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Effect of the exposure to oxidation and malondialdehyde on turkey and rabbit meat protein oxidative stability

Francesca Soglia, Giulia Baldi, and Massimiliano Petracci

Abstract: The present study aimed at evaluating the effect of the exposure to a strong oxidative environment (100 mM NaClO) and the concurrent incubation with different malondialdehyde (MDA) concentrations (0 to 5 mM) on protein carbonylation, free thiol groups, total heme pigments, and on the relative concentration of the different myoglobin (Mb) derivatives in turkey thigh and rabbit hind leg meat to elucidate their eventual role in inducing oxidative modifications on the protein fraction. With regard to turkey meat, the addition of a strong oxidant resulted in remarkably higher ($P < 0.001$) carbonyls along with a reduction in free thiol groups (which become undetectable). The relative concentration of MbO₂ and MetMb was significantly affected ($P < 0.001$) and total heme pigment was reduced by 62% when fresh meat is compared to its oxidized counterpart. The addition of MDA 2.5 mM (or greater) resulted in a 1.4-fold increase in carbonyls and a tendency ($P = 0.07$) has been observed for free thiol groups whose content decreased as the MDA level increased. Overall, the aforementioned results are in agreement with those obtained on rabbit hind leg meat in which a three-fold increase ($P < 0.001$) in carbonyls and a remarkably higher ($P < 0.001$) MetMb content coupled with a 53% reduction in free thiol groups were found. No significant differences ascribable to MDA addition were found. In conclusion, exposing turkey and rabbit meat to the same environment resulted in distinctive oxidative changes that might be related to the compositional traits and a species-specific susceptibility of the meat toward oxidation.

Keywords: malondialdehyde, meat, oxidation, protein, turkey and rabbit

Practical Application: Protein and lipid oxidation have been investigated in different meat types; nevertheless, the knowledge concerning the interaction between these two phenomena is limited. According to their compositional traits (high PUFA and heme pigment content), turkey and rabbit leg meats are particularly susceptible to oxidation. Providing information concerning the oxidative stability of turkey thigh and rabbit hind leg meat exposed to strong oxidative conditions and to a concurrent increasing content of a secondary product of lipid oxidation, the findings of the present study can be useful when proper processing strategies and storage conditions have to be implemented for manufacturing processed products.

1. INTRODUCTION

The oxidative processes affecting protein and lipid in meat and meat products are free radical-mediated chain reactions initiated by the abstraction of a hydrogen atom by reactive oxygen species (ROS) that results in the generation of a protein carbon-centered and a lipid radical, respectively (Estévez, 2011; Guyon, Meynier, & de Lamballerie, 2016; Lund, Heinonen, Baron, & Estévez, 2011; Min & Ahn, 2005). These unstable radicals further react, thus resulting in the development of final oxidation products, which differ depending on the target molecules, as well as on the oxidative conditions (Estévez, 2011). In detail, after exposing the meat to strong oxidative conditions, radicals reacting with proteins and peptides may result in alterations involving both the backbone of the polypeptide chains and the amino acids side chains. Indeed, the cleavage of peptide bonds resulting in protein fragmentation and intermolecular covalent cross-linked derivatives may occur to-

gether with the formation of carbonyl groups and hydroperoxides and the loss of cysteine and tyrosine residues (Lund et al., 2011). On the other hand, lipid oxidation results in the development of hydroperoxides, which can further react thus leading to a wide range of secondary products, including pentanal, hexanal, 4-hydroxynonenal, and malondialdehyde (MDA) (Fernández, Pérez-Álvarez, & Fernández-López, 1997; Lynch & Faustman, 2000). Being the main nonmicrobial phenomenon responsible for meat deterioration (as it affects color, texture, flavor, loss of nutritional value, and development of harmful products), lipid oxidation has been extensively studied in meat and meat products (Domínguez et al., 2019). On the other hand, the chemical mechanisms resulting in protein oxidation and its impact on meat quality have raised the attention of the meat scientists later on with the discovery that also the polypeptide chains may undergo oxidative reactions during meat storage and processing. Interestingly, as radicals can be reciprocally transferred between lipids and proteins, lipid-derived ROS and several lipid oxidation products might promote protein oxidation and catalyze the oxidative modifications taking place on their polypeptide chains (Estévez, 2011). Within this context, although the concomitant occurrence of lipid and protein oxidation in meat systems may suggest a possible interaction between these

