

Relationships between loin physical, meat quality and intramuscular collagen characteristics of carcasses from crossbred swine populations

by

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A thesis submitted in partial fulfillment of the requirements for the degree of

Master of Science

in

Animal Science

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ABSTRACT

Carcass and meat characteristics are of great importance to the pork industry as they determine the ultimate returns from animals produced. Making genetic progress in these traits requires them to be measurable, heritable and have adequate additive genetic variability. Estimates of genetic parameters must therefore be assessed to determine heritability and level of additive genetic variation for use in selection programs. In the first study, phenotypic data from 500 pedigreed crossbred pigs from Duroc sires and hybrid Large White \times Landrace sows from two different swine genetics companies were used. Fixed effects (slaughter batch, sex and company), and a random additive effect were fitted in bivariate animal models to estimate phenotypic and genetic parameters in ASReml. Moderate heritabilities were obtained for loin compositional traits ranging from 0.21 ± 0.10 for bone weight to 0.44 ± 0.11 for loin eye weight with low estimates of 0.09 ± 0.08 , 0.19 ± 0.08 , 0.13 ± 0.09 obtained for bone percent, loin weight and fat trim weight, respectively. Meat quality traits were low to moderately heritable with the highest estimate being found for intramuscular fat (0.42 ± 0.13). The heritability estimates for percentages of heat soluble and insoluble collagen were 0.12 ± 0.09 and 0.15 ± 0.09 , respectively, while 0.33 ± 0.12 was found for total collagen content. Moderate heritability implies the possibility of improving these traits through selective breeding. Strong negative genetic correlations between moisture and fat traits and a further negative correlation between fat and muscling traits confirmed that selection for improved muscling over time negatively affects fat traits and may then decrease meat eating quality. The strong genetic correlations of lightness (L^*) with soluble collagen (0.68 ± 0.30) and insoluble collagen (-0.71 ± 0.35) suggest possible pleiotropic gene effects on these traits. Warner-Bratzler shear force (WBSF) had moderate genetic correlations with insoluble collagen (0.42 ± 0.16) and soluble collagen (-0.38 ± 0.10), suggesting a potential relationship between some

of the genes impacting these traits. Genetic correlations between WBSF and collagen characteristics indicated that despite the relative youthfulness of pigs at slaughter, genetic selection for collagen solubility may decrease WBSF.

In the second study, the contributions of company, sex and loin physical and collagen characteristics to measures of pork carcass and meat quality in a Canadian context were examined. Results from this study indicated that intramuscular collagen characteristics influenced loin cooking loss and further revealed that Warner-Bratzler shear force (WBSF) values were driven more by denaturation of myofibrillar proteins than by intramuscular fat content. Means separation analysis also showed WBSF to increase with increased muscularity and reduced backfat which suggests that selection for rapid growth rate with its associated reduction in calpain activity may decrease pork tenderness. This study further implicates pH and intramuscular fat content as part of the traits influencing insoluble collagen percent indicating that deposition of fat in the perimysium may be a potential disruptor of crosslink formation. Step- wise regression indicated that pH is not a reliable sole predictor of meat quality in loin that has been frozen and then thawed and thus other variables in addition to pH are needed to reliably predict frozen/thawed pork quality. Drip loss predicted most of the variation in tenderness which indicated that pork samples in the present study may have undergone myofibrillar protein oxidation during frozen storage or dehydration during cooking. The results indicated that sex has no significant influence ($p>0.05$) on loin compositional and meat quality traits while source of animal has a significant ($p<0.01$) influence on loin composition, pH, drip loss (%), cooking loss (%) and collagen characteristics.

In summary, the genetic parameters estimated in this study will make it feasible to predict response to genetic selection on loin and collagen characteristics and further add to the already

established research database on meat quality traits. The results of this study also provide insights on variables with potential for predicting pork quality and suggest that although the populations studied were obtained from the same breed combinations differentiation in terms of loin compositional, meat quality and collagen characteristics would not be expected based on sex but would be expected in terms of animal origin.

Acknowledgements

Throughout the writing of this thesis I have received a great deal of assistance and support. I would first like to thank the Natural Sciences and Engineering Research Council (NSERC), Hypor Incorporated and Genesis Genetics for generously supporting this research.

Further, I would like to express my utmost gratitude to my supervisor Dr. Heather Bruce for giving me the opportunity to study at the University of Alberta, and for the time, effort and academic advice she provided throughout this study. The door to Dr. Bruce's office was always open whenever I had a question about my research or ran into a trouble spot. She consistently allowed this thesis to be my own work but steered me in the right the direction whenever she thought I needed it.

I would also like to express gratitude to Dr. Leluo Guan for the suggestions and guidance she provided me for the completion of this work. I further thank Professor Graham Plastow for his constructive comments and criticisms. I further thank Dr. Changxi Li and Dr. Arthur Gilmour whose expertise was invaluable during my statistical analysis.

Also, I would like to extend my thanks to Dr. Bimol Roy for his guidance during my laboratory work, Dr. Huaigang Lei for help during my statistical analysis and Dr. Robert Mukiibi who have provided me with invaluable assistance and support throughout the course of my studies.

Finally, I would really like to thank Rabaa Hamed, Jiyuan Li, Patience Coleman and all my lab mates and friends for their support and for providing happy distractions to rest my mind outside of my research. This experience would not have been the same without them.

In addition, I would also like to thank my family for their wise counsel, sympathetic ear, encouragement and their many sacrifices that have allowed me to reach this stage in my life. My father, Kwame Panin, My mother, Akua Konadu, my sisters, Diviana Amadie-Panin and Munich Konadu-Panin and Niece, Elsie Nana Achia Panin. I thank God for making everything come together in the end.

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List of Abbreviations

BP	Bone percent
BWT	Bone weight
CL	Cooking loss percent
ECM	Extracellular matrix
FOF	Foss crude fat
FOM	Foss crude moisture
FOP	Foss crude protein
FTW	Fat trim weight
IMCC	Intramuscular connective tissue characteristics
IMF	Intramuscular fat
ISOLC	Insoluble collagen percent
LEP	Loin eye percent
LEW	Loin eye weight
LOINA	Minolta a^*
LOINB	Minolta b^*
LOINC	Minolta chroma
LOINH	Minolta hue
LOINL	Minolta L^*
LWT	Loin weight
MC	Moisture content (%)
MTW	Meat trim weight
MRA	Metmyoglobin reducing activity

PSE	Pale, soft, and exudative
RTP	Rib trim percent
RTW	Rib trim weight
SOLC	Soluble collagen percent
TCOL	Total collagen (mg/g raw meat)
TFP	Thick fat percent
TFW	Thick fat weight
WBSF	Warner-Bratzler shear force (N)

Chapter 1

1.0 Introduction

Swine domestication from wild boar subspecies in Asia and Europe approximately 9000 years ago is known to have been of critical importance in establishing a secure supply of protein to various human civilizations (Bosse 2018; Guiffra et al., 2000). Indeed, pork is currently the most consumed meat worldwide accounting for about 40% of the world's meat consumption (USDA, 2017; Rothschild and Ruvinsky, 2011). Pork consumption per capita worldwide as of 1961 was 8.0 kg but saw a steady rise to 16.0 kg by 2013 (FAO, 2017; Ritchie and Roser, 2019). This increase was most likely driven by the exponential rise in the worldwide population leading to an increase in pork production from 24.8 million tons in 1961 to 112.3 million metric tons in 2013, representing an over 4-fold increase (FAO, 2017; Ritchie and Roser, 2019). In Canada, production went from 535,940 in 1961 to 2.53 million tons in 2013. Expectedly, OECD/FAO (2018) projected a 15% increase in global meat production over the next decade using the average of 2015-2017 as the base and this increase was anticipated to be driven mainly by developing countries. Concomitant with this increase, over the next ten years a rise in the global demand for mutton/lamb and beef while a decrease in the demand for pork and poultry is projected (OECD/FAO, 2018). This will most probably be due to the fact that consumers of meat in most developing countries, over the years, preferred pork and poultry due to the relatively lower market prices of their end products. However, projected growth in consumer incomes over the next decade is expected to fuel a shift in their preference to more costly meat like that from sheep and cattle (USDA, 2018; OECD/FAO, 2018). It is therefore crucial and critical that commercial pig producers and processors endeavor to boost production efficiency and most

24 importantly produce pork that meets the high standards of consumers so that they remain
25 competitive with producers of other species in the global livestock and animal protein markets
26 (Salas and Mingala, 2017).

27 Genetic improvement of animal performance traits like litter size, lean percent, carcass weight,
28 average daily gain, and growth rate that are economically vital to producers has conventionally
29 been the focus of swine breeding programs (Kanis et al., 2015). However, this trend is gradually
30 shifting as consumers are progressively allocating less importance to price in their purchasing
31 decisions and attaching more value to traits and issues like pig welfare and health, healthiness of
32 pork, organoleptic properties and overall eating quality (Tsakiridou et al., 2010; Ciobanu et al.,
33 2011). Meat quality is therefore progressively exerting much influence on processor and
34 consumer decisions and thus on the direction of the swine industry (Lonergan et al., 2001).
35 Indeed, research studies are showing that selection for traits of economic importance to
36 producers like feed efficiency, lean percent, and carcass weight could negatively affect pork
37 quality (Miar et al., 2014; Lonergan et al., 2001). The increase in demand for pork products of
38 good quality has led to an increase in research by animal breeders and meat scientists to discover
39 ways of improving meat quality (Ciobanu et al., 2001). Breeders are also increasingly becoming
40 interested in the relationships existing between and among animal performance indicators,
41 carcass, meat, and eating quality, and specific muscle histochemical and biochemical traits for
42 incorporation as traits for selection in their breeding programs (Suzuki et al., 2005; Larzul et al.,
43 1997).

44 Swine skeletal muscle tissue mass usually falls between 40 to 60% of their live weight
45 (Kauffman and Warner, 1993) and are composed of a number of tissues. These include muscle
46 fibers, vascular, nervous, adipose and connective tissues (Listrat et al., 2016). The ratio and

47 characteristics of muscle fibers, intramuscular fat and intramuscular connective tissues play key
48 roles in determining various quality dimensions (Listrat et al., 2016). Quality can be divided into
49 3 subclasses: carcass quality, meat quality and eating quality. Carcass quality usually includes
50 factors that critically affect the market price of carcasses, and some of these factors are carcass
51 fat cover, bone to meat ratio and carcass weight (Smulders, 1986) while meat quality
52 encompasses all the serviceability/processability, organoleptic, healthiness/nutritional and
53 microbiological aspects of meat at fabrication, purchase or examination (Hoffman, 1973). Eating
54 quality, a subset of meat quality, on the other hand, is defined by three cardinal traits: juiciness,
55 tenderness, and flavor (Mateescu, 2005).

56 Intramuscular fat content has been implicated in species to species flavor differences and
57 juiciness (Hornstein and Crowe, 1960) while muscle fiber density and intramuscular connective
58 tissue variability have been shown to influence meat texture (Zheng et al., 2018; Thompson,
59 2004). For many years, numerous researchers in the field of meat science have endeavored to
60 find the link between muscle texture and its composition (Dransfield, 1977). The role of
61 intramuscular connective tissues or its major protein collagen has been a topic of much debate.
62 Indeed, a critical reason for the increase in research studies focusing on the inherent
63 characteristics of intramuscular connective tissues has been because of meat texture variability
64 (Voutila et al., 2007). There is a consensus on the role of collagen in raw meat toughness but
65 some skepticism as to its role in cooked meat toughness. Proponents of the hypothesis that
66 collagen plays an only minor role on cooked meat toughness like Purslow (2018) suggest that
67 myofibrillar proteins drive meat toughness at high cooking temperatures with the collagens
68 essentially gelatinizing at these temperatures. However, other research studies like that of Light
69 et al. (1985) suggest that although collagen molecules shrink at high temperatures, meats with

70 high percentages of heat resistant trivalent cross-links tend to have stable matrices compared to
71 those with more heat labile crosslinks (which form gelatin) (Bailey and Light, 1989). Based on
72 this argument, it is suggested that the proportion of heat insoluble to heat soluble collagen
73 influences meat toughness and texture to a greater degree than previously thought (Purslow,
74 2018). Adequate additive genetic variability in collagen traits if obtained, may pave way to the
75 manipulation of its characteristics through selective breeding, as this is yet to be undertaken.
76 Indeed, studies (Brock et al., 2000) on the effects of selection for increased leanness on muscle
77 fiber type characteristics has shown that selection for reduced backfat thickness results in an
78 increase in glycolytic muscle fibers and a decrease in oxidative fibers, and the magnitude of
79 these changes is dependent on generation (selection period duration) and muscle type. This trend
80 is problematic as these changes tend to affect the metabolic properties of muscles (Brock et al.,
81 2000) and thus, muscles with higher glycolytic fibers are known to yield poor quality meat. It
82 therefore is crucial to undertake a study to characterize the genetic relationships among V, meat
83 quality and intramuscular collagen characteristics to test the hypothesis that because genetic
84 relationships exist amongst them, a genetic approach may be a better tool for managing meat
85 quality. Characterization of pork collagen has been limited most likely because pigs are
86 relatively immature (slaughtered based on physiological age rather than chronological age) at
87 slaughter and thus the proportion of heat-resistant collagen cross-links which would contribute to
88 cooked pork toughness is expected to be low. However, some studies still suggest that
89 intramuscular collagen contributes to immutable background toughness in meat irrespective of
90 maturity level (Purslow, 2005).

