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## Freezing of meat and aquatic food : Underlying mechanisms and implications on protein oxidation

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2021-11

Bao , Y , Ertbjerg , P , Estévez , M , Yuan , L & Gao , R 2021 , ' Freezing of meat and aquatic food : Underlying mechanisms and implications on protein oxidation ' , Comprehensive Reviews in Food Science and Food Safety , vol. 20 , no. 6 , ARTN 1-22 , pp. 5548-5569 . https://doi.org/10.1111/1541-4337.12841

http://hdl.handle.net/10138/348351 https://doi.org/10.1111/1541-4337.12841

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### Freezing of meat and aquatic food: Underlying mechanisms and implications on protein oxidation

Journal:	Comprehensive Reviews in Food Science and Food Safety
Manuscript ID	Draft
Manuscript Type:	Comprehensive Review
Date Submitted by the Author:	n/a
Complete List of Authors:	Bao, Yulong; Jiangsu University, School of Food and Biological Engineering Ertbjerg, Per; University of Helsinki, Food and Nutrition Estévez, Mario; Faculty of Veterinary (University of Extremadura), Food Technology Yuan, Li; Jiangsu University, School of food science and biological engineering Gao, Ruichang; Jiangsu University, School of food science and biological engineering
Keywords:	





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March 27th 2021

Dear Editor

Please find the attached manuscript entitled: **Freezing of meat and aquatic food: Underlying mechanisms and implications on protein oxidation**. We would like to have this manuscript considered for publication in Comprehensive Reviews in Food Science and Food Safety.

Over the recent decades, protein oxidation in muscle foods has gained increasing research interests. Protein oxidation occurs during freezing/thawing and frozen storage of muscle foods, leading to irreversible physicochemical changes and impaired quality traits. The submitted review paper focus on key physicochemical factors in freezing/thawing and frozen storage of muscle foods, such as formation of ice crystals, freeze concentrating and macromolecular crowding effect, instability of proteins at the ice-water interface, freezer burn, lipid oxidation, etc. Possible relationships between these physicochemical factors and protein oxidation are thoroughly discussed. This review will shed light on the complicated mechanism of protein oxidation in frozen muscle foods.

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We upload in separate files: Manuscript Tables 1 to 2 Figures 1 to 5

Yours sincerely

Ruichang Gao, Professor School of Food and Biological Engineering Jiangsu University

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1	Freezing of meat and	aquatic food:	Underlying	mechanisms a	nd implications on

- 2 protein oxidation
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#### 22 Abstract

23 Over the recent decades, protein oxidation in muscle foods has gained increasing 24 research interests as it is known that protein oxidation can affect eating quality and nutritional value of meat and aquatic products. Protein oxidation occurs during 25 26 freezing/thawing and frozen storage of muscle foods, leading to irreversible physico-27 chemical changes and impaired quality traits. Controlling oxidative damage to muscle 28 foods during such technological processes requires a deeper understanding of the 29 mechanisms of freezing-induced protein oxidation. This review focus on key 30 physicochemical factors in freezing/thawing and frozen storage of muscle foods, such 31 as formation of ice crystals, freeze concentrating and macromolecular crowding effect, 32 instability of proteins at the ice-water interface, freezer burn, lipid oxidation, etc. 33 Possible relationships between these physicochemical factors and protein oxidation are thoroughly discussed. In addition, the occurrence of protein oxidation, the impact on 34 eating quality and nutrition, and controlling methods are also briefly reviewed. This 35 36 review will shed light on the complicated mechanism of protein oxidation in frozen muscle foods. 37

38 Keywords: Protein oxidation, protein denaturation, ice-water interface, freezing
39 potential, freezer burn, macromolecular crowding

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#### 43 1. Introduction

44 Meat and aquatic products are, in a global perspective, among the most important 45 foods consumed. Muscle foods provide high-value animal proteins and a wide range of essential micronutrients to a large majority of people in the world. Muscle-based foods 46 47 are nutritious and highly appreciated by their sensory properties. However, fresh meat 48 and aquatic foods are perishable and freezing is widely used for preservation purposes. 49 The effects of freezing and thawing on muscle foods have been extensively studied and 50 many excellent reviews have been available on this topic over the years. Mazur (1970) 51 thoroughly reviewed the major effects of freezing on biological systems, and 52 emphasized that the properties of cell membrane were closely related to the responses 53 of cells to freezing. Many of the theories/hypothesis discussed in that paper benefited 54 the research area of meat freezing. In due course, Mackie (1993) reviewed the effects of freezing on muscle proteins and focused on similarities and differences between 55 proteins from different animals. Possible mechanisms of protein denaturation in muscle 56 57 during freezing and frozen storage were introduced. More recently, Leygonie et al. 58 (2012) reviewed the effects of freezing and thawing on major quality parameters of 59 meat, such as moisture loss, color, pH, protein denaturation, lipid oxidation, tenderness 60 and microbial spoilage, while Nakazawa & Okazaki (2020) introduced various factors 61 that influence quality of frozen seafoods. In recent years, advanced tools such as proteomics (Men et al., 2020; Shi et al., 2018) and molecular dynamic simulations 62 63 (Zhang et al., 2018) have been used to provide further insights into the underlying 64 biochemical processes occurred during frozen storage of muscle foods.

65 It is well known that textural properties of frozen foods will generally be compromised after thawing, and ice crystals play a key role. The biochemical 66 67 components are much less affected by frozen storage though no foods could remain unmodified during long-term storage. Native proteins are just marginally stable as the 68 69 thermodynamic stability of folded proteins is often 1 to 4 kcal/mol higher than that of 70 the corresponding unfolded ones (Sanfelice & Temussi, 2016). Freezing leads to a 71 series of physicochemical changes and therefore pronounced denaturation of muscle proteins are often observed after frozen storage. In addition to protein denaturation, 72 proteins can be oxidized during frozen storage as well. Increased protein oxidation with 73 74 frozen storage has been ascribed to catalytic iron release after membranes being 75 disrupted, and freeze-concentrating of pro-oxidants and protein molecules (Estévez et 76 al., 2011; Leygonie et al. 2012; Soladoye et al., 2015). Unlike protein denaturation, protein oxidation has been less studied in frozen meat, but it is receiving increased 77 research interest. So far, the mechanisms of freezing-induced protein oxidation are not 78 well-understood. Therefore, in this paper, we focus on the physiochemical changes 79 80 occurring in frozen muscle foods and discuss their implications on protein oxidation.

81

#### 82 2. Major physicochemical changes in frozen muscle foods

#### 83 2.1 Physicochemical environment in muscle foods prior to freezing

84 Muscle foods are derived from animal flesh. Fundamental aspects such as85 chemical composition and structural organization of meat is well known. In brief, lean

86	meat is mainly composed of water and proteins and both pro-oxidant (myoglobin, lipids,
87	etc.) and antioxidant components (reducing enzymes, glutathione, etc.) naturally occur
88	in muscle tissue (Lawrie & Ledward, 2006). The final pH value of postmortem muscle
89	is of critical importance to reactions in meat and thus affects the final quality of muscle
90	foods. Ions can be bound to or tightly associated with proteins, or they can loosely
91	associate to proteins or exist as free ions (Hamm, 1986). Like many other cellular
92	tissues, muscle contains a very high concentration of proteins (~ 20%), along with other
93	components, such as nucleic acids, lipids and sugars. They cumulatively occupy a large
94	fraction of the total cell volume since macromolecules generally cannot interpenetrate
95	and this is called macromolecular crowding. This crowding effect could affect many
96	reactions, yet many biochemical studies of macromolecular properties are carried out
97	in dilute solutions in which crowding do not occur (Minton, 2006). Much of the cellular
98	water exists between a variety of fibrous and membranous structures. Between 10%
99	and 100% of the fluid lies within one macromolecular diameter of the surface of these
100	fibrous or membranous structures. As these structures do not move, this leads to a
101	situation named as macromolecular confinement. Both crowding and confinement are
102	expected to increase the stability of folded proteins. When reactants bear net charges
103	opposite to that of the surface of nearby fibers or membranes, adsorption may occur
104	and thus change the reaction kinetics. The abovementioned macromolecular crowding,
105	confinement, and adsorption can greatly influence the equilibria and reaction rates.
106	