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two phenomena (Lund et al., 2011), the knowledge concerning the effect of the concomitant exposure to oxidative conditions and secondary products of lipid oxidation on protein oxidative stability is limited. In detail, a recent study evaluated the effect of MDA exposure only on the oxidative stability of the myofibrillar proteins extracted from rabbit meat (Wang, He, Emara, Gan, & Li, 2019), whereas no information is available on meat as raw material. Although commonly referred to as “white meats,” poultry and rabbit leg meats have relatively high PUFA and heme pigment contents, which make them particularly susceptible to the development of oxidative reactions (Hernandez & Gondret, 2006; Dalle Zotte & Szendro, 2011; Estévez, 2015; Mercier, Gatellier, Viau, Remignon, & Renner, 1998; Petracci & Cavani, 2013). Recently, the growing interest of both the rabbit and turkey industries in developing processed meat products (Barbut, 2015; Cullere & Dalle Zotte, 2018; Petracci, Soglia, & Leroy, 2018; Soglia et al., 2018) shed the light on the need to improve the knowledge concerning the impact of different processing steps (i.e., grinding and cooking) on the oxidative stability of the lipid fraction whose secondary products might promote oxidative modifications on proteins' polypeptide chains and affect their functional and technological properties (Estevez, 2015; Petracci & Cavani, 2013). Within this context, the present study aimed at evaluating the effect of the exposure to oxidative conditions and the concurrent incubation with different MDA concentrations (from 0 to 5 mM) on protein carbonylation, free thiol groups, total heme pigments, as well as on the relative concentration of the different myoglobin derivatives in turkey thigh and rabbit hind leg meat in order to elucidate their eventual role in inducing oxidative modifications on the protein fraction.

2. MATERIALS AND METHODS

2.1 Sample preparation

Five turkey carcasses were collected 24 hr post-mortem from the same flock (BUT BIG 6 hybrid, 100 days-old, average live weight of 9.7 kg) farmed, and processed under commercial conditions. After dissection, the *Extensor iliobtibialis lateralis* muscles were finely minced with a grinder and divided into 12 aliquots/each: six to be added with NaClO to a final concentration of 100 mM (to obtain a strong pro-oxidative environment), whereas the same volume of distilled H₂O was added to the others (in order to achieve the same dilution of the protein fraction). Then, fresh and oxidized samples were added with different final concentrations of MDA (i.e., 0, 0.25, 0.5, 1.0, 2.5, and 5.0 mM) and incubated in the dark at 37 ± 1 °C for 24 hr. No antimicrobial agents were added to the samples in order to inhibit microbial growth. However, the eventual changes in meat quality ascribable to microbial growth during the 24 hr incubation time would definitely be of a lower extent if compared to those related to either the addition of different MDA levels or to the exposure to oxidative conditions, considered as main factors in the present research.

After that, samples were stored at -80 °C until analyses carried out to assess the occurrence of protein oxidation, through the quantification of carbonyls and free thiol groups, total heme pigments content, and the relative concentration of the different myoglobin derivatives, such as deoxymyoglobin (DeoxyMb), oxymyoglobin (MbO₂), and metmyoglobin (MetMb). The same experiment was performed on rabbit hind legs meat by collecting 24 hr post-mortem five carcasses belonging to the same batch (Martini line, 10 weeks old, average live weight of 2.7 kg) farmed, and slaughtered in a commercial abattoir. All the muscles belong-

ing from this anatomical region were finely minced and samples were prepared and treated as described before.

2.2 Total heme pigments and myoglobin derivatives

The concentration of total heme pigments as well as the relative concentration of the different myoglobin derivatives (DeoxyMb, MbO₂, and MetMb) was determined following the procedure described by Krzywicki (1982) with slight modifications.

Briefly, 2.5 g of sample was homogenized (13,500 rpm for 10 s) with an Ultra-Turrax T25 (IKA – WERKE, Labor Technik, Staufen, Germany) in 12.5 mL of 40 mM phosphate buffer (pH 6.8). Following incubation for 1 hr at 4 ± 1 °C, samples were centrifuged (Beckman Coulter JP-26, Indianapolis, IN, USA) (15,000 × g for 30 min at 4 °C) and the supernatant filtered through filter paper (Whatman 1, GE Healthcare) and through cellulose acetate syringe filter (20 μm). Then, the absorbance of the samples was read against blank at 730, 525, 545, 565, and 572 nm and total heme pigment as well as the relative concentration of DeoxyMb, MbO₂, and MetMb was calculated as follows, where R1 = Abs 572/Abs 525, R2 = Abs 565/Abs 525, and R3 = Abs 545/Abs 525. All the absorbances were corrected taking into account the turbidity of the sample (Abs 730 nm).

