

## Influence of cutting room temperature on the microbiological quality of chicken breast meat

[Influência da temperatura da sala de cortes sobre a qualidade microbiológica da carne de peito de frango]

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### ABSTRACT

The temperature control in the processing room is one of the major factors associated with the production of safe food with a satisfactory microbiological quality. A total of 288 samples of skinless chicken breast meat were placed in a cutting room, subjected to four different temperatures (12°C, 14°C, 16°C and 18°C) and collected to evaluate the influence of the room temperature on the microbiological quality during the cutting and boning of chicken breasts. Aerobic mesophilic microorganisms were counted to evaluate the environmental contamination. In addition, coliforms at 35°C and 45°C and *Staphylococcus* spp. were counted, and an analysis for the presence of staphylococcal enterotoxins and *Salmonella* spp. was performed to determine the microbiological quality of the meat. The results showed an increase in environmental contamination ( $P=0.01$ ) with an increase in room temperature. However, no significant differences ( $P>0.05$ ) were observed in the meat cuts regarding the counts of coliforms at 35°C and 45°C, the count of *Staphylococcus* spp. and the presence of *Salmonella* spp. Moreover, no staphylococcal enterotoxins were detected in the samples analyzed. Thus, despite increasing the environmental contamination, the increase in the cutting room temperature did not affect the microbiological quality of the final product.

Keywords: Microbiological quality, environmental contamination, chicken breast meat, cutting room temperature

### RESUMO

O controle da temperatura do ambiente de processamento é um dos principais fatores relacionados à produção de alimentos seguros e com qualidade microbiológica. Com o objetivo de avaliar a influência da temperatura ambiente durante o corte e a desossa da carne de frangos sobre a qualidade microbiológica dos produtos finais, foram coletadas 288 amostras de carne de peito de frango sem pele, obtidas em uma sala de cortes climatizada submetida a quatro diferentes temperaturas ambientes (12°C, 14°C, 16°C e 18°C). Para avaliação da contaminação ambiental, foi realizada a contagem de microrganismos mesófilos aeróbios e, para a avaliação da qualidade microbiológica da carne, foram realizadas a contagem de coliformes totais e termotolerantes, a contagem de *Staphylococcus* spp., a pesquisa de enterotoxinas estafilocócicas e a pesquisa de *Salmonella* spp. Os resultados encontrados demonstraram um aumento da contaminação ambiental ( $P=0,01$ ) à medida que a temperatura da sala foi aumentada. Porém, nos cortes cárneos, não foram observadas diferenças significativas ( $P>0,05$ ) na contagem de coliformes totais e termotolerantes, na contagem de *Staphylococcus* spp. e na pesquisa de *Salmonella* spp. Também não foi detectada a presença de enterotoxinas estafilocócicas nas amostras analisadas. Foi concluído que, apesar da elevação da contaminação ambiental, o aumento da temperatura ambiente da sala de cortes não comprometeu a qualidade microbiológica do produto final.

Palavras-chave: qualidade microbiológica, contaminação ambiental, carne de peito de frango, temperatura da sala de cortes

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## INTRODUCTION

Chicken meat is an excellent source of protein, B-complex vitamins and minerals. Due to its intrinsic characteristics, such as its high nutrient components, high water activity and pH close to neutral, chicken meat also provides an excellent environment for the growth of microorganisms.

The microorganisms that contaminate chicken meat derive from the microbiota of the skin and feathers of these animals and can also originate from the nasal cavity, mouth, skin and gastrointestinal tract of the handlers. In addition, bacteria present on the surfaces of equipment, utensils and containers can be sources of food contamination. The air is another important contaminating factor because microorganism-containing aerosols and particles are spread through the air (Jay *et al.*, 2005; Silva Junior, 2007).

