TUENCE OF SEQUENCE OF TREATMENTS ON OXIDATION OF PORK MUSCLE TISSUE.

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## mary

influence on lipid oxidation of different sequences of treatments such as sample size to location of ground samples), cooking, refrigerated storage and frozen storage was intigated. A modified TBA method was used to measured oxidation on porcine longissimus muscle samples. Results indicate that, the sequence in which these treatments were applied to muscle samples has a significant (P<0.05) effect on TBA values.

## troduction

investigations have been conducted to determine the influence of tissue size reduction. cooking, storage conditions and their cumulative effect on oxidation and consequently med-over flavor (WOF) (Chang et al., 1961; Keskinel et al., 1964; Huang and Greene, 1978; he et al., 1981; Pikul et al., 1984a). However, the effect on lipid oxidation of the wence in which these processes are conducted has not been systematically studied. The 2-lobarbituric test (TBA) has been mostly used to measure oxidation in meat, but it not sible to determine the sequence effect by comparing studies performed in different contains an analysis because TBA number varies when different experimental conditions or diffications of the method are utilized (Newburg and Cocon, 1980; Melton, 1983; Igene et 1985; Willemot et al., 1985; Williams et al., 1983; Pikul et al., 1984b). Considering the objective of the present research is to evaluate the influence that different mences of common meat industry procedures (grinding, slicing, cooking, and refrigerated frozen storage) have on oxidation of pork muscle tissue as measured by TBA method.

# terials and Methods

Treatments: As it is shown in Figure 1, the samples were randomly distributed among the preparation (prep.) treatment and thirteen time-temperature (tm-tp) treatments.

Source and Sample Preparation: Pork loins from both sides of an individual pork car-Were purchased from a local packer. At the laboratory, the loins were deboned and the Rissimus dorsi muscles (LD) were excised and trimmed. The LD's were arbitrarily divided eight pieces between the proximal and distal ends and randomly distributed among the treatments. The sizes of the portions were calculated considering that the samples epared from them must be the necessary to be distributed among the tm-tp treatments. plicate patties and slices were prepared for each tm-tp treatment. For the cooked samples extra 35% more of sample was calculated to prevent cooking loss. Whole samples of .5cm thick, slices samples of 0.5±0.1cm thick and 20±0.5g patties of 5.5cm diameter and thick were prepared. The patties were made from meat that was cubed (4x4x4cm) and then through a 0.48 cm plate. The slices were sliced with an electric slicer (Hobart Model After being prepared all samples were individually wrapped in aluminum foil and taped hasking tape and then cooked or storage. The cooked samples were cooked in a convection (The G. S. Blodgett Company, Inc.) at 93.3°C and 1125 rpm blower speed to a final intemperature of 70±2°C. The refrigeration and freezing temperatures were 4±2°C and respectively. Each individual sample was weighed and its weight recorded before and ter being treated.

method of Witte et al. (1970) as modified by Pensel (1990). Blanks and standard curve ing 1,1,3,3-Tetraethoxypropane (TEP) were run simultaneously. The TEP recovery was 91% and value was 10.64, similar to the value reported by Witte et al. (1970) considering that of sample was used in this case. The results were expressed as TBA numbers in terms of malonaldehyde per 1 kilogram of meat. TBA values reported are the mean of

five replicates. To consider the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples, it was never the cooking drip and storage loss of the samples and storage loss of the sampl sary to correct the weight to a standard fresh weight (Pensel, 1990).

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Data Analysis: The statistical analysis was completed using a two-factor (prep. and to the statistical analysis was completed using a two-factor (prep. and to the statistical analysis was completed using a two-factor (prep. and to the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. and the statistical analysis was completed using a two-factor (prep. analysis) was complete treatments) fixed effect model with blocking (replications). Due to the interaction between main effects, multiple comparisons were conducted based on the "marginal" multiple TON parisons (Pensel, 1990). For this reason, WU, SU, CS and 0 time treatment were not includes. in Table 1. Analysis of variance and Duncan multiple range test were performed using BURG (1985) procedures.

### Results and Discussion

It can be seen from the statistical comparison in Table 1, that TBA numbers for CG samp were significantly higher (P<0.05) than the comparable TBA numbers obtained from SC,  $G^{C}$  oker and GU samples at all Tm-Tp treatments. Except at 0 time, the GU samples had significantly because of the comparable of the compa lower (P<0.05) TBA numbers when compared with CG, SC, GC and WC samples. The length frozen storage did not produce significant differences whatever the size of the samples. 6F samples were not significantly different from 2F samples, neither 2R-2F or 2R-6F significantly different from 6R-2F and 6R-6F respectively. Contrarily, TBA values of 68 samples were higher than TBA values of 6F-6R samples. In general, refrigeration and  $co^{00}$ , 19 lead increases in TBA values (Dawson and Schierholz, 1976) independently of the sample However, the most detrimental combination of treatments was reducing the sample size cooking. This can be seen in Figure 2, where the TBA numbers for the samples that we cooked and then ground or sliced (CG, CS) presented always the highest values, the cooked samples had intermediate values and the cooked always the highest values. This det samples had intermediate values and the uncooked samples had the lowest TBA values. shows the importance of the sequence in which the processes are carried out. For example, and SC samples were subjected to the same processes but in reverse sequence, however results were significantly different with CS samples resulting in considerable more  $o^{\chi_{i}^{j,j}}$ tion. Despite the disruption or denaturation of the cell membrane with the concomit liberation of phospholipids, release of iron from the heme and incorporation of oxygen covered be contributing factors for the increase of TBA numbers (Gray and Pearson, 1987: 1988), the sequence in which these processes occur also should be considered a factor.

which BF= Based on the results of this study, it is possible to conclude that the sequence in the processes were carried out had a significant influence on TBA numbers. Cooking follow by grinding and then six days of frozen storage and six additional days of refrigeral storage was the most detrimental combination of treatments. Then to predict the effect certain treatments have on oxidation using TBA analysis, it is necessary to sequence and storage history of the sample.