Total heme pigments (mg Mb/kg of meat)

$$= \frac{[(0.661 R1 + 0.086 R2 + 0.088 R3 + 0.090) \times \text{Abs}525] \times 17,500}{\text{sample weight (g)} \times \text{dilution factor}}$$

$$\text{DeoxyMb (\%)} = (0.369 R1 + 1.140 R2 - 0.941 R3 + 0.015) \times 100$$

$$\text{MbO}_2 (\%) = (0.882 R1 - 1.267 R2 + 0.809 R3 - 0.361) \times 100$$

$$\text{MetMb (\%)} = (-2.541 R1 + 0.777 R2 + 0.800 R3 + 1.098) \times 100$$

2.3 Carbonyl content

Protein carbonylation level was assessed in triplicates following the procedure proposed by Levine et al. (1990) and recently modified by Soglia, Petracci, and Ertbjerg (2016). Briefly, 1.0 g of sample was homogenized (9,500 rpm for 30 s) in ice by an Ultra-Turrax T25 (IKA – WERKE, Labor Technik). Five aliquots (100 μL/each) of the homogenate were added with 1 mL 10% TCA. Following centrifugation (5,000 × g for 5 min) (Beckman Coulter microfuge 16), the supernatant was discarded and 400 μL of 5% sodium dodecyl sulfate (SDS) solution was added to the resulting pellet. The samples were subsequently heated at 100 °C for 10 min and ultrasonicated (Argo-lab DU-45, Modena, Italy) at 40 °C for 30 min. Then, the samples (three replicates) were incubated (30 min) with 800 μL of 0.3% (w/v) 2,4-Dinitrophenylhydrazine (DNPH) in 3 M HCl, while the same volume of 3 M HCl was added to the blank (two replicates). Following incubation, 400 μL of 40% TCA was added and the supernatant removed by centrifugation (5,000 × g for 5 min). The pellet was washed three times with 1 mL of ethanol-ethyl acetate (1:1, v:v) solution by centrifugation at 10,000 × g for 5 min. After the final wash, the pellet was dried, resuspended in 1.5 mL of 6 M guanidine hydrochloride in 20 mM NaH₂PO₄ (pH 6.5), and the samples were incubated overnight at 4 °C. Then, the absorbance at 370 and 280 nm was measured with a UV-VIS 1800 spectrophotometer (Shimadzu Corporation, Japan) at 25 °C in order to quantify carbonyls and the protein concentration within the samples, respectively. In detail, carbonyl content, expressed as nmol/mg of protein, was calculated according to the

equation:

$$\text{carbonyl content (nmol/mg of protein)} = \frac{\text{Abs 370} - \text{Abs 370 (blank)}}{22,000 \times [\text{Abs 280} - (\text{Abs 370} - \text{Abs 370 (blank)}) \times 0,43]} \times 10^6$$

2.4 Free thiol groups

Free thiol groups were quantified following their reaction with 5,5'-Dithiobis(2-nitrobenzoic acid) (DTNB) as proposed by Ellman (1959) with some modifications (Lund, Lametsch, Hviid, Jensen, & Skibsted, 2007). Briefly, 1.0 g of samples was homogenized (13,500 rpm for 30 s) with an Ultra-Turrax T25 (IKA – WERKE, Labor Technik) in 25 mL 5% (w/v) SDS in 0.1 M Tris-HCl (pH 8.0). After heating in a water bath at 80 °C for 30 min, the homogenate was filtered through filter paper (Whatman 1, GE Healthcare) and the protein concentration was assessed by reading the absorbance at 280 nm and calculated through a standard curve prepared from 0 to 2 mg/mL bovine serum albumin. Then, free thiol groups were quantified by mixing 0.5 mL filtrate, 2 mL of 0.1 M Tris-HCl (pH 8.0), and 0.5 mL 10 mM DTNB in 0.1 M Tris-HCl (pH 8.0) and reading the absorbance of the samples at 412 nm after 30 min incubation in the dark. Indeed, forming a disulfide bond with free thiol groups, DTNB releases a thiolate ion with a maximal absorbance at 412 nm (Estévez, Morcuende, & Ventanas, 2009). Thiol groups content was calculated by using a standard curve prepared from L-cysteine and expressed as nmol/mg protein.