91 The objective of this chapter was to explore the contribution of genetics, breeding and
92 management practices to pork quality, and how collagen contributes to meat quality and is

93 affected by genetics and management practices. To understand this contribution, the structure
94 and composition of pork muscle and how it relates to pork quality must first be considered and
95 appreciated.

96 **1.1 Chemical composition and structural features of muscles**

97 Generally, post-rigor muscle that is yet to undergo its postmortem degradative transformations
98 will usually be comprised of 75% water, approximately 19% protein, about 2.5% fat and 3.5%
99 non-protein soluble substances (Voyle, 1979; Mateescu, 2015). The structural integrity of muscle
100 is maintained by 3 layers of connective tissues: there is the epimysium which sheathes the whole
101 muscle, the perimysium which surrounds bundles of muscle fibers and the endomysium which
102 surrounds each individual muscle fiber (Bailey and Light, 1989).

103 **1.1.1 Muscle water**

104 About 5-12% of the overall water content of muscle is in the space of the extracellular matrix
105 while the rest (88-95%) is held in the myofibers (between the myofibrils, within the myofibrils,
106 and between the myofibrils and cell membrane) (Voyle, 1979). Approximately 5% of the total
107 water volume in muscle is bound to hydrophilic sites on myofibrillar proteins (Lee, 2012). The
108 rest is categorized either as free water (immobilized due to myofibrillar protein configuration) or
109 loose water, which is the water expressed during water holding capacity measurements (Hamm,
110 1961).

111 **1.1.2 Muscle protein**

112 There are three basic classifications of muscle proteins. These are myofibrillar, sarcoplasmic and
113 stroma proteins (Xiong, 1997). This classification is based on their solubility characteristics
114 (Xiong, 1997). Myofibrillar proteins are salt-soluble and sarcoplasmic proteins water soluble

115 while stroma proteins are insoluble (Greaser and Guo, 2015). Figure 1.1 depicts the muscle
116 protein fractionation into these three solubility categories.

117 Myofibrillar and sarcoplasmic proteins tend to be intracellular while stroma proteins are usually
118 extracellular (Greaser and Guo, 2015). The major myofibrillar proteins by percent composition
119 are myosin (43%), actin (22%) and titin (8%) (Greaser and Guo, 2015) while myoglobin and
120 hemoglobin are the most important sarcoplasmic proteins (Papadopoulos, 2000). Stroma proteins
121 fundamentally constitute the muscle connective tissue component and include elastin, collagen,
122 and reticulin as well as the proteins in myocyte organelle membrane systems (Smulders, 1986).

123 **1.1.3 Muscle fat**

124 Fat reserves in mammals can be found in a number of internal and external anatomical locations
125 where they serve as energy depots (Lebret, 2008). Two main types of adipose cells are known so
126 far, these are white and brown adipose cells. The main differences between these two cells are
127 that brown adipose cells besides their brown color, tend to have comparatively higher
128 mitochondria, cytoplasm, and a greater level of vascularization (Swatland, 1994). Nascent fat
129 tissues tend to have a higher proportion of connective tissue, water and a lower proportion of
130 lipid; however as animals age dietary energy is diverted increasingly into fat growth, and the
131 lipid portion increases while the proportions of connective tissue and water decrease (Aberle et
132 al, 1977). The main systemic locations of adipose tissue with respect to muscles are between
133 (intermuscular) and within muscle (intramuscular). Intramuscular fat content varies with breed,
134 genotype, growth rate and nutrition (Lebret, 2008). For instance, Duroc and Meishan pigs have
135 higher fat contents than Landrace or Large White pigs (Shingfield et al., 2013). Castrated males
136 in the studies of Franco et al. (2014) and Alonso et al. (2009) had higher intramuscular fat
137 contents compared to entire females. Ramirez and Cava (2007), Kim et al. (2018) and Sundrum

138 et al. (2011) however found no effect of sex on intramuscular fat content. These differences in
139 the results from these studies underscore the need to take into consideration how sex influences
140 this trait in swine populations.

141 **1.2 The extracellular matrix (ECM)**

142 The extracellular matrix is known to be a sophisticated 3-dimensional network of
143 macromolecules found in the extracellular space (Kaushal et al., 2014). It is the non-cellular
144 component found within all organs and tissues, where it offers not only vital physical scaffolding
145 for the cellular constituents but also launches crucial biomechanical and biochemical cues (Kular
146 et al., 2014). It provides structural support for all mammalian tissues and organs, playing crucial
147 roles in cell proliferation, migration, differentiation, signaling, and survival (Hynes, 2002; 2009).
148 The importance of the ECM can be seen from the number of syndromes that occur from genetic
149 abnormalities in certain ECM proteins (Jarvelainen et al., 2009). These include macromolecular
150 network is mostly composed of laminins, proteoglycans, elastin, glycoproteins, collagens and
151 hyaluronan (Sainio and Järvelainen, 2014; Ruoslathi, 1988) secreted by a wide array of cell types
152 such as chondrocytes, osteoblasts, fibroblasts, endothelial cells, epithelial cells, adipocytes and
153 tenocytes (Kaushal et al., 2014; Sainio and Järvelainen, 2014; Milz et al., 2009). Extracellular
154 matrix components can be divided into 2 subclasses, specifically fiber-forming and interfibrillary
155 molecules. Laminins, elastin, fibronectins and certain collagen types are typical fiber-forming
156 ECM molecules, whereas the glycoproteins and proteoglycans are generally regarded as
157 interfibrillary ECM molecules (Frantz et al., 2010). The molecular components of the ECM are
158 subject to numerous post-translational modifications (Kassianidou et al., 2019) and by means of
159 these biochemical and physical characteristics, the ECM creates the mechanical and biochemical
160 properties of each organ, such as its elasticity, compressive and tensile strength, and also

161 contributes to protection through a buffering action that supports extracellular water retention
162 and homeostasis (Frantz et al., 2010). The protective, biomechanical, biochemical and
163 organizational properties of the ECM can vary from tissue to tissue (e.g. lungs versus bone) and
164 even within one tissue (e.g. renal medulla versus renal cortex) as well as from one physiological
165 state to another (cancerous versus normal) (Jarvelainen et al., 2009).

166 **1.3 Collagen structure and physiology**

167 **1.3.1 Collagen**

168 Collagens are extracellular matrix proteins that are vital in supporting the structures of many
169 tissues in the mammalian body (Myllyharju and Kivirikko, 2004). They are present in almost all
170 tissues and organs of mammals and constitute between 25-30% of the mammalian body protein
171 content, making it the most abundant protein in mammals (Langrock and Hoffmann, 2012;
172 Bailey and Light, 1989). They are key protein components of tissues of cartilage, teeth, tendons,
173 bones, skin, muscle connective tissues and blood vessels (Fratzl, 2008). Collagens have been
174 found to play important roles in pathways and processes such as cell and muscle morphogenesis,
175 cell signaling mediation, chemotaxis and even wound healing in metazoans (Myllyharju and
176 Kivirikko, 2004; Gjaltema and Bank, 2017). All collagen molecules regardless of their types
177 exhibit a repeating Gly-Xaa-Yaa motif (Exposito et al., 2010) in which the Xaa and Yaa
178 positions can be any amino acid except glycine and are usually occupied by proline and 4-
179 hydroxyprolines respectively (Bella et al., 1995). This highly ordered structure of collagen
180 results in its characteristic inextensibility and rigidity (Parvizi and Kim, 2010). The combination
181 of hydroxyproline, proline, and glycine accounts for approximately 57% of the total amino acid
182 makeup of collagen (Li and Wu, 2018).

183 **1.3.1.1 Hydroxyproline**

184 Hydroxyproline is an amino acid post-translationally derived from the hydroxylation of proline
185 by 4-prolyl hydroxylase (Cundy et al., 2014) with ascorbic acid being its cofactor (Stoilov et al.,
186 2018). At the Y-position, it forms hydrogen bonds with α chains within the triple helix thereby
187 stabilizing the trimeric structure. Hydroxyproline is not exclusive to collagen as it is also found
188 in elastin (Schmelzer et al., 2005), it uniquely constitutes approximately 14% of collagen's
189 overall imino and amino acid content (Neuman and Logan, 1950). It is therefore quantified as a
190 marker amino acid in collagen assays (Stoilov et al., 2018).

191 **1.3.2 Collagen nomenclature**

192 Roman numerals are used to number collagens in the sequence of their discovery (Myllyharju
193 and Kivirikko, 2004). Collagen molecules are composed of 3 left-handed alpha (α) chains
194 (polypeptide chains) identified with Arabic numerals. The type of collagen depends on whether
195 its component molecules are genetically distinct (heterotrimeric) or not (homotrimeric)
196 (Gjaltema and Bank, 2017); that is whether different genes express them or they are expressed by
197 a single gene (see Table 1.1).

198 This nomenclature identifies the individual α chains of the collagen molecule: $\alpha_n(N)_p$, where n
199 is the identifying number of the α chain, N is the roman numeral that indicates the collagen type
200 and p is the identifier for the polypeptide chain (Li and Wu, 2018; Hulmes, 2018). Thus, $\alpha_1(II)_3$
201 denotes a homotrimer of type II collagen consisting of 3 α_1 chains. The genes that encode
202 collagen alpha chains are prefixed by the letters COL, then the collagen type, followed by the
203 letter A-which means alpha chain and lastly the chain number (Table 1.1). For example,

204 *COL1A1* is the gene coding type 1 collagen alpha 1 chain while *COL1A2* is the gene coding for
205 type 1 collagen alpha 2 chains.

206 **1.3.3 Types of collagen**

207 Two main categories for grouping collagen types are fibrillar and non-fibrillar collagens (Table
208 1.1).

209 **1.3.3.1 Fibrillar collagens**

210 Fibril-forming collagens are the most abundant collagen types comprising about 90% of the total
211 collagens (Gelse et al., 2003). These collagens provide support, tensile and mechanical strength
212 to tissues and organs like skeletal muscles, skin, ligaments, and tendons in the mammalian body
213 (Kaushal et al., 2014). The regions of the gene that encode their triple helical chains have almost
214 identical exon and intron structures for all mammalian fibrillar collagen genes (Sutmuller et al.,
215 1997) which may indicate common evolutionary pressures and ancestry at the level of the gene
216 (Takahara et al., 1995). Collagens molecules in this category tend to be staggered (quarter-
217 staggered) with respect to neighboring molecules by integral multiples of D, where $D=234\pm 1$
218 amino acid residues or 67nm (Hulmes et al., 1973; Meet et al., 1979; Hulmes, 2008) known as
219 the Hodge-Petruska fibrillar collagen model (Petruska and Hodge, 1964) resulting in the banded
220 appearance of collagen fibrils (Kaushal et al., 2014) (Figure 1.2). Collagens types forming
221 quarter staggered fibrillar structures include collagen types I, II, III, V, XI, XXIV, and XXVII
222 (Table 1.1). This structure provides fibrillar collagen with its rope-like structure, conferring
223 substantial strength to connective tissues in which it is found.

224 **1.3.3.1.1 Some fibrillar collagens in skeletal muscles**

225 **1.3.3.1.1.1 Collagen type I**

226 Type I collagen is the most abundant and widely distributed fibrillar collagen in mammalian
227 species, and it provides strength and stability to a wide array of tissues (Fratzl, 2008; Kaushal et
228 al., 2014). Its precursor, type I procollagen is synthesized in the rough endoplasmic reticulum by
229 the combination of 1 pro α 2(I) peptide chain and two pro α 1(I) to form a triple helix with these α
230 chains initially interacting at their carboxyl-terminal propeptides (Cundy et al., 2014). It
231 constitutes about 90% of the protein in the bone matrix (Cundy et al., 2004; Gelse et al., 2003)
232 and it is also in tissues of the vasculature, lungs, skin, tendons, cornea (Humes, 2008). Thomas et
233 al. (2000) reported significant decreases in collagen type I and III mRNA expression with
234 increasing age in mice left ventricle. This observation implies that most likely, it is not the
235 synthesis of new collagen molecules that form heat stable crosslinks but the transformation of
236 existing heat labile ones. Studies by Light and Champion (1984) showed that type I collagen is
237 the major component of the epimysium and perimysium of skeletal muscles where it supports the
238 structural integrity of the muscle and thus an important component for muscle physiology.