#### 107 **2.2 Decrease in temperature**

108	Temperature is one of the most important factors in determining the rates of
109	chemical reactions (Boekel, 2008). Frozen storage of muscle foods usually takes place
110	at temperatures ranging from -10 °C to -30 °C, while storage of some precious muscle
111	foods such as tuna often requires -60 °C or lower. Frozen temperatures from -20 to -30 °C
112	are generally satisfactory in maintaining quality of muscle foods. The marginal gain in
113	quality of storage at -60 °C may be small, but can be critical for demanding consumers
114	(Ji et al., 2021). During freezing (and thawing), if heat removal (supply) is slow, muscle
115	foods will stay at temperatures around -1 °C for quite long time. During frozen storage,
116	the temperature fluctuations in the freezer will lead to fluctuations in the muscle foods
117	as well, but to a lesser extent. A storage temperature well below the freezing point of
118	muscle is expected to slow down chemical reactions, as the migration of molecules is
119	largely suppressed in the solid phase. However, some reactions may be greatly
120	enhanced by freezing or a lower storage temperature. For example, freezing was found
121	to accelerate the reaction between nitrite and dissolved oxygen by a factor of $10^5$
122	(Takenaka et al., 1992).

123

#### 124 **2.3 Formation of ice crystals**

Upon freezing, the extracellular solution first crystallizes as it has a higher freezing point due to a lower content of solutes. The extracellular water in muscle has a freezing point around -1.2 °C, while the intracellular water starts to freeze around -1.6 °C. Once ice crystals form in the extracellular space, solutes get concentrated and therefore there will be osmotic pressure across the sarcolemma (i.e. cell membrane). To achieve a 130 balanced state again, cells can response in two ways: cellular dehydration or 131 intracellular ice formation (Fig. 1a). These subsequent events during cooling depend 132 mainly on the cooling rate and membrane permeability to water (Mazur, 1970). The cooling rate can be assessed by the characteristic freezing time  $(t_c)$ , and  $t_c$  is defined as 133 time needed for temperature to decrease from -1°C (beginning of freezing) to -7°C 134 135 (freezing of 80% of the water present). Bevilacqua & Zaritzky (1980) found that 136 intracellular ice formation in beef only happened when t<sub>c</sub> was less than 5 min when freezing is transverse to the fibers. When freezing is longitudinal to the fibers, 137 intracellular ice crystals can advance in the direction of heat flow. Therefore, 138 139 intracellular ice can be found with a t<sub>c</sub> up to 20 min in this situation. The longer the t<sub>c</sub>, the larger the size of the crystals formed, and larger extracellular ice crystals lead to 140 141 increased distortion of the muscle microstructure. As for membrane permeability, not enough information is available on how the low temperature and high electrolyte 142 143 concentrations in frozen condition will affect the permeability. It has been suggested that a high concentration of electrolytes makes the sarcolemma leaky. According to 144 145 Mazur (1970), the surface membranes of cells prevent extracellular ice from entering cells at temperatures about -10 °C to -15 °C. One possible explanation is based on the 146 147 assumption that the membranes contain water-filled pores. At temperatures of -10 °C to -15 °C, ice crystals are so large that they cannot pass through pores in the membranes. 148 149 Cooling may result in a transition of membranes from the liquid crystalline to gel phase 150 (Crowe et al., 1989), which may lead to leak of cell contents.

151 During storage, ice crystals grow larger at the expense of smaller ones (Fig. 1b), 152 and increased ice crystals may lead to increased physical damage of the muscle structure. The driving force for the process of recrystallization is that small crystals 153 have higher vapor pressure and water diffuse down the pressure gradient. Temperature 154 155 fluctuation tend to accelerate the recrystallization process and hence need to be 156 controlled during frozen storage. Recently, Nakazawa & Okazaki (2020) summarized 157 factors influencing the quality of frozen seafood and they argued that the formation of ice itself may not be detrimental to muscle structure. In the case of low level of protein 158 denaturation, thawing-induced free water is absorbed and the muscle cell can be largely 159 160 recovered. If the denaturation level is high, then the released water is not absorbed and the muscle cell is not recovered (Fig. 1c). If the thawing rate is high, the melted water 161 162 does not have enough time to be re-absorbed by the muscle fibers and part of it is eliminated as exudate. 163

As volume change occurs along with the formation of ice, muscle structure may 164 165 be distorted by ice crystals. It is speculated that the rigor bond and myosin filament are 166 among the sites of distortion (Fig. 2). It is known that myosin is prone to denaturation 167 during frozen storage. And structure distortion of muscle by changing the sarcomere 168 length was found to influence protein denaturation during heating (Findlay & Stanley, 169 1984). Therefore, freezing-induced distortion of muscle structure may affect protein 170 stability. Mechanisms of ice formation and controlling methods of ice crystallization has been studied extensively. Regarding the freezing of food, modelling approaches of 171 172 water crystallization, evaluation methods for the process and different novel methods

for improving water crystallization were presented in an excellent review by Kiani &
Sun (2011). Antifreeze proteins (AFPs) can protect living organisms from freezinginduced damage via tuning ice nucleation, shaping ice crystals, and inhibiting ice
growth and recrystallization (He et al., 2018).

177

178 2.4 Concentrating of solutes

179 During freezing, a large proportion of water turns into ice and thereby reduce the 180 amount of available liquid water. At the same time, the remaining solutes (including soluble proteins, pro-oxidants, and anti-oxidants) get concentrated in the unfrozen 181 fraction of water. Freezing has been found to accelerate oxidation of some small 182 183 molecules, such as nitrite, sulfite, and ascorbic acid (Hambly & Gross, 2009). However, the extent of acceleration was much higher than what would be expected from the 184 185 concentrating effect only. Takenaka et al. (1996) proposed a mechanism for the 186 accelerated reaction between nitrous acid and oxygen (Fig. 3). With the formation of ice crystals, dissolved oxygen and nitrite are excluded from ice and concentrated at the 187 interface of each ice crystal (Fig. 3A). As ice crystals continue to grow, concentrated 188 solutions are confined in the unfrozen solution surrounded by walls of ice crystals (Fig. 189 3B). When crystals grow further, solutes become extremely high in the solutions (Fig. 190 3C) and therefore the reactions can be accelerated. 191

192 **2.5 Ice-water interface** 

193 The presence of ice is critical for the instability of proteins subjected to freezing 194 treatment. For example, lactate dehydrogenase lost its activity in frozen state, however, 195 concentrated solutions at the same temperature and composition maintained its activity (Bhatnagar et al., 2008). Both water and ice are composed of a tetrahedral network of 196 197 hydrogen bonds, and the structural difference are minimal. The ice-water interface has 198 been suggested as a major source of protein denaturation during freezing (Arsiccio & 199 Pisano, 2020). Interestingly, the nature of a growing ice surface can be different from a melting ice surface (Takenaka et al., 1996). Possible mechanisms of the ice-water 200 interface-induced protein denaturation can be summarized as follows: 201

#### 202 2.5.1 Changes in microenvironment

Generally, solutes cannot be incorporated into the crystal lattice of ice and the 203 204 expelled solutes thus get concentrated in a quasi-liquid layer. A quasi-liquid layer represents an intermediate state between the solid ice crystal and bulk liquid. 205 Concentrated protons and ions would change the pH and ionic strength and likely to 206 207 affect protein stability (Arsiccio & Pisano, 2020) (Fig. 4a). Zhang & Ertbjerg (2019) suggested that pH of pork meat decreased due to freeze-induced concentrating of 208 protons. Difference in freezing rate may resulted in different degree of proton 209 210 entrapment by ice crystals, and therefore the difference in pH values of the remaining unfrozen liquid. It is well-known that pH is an important factor for protein stability. 211

212 2.5.2 Interface adsorption

213 As a biological material is frozen, formation of ice crystals leads to an increase in 214 the concentration of all solutes, including dissolved air, which is mainly composed of nitrogen and oxygen. In addition, the solubility of oxygen increases at lower 215 temperatures. The increase in oxygen concentration can be hypothesized to accelerate 216 oxidation during freezing as follows. Once the effect of freeze concentrating on air 217 218 exceeds its solubility, air bubbles are formed (Authelin et al., 2020). Air bubbles trapped between ice crystals and the hydrophobic air-water interface denature proteins 219 (Fig. 4b). In addition to the adsorption to the air-water interface, proteins can be 220 adsorbed directly to the ice surface and unfold (Fig. 4c), such as antifreeze proteins (He 221 222 et al., 2018)

#### 223 2.5.3 Mechanical stress

Volume expansion occurs during growth of ice crystals. Water-to-ice transformation leads to approximately a 9% increase in volume, and such expansion results in great shear stress and local pressure (Fig. 4d). It has been estimated that freeze-induced local pressure can reach 200 MPa (Authelin et al., 2020). It is known that high pressure treatment led to protein modifications in muscle foods, which mainly occur in the tertiary and quaternary structures (Cruz-Romero et al., 2004; Shi et al., 2020; Xue et al., 2018).

