

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