2.5 Statistical analysis

Data were analyzed according to a 2 × 6 factorial design using the GLM procedure of Statistica (StatSoft Italy srl, Vigonza, Italy). In detail, within the same experiment (turkey and rabbit meat), a factorial ANOVA was performed to investigate the main effects of the exposure to oxidative conditions (Ox, fresh vs. oxidized) and MDA level (MDA, 0, 0.25, 0.5, 1.0, 2.5, and 5.0 mM) as well as their interaction term “Ox × MDA” on protein carbonylation, free thiol group, total heme pigments, and on the relative concentration of the different myoglobin derivatives. Mean values were subsequently separated through the parametric Tukey-HSD test. In addition, Student's *t*-test was carried out with the aim to compare protein oxidation and heme pigments in meat belonging from turkey thigh and rabbit hind leg meat without any MDA added and evaluate whether an effect associated with the meat type can be observed. All statistical differences were considered significant at a level of $P \leq 0.05$.

3. RESULTS AND DISCUSSION

The results concerning the impact of the exposure to strong oxidative conditions and the concomitant addition of increasing MDA levels on protein oxidation and heme pigments in turkey thigh meat are reported in Table 1.

As expected, the addition of a strong oxidant (100 mM NaClO) to the system resulted in remarkable oxidative modifications affecting the proteins' polypeptide chains. Indeed, if compared to its fresh counterpart, a significant increase in protein carbonylation (3.49 vs. 7.58 nmol/mg pf protein; $P < 0.001$) along with a concomitant decrease in free thiol groups was observed in oxidized meat. In detail, free SH groups became readily undetectable after incubating turkey meat in the presence of NaClO, meaning that the oxidative environment was strong enough to induce oxidation of the cysteine residues. Indeed, following a one- or two-electron

oxidation reaction, thiol groups in cysteine can result in the formation of thiyl radicals or sulfenic acids. Because of their unstable nature, these species can further react with other thiol/thiolate to generate disulfide or result in the formation of a disulfide bond, respectively (Zhang, Xiao, & Ahn, 2013). Thus, the loss of free thiol groups can be considered a reliable marker for protein oxidation (Winther & Thorpe, 2014) and suggests the occurrence of oxidative modifications in turkey meat involving the formation of disulfide bonds. Overall, this reduction in free thiol groups is in agreement with the findings of previous studies carried out on oxidized myofibrils (Frederiksen, Lund, Andersen, & Skibsted, 2008; Morzel, Gatellier, Sayd, Renner, & Laville, 2006).

As for heme pigments, exposing turkey thigh meat to a strong oxidant significantly affected the content as well as the relative concentration of the oxidized and the oxygenated myoglobin derivatives, whereas the reduced one (DeoxyMb) was not affected ($P > 0.05$). In detail, the content of total heme pigments was found to be decreased by 63% by comparing fresh meat with its oxidized counterpart (2.04 vs. 0.77 mg Mb/kg of meat; $P < 0.001$). This result might be ascribed to a possible damage to the globin molecule that, affecting the stability of the heme group itself, ultimately results in the release of iron (Estévez & Cava, 2004). The same authors hypothesized a possible relationship between the release of iron from the heme group and the occurrence of protein oxidation, which seems to be in agreement with the results of our study. Concurrently, a 67% reduction ($P < 0.001$) in the relative concentration of MbO₂ matching with a two-fold increase ($P < 0.001$) in the proportion of MetMb was found in oxidized in comparison with fresh meat. Undeniably, the exposure to pro-oxidant conditions can lead to the destabilization of the myoglobin molecule and to a subsequent meat discoloration (Suman & Joseph, 2013). Indeed, the oxidation of DeoxyMb leads to the formation of MetMb and superoxide that subsequently dismutates to generate H₂O₂ and O₂ (Chan, Faustman, Yin, & Decker, 1997). As previously hypothesized by Thiansilakul, Benjakul, and Richards (2011), the H₂O₂ thus formed might have induced conformational changes in the globin structure and, affecting myoglobin redox state, ultimately resulted in the release of iron from the porphyrin ring. Intriguingly, it is noteworthy to mention the absence of significant differences in the relative concentration of the reduced pigment form (DeoxyMb). It seems reasonable to hypothesize that this result might be ascribed to the structure of the DeoxyMb molecule itself, in general, and to the absence of ligands at the sixth coordinate of the heme iron in this myoglobin derivative, in particular. Indeed, this may have helped in maintaining the hydrophobic pocket-like structure of the globin thus protecting its heme group from the oxidative environment developed in our experiment.