239 **1.3.3.1.1.2 Collagen type III**

240 This collagen type along with type 1 collagen forms a significant component of the interstitial
241 matrix (Nielsen and Karsdal, 2016). Collagen type III was found to be a minor component of the
242 epimysium and perimysium (Light and Champion, 1984). Apart from its role in structural
243 support, it also acts as a ligand for numerous proteins including von Willebrand factor, G
244 protein-coupled receptor-56 and integrin α 2 β 1 (Nielsen and Karsdal, 2016). It can also be found
245 in comparatively elastic mammalian tissues like the lungs, blood vessels and the skin of an
246 embryo (Hulmes, 2008). Fugii et al. (1976) found lysinonorleucine to be the predominant

247 crosslink in fibrils of collagen type III while collagen type I mainly formed
248 hydroxylysinoxidation crosslinks. Shuttleworth and Forrest (1975) observed much higher
249 collagen type III-specific activity than collagen type I in guinea pig fetal dermal tissues although
250 it declined dramatically with increasing fetal age until the specific activity of type I collagen
251 surpassed that of Type III at day 70 of gestation. Based on these observations, they concluded
252 that maximum synthesis of collagen type III occurs at the fetal stage.

253 **1.3.3.1.3 Collagen type V**

254 This collagen type is widely distributed in various tissues of the body and is critical to the
255 formation of collagen type I and III tissue-specific matrices (fibrillary formation) (Leeming and
256 Karsdal, 2016). Indeed, Light and Champion (1984) found traces of this collagen type in the
257 perimysium. It supports the integrity of interstitial muscle matrices, corneal stroma, and bone
258 matrices. It forms heterotypic fibrils with type I collagen especially in the cornea (Birk et al.,
259 1988). Birk et al. (1986) have chronicled a relationship between type V and type I in which
260 quantitatively increasing type V collagen ratio relative to type I decreased the ultimate collagen
261 fibril diameter in heterotypic corneal fibrils. The relationship between type I and V collagens
262 illustrates that some collagens, like type V, exist in muscle to cooperatively modify the
263 quaternary structure of other collagens.

264 **1.3.3.2 Non-fibrillar Collagens**

265 Non-fibrillar collagens are predominantly lattice-forming collagens which tend to be important
266 structural components of all basement membranes like Type IV collagen (Ricard-Blum, 2011).
267 Their triple helical chains are of varying lengths, interrupted by non-helical sections (Kaushal et
268 al., 2014). This category of collagen includes collagens with multiple triple helical domains with

269 interruptions (multiplexins)(Types XV and XVIII), basement membrane collagens (Types XIII,
270 XVII, XXIII, and XXV), network collagens (Types IV, VIII, and X), and fibril-associated
271 collagens with interrupted triple helices (Types IX, XII, XIV, XVI, XIX, XX, XXI and
272 XXII)(Ricard-Blum, 2011).

273 **1.3.3.2.1 Some non-fibrillar collagens in skeletal muscles**

274 **1.3.3.2.1.1 Collagen type IV**

275 This structural collagen is regarded as the collagen type most vital to the integrity of muscle fiber
276 basement membranes (Sand et al., 2016). It possesses several discontinuities in its Gly-Xaa-Yaa
277 motifs due to the presence of non-collagenous sections (Dolz et al., 1988; Hulmes, 2008).
278 Collagen type IV exists in all known metazoans (Sundaramoorthy et al., 2002) especially in their
279 glomerular basement membranes (Hulmes, 2008). The C-terminal non-collagenous domains of
280 type IV collagen prevent angiogenesis (Petitclerc et al., 2000). Expression of this collagen type
281 has been reported (Feru et al., 2016) to decrease with age in some tissues while transforming
282 growth factor beta 1 (TGF- β 1) treatment causes a significant rise in its expression in fibroblasts
283 indicating a possible influence of TGF- β 1 on collagen IV expression and indirectly, basement
284 membrane integrity. Future studies may need to look at the effect of these hormonal treatments
285 on the expression and synthesis of collagen molecules in livestock species as manipulation of
286 these hormones could lead to a lower collagen synthesis to turnover ratio and probably a
287 weakened muscle structure.

288 **1.3.3.1.1.2 Collagen type VI**

289 Tetramers of this collagen type aggregate to form beaded filament structures in most
290 extracellular matrices as it forms microfibrillar networks between the interstitial matrix and

291 basement membrane interfaces (Sun and Karsdal, 2016). In humans, mutations in its alpha chain
292 coding genes lead to muscular defects like Bethlem myopathy and Ullrich disease (Lampe and
293 Bushley, 2005). Bonaldo et al. (1998) in their study of the role of collagen VI in muscle fiber
294 integrity found that *COL6A1* gene knockout mice exhibited pronounced muscle fiber
295 phagocytosis and necrosis as well as fiber diameter variations. These myopathic characteristics
296 suggest that collagen VI is crucial to muscle fiber integrity.

297 **1.3.3.1.3 Collagen type XXI**

298 In the extracellular matrix, collagen XXI acts as molecular bridges by enhancing protein-protein
299 interactions using their non-collagen N-terminal domains (Kehlet and Karsdal, 2016). The
300 marked expression of *COL21A1* in smooth muscle cells of the aorta after introducing platelet-
301 derived growth factor led Cho and Li (2002) to suggest that it may be involved in extracellular
302 matrix arrangement during formation of blood vessels. Furthermore, its co-expression with type
303 1 collagen in skeletal muscles may play an important role in interstitial collagen fibril
304 organization (Fitzgerald and Bateman, 2001) and thus an important collagen for extracellular
305 matrix integrity.

306 **1.3.4 Collagen biosynthesis**

307 All proteins belonging to the collagen superfamily are secretory proteins that are synthesized on
308 bound ribosomes (protein synthesis site in cells), secreted into the endoplasmic reticulum lumen
309 for post-translational modifications, transported to the Golgi apparatus for further modifications
310 and finally secreted into the extracellular space (Myllaharju, 2005; Koide and Nagata, 2005;
311 Cundy et al., 2014). Figures 1.3 and 1.4 depict the steps involved in collagen synthesis and
312 subsequent post-translational modifications.

313 **1.3.4.1 Ribosomal stage**

314 The genes coding for their amino acid sequences essentially control the basic structure/template
315 of collagens (Di Pasquale, 1997). In the ribosome, triplet codons for particular amino acids on
316 the mRNA are recognized and read by the anti-codon on the cognate amino-acyl tRNA (Zaher
317 and Green, 2009). This process occurs at the 80S when the 40S and 60S subunits combine to
318 initiate translation. Uncharged amino acids, at the onset, interact with Mg^{2+} and ATP forming an
319 amino acid adenylate (enzyme bound) catalyzed by amino acyl-tRNA synthetase (Coulson,
320 1994). This aminoacylation guarantees that only matching amino acids bind to the tRNA's. A
321 complex composed of amino-acyl tRNA, guanosine triphosphate and eEF1A elongation factor is
322 conveyed to the aminoacyl site. This ternary complex interacts with peptidyl tRNA at the
323 peptidyl site, elongating the forming polypeptides (Zaher and Green, 2009). Synthesis of the
324 nascent polypeptide is terminated when a stop codon at the Aminoacyl site is detected by
325 eukaryotic translation termination factor 1.

326 In summary, polysomes control the inclusion of amino acyl groups to the increasing peptidyl
327 tRNA forming peptide bonds through condensation reactions (Coulson, 1994). Pre-procollagen,
328 a protein precursor, translated from mRNAs on the ribosomes is formed.

329 **1.3.4.2 Endoplasmic reticulum stage**

330 Pre-procollagen, a newly formed polypeptide chain, is synthesized with a hydrophobic leader
331 peptide that enhances the binding of ribosomes to the endoplasmic reticulum and directs the
332 lengthening polypeptide chain (pre-procollagen) into the lumen of the endoplasmic reticulum
333 (Kaushal et al., 2014). Thus, the polypeptide chains undergo co-translational translocation across
334 the endoplasmic reticulum lumen (Bullied et al. 1997). Initiation of post-translational

335 modifications of pre-procollagen occurs upon cleaving of the leader peptide in the endoplasmic
336 reticulum coupled with the addition of hydroxyl groups to lysine and proline residues (Kaushal et
337 al., 2014).

338 Proline residues at the Y-position are mostly modified to 4-hydroxyproline by prolyl-4-
339 hydroxylase (Cundy et al., 2004). When proline residues are not fully hydroxylated, collagen
340 molecules tend to have comparatively lower thermal stability as well as abnormal assemblage
341 (Gjaltema and Bank, 2017) indicating the importance of this modification in the stability of the
342 whole structure. Lysyl hydroxylase catalyzes the conversion of lysine residues into 5-
343 hydroxylysine on the individual α chains of a procollagen molecule (Gjaltema and Bank, 2017).
344 Proline residues in the X-positions convert to 3-hydroxyproline catalyzed by prolyl-3-
345 hydroxylases (Koide and Nagata, 2005). Prolyl-4-hydroxylases, lysyl hydroxylases, and prolyl-
346 3-hydroxylases belong to the alpha-ketoglutarate-dependent hydroxylase enzyme family and thus
347 require oxygen, vitamin C (ascorbate), 2-oxoglutarate and iron (Fe^{2+}) to perform their functions
348 (Gjaltema and Bank, 2017). The hydroxylation of these residues is a necessary modification
349 leading towards the formation of intermolecular crosslinks that are stable.

350 During the actual triple helix formation, the alpha chain association is initiated at the carboxyl
351 (C)-terminal propeptide propagating the triple helix formation in a zipper-like folding to the
352 amino (N)-termini (Engel and Prockop, 1991). The formation of the triple helix is completely
353 driven by the initial association of the alpha chains through their C-terminal poly-peptides, as
354 Bullied et al. (1997) found that helix formation was not initiated among monomeric chains
355 lacking the carboxyl-terminal propeptides.

356 Proper trimerization and folding of collagen are achieved through the assistance of various
357 enzymes (Koide and Nagata, 2005). Involved enzymes include protein-disulfide isomerase,
358 which catalyzes the formation of inter and intrachain disulfide bonds at the carboxyl-terminal
359 region, allowing for the folding and association of peptide chains into a triple helical structure
360 (Okumura et al., 2015; Cundy et al., 2014). Prolyl peptidyl cis-trans isomerase B accelerates the
361 folding around prolyl residues by catalyzing isomerization of cis-peptide bonds in pro α -chains
362 into trans-peptide bonds (Cundy et al., 2014). Lysyl hydroxylases-2 aids the hydroxylation of
363 lysine residues at the exterior of the helix (telopeptides) to 5-hydroxylysine. Some lysine
364 residues in the Y-position inside the triple helical domain undergo hydroxylation catalyzed by
365 the enzyme lysyl hydroxylase-1. Galactosyltransferase catalyzes the occurrence of O-linked
366 glycosylation by adding galactosyl residues to hydroxylysine. Glucose groups are further added
367 to the galactosyl hydroxylysine to form a disaccharide catalyzed by glucosyltransferase.
368 Glycosylation occurs only on non-collagenous domains by these enzymes (Kaushal et al., 2014).
369 In the non-fibrillar domains, oligosaccharides are added (N-linked glycosylation) on specific
370 asparagine residues. Non-fibrillar collagens tend to be more glycosylated compared to their
371 fibrillar counterparts most likely due to the higher occurrence of non-helical domains in them.
372 These hydroxylated residues are vital in the formation of covalent crosslinks and subsequent
373 thermal stability (Gjatema and Bank, 2017; Cundy et al., 2014).

374 The multiple enzymes involved mean that the formation of collagen structures is a metabolically
375 intensive process, with ample potential for error. As a result, chaperone proteins are essential in
376 the processes leading to collagen formation (Koide and Nagata, 2005). These include the heat
377 shock protein 47 (HSP 47), a glycoprotein found to aid triple helix propagation by binding
378 procollagen and to prevent collagen aggregation in the endoplasmic reticulum (Nagata and

379 Yamada, 1986). FK-binding protein 65 (FKBP 65) is a chaperone that has been found to be
380 involved in procollagen synthesis and regulation (Knüppel et al., 2018) and mutation in this
381 protein has been reported to cause a decline in extracellular matrix rigidity (Staab-Weijnitz et al.,
382 2015).

383 **1.3.4.3 Golgi apparatus stage**

384 After triple helix formation in the rough endoplasmic reticulum, the procollagen molecule is
385 transported through the Golgi apparatus (Koide and Nagata, 2005). Here, it undergoes packaging
386 into cylindrical aggregates in membrane-bound secretory organelles (Myllyharju, 2005). At this
387 stage, the initial formation of fibrils occurs in the case of fibril-forming collagens (Cundy et al.,
388 2014). These secretory organelles are then conveyed into the extracellular space via exocytosis.
389 During the secretion of these procollagen molecules, further modifications occur as specific C-
390 and N- terminal procollagen proteinases cleave off the non-helical domains of the molecule
391 (Prockop et al., 1998). Disintegrin and metalloproteinase with thrombospondin motifs 2
392 (ADAMTS-2), also known as procollagen N-proteinase, cleave the N-terminal propeptides while
393 bone morphogenetic protein 1 (BMP 1) also called procollagen C-proteinase cleaves the C-
394 terminal propeptides (Prockop et al., 1998). The resulting molecule is termed tropocollagen.