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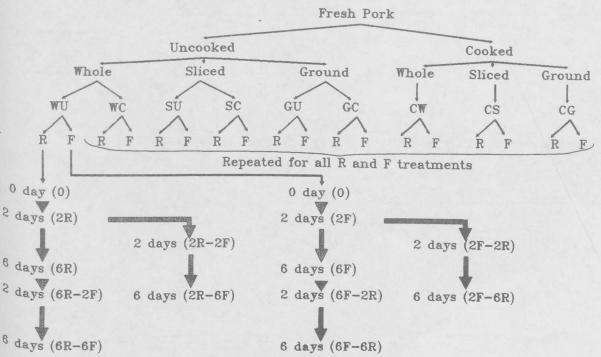
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# ure 1: Treatments Design

ted, GU = ground uncooked, GC = ground and then cooked, CW or WC = whole cooked, CS = cooked and then then cooked and then cooked and then cooked and then ground.

temperature treatments: R= refrigerated storage, F= frozen storage, = indicates refrigerated = indicates frozen storage, 0= zero day, 6F-6R= 6 days F and then 6 days R, 6R-2F= 6 and then 2 F, 6R= 6 days R, 2F-6R= 2 days F and then 6 days R, 6R-6F= 6 days R and then 6 days F, 6F-2R= 6 days F and 2 days R, 2F-2R= 2 days F and then 2 days R, 2F-2R= 2 days R and then 6 days F, 2R= 2 days R, 2F= 2 days F, 6F= 6 days F.



## Table 1. Effect of Different Sequences of Treatments on TBA Numbers a,b

a: Cells in the same column or row, bearing the same letter are not significantly different (P<0.05)

b: TBA Number is expressed as milligrams malohaldehyde/kilogram muscle. Each value represents a mean of five replicates (±SD).

CG= cooked and then ground, SC= sliced and then cooked, GC= ground and then cooked, WC= whole cooked. GU= ground uncooked, R= refrigerated storage, F= frozen storage, 6F-6R= 6 days F and the 6 days R, 6R-2F= 6 days R and then 2 F, 6R= 6 days R, 2F-6R= 2 days F and then 6 days F, 6R-6F= 6 days R and then 6 days F, 6F-2R= 6 days F and 2 days R, 2F-2R= 2 days F and then 2 days R, 2R-6F= 2 days R and then 6 days F, 2R= 2 days R, 2F= 2 days F, 6F= 6 days F.

	6F-6R	6R-2F	6R	2F-6R	6R-6F	2R-2F	6F-2R	2F-2R	2R-6F	2R	2F	6F
C G	A 1.783 ± 0.12	A B 1.779 ± 0.10	A B C 1.742 ± 0.12	A B C 1.743 ± 0.15	B C D 1.772 ± 0.15	C D E 1.707 ± 0.11	C D E 1.732 ± 0.09	DEF 1.772 ± 0.11	E F G 1.536 ± 0.07	F G 1.585 ± 0.07	G 1.480 ± 0.17	G 1.418 ± 0.10
SC	H 0.652 ± 0.25	H I 0.761 ± 0.36	H I J 0.657 ± 0.38	HIJ 0.450 ± 0.23	I J K 0.471 ± 0.16	J K L 0.617 ± 0.26	J L K 0.354 ± 0.20	K L M 0.368 ± 0.22	L M N 0.274 ± 0.07	M N 0.316 ± 0.16	N 0.235 ± 0.08	N 0.227 ± 0.18
G C	H 0.577 ± 0.39	H I 0.560 ± 0.30	HIJ 0.308 ± 0.05	HIJ 0.641 ± 0.38	IJK 0.590 ± 0.28	JKL 0.493 ± 0.14	J K L 0.481 ± 0.18	KLM 0.351 ± 0.17	L M N 0.438 ± 0.14	M N 0.251 ± 0.08	N 0.324 ± 0.08	0.203 0.07
W C	H 0.761 ± 0.29	H I 0.549 ± 0.34	HIJ 0.623 ± 0.50	HIJ 0.472 ± 0.22	I J K 0.354 ± 0.15	J K L 0.248 ± 0.09	J K L 0.462 ± 0.14	K L M 0.315 ± 0.13	L M N 0.285 ± 0.05	M N 0.213 ± 0.05	N 0.064 ± 0.05	0.190 ± 0.21
G U	0 0.217 ± 0.15	0 P 0.225 ± 0.17	0 P Q 0.299 ± 0.27	0 P Q 0.154 ± 0.20	P Q R 0.187 ± 0.08	Q R S 0.116 ± 0.07	Q R S 0.079 ± 0.07	R S T 0.118 ± 0.08	S T U 0.170 ± 0.08	T U 0.120 ± 0.07	U 0.068 ± 0.06	0.109 ± 0.05

Figure 2: Effect of Treatment Combinations on TBA Numbers

a- 0, 2R, 6R, 6R\_2F, 6R-6F Tm-Tp Treat. c- 0, 2F, 6F, 6F-2R, 6F-6R Tm-Tp Treat.

b- 0, 2R, 2R-2F, 2R-6F Tm-Tp Treat. d- 0, 2F, 2F-2R, 2F-6R Tm-Tp Treat. OSPI

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