#### 231 2.5.4 Cold denaturation

Cold denaturation of proteins is linked to specific and strongly temperature-dependent interaction of protein nonpolar groups with water. There are many nonpolar

234 groups in proteins and the hydrophobic forces contribute to stabilization of the native protein structure. At sufficiently low temperature, the interaction between nonpolar 235 236 residues with water becomes less unfavorable (Fig. 4e). This may promote exposure of the hydrophobic core of the protein, and this phenomenon is called as cold unfolding 237 or cold denaturation. For example, Arsiccio et al. (2020) found that protein L was 238 239 destabilized in the presence of ice, and they explained that ice slowed down nearby 240 water molecules, and theses water molecules could form a larger number of hydrogen bonds with the protein, thus promoting hydration of nonpolar patches and ultimately 241 leading to exposure of the hydrophobic core. 242

243

#### 244 **2.6** pH change

As aforementioned, pH might change in the quasi-liquid layer under frozen 245 246 conditions (section 2.5.1). One hypothetical explanation is related to the freezing 247 potential. When a dilute electrolyte solution is frozen, an electric potential is generated between the ice and the solution. This freezing potential is also known as the Workman-248 249 Reynolds Effect (Workman & Reynolds, 1950). The sign and value of the freezing potential mainly depend on the kinds of solutes and the freezing rate, which lead to 250 251 selective segregation of ions with opposite sign into ice phase and the quasi-liquid layer (Levi & Milman, 1966). In extensively frozen solutions, the quasi-liquid layer may act 252 253 as a microscopic film wetting polycrystalline ice. Ions are marginally, but selectively, incorporated in ice as substitutional dopants during the freezing of electrolyte solutions. 254

255 For example, more Cl<sup>-</sup> is incorporated into the ice than Na<sup>+</sup> during freezing of aqueous 256 solution of NaCl, the charge of the ice will be negative; while in the case of NH<sub>4</sub>Cl, more NH<sub>4</sub><sup>+</sup> is incorporated than Cl<sup>-</sup> and therefore produces positive potential (Takenaka 257 et al., 1996). The potential is then neutralized by highly mobile  $H_3O^+$  or  $OH^-$ , therefore 258 pH in the quasi-liquid layer is varied (Fig. 5). For solutions of 2-(N-morpholino) 259 260 ethanesulfonic acid zwitterion, the acidity barely changes upon freezing. This is because the covalently linked cationic and anionic centers cannot be separated across 261 the ice/solution interface (Robinson et al., 2006). Freezing potential may affect 262 chemical reactions. In the reaction of nitrite with oxygen, addition of a stainless-steel 263 264 wire depressed the oxidation rate, suggesting the acceleration of the reaction by freezing is an electrostatic effect rather than an electrochemical reaction (Takenaka et al., 1996). 265 266 Those authors further investigated the electrostatic effect in the freezing process on the reaction. Different salts were added to the reaction mixture, and results showed that the 267 formation of nitrate is affected by the sign of the freezing potential, rather than by the 268 kind of ions. 269

Besides the above-mentioned mechanisms, pH change in frozen meat may be affected by other factors. At the generally used frozen storage temperature of -20 °C, around 90% water are frozen in beef and haddock muscle (Mackie, 1993). If protons remained in the liquid water during slow freezing, 90% frozen water would theoretically cause a 10-fold increase in proton concentration which equals to a drop of 1 pH unit. Although the buffering capacity of muscle would counteract the increase of protons, freezing could plausibly lead to a lower pH in vicinity of the protein surfaces

277	of frozen meat. Zhang & Ertbjerg (2019) suggested that fast freezing entraps more
278	protons in ice crystals, therefore less proton will remain in solution which leads to a
279	higher pH. At certain temperatures, solutes may exceed their solubilities and precipitate.
280	Precipitation of solutes may shift pH during freezing (Pikal-Cleland et al., 2000).
281	
282	2.7 Occurrence of 'Freezer burn'
283	'Freezer burn' describes the dehydration and associated changes in color, texture
284	and flavor that may occur on the surface of foods during frozen storage. The
285	dehydration is caused by sublimation of ice when the vapor pressure of ice at the surface
286	is greater that the vapor pressure of water in the air (Schmidt & Lee, 2010). A lower
287	storage temperature leads to lower vapor pressure, for example the vapor pressure of
288	ice is about 100 Pa at -20 °C, but reduces to around 4 Pa at -50 °C. Surface dehydration
289	leads to a local concentrating effect, where endogenous pro-oxidants and soluble
290	proteins get concentrated, and the oxidation process may be enhanced. Surface color of
291	frozen beef gradually change from red to brown due to the oxidation of myoglobin to
292	the brown colored metmyoglobin. Sublimation from the surface of frozen meat may
293	cause small air pockets to develop, forming a honeycomb structure. Modern self-
294	defrosting freezers may contribute to an increase in freezer burn. In a traditional manual
295	defrost freezer, accumulation of ice in the freezer compartment serves to increase the
296	vapor pressure of the air in the freezer, and thereby decreases the sublimation. However,

297 in self-defrost freezers, the removal of frost keeps the vapor pressure in the air at a low 298 level and therefore promotes sublimation from the food surface.

299 2.8 Lipid oxidation and hydrolysis

300 Lipids are chemically unstable food components and readily undergo oxidative reactions. Intramuscular fat, especially the lipids in fish tissues, are of high unsaturation 301 level. Therefore, lipid oxidation progresses to a noticeable extent during long-term 302 frozen storage. According to Leygonie et al. (2012), lipid oxidation mainly occurs at 303 the cellular membranes, therefore, lipid oxidation was observed both in lean and fatty 304 meats. In addition to oxidation, lipids may undergo hydrolysis during frozen storage 305 due to enzymatic hydrolysis, and free fatty acids will create more hydrophobic regions 306 in protein and promote denaturation (Mackie, 1993). 307

308

### revie 3. Protein oxidation in frozen muscle foods 309

3.1 Protein oxidation: chemistry and consequences 310

Meat, being a protein-rich food and containing pro-oxidants, such as lipids and 311 312 myoglobin, is susceptible to oxidation. One pathway of protein oxidation is via the 313 abstraction of a hydrogen atom in the protein by reactive oxygen species (ROS), which 314 is a collective term that includes oxygen-containing radicals  $(O_2^{\bullet}, OH, RO, RO_2^{\bullet}...)$ and also some non-radical derivatives of oxygen (H<sub>2</sub>O<sub>2</sub>, HClO, O<sub>3</sub>...) (Bao & Ertbjerg, 315 316 2019). In addition, reactive nitrogen species may also induce oxidative stress in muscle proteins (Skibsted, 2011). The detailed aspects of the chemistry of protein oxidation are 317

318	out of the scope of this paper, and they have been covered in a number of previous
319	review papers (Hawkins & Davies, 2001; Hellwig, 2019, 2020). Oxidative conditions
320	readily occur in post-mortem muscle and protein oxidation products may accumulate
321	in food during processing, storage and subsequent food intake (Estévez & Luna, 2017).
322	Oxidative modifications of muscle proteins include carbonylation, depletion of thiols
323	and tryptophan, formation of crosslinks, etc. (Estévez, 2011; Lund et al., 2011). Various
324	methods can be applied to the quantification of protein oxidation in biological systems.
325	Those methods are generally based on monitoring changes in parent amino acid
326	residues (Cys, Met, His, Lys etc.), detection of radical and non-radical intermediates,
327	or formation of products (e.g. formation of carbonyls, protein cross-linking).