With regard to the effect of MDA, the achievement of a concentration 2.5 mM (or greater) resulted in a significantly ($P < 0.001$) higher carbonyl content (1.4-fold increase) if compared to the control group (0 mM MDA). This result corroborates the hypothesis that MDA, as a secondary product of lipid oxidation, may trigger an increase in carbonyl groups (Burcham & Kuhan, 1996; Estévez et al., 2019; Wang et al., 2019), which represents one of the most common consequences of protein oxidation (Estévez, 2011; Lund et al., 2011). However, it is interesting to highlight that, aside from the in-site formation of carbonyl groups following the attack of ROS to the side chain of susceptible amino acids (i.e., proline, lysine, and arginine), protein carbonylation can also result from the addition of secondary preformed carbonyls including those generated from lipid oxidation (among the others, also MDA) (Estévez et al., 2019). Thus, it might be argued whether the increased

Table 1—Effect of the exposure to oxidative conditions (100 mM NaClO) and the concurrent incubation with different MDA levels (0, 0.25, 0.5, 1.0, 2.5, and 5.0 mM) on protein oxidation, heme pigments content, and relative concentration of myoglobin derivatives in minced turkey thigh meat.

Treatment (Ox) N = 30/group MDA level (MDA) N = 10/group	Fresh Oxidized 0 mM 0.25 mM 0.50 mM 1.00 mM 2.50 mM 5.00 mM	Free thiol groups (nmol/mg of protein)			Heme pigment content (mg Mb/kg of meat)	DeoxyMb (%)	MbO ₂ (%)	MetMb (%)
		Carbonyls (nmol/mg of protein)	Free thiol groups (nmol/mg of protein)	Heme pigment content (mg Mb/kg of meat)				
Ox × MDA N = 5/group	Fresh	3.49	141.9	2.04	23.9	41.1	34.9	
	Oxidized	7.58	b.d.	0.77	24.2	13.5	62.3	
	0 mM	4.53 z	75.5	1.42	24.5	28.3	47.2	
	0.25 mM	5.32 y	68.8	1.41	24.4	24.0	51.6	
	0.50 mM	5.20 yz	75.4	1.48	23.4	27.2	49.3	
	1.00 mM	5.57 y	74.1	1.41	25.1	27.2	47.6	
	2.50 mM	6.31 x	71.9	1.32	22.8	28.0	49.2	
	5.00 mM	6.30 x	60.2	1.37	24.0	29.1	46.9	
	Fresh	2.17 d	150.9	2.13	25.2	46.4	28.4	
	0.25 mM	3.17 d	137.6	2.14	24.5	37.4	38.1	
	0.50 mM	3.20 d	150.9	2.13	23.0	39.2	37.8	
	1.00 mM	3.26 d	148.2	2.10	24.4	38.5	37.2	
	2.50 mM	4.54 c	143.8	1.83	24.2	43.1	32.7	
	5.00 mM	4.61 c	120.4	1.92	22.2	42.0	35.8	
Oxidized	6.89 b	b.d.	0.71	23.9	10.2	65.9		
0.25 mM	7.48 ab	b.d.	0.69	24.3	10.6	65.1		
0.50 mM	7.19 ab	b.d.	0.83	23.9	15.2	60.9		
1.00 mM	7.89 ab	b.d.	0.73	25.9	16.0	58.1		
2.50 mM	8.08 a	b.d.	0.81	21.5	12.8	65.7		
5.00 mM	7.98 ab	b.d.	0.83	25.8	16.2	58.0		
Sem		0.29	9.4	0.10	0.6	2.2	2.2	
P-value		***	***	***	ns	***	***	
Ox		***	Ns	Ns	ns	ns	ns	
MDA		***	Ns	Ns	ns	ns	ns	
Ox × MDA		**	Ns	Ns	ns	ns	ns	

b.d., below detection limit; ns, not significant.
 ** = $P < 0.01$; *** = $P < 0.001$.
 x to z = Mean values followed by different letters significantly differ among the groups treated with increasing MDA level.
 a to d = Mean values followed by different letters significantly differ among the groups exposed to oxidative conditions and treated with increasing MDA level.