395 **1.3.4.4 Extracellular stage**

396 In the extracellular space, tropocollagen trimers are assembled into collagen fibers and fibrils.
397 Intermolecular aldehyde-derived crosslinks stabilize these structures. Also, oxidative
398 deamination of the amino groups from some hydroxylysine and lysine residues side chains,
399 catalyzed by lysyl oxidase results in the production of hydroxyallysine and allysine (reactive
400 aldehyde derivatives) (Fujii et al., 1976). Subsequent formation of aldol condensation products

401 by these aldehyde groups results in crosslinks between and within the collagen molecules. Imine
402 (Schiff base) crosslinks may also be formed when these aldehyde groups react with amino
403 groups from unoxidized hydroxylysine and lysine residues. Stable cross-links like
404 lysinonorleucine may be formed from the rearrangement, reduction or dehydration of the above
405 products. Other non-reducible cross-links include lysylpyridinoline and
406 hydroxylysylpyridinoline. Lysylpyridinoline crosslinks are composed of 2-hydroxyallysine
407 residues in the telopeptide section and a specific lysine residue in the helical domain while
408 hydroxylysylpyridinoline is made from 2 hydroxyallysine residues in the telopeptide region and
409 a specific 5-hydroxylysine in the triple helical domain (Cundy et al., 2014). Research on the
410 organic compound β -aminopropionitrile, which inhibits the activity of lysyl oxidase, strongly
411 suggests that the formation of collagen crosslinks is important for tissue strength and mechanical
412 properties as a significant reduction in pyridinoline cross-link content and bone strength was
413 observed after treating mice with β -aminopropionitrile (McNerny et al., 2015). Thus, reducibility
414 and density of collagen crosslinks likely affect meat toughness. Ehrlich chromogen is another
415 crosslink proposed by Scott et al. (1983). It is a trifunctional pyrrolic crosslink derived from a
416 Knorr reaction between amino-ketones to produce pyrroles (Scott et al., 1981).

417 **1.4 Pork quality**

418 **1.4.1 Definition of pork quality**

419 Pork quality is dictated mainly by muscle metabolism peri- and post-mortem and is dependent on
420 a pig's genetics and environmental factors (Lebret, 2008; Enfält et al., 1997). Some of these
421 environmental factors include management practices, animal nutrition, slaughter practices and
422 carcass management (Salas and Mingala, 2017). Pork quality is defined by Hoffman (1973) as
423 encompassing all the toxicological, technological, food physiological and sensorial attributes of

424 pork (Figure 1.5). A similar definition by Ciobanu et al. (2011) described pork quality as
425 consisting of a class of sensorial and meat processing traits critical for competition and profit-
426 making by the pork industry in the ever-dynamic food sector. Sensorial traits were explained to
427 include eating quality and appearance as assessed by sensory panelists while meat processing
428 traits included cooking loss, water-holding capacity, color, and firmness. Further, a typical meat
429 scientist according to Bray (1966) will define the quality of fresh meat to include the
430 components/aspects linked to the palatability of cured and fresh meat products as well as the
431 associated financial losses incurred during its processing and subsequent distribution.

432 From the above definitions, it is clear that different definitions exist in the pork industry to
433 describe quality. However, the one that should carry the most weight is that one given by
434 consumers. The general view on meat quality of consumers is summed up as meat products that
435 are wholesome, safe, nutritious, aesthetically pleasing, worth its cost and produced with
436 procedures that are morally admissible (Eikenboom, 1983 as cited by Smulders, 1986). The
437 perceptible quality factors are especially of economic significance as consumers experience these
438 themselves (Mir et al., 2017).

439 **1.4.1.1 Meat quality versus carcass quality**

440 As is often the case, people tend to confuse the term carcass quality with meat quality. However,
441 there are some distinctions between those two terms (Smulders, 1986). Carcass quality usually
442 subsumes factors that critically affect the market price of carcasses, and some of these factors are
443 carcass fat cover, bone to meat ratio and carcass weight (Smulders, 1986). In most countries,
444 carcass grading policies, pricing, and procedures are designed in such a way as to allow carcass
445 quality characteristics/attributes to, for the most part, harmonize with an improved meat quality

446 derived from the carcass (Smulders, 1986). However, there are numerous factors affecting meat
447 quality and these determinants justify the distinction between carcass and meat quality.

448 Different players in the food sector such as consumers, microbiologists, nutritionists,
449 veterinarians, and meat technologist, at times, append similar weights to some subcomponents of
450 meat quality. However, now and then, their interests conflict (Smulders, 1986). A classic
451 example is that of a dark-cutting (dark, firm and dry) meat. The microbiologist will consider
452 such meat as a possible hazard due to its unusually high pH; however, the meat technologist or
453 processor, realizing that such high pH will likely equate to outstanding water holding capacity
454 will see such meat as particularly suitable for certain sausage types.

455 **1.5 Eating quality**

456 A major aim of the current pork industry is to offer end-products that will achieve consumer
457 satisfaction at equitable costs (Drescher et al., 2012). From their principal component analysis,
458 Dransfield et al. (1984) reported that overall acceptability of bovine longissimus muscle steaks
459 was determined primarily by juiciness, flavor, and tenderness. Similarly, eating quality attributes
460 of pork are described by consumers with respect to its juiciness, tenderness, and flavor (Aaslyng
461 et al. 2003). It is important therefore to understand how the characteristics of pork affect pork
462 juiciness, tenderness and flavor.

463 **1.5.1 Juiciness**

464 Juiciness, a subjective sensory trait (Warner, 2017) is one of the most significant factors
465 influencing meat palatability (Tannor et al., 1943). It is the sensation of lubrication and wetness
466 (moisture) experienced during mastication of meat (Warner, 2017). Thus, it provides an
467 indication of the amount and level of moisture in meat that is released during chewing (Heymann

468 et al., 1990). Juiciness of meat can be divided into two aspects: the sensation of wetness
469 perceived during initial mastication due to the expression of meat fluids; and the perceived
470 juiciness experienced during sustained mastication. The latter aspect may be related to the
471 stimulating influence of muscle fats on salivation (Winger and Hagyard, 1994; Aaslyng et al.,
472 2003).

473 Well-marbled pork gives a better impression of juiciness and tenderness than pork devoid of
474 marbling (Thompson, 2004). This could be due to the ECM remodeling theory proposed by
475 Nishimura (2010) in which deposition of fat in the perimysium results in a remodeling of the
476 extracellular matrix leading changes in the mechanical properties (weakening) of the
477 intramuscular connective tissue. Fernandez et al. (1999) found that rising levels of intramuscular
478 fat were significantly linked to juiciness in porcine Longissimus muscles. Kempster et al. (1986)
479 made similar observations, although in their study the level of leanness did not affect overall
480 product acceptability. These studies indicated that there is an intimate relationship between
481 intramuscular fat content and juiciness in meat. Wood et al. (2004) found that feeding intact male
482 pigs low protein diets of about 16% crude protein content with higher lysine to energy ratio
483 resulted in a much juicier and tender longissimus muscle compared to a high protein diet of 20%
484 crude protein and these results showed that certain quality traits can be manipulated through
485 animal nutrition.

486 Swine breed has also been shown to affect pork sensory quality. Cameron et al. (1990) in
487 comparing the British Landrace with Duroc found that pork from Durocs had higher scores for
488 juiciness than Landrace and this was subsequently attributed to the greater levels of
489 intramuscular fat in the Duroc pork compared to that of the Landrace. Ventanas et al. (2007) also
490 reported that purebred Iberian pigs had higher scores for juiciness than Duroc × Iberian pigs in

491 their study on dry-cured loins. Serra et al. (1998) found a positive correlation between
492 intramuscular fat percent and type I muscles fibers. In that study, Serra et al. (1998) compared
493 quality and histochemical characteristics between Landrace and Iberian pigs and found the latter
494 to have twice the backfat depth and 5 times the intramuscular fat percent and a significantly
495 higher proportion of oxidative type I fibers. Similar results were obtained by Lee et al. (2012) as
496 their results showed that breeds with higher fat content like Berkshires and Duroc also had
497 higher Type I and lower Type IIB muscle fibers compared to the Landrace and Yorkshire who
498 had comparative lower fat content. Thus, a plausible underlying driving factor for these breed
499 differences in juiciness is most likely muscle fibre type (oxidative vs glycolytic). Interestingly,
500 this trend may not be species specific as it has also been reported in Hanwoo beef steers by Joo
501 et al. (2017). In their study, they found a strong positive correlation between intramuscular fat
502 content and the proportions of Type I muscle fibers but a negative correlation of similar
503 magnitude with the proportion of Type IIB muscle fibers.

504 A relationship between ultimate pH and juiciness has also been established by Eikelenboom et
505 al. (1996), who reported a positive correlation between these traits and subsequently suggested
506 that selection of carcasses for juiciness should be based on ultimate pH. Jonsall et al. (2001)
507 reported juicier pork from pigs reared indoors compared outdoors which may have been due to
508 the outdoor pigs requiring more energy for locomotion and maintaining body temperature thus
509 resulting in less fat deposition. Channon et al. (2016) reported higher juiciness scores for
510 castrates compared to intact male pigs. This is not surprising as castrates are known to have a
511 higher propensity for fat deposition (Bonneau, 1998). The presence or absence of testosterone is
512 the main difference between intact male pigs and castrates. Indeed, low testosterone levels have
513 been associated with metabolic syndrome and obesity (Wang et al., 2011). Wang et al. (2017)

514 speculated that the *MARK4* gene might play a vital role in obesity as their results showed that fat
515 accretion in castrates may be regulated by ssc-miR-7134-3p by acting as a translational inhibitor
516 of *MARK4*. Thus, castrates had lower levels of ssc-miR-7134-3p and increased MARK4 protein
517 levels in their fat tissues than intact males. Muscle fiber type may also affect juiciness, as Shen et
518 al. (2005) found that pork with proportionally more glycolytic muscle fibers had a reduced pH
519 and light colour, which is associated with pale, soft and exudative pork (Lee et al., 2012).

520 End-point temperature has also been found to significantly influence juiciness ratings, as higher
521 juiciness ratings have been reported (Channon et al., 2016) for pork cooked to an internal end-
522 point temperature of 70°C relative to those cooked to 75°C. This is probably because with
523 increasing temperature comes a concomitant increase in loss of fluids and volatile fatty acids
524 which may decrease the sensation of juiciness during mastication. Sheard et al. (1999) reported
525 significantly improved juiciness in polyphosphate-injected pork loins. This is probably because
526 phosphates are known to increase meat pH and thus the water holding capacity of meat (Hamm,
527 1961). All the above studies show that animal genetics, management practices and processing
528 techniques can be manipulated to affect meat juiciness.

529 **1.5.2 Tenderness**

530 Tenderness is considered the most important eating quality trait influencing consumer
531 satisfaction and acceptance (Suman, 2012; Williams., 1986). It can be defined as the ease with
532 which meat fibers are cut through by the teeth during chewing (Coggins, 2012), thus simulating
533 the effort required to masticate. In general, three methods have been used to evaluate the
534 differences in meat tenderness. The first method is sensory evaluation in which trained panelists
535 are used to describe the meat product of interest under standardized conditions. Another method
536 employed by some researchers to evaluate tenderness is chemical analysis through the

537 determination of elastin and collagen contents of meat samples. A third method widely accepted
538 in most sectors in the food industry is the mechanical method, that is, an instrumental
539 measurement which assesses the shear force strength or resistance to shearing of a meat product
540 (Bratzler, 1930). Dissatisfaction among consumers could arise due to variations in the tenderness
541 of meat products (Mane et al., 2014). As illustrated in Figure 1.6 numerous factors affect pork
542 tenderness. Some of these factors are highlighted below. The two most important groups of
543 proteins determining tenderness are connective tissue proteins and myofibrillar proteins (Mane et
544 al., 2014; Bouton et al., 1974).

545 In cooked meats (Purslow, 2018), connective tissues are thought to contribute to an immutable
546 background toughness, although they are still implicated in muscle to muscle and animal to
547 animal variation while myofibrillar contraction state influences the sensation of toughness for the
548 most part (Bouton et al., 1974; Huff-Lonergan et al., 2010; Sikes et al., 2010). The statement
549 about the immutable background role of collagen is still subject to some debate as discussed later
550 in section 1.5.5.

551 The mean adhesion value of meat cooked to a temperature of 50°C according to Bouton et al.
552 (1974) was approximately five times that cooked at 90°C for 1hr. Because these values represent
553 the strength of connective tissues among the meat fibers, these results suggest a major influence
554 of connective tissues at low temperatures. Williams et al. (1986) reported a rise in Z-disc
555 fragmentation and a decrease in sarcomere length in poultry breast muscle cooked to an internal
556 temperature of 80°C. They further observed during their sensory tests that samples with longer
557 sarcomere lengths scored as more tender, adding to the well-established effect of sarcomere
558 length on meat toughness. Feldhusen and Kuhne, (1992) observed an increase in sarcomere
559 length and tenderness of pork loins after aging for two days. Channon et al. (2004) further looked

560 at a greater period of 7 days with vacuum bags and found that pork loins aged for that length of
561 time were significantly more tender than those aged for two days. Wood et al. (1996) also
562 reported an increase in myofibrillar fragmentation index with increasing aging time. These
563 results suggest that as aging period increases, so too does tenderness probably due to an increase
564 in proteolytic enzyme activity.