Physicochemical properties of proteins are greatly affected by mobile electrolyte 328 329 ions and by ionized amino acid residues attached to the backbones of protein molecules. Oxidative modifications of those ionized amino acids are expected to affect the protein 330 functionality in muscle food. Histidine, for example, mainly exits as a positively 331 charged amino acid when pH is lower than its pKa ( $\sim 6.5$ ). And it has been suggested 332 to be oxidized into 2-oxo-histidine, which carries no charges. This loss of charge will 333 subsequently affect protein net charges and ultimately the filament net charges (Bao et 334 335 al., 2018). The oxidation-induced modification of protein net charges offered a novel perspective in understanding oxidation-induced functionalities change of muscle 336 337 proteins. Protein oxidation also affects eating quality of muscle foods, such as color, water-holding and texture. Many reviews have covered those topics in great depth (Bao 338

339 & Ertbjerg, 2019; Estévez, 2011; Hematyar et al., 2019; Lund et al., 2011; Zhang et al.,

340 2013). In general, protein oxidation has a negative impact on the eating quality of meat.

341 Relationship between dietary protein oxidation and human health has received great interest. As pointed out by Hellwig (2019), several aspects need to be considered 342 343 on the relation between protein oxidation and human health. Those aspects include the 344 loss of essential amino acids, the influence on protein digestibility, the formation of potentially toxic compounds and possible transfer of oxidative damage to body proteins. 345 Oxidative modification of essential amino acids limits their bioavailability. The 346 347 mechanism by which oxidation modulates digestibility is complex. Mild oxidation of 348 many proteins increases their susceptibility to proteolysis. When oxidation proceeds to 349 a certain degree, excessive protein polymerization and aggregation occur, which 350 subsequently impairs protein digestibility (Li et al., 2017; Santé-Lhoutellier et al., 2007; Santé-Lhoutellier et al., 2008). Other than lowered nutritional value, protein oxidation 351 may be potentially toxic as the gastrointestinal tract and internal organs are exposed to 352 oxidation products. Oxidatively modified amino acids and peptides which resist 353 digestion may enter the large intestine and be utilized by microbiota and turn into 354 355 mutagenic compounds such as biogenic amines, ammonia, cresol and indole (Hellwig, 356 2019). Some pathologies are accompanied by the presence of oxidized amino acids at 357 levels beyond the physiological situation, indicating a toxic effect of oxidized proteins. 358 Oxidative damage may transfer from proteins to other targets such as DNA and lead to malfunction of gene expression. Furthermore, there is a strong link between protein 359 360 oxidation and aging (Davies & Dean, 1997)

# 361 3.2 Possible links between physicochemical changes and protein oxidation in 362 frozen muscle foods

363	As discussed in section 2, freezing leads to a series of physicochemical changes in
364	muscle foods, including temperature drop, formation of ice, concentrating of solutes,
365	alteration of ionic strength and pH, freezer burn, lipid oxidation, protein denaturation,
366	etc. These physicochemical changes may have significant impacts on protein oxidation.
367	Environmental factors such as pH and ionic strength are expected to affect many
368	chemical reactions, including protein oxidation. Bertram et al. (2007) investigated the
369	oxidation of myofibrillar proteins by hemoglobin and $H_2O_2$ under different
370	combinations of pH and ionic strength, the results showed that pH had a great impact
371	on the formation di-tyrosine, a general maker of protein oxidation. The formation of di-
372	tyrosine was much greater in pH 5.4, as compared to pH 6.2 or pH 7.0, and di-tyrosine
373	appeared to increase with increased ionic strength. Several possible explanations were
374	proposed for the pH-dependent formation of di-tyrosine including i) The activation and
375	pro-oxidant actions of hemoglobin is facilitated at acidic pH ii) lower pH induces
376	denaturation of hemoglobin, leading to iron release which, in turn, initiates a Fenton
377	reaction and iii) elevated pH leads to longer distances between tyrosine residues and
378	limits the effective cross-linking reaction. Other than directly affecting protein
379	oxidation, changes in pH and ionic strength may lead to protein denaturation (Zhang et
380	al., 2021). Denatured proteins may expose the previously buried oxidation-susceptible
381	amino acid residues, and thereby enhance protein oxidation. Other factors, such as the
382	formation of ice-water interface, distortion of myofibrillar proteins caused by ice

383	crystals, and generation of free fatty acids, all lead protein denaturation in frozen meat.
384	Therefore, those factors may promote protein oxidation as well. In the study of Zhang
385	et al. (2021), they used different combinations of pH and ionic strength to mimic the
386	condition of frozen-thawed meat. Results showed that both a lower pH and a higher
387	concentration of KCl increased the surface hydrophobicity of myofibrillar proteins,
388	which indicated a greater protein denaturation. It should be noted that both Bertram et
389	al. (2007) and Zhang et al. (2021) used model systems, where extracted myofibrillar
390	proteins were incubated with solutions containing different concentrations of salt and
391	protons. The reactions in the meat matrix can differ significantly, due to reasons like
392	macromolecular crowding, confinement or adsorption as discussed in section 2.1.

A portion of water remains as liquid in frozen muscle foods and it is concentrated 393 394 with solutes (most of them prooxidants). Such unfrozen water is normally surrounding muscle proteins and therefore molecular crowing takes place during frozen storage may 395 promote the contact between pro-oxidants and susceptible protein molecules, making 396 397 of frozen meat a pro-oxidative environment despite of the reduced movement of molecules due to low temperatures. For example, freezing-induced dehydration caused 398 399 close approach of the protoplasmic proteins in leaves, which promoted the formation 400 of disulfide bonds (Levitt, 1962), a general marker of protein oxidation. Oxygen exposure is obviously one of the important factors for the development of protein 401 402 oxidation. Therefore, membrane rupture caused by ice crystals can accelerate oxidation via allowing more diffusion of oxygen into close contact with cell interior. In the case 403 404 of severe freezer burn, sublimation-induced honeycomb structures in the surface of 405 meat will greatly facilitate the penetration of oxygen into a deeper position, and thereby406 accelerate oxygen-induced protein oxidation.

407 Hambly & Gross (2009) aimed to address the question of whether protein oxidation occurs primarily when its solution is frozen or during freezing, and they found 408 409 that freezing of apomyoglobin in peroxide solution led to protein oxidation. 410 Furthermore, the oxidation was dependent on incubation temperature and time, but not as a result of freeze or thaw. After 2 hour, protein oxidation is less at 4 °C or 22 °C, as 411 compared to freezing at -15 °C or -80 °C. At temperatures above the freezing point, 412 H<sub>2</sub>O<sub>2</sub> molecules are moving in three dimensions and they are constantly moving away 413 414 from the protein. Therefore, the oxidation reaction proceeds slowly. In contrast, when the solution is frozen, the few  $H_2O_2$  molecules are trapped and cannot diffuse away 415 416 from the protein. Oxidant diffuses over some portion of the protein surface in the solid state and oxidize the most reactive amino acid residues. Buried residues are also 417 418 oxidized suggesting electron transfer between the solvent-exposed residues and the interior of the protein in the solid state. 419

Theoretically, oxidative reactions could be transferred between lipids and proteins. This was supported by Burcham, & Kuhan (1996) who showed that incubation with MDA introduced carbonyl groups into BSA. In agreement, Zhang et al. (2011) found that consumption of oxidized oil was related to higher protein carbonyl content in breast meat of broiler chickens. There were also evidences of lipid-induced protein oxidation in frozen meat. For example, Utrera et al. (2014a) observed more intense protein oxidation in frozen beef patties with higher fat content. Many studies found that protein 427 oxidation was accompanied with lipid oxidation in frozen muscle foods. However,
428 other than lipid oxidation, the relationships between physicochemical factors (muscle
429 structure distortion, macromolecular crowding, generation of freezing potential, altered
430 pH, etc.) and protein oxidation in frozen muscle foods require further deep
431 investigations.