The growth of microorganisms in food depends on extrinsic and intrinsic factors; thus, a series of standards and techniques must be followed within the slaughter and processing industries of broilers to minimize contamination of the carcasses. The room in which the meat cutting is performed is one of these environments that requires monitoring, and the room temperature must be kept below 12°C, according to Regulation (EC) 853/2004 (European Commission, 2004). The rationale for the use of lower temperatures is to reduce microbial, chemical, and enzymatic activities that can alter overall food quality because the rate of such changes is directly proportional to the temperature. Thus, the surface temperature of the meat may also favor the growth of microorganisms and affect the quality of the final product (Borges and Freitas, 2002).

There are few reports in the literature about this subject. The aim of this study was to evaluate the influence of the cutting room temperature on the microbiological quality of chicken breast meat.

## MATERIALS AND METHODS

The experimental samples were collected directly from a meatpacking plant under official

veterinary inspection; this plant had implemented quality control systems (Hazard Analysis and Critical Control Points - HACCP).

To perform the microbiological analyses of the meat, 288 whole broiler carcasses and 288 skinless chicken breast cuts were collected. The carcasses were collected shortly after leaving the pre-cooling area and before entering the cutting room, and the skinless chicken breast cuts were collected immediately after their production, which was performed in a processing room with a controlled temperature. The microbiological analyses of the carcasses were performed to evaluate the initial contamination and assess their uniformity.

For the analyses of the chicken breast cuts, four treatments, corresponding to the four temperatures (12°C, 14°C, 16°C and 18°C) to which the cutting room was subjected were evaluated. The same cutting room was used for all treatments and samples were only collected after stabilization of the temperature according to the each treatment. Twelve replicates were used per treatment. Each work shift was considered a replicate, and each replicate consisted of a pool of six skinless chicken breast cuts, resulting in a total of 48 samples analyzed.

For the analyses of whole broiler carcasses, pools of six carcasses from each of the 48 experimental batches that went to the cutting room were used. The temperatures of the carcasses (*Pectoralis major* muscle) before entering the processing room and of the chicken breast collected after the cutting and boning were measured. The time between the entry of the carcasses in the processing room and obtaining the chicken breast cuts was three minutes (according to the routine in the industry); this time was similar for all of the treatments evaluated.

Throughout the experiment, the maximum and minimum temperatures of the cutting room were monitored for the different treatments and are recorded in Table 1.

Table. 1. Mean temperatures (°C) and standard deviations measured in the cutting and boning room for the different treatments

Treatments	Mean temperature (°C)
12°C	12.02 ± 0.11
14°C	14.05 ± 0.10
16°C	16.10 ± 0.10
18°C	18.02 ± 0.15

The environmental contamination of the air-conditioned cutting room in the different treatments was assessed by counting the aerobic mesophilic microorganisms (Marshall, 1992). For these analyses, Petrifilm™ Aerobic Count (AC) plates (3M™, Maplewood, MN, USA) were hydrated with 1mL of 0.1% saline solution and exposed in different places in the room for 15 minutes. Then, the plates were incubated at 35±1°C for 48 hours, after which the colonies were counted. The result was expressed in microbial density (CFU/m<sup>2</sup>/hour), which was calculated based on the following formula:

$$\text{Microbial density} = \text{CFU} \times 60\text{min} / 15\text{min} \times 10000\text{cm}^2 / 20\text{cm}^2$$

The microbiological quality of the samples was evaluated by counting coliforms at 35°C and 45°C, *Staphylococcus aureus* count, presence of staphylococcal enterotoxins and *Salmonella* spp.

Coliforms at 35°C and 45°C in the carcasses and chicken breast cuts (CFU.g<sup>-1</sup>) were counted using the Petrifilm™ EC method (Curiale et al., 1991). For these analyses, 25-g aliquots of the samples were added to 225mL of 0.1% peptone salt solution and homogenized with a Stomacher (Interscience, Saint Nom La Breteche, France) for 60 seconds. The readings were performed after incubation of the plates at 35±1°C for 48 hours.