565 Guzek et al. (2016) reported no significant differences in Longissimus lumborum shear force
566 values and sarcomere lengths for crossbred Polish Landrace× Duroc and purebred Landrace pigs.
567 Wood et al. (1996) also reported no difference in tenderness between the Duroc and Large White
568 pigs. These results partially question the influence of breed on these pork quality traits. Carlson
569 et al. (2017) implicated proteolytic degradation and lipid content in the significant variations they
570 observed in their instrumental meat tenderness measurements, as the groups of samples
571 determined as tender tended to have high lipid and marbling scores as well as significantly
572 greater degradation of filamin, troponin, and desmin. Wheeler et al. (2000) stated that enzymatic
573 proteolysis, sarcomere length, variations in collagen content and the extent of their interactions
574 dictates individual pork muscle tenderness.

575 **1.5.3 Flavor**

576 For us to generate the numerous perceptual responses of food during mastication, numerous
577 chemicals are detected and then turned into nerve impulses (Reineccius, 2016). Flavor according
578 to Lawrie (1966) is essentially a complex sensation experienced by humans that involves taste,
579 pH, temperature, texture, and odor. It is regarded as one of the factors affecting consumer
580 purchasing decisions (Spanier et al., 2004).

581 Flavor formation and subsequent deterioration in meat products are said to be a process that is
582 continuous with respect to the formation and subsequent deterioration of some desirable flavor
583 compounds as well as the development of some off-flavor components (Spanier et al., 1992).
584 Formation of these flavor generating components has been reported to be correlated with some
585 factors like end-point cooking temperature, post cooking storage and aging period postmortem
586 (Yang et al., 2016; Cambero et al., 1992; Spanier et al., 2004). Factors influencing pork flavor
587 are illustrated in Figure 1.7 and indicate the complexity of flavor development.

588 Some descriptors used by sensory analysts to describe pork flavor include metallic, serum,
589 musty, boar odor, bloody, liver-like, porky, piggy, cardboard, rancid, briny, fatty and browned
590 (Hayes, 2010). More than 1,000 volatile compounds that dictate aroma characteristics and meat
591 flavor have been isolated from cooked meats (Mottram, 1994). Maillard reactions between some
592 reducing sugars and amino acids result in the formation of aroma volatiles during cooking
593 (Mottram, 1994). This discovery was originally made by Morton et al. (1960) after noticing that
594 at elevated temperatures, reacting cysteine with pentose monosaccharides like ribose in the
595 presence of water led to the production of meat flavors and the author subsequently patented this
596 discovery. Some other reactions that may occur during cooking of meat and which could
597 ultimately result in flavor development are caramelization of some carbohydrates, thiamine and
598 ribonucleotide degradation, peptides and amino acids pyrolysis, and lipid degradation (MacLeod
599 and Seyyedain-Ardebili, 1981).

600 Hornstein and Crowe (1960) in their research on pork and beef flavor reported that odor
601 characteristics of heated lean pork and beef aqueous extracts were identical as sensory panelist
602 could not differentiate the meats of those two species. Further, they observed a striking
603 resemblance in the spectrophotometric and gas chromatographic behaviors of their compound

604 isolates. However, flavor difference existed in their heated fat portions, producing species-
605 specific flavors. Indeed, Fernandez et al. (1999) found that flavor scores increased with
606 increasing intramuscular fat content. Wood et al. (1996) reported higher flavor intensity ratings
607 for Duroc pigs compared to Large Whites which was not surprising as they also observed higher
608 marbling fat content in the Durocs.

609 Sheard et al. (1999) found that injection of polyphosphates in porcine longissimus muscle
610 reduced flavor intensity. This is not surprising as these polyphosphates are known to have water-
611 binding properties and therefore may increase the water ratio of the muscle leading to a decrease
612 in the concentrations of flavor compounds. It was also not surprising that after heating their
613 samples to an internal temperature of 80°C, Sheard et al. (1999) recorded an increase in flavor
614 intensity probably because evaporation of moisture might have caused an increase in flavor
615 compound concentration suggesting a possible negative relationship between increasing moisture
616 content and flavor.

617 **1.5.4 Intramuscular fat and pork eating quality**

618 Pork intramuscular fat, in general, has been known to influence the three cardinal eating quality
619 traits, specifically juiciness, tenderness, and flavor (Smith and Carpenter, 1974). The notion that
620 fat deposition improves the worth of meat has been recognized since the beginning of written
621 history (Smith and Carpenter, 1974). This belief that fatness is positively correlated with
622 palatability and eating quality of meat products led famous 18th-century British animal breeding
623 pioneers like Robert Bakewell to improve a number of livestock species during the agricultural
624 revolution of the mid-18th century (Philips, 2001). For instance, Robert Bakewell transformed the
625 Leicestershire sheep, a previously heavily boned breed with little disposition to fatten into a fast
626 fattening, fine-boned breed also known as the Bakewell Leicester (Bulliet, 2005). It may

627 however be worth point out that during that period, many workers expended lots of energy on
628 tasks presently performed by machines and as fat has comparatively larger amounts of stored
629 energy than lean, the meat of those times supplied a significant proportion of their daily energy
630 requirements (Swatland, 1994).

631 Hall (1910) observed that marbling, in addition to its contribution to juiciness and flavor, also
632 increased meat tenderness, through a reduction in connective tissue elasticity as a result of the fat
633 deposition in it. Nishimura et al. (1999) studied the relationship between marbling and beef
634 tenderness in Japanese black cattle, and these authors reported that shear force values obtained
635 from the Longissimus lumborum muscle increased by about 50N between 9 to 20 months of age
636 but saw a reduction of about 45N from 20 months to 32 months of age. Interestingly, this is the
637 final fattening period of those animals. They, therefore, concluded that increased fat deposition
638 might increase affect tenderness by altering intramuscular connective tissue structures in breeds
639 with predispositions to increased marbling.

640 **1.5.5 Collagen and pork eating quality**

641 Light et al. (1985) proposed two theories of the impact of collagen on meat toughness and
642 texture. Firstly, they proposed that shrinkage of collagen fibers in the connective tissues results
643 in muscle bundle compression and moisture loss which ultimately affects texture. Secondly, they
644 proposed that the amount of insoluble cross-linked collagen which could offer some resistance to
645 shear force could also affect meat toughness. Researchers have identified two major
646 consequences of temperature (heat) on collagen characteristics intimately linked to the theories
647 proposed by Light et al. (1985) as these effects can significantly affect cooked meat texture.
648 Firstly, at approximately 65°C (shrinkage temperature of mammalian collagen), collagen fibers
649 undergo significant denaturation resulting in the collapse of its structure, leading to shrinkage by

650 as much as one fourth its original length (Bailey and Light, 1989) and this ultimately results in
651 toughening of the meat (Voyle, 1979). Secondly, most collagen fibers solubilize (Figure 1.8) at
652 temperatures around 80°C. However, the degree of solubilization is dependent upon the
653 proportions of heat resistant intramolecular and intermolecular crosslinking (Hill, 1966) resulting
654 in lower intrafibrillar water content and this decreased hydration leads to an increase in thermal
655 stability (Miles et al., 2005). Research by Purslow et al. (2016) however suggested that collagen
656 fiber shrinkage may not be the cause of fluid losses in cooked meat. After comparing the
657 dimensional variations in muscles with intact perimysium and endomysium with those with only
658 endomysium and those without connective tissues, they attributed transverse shrinkage at 50-
659 65°C to myosin denaturation while actin denaturation caused longitudinal shrinkage at 70-75°C.
660 However, comparative analysis of fluid or weight loss between whole meat blocks and
661 myofibrils devoid of connective tissue was not provided by Purslow et al. (2016); this
662 information however would have been very helpful for drawing concrete conclusions.
663 Furthermore, Purslow et al. (2016) premised the hypothesis of their study on research by
664 Tornberg (2005) who compared cooking loss (%) between whole meat, hamburger and
665 emulsion-type sausages. It would have been more informative for Purslow et al. (2016) to
666 indicate that for all the temperatures examined by Tornberg (2005), whole meat had the highest
667 mean cooking loss (%) of the 3 products and it was especially significant at 65°C, the
668 temperature at which collagen fiber shrinkage occurs. In fact, addition of salt (0.4 and 4% for the
669 hamburger and emulsion-type sausage respectively) and comminution of meat leads to meat
670 protein extraction which form protein gel networks upon heat treatment, which is what would be
671 expected to occur in emulsion-type sausages and to an extent, hamburger patties and thus one
672 would expect water holding capacity of hamburgers to be higher than that of whole meat. The

673 evidence suggests that fluid losses may be driven by connective tissue shrinkage up to 65°C
674 above which shrinkage of muscle proteins dominates, as connective tissue fibers will start
675 gelatinizing at higher temperatures. Future studies should look at whether the water released is
676 bound or loose by comparing the shrinkage of connective tissue-bound muscle fibers with
677 muscle fibers devoid of connective tissue when heated.

678 There is disagreement on the effect of the presence and level of collagen crosslinking on cooked
679 meat texture. Avery et al. (1996) concluded that perimysial collagen crosslinks were not related
680 to cooked meat texture obtained either subjectively or objectively. Ngapo et al. (2002) also found
681 a weak and insignificant relationship between crosslink amounts and shear force values of
682 cooked meat. In contrast, researchers like Jeremiah et al. (2003) reported that the amount of
683 insoluble collagen negatively affected meat texture. Dransfield (1997) also chronicled a
684 relationship between muscle toughness and total collagen content especially at lower
685 temperatures (60°C for 20 mins).

686 **1.5.5.1 Animal age**

687 Irrespective of species and muscle type, the amount of heat-soluble, reducible collagen crosslinks
688 decreases with increasing age (Figure 1.9) (Shimokomaki et al., 1972); implying that these
689 reducible cross-links may be intermediates which with time most likely become stabilized during
690 the crosslinking (Bailey and Shimokomaki, 1971). Indeed, Goldspink et al. (1994) reported an
691 age-induced decline in collagen expression in skeletal muscles and Khorramiza et al. (1999)
692 further reported an increase in collagenase mRNA expression in older fibroblasts.

693 A greater amount of collagen is solubilized in cooked meat from young animals compared to
694 matured ones, and this, according to Hill (1966) is most likely a major contributing factor to the
695 comparatively increased sensation of tenderness experienced during mastication of meat

696 products from young animals. The decrease in solubility that occurs with increasing age (Figure
697 1.9) is most likely due to the decline in new collagen synthesis, thus lending time for the heat-
698 soluble crosslinks to stabilize and convert into matured heat-stable crosslinks (Shimokomaki et
699 al., 1972). Hill (1966) reported approximately five times more soluble collagen percent in young
700 hogs compared to old boars and sows although the hogs had a greater percentage of total
701 collagen in muscle. Hill (1966) attributed the comparatively higher total collagen content in
702 young animals versus old ones to muscles not being fully developed at young ages.
703 Shimokomaki et al. (1972) further implicated rapid growth rate during the formative months and
704 therefore the presence of more intermediate heat labile crosslinks. This conflicts with the
705 findings of Berge et al. (1997) who found significant positive correlations between crosslink
706 concentrations and total collagen in their study of the crosslink profile of the semitendinosus and
707 longissimus muscles in three bovine breeds types (Angus × Hereford, Piedmontese × Hereford,
708 pure Brahman). It is therefore contentious for Shimokomaki et al. (1972) to suggest that it is the
709 ratio of heat resistant to heat labile intermolecular crosslinks and not total collagen that is of
710 prime importance.

711 **1.5.5.2 Muscle Location**

712 Intramuscular connective tissues as has been previously elucidated, supports the structural
713 integrity of muscles. They comprise about 2-10% of skeletal muscles on a dry weight basis
714 (Bendall, 1967) and the more active a muscle is, the more connective tissues and mature
715 crosslinks it tends to possess (Provost et al., 2016). In addition, muscles with immediate or close
716 anatomical connections with various skeletal structures tend to possess more connective tissue
717 amount than those further away (Mackenzie, 1919). By way of some research studies supporting
718 the above statements, Kim et al. (1967) found significantly higher amounts of collagen nitrogen

719 in the biceps femoris compared to the longissimus dorsi. Furthermore, their trained sensory
720 panelists rated the longissimus muscle as softer with comparatively lower connective tissue
721 content compared to the bicep femoris. Torrescano et al. (2003) in their study of 14 muscles from
722 Swiss Brown bulls found that the psoas major had the lowest total collagen content (0.31 ± 0.04
723 mg hydroxyproline/g wet tissue) and insoluble collagen concentration (0.18 ± 0.01 mg
724 hydroxyproline/g wet tissue) while the infraspinatus had the highest total and insoluble collagen
725 contents: 0.98 ± 0.05 mg hydroxyproline/g wet tissue and 0.76 ± 0.08 mg hydroxyproline/g wet
726 tissue respectively. Light et al. (1985) also found that tougher bovine muscles like the
727 gastrocnemius and sternomandibularis tended to have greater fractions of perimysium,
728 epimysium and total collagen in comparison with more tender cuts like the longissimus dorsi and
729 psoas major. Based on these observations, one can conclude that intramuscular connective
730 tissues characteristics influence textural incongruities observed between tough and tender
731 muscles and ultimately meat tenderness.