carbonylation level found in association with higher MDA concentration might be attributed to the formation of adducts with this compound rather than to the occurrence of protein oxidation. Having this in mind, in order to get more information concerning the effect of an increasing MDA content on the oxidative damage as well as on the modifications affecting the proteins' structure, free thiol groups were also quantified. Interestingly, a tendency ($P = 0.07$) was observed for free thiol groups whose content was found to decrease as the MDA level added to the system increased. This finding corroborates the hypothesis of the existence of a possible interaction between the aldehydes generated from lipid oxidation and the oxidative modifications affecting proteins' polypeptide chains. In detail, as observed in previous studies, MDA might be involved in the formation of protein aggregates through the generation of cross-linking (Buttkus, 1967; Xiong, Park, & Ooizumi, 2009). Therefore, it might be speculated that, incubating the samples with increasing MDA levels may have resulted in a conversion of the sulfhydryl groups into disulfide and in the formation of disulfide bonds in the myofibrils thus accounting for the reduction in free thiol groups content. Analogously, a sharp decrease in the sulfhydryl content matching with an increase in the amount of disulfide bonds was previously reported by Liu, Xiong, and Butterfield (2000) in oxidized myofibrillar proteins. At last, the presence of increasing MDA level did not exert any relevant effect on total heme pigments as well as on the relative concentration of the different myoglobin derivatives in turkey meat.

The significant interaction term "Ox \times MDA" observed for carbonyls in turkey meat might be ascribed to the different carbonylation levels observed as a consequence of the increasing MDA concentration in the system and the concurrent exposure to oxidative conditions. In detail, a relevant increase in carbonyl content was found as MDA concentration attained 2.5 mM in both fresh and oxidized meat. However, in the case of fresh meat, a further increase in MDA level (to 5 mM) led to the same result, whereas an opposite trend was found in oxidized meat in which the highest MDA concentration added to the samples did not result in a significant increase in carbonyls ($P > 0.05$).

Overall, the aforementioned results are in agreement with those obtained for rabbit hind leg meat subjected to the same experimental design, which are reported in Table 2. Exposing rabbit meat to oxidative conditions significantly affected all the parameters considered within this study with the only exception being the relative concentration of DeoxyMb and MbO₂ that did not differ between fresh meat and its oxidized counterpart ($P > 0.05$). Indeed, a three-fold increase ($P < 0.001$) in carbonyl content coupled with a one-half reduction (-53% ; $P < 0.001$) in free thiol groups was found in oxidized rabbit meat. These findings suggest the occurrence of intense oxidative modifications to the proteins' polypeptide chains. In detail, although it is not clear whether the hydrazones (DNPH-derivatized carbonyls) measured are the result of an in-site formation of carbonyls mediated by ROS or are rather a consequence of adducts with MDA or preformed carbonyl compounds, the remarkable increase in protein carbonylation level associated with a sharp reduction in free thiol groups is considered a good marker of oxidation (Lund et al., 2011). It is noteworthy that the exposure of turkey thigh and rabbit hind leg meat to the same environment affected the protein fraction with a different extent and might have resulted in different oxidative changes. Indeed, if compared to turkey thigh exposed to oxidative conditions, a more intense protein carbonylation seems to take place in rabbit meat (two-fold vs. three-fold increase, in absolute terms) in which, simultaneously, the development of disulfide bonds seems to oc-