432

#### 433 **3.3 Factors affecting protein oxidation in frozen muscle foods**

Protein oxidation occurs readily in frozen muscle foods. Studies on protein
oxidation in frozen muscle foods are summarized in Table 1 (livestock meat) and in
Table 2 (aquatic products) with a focus on recent progress. Various factors may affect
protein oxidation in frozen muscle foods as outlined below.

438 3.3.1 *Types of muscle foods* 

Major categories of muscle foods include livestock animals and aquatic animals. 439 Chemical composition of muscles from mammals, domestic birds and aquatic animals 440 can be greatly influenced by type of animal, feeding, physiological status, slaughter 441 442 process, muscle cuts, etc. (Lawrie & Ledward, 2006). Such factors are expected to influence protein oxidation as they determine the concentration of pro- (heme iron, 443 444 unsaturated lipids, etc.) and antioxidant (tocopherols, carotenoids, etc.) components in muscle tissue. For example, Benjakul et al. (2003) compared the stability of muscle 445 proteins from some tropical fish during frozen storage, and they found that sulfhydryl 446 447 content in lizardfish decreased quickly and was lower compared to other species studied.

This was attributed to a higher level of formaldehyde formed in lizardfish. Estévez et 448 al. (2011) found a significant effect of muscle type on the formation of specific 449 carbonyls ( $\alpha$ -aminoadipic and  $\gamma$ -glutamic semialdehydes) in pork subjected to frozen 450 storage. And the different susceptibility of muscle cuts to oxidative reactions during 451 frozen storage was ascribed to the variations between muscle fibers (oxidative vs. 452 453 glycolytic), which largely determines the overall chemical composition of the muscles. 454 Utrera et al. (2014b) observed the oxidative stability of patties made from different muscle cuts, and the heme-iron content, antioxidant enzyme activity and PUFA content 455 were suggested to play a major role. Myofibrillar proteins are susceptible to oxidation 456 457 with myosin being the most sensitive (Lund et al., 2011). Liu et al. (2011) compared the stability of fish actomyosin and pork actomyosin, and they found that pork 458 actomyosin had higher stability (less exposed thiols and hydrophobic groups) during 459 cold storage. Differences in physicochemical properties of muscle proteins and the 460 antioxidative defense systems of the muscle are likely to affect the process of oxidation 461 during frozen storage. 462

463

#### 464 3.3.2 *Freezing rate, and frozen storage and thawing conditions*

In general, longer storage time leads to increased protein oxidation in frozen muscle foods. However, the carbonyl content of amino acid side chains, a generally used marker of protein oxidation, may decrease with prolonged storage (Estévez et al., 2011; Holman et al., 2018). The decrease may be caused by cross-linking between

469	protein semialdehydes and other amine-containing amino acid residues, as suggested
470	by Estévez et al. (2011). As for storage temperature, Baron et al. (2007) found that
471	carbonyls of rainbow trout fillets increased with storage time at -20 °C, but not at -30
472	or -80 °C, indicating that temperatures lower than -30 °C can inhibit protein oxidation.
473	To similar conclusions came Utrera et al. (2014), who reported that frozen storage had
474	a remarkable impact on specific oxidative changes in meat proteins (carbonylation, and
475	tryptophan depletion) and such modifications were temperature-dependent. According
476	to these authors, the protein oxidation occurred during frozen storage at -8°C and -18°C
477	contributed to impaired quality traits in patties cooked upon thawing. Similarly, Li et
478	al. (2020a) reported that pork patties stored at -18 or -25 °C had lower content of
479	carbonyls than at -8 °C, but no difference was found between -18 and -25 °C. Qian et al.
480	(2021) investigated protein oxidation in beef stored at temperatures ranged from -1 °C
481	to -18 °C, and they found that myofibrillar protein oxidation decreased when the
482	temperature decreased from -1 to -12 °C, but there was no clear difference between -12
483	°C and -18 °C. Therefore, it appears that protein oxidation in frozen muscle foods can
484	be inhibited when the storage temperature is lower than a critical point. However, this
485	critical temperature varies between studies, and the mechanism remains to be
486	investigated. Szymczak et al. (2020) compared the effect of constant and fluctuating
487	temperatures during frozen storage on quality of herrings, and they found that herrings
488	stored at constant temperatures had lower peroxide values, suggesting constant storage
489	temperature may be beneficial in maintaining the oxidative stability.

490 Other than frozen storage, the freezing and thawing process may also affect protein 491 oxidation in muscle foods. Increased freeze-thaw cycles have been demonstrated to 492 increase protein oxidation in a variety of muscle foods (Pan et al., 2021; Shao et al., 2018; Zheng et al., 2020). Kim et al. (2018) found that the freezing rate had an impact 493 on the oxidative stability of muscle proteins, however, the aging combination had 494 495 overriding impacts over freezing rate. Hou et al. (2020) observed no difference in total 496 and free sulfhydryl groups of pork between immersion solution freezing and air blast freezing, although they found that immersion solution freezing reduced lipid oxidation. 497 Different thawing methods have been studied in meat and meat products, including 498 499 conventional thawing by water or air, or combined with microwave, ultrasound, electrostatic field, etc. Cai et al. (2020b) compared effects of different thawing methods 500 501 on the myofibrillar protein oxidation in large-mouth bass. The results showed that microwave thawing in combination with either vacuum or magnetic nanoparticles 502 resulted in a lower degree of protein oxidation as compared to microwave thawing alone. 503 Combination of far-infrared thawing with magnetic nanoparticles was found to be the 504 most effective methods in controlling protein oxidation. Protein oxidation did not 505 receive much research attention during the freezing and thawing of meat, but existing 506 507 literature clearly show that the freezing and thawing processes have significant impact on protein oxidation. 508

509

510 3.3.3 *Processing of the muscle foods* 

511 Processed meat constitutes a large percentage of muscle foods. Various processing 512 technologies affect the physicochemical properties of meat and they potentially affect protein oxidation. Soladoye et al. (2015) thoroughly reviewed the impacts of processing 513 514 on protein oxidation in meat and meat products. It is worth to point out that frozen storage may increase the susceptibility towards oxidation during successive 515 516 technological processes, and processing frozen/thawed meat compared to unfrozen one 517 has been reported to influence protein oxidation and quality traits in dry-cured and 518 cooked meat products (Utrera et al., 2012; Utrera et al., 2015; Lorido et al., 2016). Some processing technology can be applied to meats in combination with freezing. The 519 520 impacts of processing on protein oxidation in frozen muscle foods, with a focus on recent studies, are briefly summarized as below. 521

522 Packaging is widely used for meat and meat products and oxygen in the package is expected to induce protein oxidation. Vacuum packaging was found to inhibit the 523 formation of protein carbonyls in frozen pork as compared to oxygen permeable bags 524 525 (Estévez et al., 2011). Similarly, it was observed that both light exposure and increased 526 oxygen concentration in the package could significantly accelerate protein oxidation in 527 the frozen obscure pufferfish (Zheng et al., 2021). Non-thermal processing technologies 528 can be applied in muscle foods to extend shelf life. It has been reported that treatment 529 with high pressure (Prego et al., 2021), pulsed electric field (Kantono et al., 2021; Li et 530 al., 2020b), and irradiation (Arshad et al., 2020) promoted lipid oxidation in frozen muscle foods. Since protein oxidation and lipid oxidation are closely related, it is 531 532 expected that these non-thermal processes can enhance protein oxidation as well. For

533	example, Cropotova et al. (2019) found that high-pressure treated haddock and
534	mackerel minces had higher content of carbonyls in sarcoplasmic and myofibrillar
535	proteins. The relationship between lipid oxidation and protein oxidation in high
536	pressure treated meat has been thoroughly discussed by Guyon et al. (2016). During the
537	production of meat products, non-meat ingredients may be added. It has been shown
538	that the incorporation of plant extracts (Botsoglou et al., 2014; Huang et al., 2019; Ozen
539	& Soyer, 2018; Turgut et al., 2017; Wang et al., 2017), hydrolysates from surimi
540	process byproducts (Zhang et al., 2020), and antifreeze protein (Nian et al., 2020; Wang
541	et al., 2021) inhibited protein oxidation in frozen muscle foods.