732 **1.5.5.3 Breed and species**

733 Gonzalez et al. (2014) studied in cattle the influence of Brahman genetics on tenderness and
734 mRNA expression of the genes for cystatin C (*CST3*), lysyl oxidase (*LOX*) and bone
735 morphogenetic protein 1 (*BMP1*), and their association with meat quality and collagen
736 intermolecular crosslink formation. They found that cattle with 50% Brahman genetics and
737 purebred Brahmans produced higher Warner Bratzler shear force values and had higher
738 expression of *CST3* and *LOX* at weaning compared to the Angus and Brangus cattle. They
739 therefore concluded that meat tenderness is influenced by the activities of collagen crosslinking
740 enzymes in young animals. Hill (1966) found young hogs and old sows to possess more soluble
741 and total collagen than calves and cows as well as lamb and ewes, and the inclusion of the

742 percent insoluble collagen in that work would have been more informative. Porcine longissimus
743 muscle intramuscular connective tissue had the lowest thermal shrinkage temperature
744 ($60.2^{\circ}\text{C} \pm 0.4$) compared to ovine ($63.0^{\circ}\text{C} \pm 0.7$) and bovines ($63.0^{\circ}\text{C} \pm 0.4$) (McClain et al., 1971)
745 indicating that connective tissue from swine may be more heat labile compared to the other
746 livestock breeds.

747 Research by Dunner et al. (2013) with 15 European cattle breeds on genes associated by meat
748 quality traits found single nucleotide polymorphisms (SNPs) in the aralkylamine N-
749 acetyltransferase (*AANAT*) gene that were significantly associated with total collagen. These
750 authors also found that SNPs in the matrix metalloproteinase-1 (*MMP-1*) gene were associated
751 with m-calpain activity in the muscle. Taye et al. (2017) studied candidate genes related to meat
752 quality in Ankole cattle and identified *COL9A2* influenced meat tenderness. Also, genome wide
753 association studies (GWAS) by Castro et al. (2017) obtained 46 SNPs associated with tenderness
754 in polled Nellore cattle. These studies shed light on the biological mechanisms influencing
755 tenderness in various cattle breeds; however, the same cannot be said for swine breeds especially
756 with respect to genetic parameters for collagen characteristics and their genetic architecture.

757 **1.6 Major Genes affecting pork quality**

758 A major gene is defined as a single gene with a detectable and pronounced effect or influence on
759 the phenotypic expression of a quantitative trait (Bouchard et al., 1997). The halothane and
760 Rendement Napole genes are major genes identified as influencing pork eating quality as well as
761 carcass quality (Salas and Mingala, 2017; Apple and Yancey, 2013).

762 **1.6.1 Halothane gene**

763 Recognition of porcine stress syndrome susceptible pigs was made possible largely by the
764 halothane gene discovery (Fujii et al., 1991; Salas and Mingala, 2017). Eikenkelenboom and
765 Minkema (1974) observed that when exposed to the halothane gas, pigs that were stress
766 susceptible died and subsequently produced a pale, soft and exudative meat.

767 With knowledge of the halothane gene's inheritance, pigs can be placed into three possible
768 genotypes, namely: normal/halothane-negative (Hal^{NN}), heterozygote/halothane-carrier (Hal^{Nn})
769 and mutant/halothane positive (Hal^{nn}) (Apple and Yancey, 2013). When exposed to pre-
770 slaughter stress, muscles from pigs of the Hal^{nn} genotype (halothane positive pigs) are more
771 likely to produce a pale, soft and exudative meat due to an atypical metabolism of lactic acid
772 resulting in a dramatic drop in pH while the muscle is still warm, leading to the denaturation of
773 some muscle proteins (Salas and Mingala, 2017). This denaturation of muscle proteins inevitably
774 leads to the low water holding capacity evident in PSE pork (Bowker et al., 2000).

775 The porcine halothane locus is composed of 2 alleles, Hal^n - the recessive allele and Hal^N -
776 normal allele located at the 6p11-q21 region of chromosome 6 (Fujii et al., 1991). Fujii et al.
777 (1991) found that a single mutation (position 1843) in the gene encoding for ryanodine receptor
778 ($RYR1$) in the calcium release channel of the sarcoplasmic reticulum was linked to porcine stress
779 syndrome in some lean swine breeds. At the time of muscle contraction or relaxation, ryanodine
780 receptor proteins on the sarcoplasmic reticulum membrane surface regulate calcium release rate
781 or uptake (Apple and Yancey, 2013). Pigs with the recessive genotype (Hal^{nn}) have defective
782 ryanodine receptors which results in an uncontrolled Ca^{2+} release and an inability to sequester
783 Ca^{2+} in the sarcoplasmic reticulum after contraction of a muscle (Salas and Mingala, 2017;

784 Apple and Yancey, 2013). This high Ca^{2+} concentration in the muscle results in a rapid
785 postmortem metabolism which, at high muscle temperatures, leads to denaturation of some
786 muscle proteins, resulting in increased moisture losses and subsequently, an increased
787 occurrence of pale, soft and exudative pork in susceptible pigs (Apple and Yancey, 2013). Zhang
788 et al. (1992) reported that the halothane locus explained about 20-30% of the overall meat quality
789 variation in swine populations.

790 Apple et al. (2002) after randomly assigning homozygote halothane negative and heterozygote
791 carriers to diets containing varying levels of magnesium mica, found that halothane carriers
792 produced significantly heavier and leaner carcasses than halothane negative individuals. Further,
793 a higher proportion of carcasses having color scores indicative of the pale, soft and exudative
794 defect came from these carriers. Fernandez et al. (2002) found the longissimus muscle of
795 halothane positive pigs to have higher L^* and drip loss values than the other 2 genotypes. Their
796 subjective sensory test also revealed a lower color score (lighter color) for halothane positive
797 pigs, but a higher rating for tenderness was observed in the halothane negative pigs. Fisher et al.
798 (2000), studying 60 crossbred Landrace \times Large White pigs composed of the three halothane
799 genotypes, found that halothane positive pigs had less mean fat (at the 2nd-3rd last rib), larger
800 mean loin eye area and depth, longer carcasses and higher mean lean percent than halothane
801 negative genotypes. This result is similar to the findings of Aalhus et al. (1991) who also
802 observed a higher lean percent in halothane positive pig carcasses.

803 **1.6.2 Rendement Napole gene**

804 Another major gene with significant influences on pig carcass and meat quality is the
805 Rendement Napole gene also called the *protein kinase AMP-activated non-catalytic subunit*
806 *gamma 3 (PRKAG3)* gene located on chromosome SSC 15 (Monin and Sante-Lhoutellier, 2014).

807 The $\gamma 3$ isoform of *adenosine monophosphate activated protein kinase* (*AMPK*), which is highly
808 expressed in skeletal muscles (Cheung et al., 2000; Mahlapuu et al., 2004), is encoded by
809 *PRKAG3* (Ryan et al., 2012). Mahlapuu et al. (2004) found a significant correlation between the
810 protein expression of $\gamma 3$ isoforms and its mRNA expression in mice and rat gastrocnemius
811 muscles. They further found that the $\gamma 3$ isoform tended to associate with $\alpha 2$ and $\beta 2$ isoforms to
812 form $\alpha 2\beta 2\gamma 3$ complexes, the main AMPK heterotrimer in most skeletal muscles especially
813 white muscles. The AMPK enzyme is an energy sensing enzyme vital in the regulation of
814 numerous pathways including glycogen metabolism (Monin and Sante-Lhoutellier, 2014;
815 Winder, 2001). A mutation that occurs on *PRKAG3* in which guanine is substituted with adenine
816 resulting in a non-conservative substitution to glutamine from arginine has been reported to be
817 the driving force behind the low ultimate pH or acid pork characteristic of homozygous dominant
818 RN^- genotypes (Milan et al., 2000).

819 The RN^- allele has proven to be of appreciable economical relevance to the swine industry due
820 to its detrimental impact on yield after processing (Milan et al., 2000). Le Roy et al. (1990) first
821 tested the hypothesis that the presence of a major gene locus significantly influencing the Napole
822 technological yield trait- a measure of product yield after curing and subsequent cooking
823 (England et al., 2017)- using segregation analysis in the semimembranosus of pigs resistant to
824 porcine stress syndrome. They indeed found a major locus comprising of 2 alleles and these are:
825 RN^- , a completely dominant allele with detrimental effects on Napole Yield; and rn^+ , a normal
826 recessive allele. Research by Monin and Sellier (1985) had earlier revealed the influence of the
827 RN^- gene on pork quality in Hampshire pigs, which they described as the Hampshire
828 type/effect. In their study, three swine breeds were used: Large White, Pietrain (halothane
829 positive) and Hampshire. They found that although pH values 1hr post slaughter were the same

830 between the Hampshire and Large white, a lower ultimate pH was recorded for the Hampshire
831 pigs, which they suggested was most likely due to an extremely high glycolytic potential in the
832 Hampshire pigs. Estrade and Monin (1993) found that pigs with RN^- mutation had as high as a
833 70% rise in glycolytic potential/glycogen content in their longissimus muscles compared to rn^+
834 pigs. This increase in glycolytic potential leads to increased lactate production at the time of
835 postmortem glycolysis, leading to a low ultimate pH (Monin and Sante-Lhoutellier, 2014) which
836 leads to a reduction in water holding capacity of meat from RN^- genotypes. In fact, this low
837 ultimate pH is not the only cause of reduced water holding capacity in these meats as the high
838 glycogen content also influences water holding capacity. The water binding potential of
839 glycogen is approximately the same as that of proteins (Lebret et al., 1999), however, during post
840 slaughter glycolysis or meat processing, glycogen is depleted, leading to the release of water
841 bound to it, which is subsequently lost during cooking, resulting in the lower product yields
842 observed in RN^- genotypes (Lebret et al., 1999).

843 The RN^- gene is primarily found in swine populations with Hampshire ancestry (Moeller et al.,
844 2003). Its allele frequency in the United States (US) Hampshire populations has been found to be
845 0.63 (Miller et al., 2000). Considering the propensity of the RN^- genotype to produce poor
846 quality pork, this high frequency was problematic as Hampshire pigs were widely utilized as
847 terminal sires in the US. The Canadian Centre for Swine Improvement (CCSI) in conjunction
848 with 11 artificial insemination centers and the Center for Food Research and Development
849 (CFRD) reported a net loss of \$12 per hog for those carrying the RN^- gene. Further, CCSI
850 (2001) reported on a research study at the Lacombe Research Centre in which 25% of pork chops
851 obtained from various retail stores in Alberta had abnormally high glycolytic potential
852 suggesting they were likely RN^- gene carriers. Enfalt et al. (1997a) found that RN^- allele

853 carriers had significantly greater average daily gain, lean percent and hind leg muscles than non-
854 carriers, but lower Napole yield, ultimate pH, and crude protein, and higher drip loss, ash and
855 glycolytic potential. Thus, the influence of the RN^- allele may not be restricted to meat quality
856 traits alone but might affect carcass traits and even proximate composition. Estrade and Monin
857 (1993) also found a 10% reduction in protein content in the Longissimus muscle of RN^- pigs
858 compared to their normal counterparts. Lebret et al (1999) found no difference in hydroxyproline
859 content (mg/g) between the three RN genotypes. Enfalt et al. (1997b) looked at breed differences
860 and found RN^- carriers to produce leaner carcasses compared to the Swedish Landrace and
861 purebred Yorkshire breeds.

862 Both RN^- and Hal^{lm} have been reduced to non-significance in the Canadian swine populations
863 through removal of the Hampshire breed from breeding programs and coordinated testing and
864 removal of Hal positive boars and sows from nucleus breeding herds (Mathur and Liu, 2003,
865 https://www.ccsi.ca/Reports/Reports_2003/nsif_CCSI.pdf). Vigilance is still required by the
866 pork industry, however, as Nonneman et al. (2012) identified a novel stress syndrome in pigs
867 characterized by cardiac abnormalities which under stressful conditions can result in death. The
868 dystrophin gene was found to be significantly associated with this syndrome. This stress
869 susceptibility was found to be due to an arginine to tryptophan polymorphism in exon 41 of
870 dystrophin gene. Affected animals tended to have elevated plasma creatine phosphokinase (CPK)
871 levels (Nonneman et al., 2012).