*Staphylococcus aureus* analyses were performed according to the procedures described in the Bacteriological Analytical Manual (Bennett and Lancette, 2001). For this purpose, 25g of each sample were weighed, and 225mL of 0.1% peptone salt solution was added. After homogenization of the samples, serial dilutions were performed and plated on Baird-Parker agar plates, which were incubated at 36±1°C for 48 hours. The isolated colonies were inoculated into brain heart infusion (BHI) broth (Oxoid,

Hampshire, UK) and subjected to the coagulase test using rabbit plasma (Laborclin, Pinhais, PR, Brazil). In addition to the coagulase test, two complementary tests were also performed: Gram staining, to assess morphology, and the thermonuclease test using DNase Test Agar (BD, New Jersey, USA). Because no *Staphylococcus aureus* were detected in the analyses performed throughout the entire experiment, the results were expressed as *Staphylococcus* spp.

To detect the staphylococcal enterotoxins (A, B, C<sup>1</sup>, C<sup>2</sup>, C<sup>3</sup>, D, and E), the enzyme linked fluorescent assay (ELFA) technology associated with the VIDAS™30 equipment (Hennekinne et al., 2007). For this purpose, 25g of each sample were homogenized with 25mL of the extraction solution (Tris buffer, 0.3M, pH 8.0) for 3 minutes to obtain a homogeneous suspension. The samples were centrifuged for 15 minutes at 3000-5000x g, and the supernatant was filtered. Then, 500µL of the filtrate (pH 7.5 - 8.0) was transferred to VIDAS® Staph Enterotoxin - SET2 strips. The results were read 80 minutes after inserting the strips into the equipment.

The analysis of *Salmonella* spp. was performed according to the AOAC International (2005). A 25-g aliquot of each sample was added to 225mL of 1% sterile peptone salt solution and incubated at 35°C for 24 hours. After the pre-enrichment, 100µL of the culture was transferred to Salmonella Xpress broth (bioMérieux, Hazelwood, MO, USA) and incubated at 41°C for 24 hours. Then, 500 µL was transferred to specific strips for the detection of *Salmonella* spp. The strips were heated at 100°C for 15 minutes and placed in the VIDAS™30 equipment (bioMérieux, Hazelwood, MO, USA). The reading was performed after 45 minutes. The presence of *Salmonella* spp. was confirmed through biochemical tests using Vitek™2 equipment (bioMérieux, Hazelwood, MO, USA) (Biomeriëux, 2008).

The analyses were performed using a completely randomized design with the chicken breast cut as the covariate and the estimate of the regression models for the treatments. Four treatments, corresponding to the temperature to which the cutting room was subjected (12°C, 14°C, 16°C and 18°C), were evaluated using 12 replicates of six chicken breast each. The data were transformed to evaluate the microbiological

variables, and for those with a normal distribution (environmental contamination), Tukey's test was used with a significance level of 5%. Those without a normal distribution and/or homoscedasticity were evaluated by the Kruskal-Wallis test, also with a significance level of 5%. The temperature data of the carcasses and chicken breast cuts were evaluated using Tukey's test with a significance level of 5%. The results for the *Salmonella* spp. and staphylococcal enterotoxins were subjected to

descriptive analyses due to the low number of positive samples observed in the experiment.

## RESULTS AND DISCUSSION

The results of coliforms at 35°C and 45°C and *Staphylococcus* spp. in the carcasses that were collected before entering the cutting room showed no significant differences ( $P > 0.05$ ) (Table 2). Staphylococcal enterotoxins and *Salmonella* spp. were also not detected.

Table 2. Results of the microbiological analyses (CFU.g<sup>-1</sup>) of the carcasses collected before entering the cutting room according to the four treatments

Treatment	Coliforms 35°C			Coliforms 45°C			<i>Staphylococcus</i> spp.		
	Min	Max	Median	Min	Max	Median	Min	Max	Median
12°C	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	1.0x10 <sup>4</sup>	<1.0x10 <sup>2</sup>
14°C	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	7.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>
16°C	<1.0x10 <sup>2</sup>	2.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	3.0x10 <sup>4</sup>	2.2x10 <sup>2</sup>
18°C	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	5.5x10 <sup>3</sup>	<1.0x10 <sup>2</sup>

Min = minimum; Max = maximum.