cur with a lower extent. Indeed, in turkey thigh, free thiol groups became readily undetectable after the exposure to oxidative conditions, whereas a 50% reduction in their content was found in rabbit hind legs. It might thus be argued whether these distinctive oxidative changes might be related to the compositional traits (i.e., cysteine content, fat content, and fatty acids profile) of the meat considered in the present experiment that might have affected its susceptibility toward oxidation and promoted one oxidative change rather than the other. In addition, a significant reduction in total heme pigments (0.21 vs. 0.68 mg Mb/kg of meat; $P < 0.001$) associated with an increased proportion of MetMb (+9%) was found in rabbit leg meat exposed to oxidative conditions in comparison with the fresh one in agreement with the results observed for turkey thigh meat (Table 1). However, a remarkable difference in the extent of MetMb formation was found, with the increase in the relative concentration of the oxidized pigment form being of about 10% and 200% in rabbit and turkey meat, respectively. This finding might suggest a species-specific susceptibility of the myoglobin molecules to oxidation. Indeed, it might be speculated that, because of a higher susceptibility to oxidation, the myoglobin molecules in rabbit meat might have been largely oxidized (i.e., MetMb formed) already during the 24 hr incubation period at 37 °C. The same mechanism might have likely occurred for the lipid fraction, which is relatively high in rabbit meat. Within this context, myoglobin oxidation to MetMb might have been further accelerated by the reactive products of lipid oxidation, which are polar and can readily diffuse within the sarcoplasm (Esterbauer, Schaur, & Zollner, 1991; Faustman, Liebler, McClure, & Sun, 1999; Suman & Joseph, 2013).

The achievement of increasing MDA levels within the system did not exert any relevant effect neither on protein oxidation nor on heme pigments and, similarly, no significant interaction terms "Ox \times MDA" were observed. However, it has to be mentioned that a 25% reduction in free thiol groups was found by comparing the results obtained in the control group (0 mM MDA) with those observed with the addition of 5 mM MDA. Although not statistically significant, this result corroborates the hypothesis of an MDA-mediated oxidation process involving the development of disulfide bonds.

The results concerning protein oxidation, heme pigments content, as well as the relative concentration of the different myoglobin derivatives found in fresh (without any MDA added, 0 mM) turkey and rabbit meat, are reported in Table 3. Although no significant differences were found in protein carbonylation level ($P > 0.05$), a significantly higher free thiol groups content was found in rabbit meat in comparison with that observed in turkey matrix (225.0 vs. 150.9 nmol/mg of protein; $P < 0.001$). Considering that the control groups were also incubated at 37 °C for 24 hr, these findings corroborate the hypothesis of a species-related development of oxidation products. Indeed, it is likely that part of the free thiol groups in turkey meat has been readily oxidized during the incubation time and the remaining ones were quickly involved in the development of disulfide bonds after the exposure to the oxidative environment (Table 1). Aside from the amino acid composition of the polypeptide chains which vary among species, these results might be ascribed to the exposure to pro-oxidant compounds, such as hemoproteins (i.e., myoglobin) (Van Dyck, 2010), which are considerably high in turkey meat (Estévez, 2015; Pereira & Vicente, 2013). In agreement with that, a different susceptibility to oxidation was found in a previous study carried out on meat belonging from different species in which myoglobin was found to be one of the primary determinants for the differ-

Table 2—Effect of the exposure to oxidative conditions (100 mM NaClO) and the concurrent incubation with different MDA levels (0, 0.25, 0.5, 1.0, 2.5, and 5.0 mM) on protein oxidation, heme pigments content, and relative concentration of myoglobin derivatives in minced rabbit hind legs meat.

Treatment (Ox) N = 30/group	Fresh	Carbonyls (nmol/mg of protein)	Free thiol groups (nmol/mg of protein)	Heme pigment content (mg Mb/kg of meat)	DeoxyMb (%)	MbO ₂ (%)	MetMb (%)
		2.27	197.5	0.68	20.3	6.5	73.5
	Oxidized	6.58	93.3	0.21	17.7	5.3	79.9
	0 mM	4.19	159.2	0.45	20.4	3.7	75.9
	0.25 mM	4.65	135.6	0.42	18.8	9.7	79.7
	0.50 mM	4.30	144.5	0.42	17.4	3.8	78.7
	1.00 mM	4.09	158.7	0.48	18.4	5.1	76.6
	2.50 mM	4.15	154.8	0.47	17.4	6.3	76.5
	5.00 mM	4.91	124.8	0.43	21.8	6.9	72.9
Ox × MDA N = 5/group	Fresh	1.73	225.0	0.73	25.2	2.3	72.5
	0.25 mM	2.10	203.3	0.64	13.4	10.2	76.4
	0.50 mM	2.61	172.5	0.62	20.0	2.6	77.4
	1.00 mM	2.14	213.2	0.72	19.4	5.6	75.0
	2.50 mM	2.37	203.2	0.69	20.1	9.5	70.4
	5.00 mM	2.68	167.9	0.66	23.9	9.0	69.5
	Oxidized	6.64	93.4	0.17	15.6	5.2	79.2
	0.25 mM	7.20	67.8	0.20	24.2	9.2	82.9
	0.50 mM	5.99	109.5	0.23	14.9	5.0	80.1
	1.00 mM	6.54	104.3	0.24	17.4	4.6	78.1
	2.50 mM	5.93	106.4	0.24	14.6	3.2	82.6
	5.00 mM	7.14	81.7	0.21	19.6	4.7	76.3
Sem		0.31	8.5	0.03	0.7	0.7	0.9
P-value		***	***	***	Ns	ns	***
Ox		ns	Ns	ns	Ns	ns	ns
MDA		ns	Ns	ns	Ns	ns	ns
Ox × MDA		ns	Ns	ns	Ns	ns	ns