872 **1.7 Other genes affecting pork quality**

873 A number of genes with convincing association to pork quality have been found by various
874 researchers. Ciobanu et al. (2004) reported a quantitative trait locus mapped to the SSC 2 region

875 where the *calpastatin* (*CAST*) gene is found that was significantly associated with eating quality
876 traits like tenderness and juiciness. Calpastatin is a specific inhibitor of calcium-dependent
877 cysteine proteases. These proteases are known to play important roles in the tenderization of
878 meat through degradation of muscle proteins postmortem. Gerbens et al. (1998) studied the
879 association of the *adipocyte fatty acid binding protein* (*A-FABP*) gene with pork quality traits
880 and their results showed that the *A-FABP* locus on SSC4 is associated with the regulation of
881 intramuscular fat (IMF) deposition in pigs. Gerbens et al. (1999) also found increased
882 intramuscular fat content and backfat thickness to be associated with identical heart *fatty acid*
883 *binding protein restriction fragment length* polymorphism genotypes (*RFLP*) on SSC6 in
884 halothane negative Duroc pigs. Kim et al. (2004a) examined the porcine *Delta-like non-*
885 *canonical notch ligand* (*DLK-1*) *gene trap locus 2* (*GLT 2*) region in Yorkshire and Berkshire
886 pigs to assess whether the callipyge polar over-dominance was present in those breeds. They
887 found that F2 progeny with a maternally transmitted allele 1 and a paternally derived allele 2 had
888 greater lean percent and lower backfat thickness. This form of overdominance is described as
889 polar because the phenotypic expression of the heterozygote is more common than the other
890 homozygote genotypes on the same locus (Noelle et al., 1996). Parental imprinting is closely
891 related to this form of overdominance as only one parental allele is expressed in the progeny
892 (Oczkowicz, 2009). This form of overdominance was first reported in sheep after continuous
893 selection for increased muscling resulted in a ram lamb (named Solid Gold) being born with
894 pronounced muscularity in the rump and upper hind legs. It was saved for future mating and
895 descendants of this ram are known as callipyge (Greek for “beautiful buttocks”). The genomic
896 location of the gene influencing this form of muscular hypertrophy has been mapped to the distal
897 end of chromosome 18 (Murphy et al., 2006).

898 Kim et al. (2004b) reported a potential association of the *porcine agouti signaling protein (ASIP)*
899 gene, mapped to SSC17q21 to be having an effect on fat accretion and growth. These results
900 suggest that there may be numerous genes with small additive effects on quality. However, to the
901 author's knowledge, there have been no reports on genetic correlations of Warner-Bratzler shear
902 force measures with soluble and insoluble collagen concentrations. An understanding of this
903 relationship will enrich our knowledge of connective tissue characteristic and determine if
904 selection for collagen traits is possible.

905 Cameron (1990) reported that traits like pork flavor, flavor-liking, and abnormal flavor were low
906 to moderately heritable (0.16-0.31). Cameron (1990) further found a strong positive genetic
907 correlation between pork flavor and intermuscular fat weight (0.60) and subcutaneous fat weight
908 (0.85), thus selecting for these two traits over time may lead to increased pork flavor. Jonsall et
909 al. (2001) reported that ham from carriers of RN^- gene had higher scores for odor intensity than
910 non-carriers, insinuating a possible influence of major genes on this trait, although further
911 research may be needed to substantiate this observation.

912 **1.8 Heritability estimates of some pork quality traits**

913 Information on the heritabilities of meat quality traits provides tremendous insights on the most
914 probable response to selection (Mateescu, 2015). Most pork quality traits typically are low to
915 moderately heritable, as the heritabilities reported for these traits in this current review usually
916 lay within the bounds of 0.10-0.35. Mateescu (2015) also reported low to moderate means in a
917 review on the heritability estimates of beef quality (physicochemical) quality traits ranging
918 between 0.02-0.47. Also, Safari et al. (2005) reported low to moderate heritabilities for meat
919 traits in sheep, ranging from 0.04 to 0.35. Mir et al. (2017) however observed higher heritability

920 estimates for poultry, which had moderate to high heritabilities for quality traits between 0.35
921 and 0.81.

922 Table 1.2 presents a summary of the heritabilities of some selected pork and eating quality traits.
923 Ciobanu et al. (2011) reported that in the category of sensory traits, traits like tenderness
924 (measured either subjectively or through instrumental determinations) were more heritable than
925 traits like juiciness and flavor. Similar observations were made during the scan of the literature
926 for this current review (Table 1.2). Technological traits like color and shear force value were
927 more heritable than pH and cooking loss (Table 1.2). The results from the papers reviewed
928 suggest that improvements in meat quality traits like tenderness (Miar et al 2014), water holding
929 capacity and intramuscular fat are likely with genetic selection. Suzuki et al. (1995) and
930 Hovenier et al. (1993) reported low-moderate heritability estimates for total collagen which
931 suggests that improvements can be made in this trait over time.

932 **1.9 Genetic relationships among some pork quality traits**

933 Genetic correlation between two traits is basically a measure of the influence of pleiotropic genes
934 on those traits and thus, estimates the strength and direction of the relationship between the
935 breeding values of two traits under consideration while phenotypic correlations measure the
936 strength of the relationship between performance/phenotypes of two traits under consideration
937 (Bourdon, 1995).

938 Hovenier et al. (1993) reported a high negative phenotypic correlation (-0.51) between sensory
939 panelist assessment of tenderness and Warner-Bratzler shear force values and subsequently
940 recommended the use of shear force values to predict pork tenderness and possibly as an index
941 trait since the genetic correlation between those traits were also high (-0.71) (table 1.3). Further,

942 they found an almost non-existent (0.01) phenotypic correlation between collagen value and
943 subjective tenderness and a very weak phenotypic relationship with Warner-Bratzler shear force
944 values, indicating that collagen value may not be a good indicator of these two traits (Table 1.4).
945 However, given the sparseness of the information provided on how collagen determinations were
946 made, and the fact that determinations of the amounts of soluble and insoluble collagen were
947 apparently not made, further research is warranted in this regard.

948 **1.10 Conclusions and overall hypotheses**

949 Although heritability and genetic relationships between total collagen and some carcass and meat
950 quality traits have been reported by Suzuki et al. (2005), no literature examining the heritability
951 of collagen heat solubility was noted. Therefore, this thesis, will examine genetic and phenotypic
952 relationships between total, insoluble and soluble collagen and pork quality as no studies have
953 looked at these two traits especially in Canadian swine populations. Specifically, this thesis will
954 test the hypotheses that 1) pork intramuscular collagen characteristics are heritable, and 2) are
955 related to Warner-Bratzler shear force and other meat quality traits. The influence of collagen
956 content on pork texture is not yet well understood and, in light of the competition between pork
957 products and other meat products, there is a need to estimate the heritabilities and genetic
958 relationships of pork collagen characteristics.

959 The objectives of this thesis were then to:

960 1) To estimate the heritabilities and genetic and phenotypic correlations of total, heat-soluble and
961 heat-insoluble collagen from pork intramuscular tissue and compare them to those of other meat
962 quality characteristics;

- 963 2) To examine regression equations considering total, heat-soluble and heat-insoluble collagen
964 that describe pork quality indicators; and
- 965 3) To examine how intramuscular pH, fat, muscle lightness, and Warner-Bratzler shear force
966 relate to each other and other pork quality attributes.

967 **1.11 Tables**

968 Table 1.1: Classification of collagen with their tissue distributions and their chromosomal locations

Collagen Category	Structures formed/ collagen sub-family	Members/ collagen type	Molecular/alpha chain composition	Location/tissue distribution	Alpha chain coding genes (gene location) †	Chromosome number- Sus scrofa	NCBI Gene ID Homo sapiens	NCBI Gene ID (Sus scrofa)	Trimer composition	
Fibrillar	Quarter staggered fibrils	I ^{1-4,7}	$[\alpha 1(I)]_2\alpha 2(I)$	Skin, bone, ligaments, striated muscles, most connective tissues, vitreous humor, cartilage	<i>COL1A1</i> (17q21.3)	12 9	1277 1278	1007381 23 1006267 16	Heterotrimer	
		II ^{8, 10}	$\alpha 1(II)]_3$		<i>COL2A1</i> (12q13.11)	2	1280	397323	Homotrimer	
		III ⁶	$[\alpha 1(III)]_3$		<i>COL3A1</i> (2q31)	15	1281	1001520 01	Homotrimer	
	V ³			$[\alpha 1(V)]_2\alpha 2(V)$ $[\alpha 1(V)]_3$	Skeletal muscle, skin, heart, bone	<i>COL5A1</i> (9q34.3)	1 15	1289 1290	397533 397532	Heterotrimer or homotrimer
						<i>COL5A2</i> (2q32.2)	2	50509	397531	
						<i>COL5A3</i> (19p13.2)				
	XI ^{3, 9}			$\alpha 1(XI)\alpha 2(XI)\alpha 3(XI)$	Cartilage	<i>COL11A1</i> (1p21.1)	4 7	1301 1302	397175 1005209	Heterotrimer
<i>COL11A2</i> (6p21.32)						-	1280	15		
<i>COL11A3/COL2A1</i> (12q13.11)								-		
XXIV ^{7, 11}			$[\alpha 1(XXIV)]_3$	Nascent bones, cornea	<i>COL24A1</i> (1p22.3)	4	255631	1021578 45	Homotrimer	
XXVII ^{3, 7, 12}			$[\alpha (XXVII)]_3$	Cartilage, ear, eye	<i>COL27A1</i> (9q32)	1	85301	1021596 86	Homotrimer	
Non-	Basemen	IV ^{3, 13, 14}	$[\alpha 1(IV)]_2\alpha 2(IV)$	Basement	<i>COL4A1</i>	11	1282	1005153	Hetero-	

fibrillar	t membran es	$\alpha 1 - \alpha 6$	membranes of tissues	(13q34)	11	1284	36	trimer	
				<i>COL4A2</i>	15	1285	1001534		
				(13q34)	15	1286	54		
				<i>COL4A3</i>	X	1287	1006215		
				(2q36.3)	X	1288	04		
				<i>COL4A4</i>			1006214		
				(2q36.3)			03		
				<i>COL4A5</i>			1007370		
			(Xq22.3)				91		
			<i>COL4A6</i>				1006207		
			(Xq22.3)				24		
Short chain (hexagon al lattices)	VIII ¹⁵	$[\alpha 1(VIII)]_2\alpha 2(VIII)$	Descemet's membrane, striated muscles, endothelial cells	<i>COL8A1</i>	13	1295	397430	Hetero- trimer	
				(3q12.1)	6	1296	1005255		
				<i>COL8A2</i>			85		
			(1p34.3)						
	X ^{3,16}	$[\alpha 3(X)]_3$	Growth plate cartilage	<i>COL10A1</i>	1	1300	448809	Homo- trimer	
				(6q22.1)					
Beaded filaments	VI ^{7,17}	$\alpha 1(VI)\alpha 2(VI)\alpha 3(VI)$	Skeletal muscles, cornea, most connective tissues	<i>COL6A1</i>	13	1291	1006237	Hetero- trimer	
				(21q22.3)	-	1292	20		
				<i>COL6A2</i>	15	1293	1001015		
				(21q22.3)			52		
				<i>COL6A3</i>			1001015		
			(2q37.3)				51		
Fibril- associate d collagens with interrupte d triple helices (FACITs)	IX ¹⁸	$\alpha 1(IX)\alpha 2(IX)\alpha 3(IX)$	Cornea, cartilage	<i>COL9A1</i> (6q13)	1	1297	1001553	Hetero- trimer	
				<i>COL9A2</i>	6	1298	19		
				(1p34.2)				1006219	
								11	
	XX ^{5,19}	$[\alpha 1(XX)]_3$	Skin, tendon, cartilage	<i>COL20A1</i>	17	57642	1102573	Homo- trimer	
				(20q13.33)			77		
	XXI ^{3,24}	$[\alpha 1(XXI)]_3$	Blood vessel walls Skeletal muscle, heart, stomach	<i>COL21A1</i>	7	81578	1005236	Homo- trimer	
				(6p12.1)			18		
	XII ^{6,26}	$[\alpha 1(XII)]_3$	Skeletal muscle, endothelial cells, tendons	<i>COL12A1</i>	1	1303	1001566	Homo- trimer	
				(6q13-q14.1)			89		

	XXII ⁵	[$\alpha 1(XXII)$] ₃	Myotendinous junctions of skeletal	<i>COL22A1</i> (8q24.23-q24.3)	4	169044	100519955	Homotrimer
	XIV ⁶	[$\alpha 1(XIV)$] ₃	Tendon, liver, skin	<i>COL9A1</i> (6q13)	1	12839	100155319	Homotrimer
	XVI ^{6,7}	[$\alpha 1(XVI)$] ₃	Fibroblast, keratinocytes	<i>COL16A1</i> (1p35.2)	6	1307	100737666	Homotrimer
	XIX ³	[$\alpha 1(XIX)$] ₃	Skeletal muscle, spleen	<i>COL19A1</i> (6q13)	1	1310	100512751	Homotrimer
Anchorin g fibrils	VII ²	[$\alpha 1(VII)$] ₃	Skeletal muscle, skin, esophagus	<i>COL7A1</i> (3p21.31)	13	1294	100523209	Homotrimer
Multiple triple helix domains and interruptions (MULTIPLEXINS)	XV ³	[$\alpha 1(XV)$] ₃	Smooth muscle cells, fibroblasts	<i>COL15A1</i> (9q22.33)	1	1306	100620394	Homotrimer
	XVIII ⁶	[$\alpha 1(XVIII)$] ₃	Liver, lungs	<i>COL18A1</i> (21q22.3)	13	80781	100624675	Homotrimer
Transmembrane collagens	XIII ²⁰	[$\alpha 1(XIII)$] ₃	Endomysium, epidermis, intestine	<i>COL13A1</i> (10q22.1)	14	1305	100157199	Homotrimer
	XVII ²¹	[$\alpha 1(XVII)$] ₃	Eye, epithelial hemidesmosomes	<i>COL17A1</i> (10q24.3)	14	1308	414914	Homotrimer
	XXIII ^{5,7}	[$\alpha 1(XXIII)$] ₃	Skin, lung, tendon	<i>COL23A1</i> (5q35.3)	2	91522	100521775	Homotrimer
	XXV ²³	[$\alpha 1(XXV)$] ₃	Testis, heart	<i>COL25A1</i> (4q35)	8	84570	102160320	Homotrimer
Others	XXVIII ^{5, 25}	[$\alpha 1(XXVIII)$] ₃	Peripheral nerves	<i>COL28A1</i> (7p21.3)	9	340267	100510974	Homotrimer
	XXIX ^{22, 27}	[$\alpha 1(XXIX)$] ₃	Skin, lung	<i>COL29A1/COL6A5</i> (3q22.1)	-	256076	-	Homotrimer
	XXVI ^{5, 25}	[$\alpha 1(XXVI)$] ₃	Testis, ovary	<i>COL26A1</i>	3	136227	1005195	Homo-