The results were similar according to the Kruskal-Wallis test ( $P > 0.05$ ).

The mean results of the carcass temperatures measured before entering the cutting room are indicated in Table 3. The carcasses from the 48 batches used for the experimental evaluation had similar temperatures ( $P > 0.05$ ).

Table 3. Mean temperatures (°C) and standard deviations of the carcasses collected before entering the cutting room according to the four treatments

Treatments	Carcasses (°C)
12°C	5.28 ± 0.29
14°C	5.32 ± 0.22
16°C	5.16 ± 0.29
18°C	5.19 ± 0.40

The means were similar according to Tukey's test ( $P > 0.05$ ). Coefficient of variation (CV) = 5.88%.

The microbiological results of the carcasses and the assessment of their temperature showed that all of the batches used in this study had uniform conditions and were thus considered suitable for the experimental evaluation of the chicken breast cuts.

The results of the count of aerobic mesophilic microorganisms that were used to evaluate the environmental contamination of the cutting room at four different temperatures are shown in Table 4 and Figure 1. A significant difference was observed ( $P < 0.05$ ) between the microbial density and the different temperatures of the cutting room.

Table 4. Mean results of the environmental contamination analyses of the cutting room, subjected to different temperatures

Temperature (°C)	Microbial density (CFU/m <sup>2</sup> /hour)
12°C	2.0x10 <sup>4</sup> a
14°C	3.5x10 <sup>4</sup> ab
16°C	5.2x10 <sup>4</sup> ab
18°C	6.0x10 <sup>4</sup> b

The means followed by different letters in the column differed significantly according to Tukey's test ( $P = 0.01$ ). CV = 8.0%.

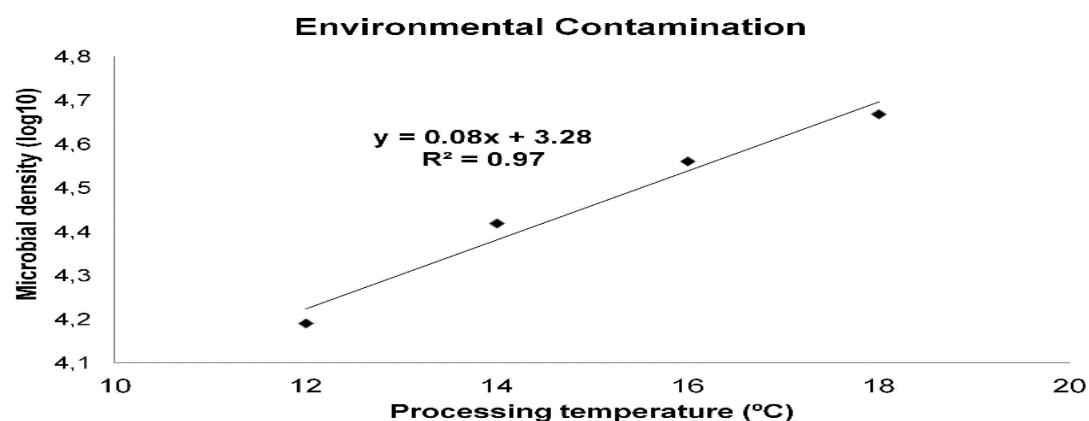


Figure 1. Linear regression plot of the environmental contamination *versus* the different temperatures.

The results demonstrated that an increase in temperature in the cutting room led to an increase in the count of aerobic mesophilic microorganisms in the environment. In fact, a significant difference ( $P=0.01$ ) was observed as the cutting room temperature was increased from 12 to 18°C. These results were expected because the conditions for the development and multiplication of aerobic mesophilic

microorganisms are more appropriate at higher temperatures.