*** = P < 0.001.
ns, not significant.

Table 3—Comparison of protein oxidation (carbonyls and free thiol groups), heme pigments content, and relative concentration of myoglobin derivatives in fresh (0 mM MDA) minced turkey thigh and rabbit hind legs meats (N = 5 samples/group).

	Carbonyls (nmol/mg of protein)	Free thiol groups (nmol/mg of protein)	Heme pigment content (mg Mb/kg of meat)	DeoxyMb (%)	MbO ₂ (%)	MetMb (%)
Turkey thigh	2.17	150.9	2.13	25.2	46.4	28.4
Rabbit hind leg	1.73	225.0	0.73	25.2	2.3	72.5
Sem	0.22	7.2	0.07	0.5	1.5	1.8
P-value	Ns	***	***	ns	***	***

*** $P < 0.001$.
ns, not significant.

ences in oxidative stability of meat (Min, Nam, Cordray, & Ahn, 2008). With regard to heme pigments, if compared to rabbit hind leg meat, turkey thigh matrix exhibited a significantly higher total heme and MbO₂ content along with a significantly lower proportion of MetMb. On the other hand, no significant differences were found in the relative concentration of DeoxyMb. The significantly lower heme pigment content observed in rabbit hind leg in comparison with turkey thigh meat might be a consequence of a partial release of iron from the heme group (discoloration) taking place during the 24 hr incubation time at 37 °C. In addition, as lipid oxidation proceed fast, the higher lipid content in rabbit hind leg meat may have resulted in a more intense development of aldehydes that, forming adducts with the histidine residue in myoglobin, can strongly affect color stability (Suman & Joseph, 2013). On the other hand, the remarkable differences in MbO₂ and MetMb might be explained by considering the different susceptibility of avian and mammalian myoglobin to oxidation hypothesized by Yin et al. (2011). Though myoglobin functions are among the highly conserved traits across poultry and livestock species, its primary structure, ultimately affecting the interactions with other molecules, profoundly differs (Suman & Joseph, 2013). Indeed, the presence of negatively charged residues was previously found to affect buffalo myoglobin stability whose meat resulted more prone to discoloration (Dosi et al., 2006). In addition, also the number of histidine residues, favorable target for the adduction of aldehydes generated by lipid oxidation, was also hypothesized to contribute to myoglobin susceptibility to oxidation (Suman & Joseph, 2013). Thus, within this context, the differences in MbO₂ and MetMb observed between turkey thigh and rabbit hind leg meat are likely ascribable to the unique primary structure of their myoglobin molecules.

4. CONCLUSION

The findings of the present study suggest the occurrence of species-specific oxidative changes affecting proteins polypeptide chains in different meat types. These may partly be attributed to the different susceptibility to oxidation of the myoglobin molecules belonging to different species that should thus be carefully considered and requires the implementation of processing strategies aiming at minimizing the occurrence of meat discoloration or, at least, reduce its impact on consumers' acceptability. In addition, as the occurrence of oxidative reactions strongly depends on the meat-cut taken into account but is also influenced by each processing step and the storage conditions as well, these aspects need to be surely considered when developing further processed products with the inclusion of turkey and rabbit meats.

AUTHOR CONTRIBUTIONS

Francesca Soglia and Massimiliano Petracchi were responsible for study conception and design. Francesca Soglia and Giulia Baldi performed the analyses and interpreted the results. All the authors contributed to draft the manuscript and critically revised it.

CONFLICTS OF INTEREST

No conflicts of interest to declare.

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