542

#### 543 **3.4** Control of protein oxidation in frozen muscle foods

Inhibition of protein oxidation in muscle foods can be achieved through animal 544 545 feeding or processing (Estévez, 2011; Lund et al., 2011). Feeding can modify the fatty acid composition of muscle tissue and also deposit anti-oxidants such as vitamin E in 546 muscle tissue, and thereby regulate the oxidative stability of raw meat. Liu et al. (2018) 547 548 investigated the effects of saturation level of added lipids on protein oxidation in minced pork, and the results showed that less saturated group had higher oxidative 549 550 stability in relation to both lipid and protein oxidation. This is in contrast to general belief and may be caused by the higher amount of antioxidant (vitamin E) in less 551 552 saturated lipid fractions obtained from pork back fat. Incorporation of antioxidants into muscle foods is an effective way of controlling protein oxidation. There is a growing 553

interest in using natural antioxidants in meats (for generally used natural antioxidants
in meat and products, see Ribeiro et al., 2019). In frozen muscle foods, it has been
demonstrated that extracts of olive leaf (Botsoglou et al., 2014), dog rose (Utrera et al.,
2015), pomegranate peel (Turgut et al., 2017), green tea, grape seed, and pomegranate
rind (Ozen & Soyer, 2018), clove and rosemary (Huang et al., 2019), nutmeg (Zhu et
al. 2020) were effective antioxidants towards protein oxidation.

560 Freezing, storage and thawing are known to affect protein oxidation, optimization of these processes may bring benefits. For examples, Hou et al. (2020) found that 561 immersion solution freezing produced smaller ice crystals in pork, and reduced lipid 562 563 oxidation was observed as compared to air blast freezing. The addition of ice structuring protein (Wang et al., 2021) or herring antifreeze protein (Nian et al., 2020) were found 564 565 to inhibit protein oxidation in frozen meats. As we discussed in this paper, a series of physicochemical factors may affect protein oxidation throughout the whole process of 566 frozen muscle foods. A better understanding of the possible mechanism of frozen-567 568 induced protein oxidation would allow to achieve a better control of protein oxidation, and ultimately improve the quality of frozen muscle foods. 569

570

571 4. Concluding remarks

572 The present review thoroughly discussed key physicochemical changes in frozen
573 muscle foods, including the formation of ice crystals and the ice-water interface,
574 concentrating of proteins and pro-oxidants, altered pH and ionic strength, freezer burn,

575 etc. These changes may contribute to the occurrence of protein oxidation in frozen 576 muscle foods. Of these key physicochemical changes in frozen muscle foods, the ice-577 water interface is believed to be a main source of protein instability. Based on the assumption that protein denaturation would expose previously buried residues which 578 are susceptible to oxidation, we suggest that freezing-induced oxidation may be 579 580 mediated by protein denaturation in frozen muscle foods. In muscle foods, protein 581 oxidation can proceed to a significant extent, which often negatively affects the eating quality and nutritional value. A deeper understanding of the mechanism for freezing-582 induced protein oxidation may lead to a better control of oxidative damage to muscle 583 584 foods.

585

#### 586 Acknowledgements

587 Yulong Bao would like to acknowledge the financial support from the National
588 Natural Science Foundation of China (31901758) and the Natural Science Foundation
589 of Jiangsu Province (BK20190591). Mario Estévez would like to acknowledge the
590 financial support from the Spanish Ministry of Economy and Competitiveness through
591 the project AGL2017-84586R.

592

#### 593 Conflicts of Interest

594 The authors confirm that there are no conflicts of interest to declare.

595

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## 973 Figure legends

974

Fig.1. Schematic illustration of ice crystals in frozen muscle foods. (a) Formation of
ice crystals during freezing (Mazur 1970); (b) Ice recrystallization during frozen
storage; (c) Effect of ice crystals on muscle structure after thawing (Nakazawa &
Okazaki, 2020).

979

980	Fig.2. Illustration of mechanical distortion of muscle structure in frozen meat. (A)
981	Under unfrozen condition, water stays in liquid form and myosin heads are bound to
982	actin filament to form rigor bonds; (B) Under normal frozen condition, a major part of
000	the material state in the former of a lidite. Due to the formation of anter called an inc

983 the water exists in the form of solid ice. Due to the formation of extra-cellular ice

984 crystals, muscle fibers are severely compressed. The fiber compression leads to

985 distortion of rigor bonds and ultimately to distortion of myosin heads.

986

Fig.3. Schematic illustration of the change in reaction conditions caused by formationof ice crystals (Adapted from Takenaka et al. 1996)

989

990 Fig.4. Possible mechanisms of ice-induced protein denaturation. (a) change of pH and

ionic strength; (b) accumulation of air bubbles at the ice interface; (c) adsorption at

992 the ice interface; (d) Pressure due to ice growth; (e), enhance cold denaturation.

993 Modified from Arsiccio & Pisano 2020.

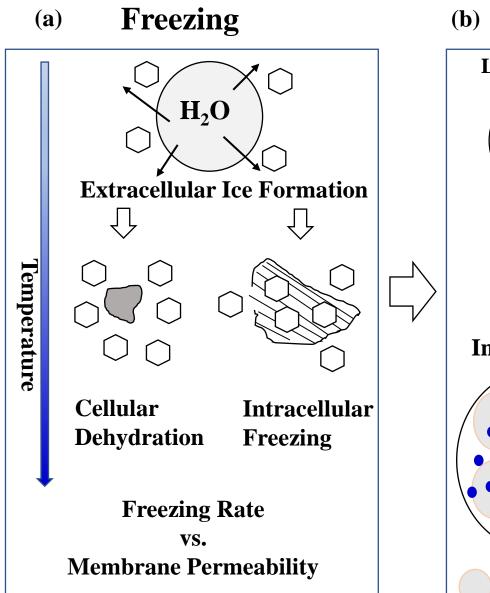
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995 Fig.5. Schematic illustration of pH changes in frozen electrolyte solutions mediated

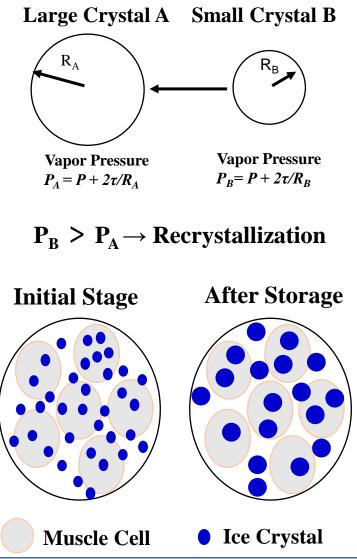
by freezing potential, based on Carmen et al., 2006.

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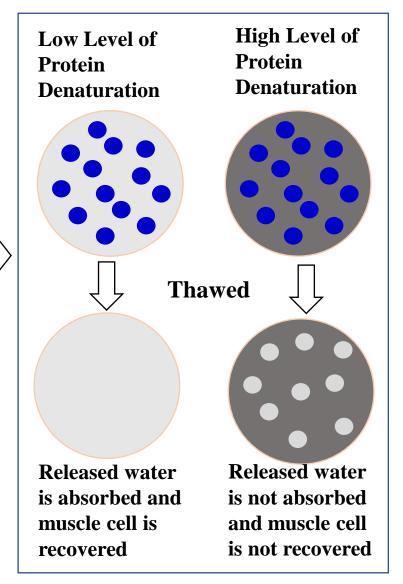
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## b) Frozen storage