The results of the microbiological evaluation of the chicken breast cuts produced in the cutting room under different temperatures are shown in Table 5. No significant differences ( $P>0.05$ ) were observed between coliforms at 35°C and 45°C counts and *Staphylococcus* spp. in the skinless chicken breast cuts.

Table 5. Results of the microbiological analyses (CFU.g<sup>-1</sup>) of the broiler chicken breast samples, according to the temperature of the cutting room

Treatment	Coliforms 35°C			Coliforms 45°C			<i>Staphylococcus</i> spp.		
	Min	Max	Median	Min	Max	Median	Min	Max	Median
12°C	<1.0x10 <sup>2</sup>	3.0x10 <sup>3</sup>	1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	4.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	1.6x10 <sup>4</sup>	7.8x10 <sup>2</sup>
14°C	<1.0x10 <sup>2</sup>	3.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	1.6x10 <sup>4</sup>	9.0x10 <sup>2</sup>
16°C	<1.0x10 <sup>2</sup>	5.0x10 <sup>2</sup>	1.5x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	5.1x10 <sup>3</sup>	6.0x10 <sup>2</sup>
18°C	<1.0x10 <sup>2</sup>	4.0x10 <sup>2</sup>	1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	2.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	<1.0x10 <sup>2</sup>	5.3x10 <sup>3</sup>	2.0x10 <sup>2</sup>

Min = minimum; Max = maximum.

The results were similar according to the Kruskal-Wallis test ( $P>0.05$ ).

Low counts of coliforms at 35°C and 45°C were observed in the breast meat samples processed in the cutting room under different temperatures (12°C, 14°C, 16°C and 18°C). The counts obtained met the requirements of the official inspection agencies, which stipulate a maximum value of 10<sup>4</sup> CFU.g<sup>-1</sup> for coliforms at 45°C (Brazil, 2001). These microorganisms are sanitary indicators, and their presence in food suggests that other fecal pathogens may be present (Jay *et al.*, 2005). Thus, the low incidence of these microorganisms represents a low risk of the presence of meat-contaminating pathogenic bacteria.

The counts of *Staphylococcus* spp. were considered low, and no staphylococcal enterotoxins (A, B, C<sup>1</sup>, C<sup>2</sup>, C<sup>3</sup>, D and E) were observed. The low counts of *Staphylococcus* spp. may explain the absence of staphylococcal enterotoxins in the chicken breast meat because this microorganism has the potential to produce toxins when present in levels exceeding 10<sup>5</sup> CFU.g<sup>-1</sup> (Valero *et al.*, 2009). Regarding *Salmonella* spp., only one (2.08%) chicken breast sample collected in the room at a temperature of 16°C was positive for this microorganism.

The results of the mean temperatures of the chicken breast cuts are described in Table 6. No significant differences ( $P>0.05$ ) were observed in the temperature of the meat cuts between treatments.

Table 6. Mean temperatures (°C) and standard deviations of the broiler breast cuts according to the treatments

Treatments	Chicken breast (°C)
12°C	5.40 ± 0.31
14°C	5.30 ± 0.24
16°C	5.38 ± 0.18
18°C	5.45 ± 0.24

The means were similar according to Tukey's test ( $P<0.05$ ) CV = 4.60%.

The time between the entry of the carcass in the cutting room and obtaining the breast cut was only three minutes. This short period can explain the consistently low temperature of the meat cuts, regardless of the variation in the room temperature.

The microbiological quality of the carcasses and chicken breast cuts reflects the efficiency of the quality control programs adopted by the company. Because of the differences in processing conditions and quality control found in slaughterhouses, further studies are needed in other processing plants to confirm or not the influence of the cutting room temperature on the microbiological quality of poultry meat.

### CONCLUSION

Despite the increase in environmental microbial contamination, increasing the temperature of the cutting room did not affect the microbiological quality of the chicken breast meat.

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