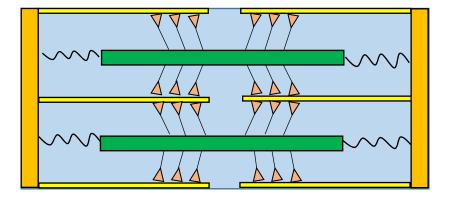


## (c) **Thawing**

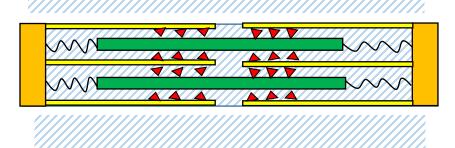


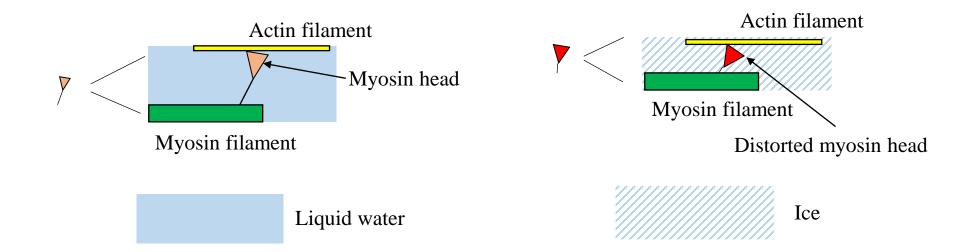
B

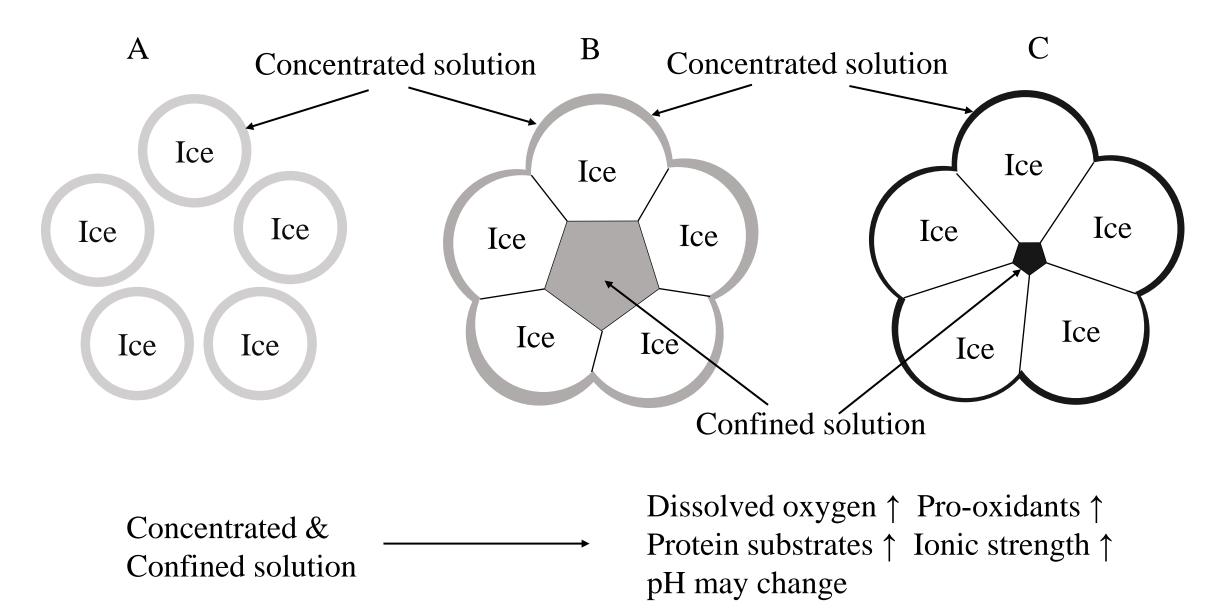
## A Unfrozen sarcomere

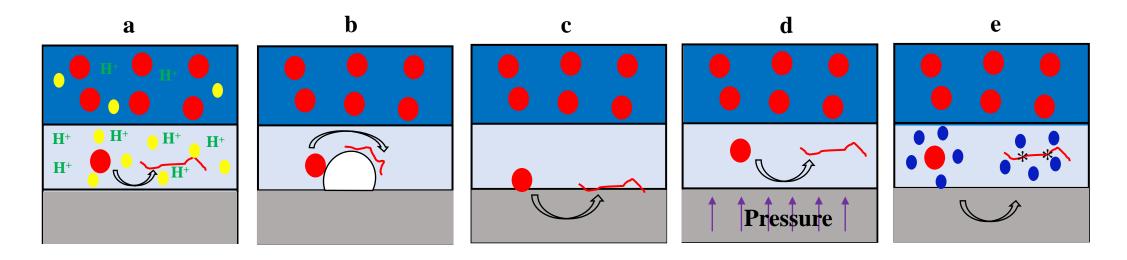


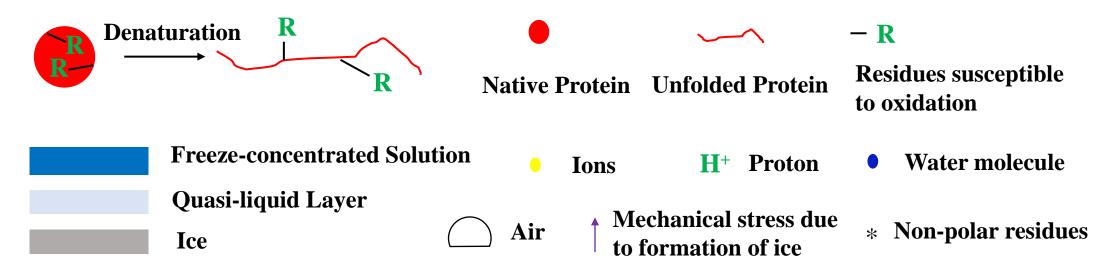












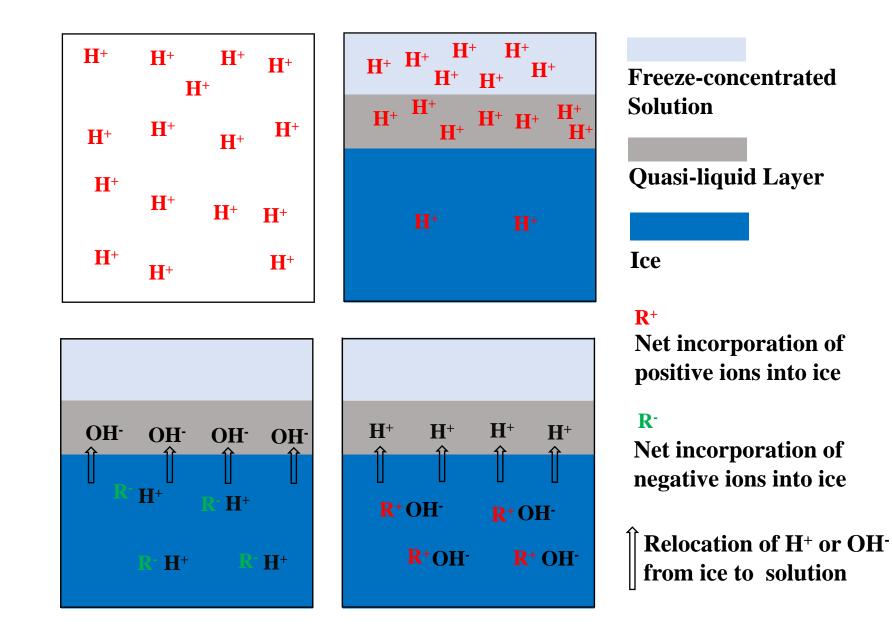


Table 1. Summary of literature on the effects of freezing on protein oxidation in livestock muscle foods

Frozen Muscle foods	Main findings on protein oxidation	Physicochemical changes relevant to protein oxidation	References
Pork patties	Carbonyls increased with increased freeze-thaw cycles and fat content of patties	Lipid oxidation increased with increased freeze-thaw cycles and fat content of patties	Pan et al., 2021
Pork patties	Addition of ice structuring protein to patties reduced protein carbonyls	Ice structuring protein limited growth of ice crystals and inhibited lipid oxidation	Wang et al., 2021
Pork patties	Storage at -18, -25, -18/-25 °C led to lower content of carbonyls than at -8 °C or - 8/-18 °C; generally, no difference among -18, -25, -18/-25 °C	Li et al., 2020a	
Pork dumplings	Carbonyls increased with storage time; incorporation of extracts of clove and rosemary led to lower carbonyls and less protein cross-linking	Huang et al., 2019	
Pork	No difference in total and free sulfhydryl groups between Immersion solution freezing (ISF) and air blast freezing (AF); sulfhydryl groups decreased with time	ISF produced smaller ice crystals and reduced lipid oxidation compared to AF	Hou et al., 2020
Pork	Freezing rate had an impact on oxidative stability of muscle proteins, however, Aging prior or after freezing increased lipid the aging process had overriding impacts over freezing rate oxidation		Kim et al., 2018
Pork patties	With increased time, carbonyls increased and sulfhydryl decreased; addition of olive leaf extract inhibited lipid oilve leaf extract inhibited protein oxidation oxidation		Botsoglou et al., 2014
Pork dumplings	Carbonyls increased with storage time; lower temperature had lower carbonyls	Free fatty acids increased with frozen storage time; lipid oxidation increased with storage time and temperature	Huang et al., 2013
	Specific carbonyls AAS and GGS first increase and then decrease with time;		
Pork	oxygen permeable bag vs. vacuum pack, minced vs. intact, <i>longissumus dorsi</i> vs. <i>psoas major</i> generally led to greater AAS and GGS	-	Estévez et al., 2011
Pork loin	Carbonyls increased and sulfhydryl decreased with increased freeze-thaw cycles	Lipid oxidation and protein denaturation increased with freeze-thaw cycles	Xia et al., 2009

	Myofibrillar protein oxidation decreased when storage temperature decreased	Protein denaturation was generally greater	
Beef	from -1 to -12 °C, but there was no clear difference between -12 °C and -18 °C	at higher temperatures, ionic- and hydrogen	Qian et al., 2021
		bonds were greater at lower temperatures	
Beef loin	Frozen storage first led to increased carbonyls and then decreased	-	Holman et al., 2018
Beef	With increased time, carbonyls increased and sulfhydryl decreased; addition of	Pomegranate peel extract reduced lipid	Turgut et al., 2017
meatballs	pomegranate peel extract greatly inhibited the change in oxidation marker	oxidation	Tulgut et al., 2017
	Frozen storage increased carbonyl compound (AAA and AAS), and also		
Beef patties	increased the susceptibility towards oxidation during successive technological	Rose extract inhibited lipid oxidation	Utrera et al., 2015
	process; addition of rose extract inhibited the formation of AAS and AAA,		
Beef patties	Frozen storage increased the specific carbonyl (AAS) and Schiff base, higher fat		Utrera et al., 2014a
beel patties	content in patties led to increased AAS and Schiff base	-	
Beef patties	Different muscle cuts had different oxidative stability	Heme-iron content, antioxidant enzyme	Utrera et al., 2014b
-		activity and PUFA content differ	
Lamb loin	Frozen storage did not affect carbonyls		Coombs et al., 2018
Emulsified	Carbonyls increased with storage time; no difference due to types of used fat	Use of skin fat lower lipid oxidation	Santos et al., 2020
chicken	(skin and abdominal fat)		
patties			
Chicken	After 3 freeze-thaw cycles, carbonyls increased and sulfhydryl decreased	Lipid oxidation increased after 3 cycles	Ali et al., 2015
breast			· · · · · · · · · · · · · · · · · · ·
	Carbonyls in chicken leg and breast increased with time; lower freezing		
Chicken	temperature led to lower carbonyls; sulfhydryl decreased with time; lower	Lipid oxidation increased with time	Soyer et al., 2010
	freezing temperature led to higher content of sulfhydryl in leg meat		

Table 2. Summary of literature on the effects of freezing on protein oxidation in aquatic muscle foods

Frozen Muscle foods	Main findings on protein oxidation	Physicochemical changes relevant to protein oxidation	References
Puffer fish	Light exposure, oxygen concentration or increased	Changes of protein structure were in line with the	Zheng et al., 2021
	freeze-thaw cycles accelerated protein oxidation	extent of protein oxidation	-
Shrimp	Protein oxidation increased with time and storage temperature, and	Protein denaturation followed similar pattern with	Ji et al., 2021
	temperature had a greater impact	protein oxidation	
Puffer fish	Increased freeze-thaw cycles resulted decrease in sulfhydryl content and antioxidant enzyme activity, and increase in carbonyls	Increased freeze-thaw cycles led to increased	Zheng et al., 2020
i unci nisii	and protein cross-linking	lipid oxidation	Zheng et al., 2020
	Fish samples soaked in solution containing herring antifreeze		
Largemouth bass	protein (hAFP) had lower content of dityrosine, carbonyls, and	hAFP inhibited recrystallization and decreased	Nian et al., 2020
	higher total sulfhydryl	freezing point	
Red sea bream	Fish samples soaked in solution containing herring antifreeze	hAFP stabilized secondary and tertiary	Cai et al., 2020a
Keu sea Dream	protein had lower content of dityrosine, carbonyls, and disulfide	conformation of myofibrillar proteins	Cal et al., 2020a
	Microwave thawing in combination with either vacuum or	Protein denaturation level was similar among different thawing methods	
Largemouth bass	magnetic nanoparticles, and far-infrared thawing with magnetic		Cai et al., 2020b
	nanoparticles had lower degree of protein oxidation		
Silver carp	Hydrolysates via trypsin- and alcalase-treated surimi process	Hydrolysates delayed the destroy of myofibrillar protein structural integrity	
surimi	byproducts delayed the oxidation of cysteine, and lowered protein		Zhang et al., 2020
	carbonylation Carbonyls of sarcoplasmic and myofibrillar proteins increased with	-	
Mackerel fillets	storage time		Cropotova et al., 2019
Large-mouth	Total sulfhydryl decreased with freeze-thaw cycles; fermented		
bass	soybean delayed the decline of reactive and total sulfhydryl	-	Shao et al., 2018
Mackerel mince	Carbonyls increased with time; addition of extracts from green tea,	Lipid oxidation increased with time; addition of	Ozen & Soyer, 2018

	grape seed, pomegranate rind delayed the increase of carbonyls, but	extracts inhibited lipid oxidation	
	no difference among antioxidant sources; similar trend for the		
	decrease of sulfhydryl		
Haddock and	High-pressure treated samples had higher content of carbonyls in	Lipid oxidation was higher in 200 MPa treated	Cropotova et al., 2020
mackerel minces	sarcoplasmic and myofibrillar proteins; higher pressure led to		
mackerer minices	higher carbonyls	mackerel minces, but lower in 300 MPa	
	Total sulfhydryl decreased with time, and $\alpha$ -tocopherol, ascorbic		
Red sea bream	acid, brown seaweed polyphenol inhibited the loss of sulfhydryl;	All added antioxidants inhibited linid avidation	Wong et al. 2017
mince	carbonyls increased, and $\alpha$ -tocopherol inhibited the formation of	All added antioxidants inhibited lipid oxidation	Wang et al., 2017
	carbonyl		
	Incorporation of carrageenan oligosaccharides prior to frozen		
Peeled shrimp	storage decreased protein carbonyl and dityrosine, maintained a	-	Zhang et al., 2018
	higher total sulfhydryl content when compared with control		
Cod mince	Frozen storage increased carbonyls, the addition of caffeic acid had	affair and reduced limit evidetion	Lorgon & Undeland 2010
Cod mince	no protective effect on protein carbonylation	caffeic acid reduced lipid oxidation	Larsson & Undeland, 2010
Rainbow trout	Carbonyls increased with time at -20 °C, but not at -30 or -80 °C	Lipid oxidation and free fatty acids increased	Baron et al., 2007
fillets	Carbonyis increased with time at -20°C, but not at -30°C	with time and frozen storage temperature	Baron et al., 2007
	Carbonyls were higher at -20 °C in the total meat homogenate and		
Rainbow trout	low-salt soluble fraction; carbonylated proteins include nucleoside		
fillets	diphosphate kinase, adenylate kinase, pyruvate kinase, actin,	-	Kjaersgard et al., 2006
miets	creatine kinase, tropomyosin, myosin light chains 1 and 2, and		
	myosin heavy chain		