969 1-MGC (2002); 2-Sutmuller et al. 1997; 3-Gelse et al. (2003); 4-Henriksen and Karsdal (2016); 5-Myllyharju and Kivirikko (2004); 6-
970 Kovanen (2002); 7- Uhlen et al. (2015); 8-Gudman and Karsdal, (2016); 9-Morris and Bachinger (1987); 10-Keigo et al. (2017); 11-
971 Matsuo et al. (2008); 12- Plumb et al (2011); 13- Sand et al. (2016); 14-Khoshnoodi et al. (2008); 15- Shuttleworth (1997); 16-
972 Gudmann and Karsdal (2016); 17- Cescon et al. (2015); 18-Olsen (1997); 19-Koch et al. (2001); 20-Siebuhr and Karsdal (2016); 21- Li
973 et al. (1993); 22- Wefel et al. (2011); 23-Keld and Karsdal (2016); 24- Fitzgerald and Bateman (2001); 25- Gordon and Haln (2010);
974 26- Dublet et al. (1989); 27-Söderhäll et al. (2007) †-Homo sapiens.

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989 Table 1. 2: Heritability estimates of some meat quality traits

Trait	Heritability (h^2) [†]	Number of estimates	Range
Cooking loss	0.13	5	0.06-0.20
Shear force value	0.25	6	0.04-0.45
Ultimate pH	0.18	11	0.00-0.39
Drip loss	0.17	8	0.08-0.30
Intramuscular fat	0.40	12	0.19-0.67
Minolta <i>L</i> *	0.26	6	0.16-0.44
Minolta <i>a</i> *	0.29	2	0.21-0.36
Minolta <i>b</i> *	0.18	2	0.15-0.20
Tenderness [†]	0.34	2	0.23-0.45
Juiciness [†]	0.16	3	0.12-0.19
Overall acceptability [†]	0.16	1	-
Water holding capacity	0.63	1	-
Pork flavor [†]	0.28	2	0.13-0.14
Collagen value	0.27	2	0.23-0.30

990 † obtained through a subjective sensory evaluation with panelists. †- The heritabilities hereby presented were average from the
 991 following papers: Allen et al. (1966); Cai et al. (2008); Cameron (1990); DeVries et al. (1994); Cabling et al. (2015); Hovenier et al.
 992 (1992); Jensen et al. (1967); Knapp et al. (1997); Larzul et al. (1997); Lo et al. (1992); Mair et al. (2004); Suzuki et al. (2005); Van
 993 Wijk et al. (2005); Hovenier et al. (1993).

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1000 Table 1. 3: Genetic and phenotypic correlations of cooking loss, shear force and intramuscular fat with some meat quality traits

Traits	Cooking loss		Intramuscular fat		Shear force		Authors
	r_g^1	r_p^1	r_g^2	r_p^2	r_g^3	r_p^3	
Subjective tenderness			0.31	0.05	-0.73	-0.40	Cameron (1990) ² , Malmfors and Nilsson (1978) ³
pH			-0.18	-0.09			Hovenier et al. (1992) ²
Moisture Content			-0.72	-0.50			Malmfors and Nilsson (1978) ²
Drip loss	0.66	0.19	-0.59	-0.35			Malmfors and Nilsson (1978) ¹ , Jensen et al. (1967) ²
Ether extract				0.70			Jensen et al. (1967) ²
Water holding capacity			-0.14	-0.25			Jensen et al. (1967) ²
Lean percent	-	0.10	-0.37	-0.23			De Vries et al. (1994) ^{1,2}
	0.06						

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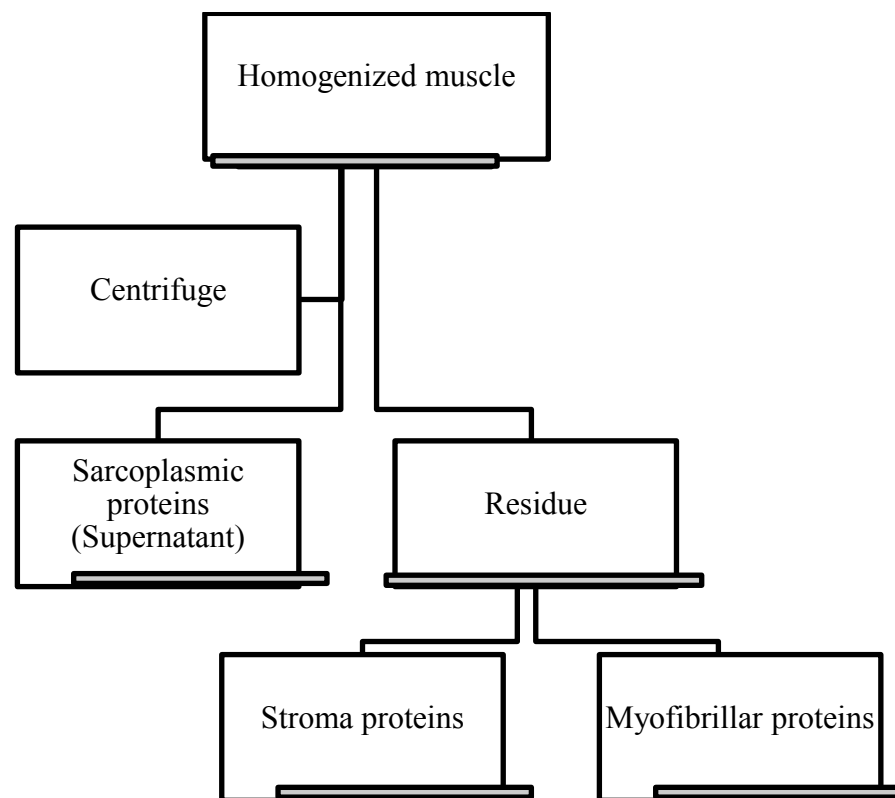
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1011 Table 1. 4: Genetic and phenotypic correlations of total collagen with some pork quality traits

Traits	Total collagen		Author
	r_g	r_p	
Tenderness		0.01	Hovenier et al. (1993)
Intramuscular fat	0.43	0.12	Suzuki et al. (2005)
pH	-0.42	0.03	Suzuki et al. (2005)
Drip loss	-0.09	-0.26	Suzuki et al. (2005)
Cooking loss	-0.64	0.06	Suzuki et al. (2005)
L^*	-0.17	0.18	Suzuki et al. (2005)
a^*	-	0.11	Hovenier et al. (1993)
Shear force value	0.26	-0.10	Suzuki et al. (2005)
Longissimus muscle area	0.19	0.07	Suzuki et al. (2005)
Back fat	-0.35	-0.07	Suzuki et al. (2005)

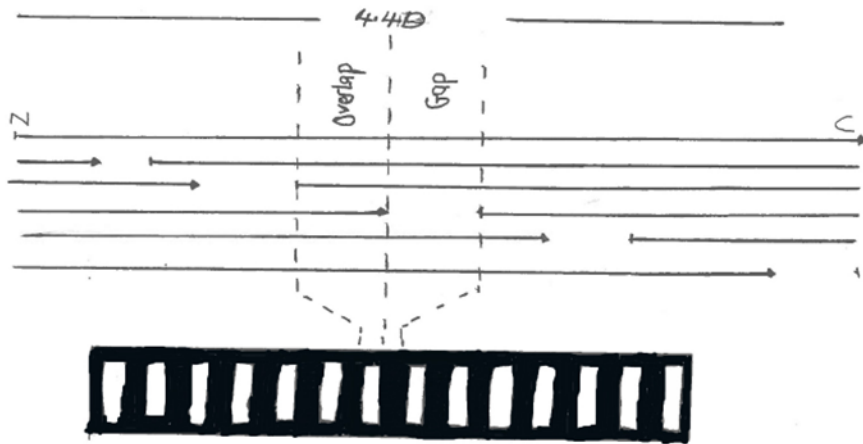
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1013 1.12 Figures



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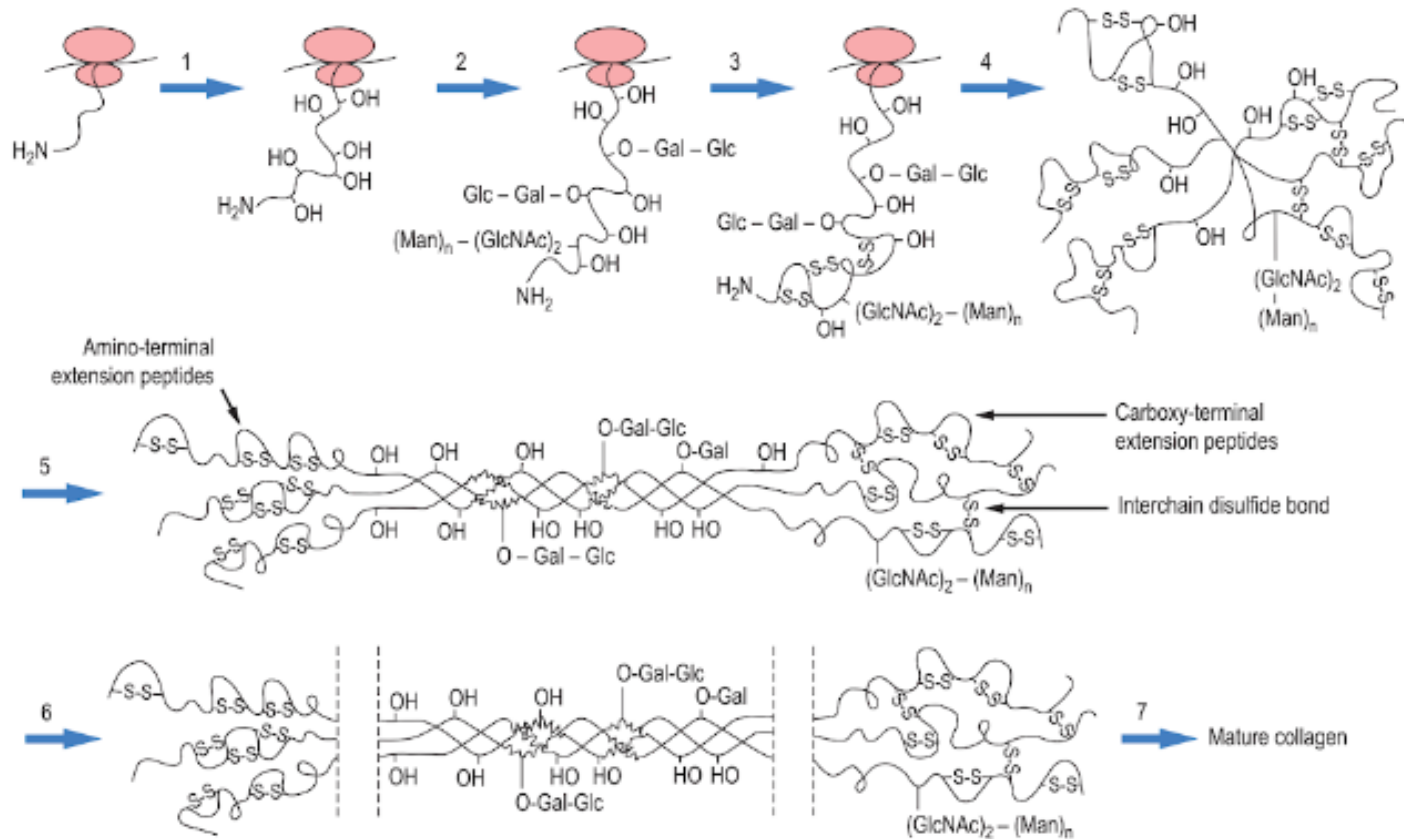
1015 Figure 1. 1: Muscle protein fractionation into its solubility categories. Adapted from Greaser and Guo (2015) with modifications.



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1017 Figure 1. 2: A Schematic representation of the 4.4 D overlap conformation of negatively stained collagen fibers under electron
 1018 microscope showing the banding patterns.

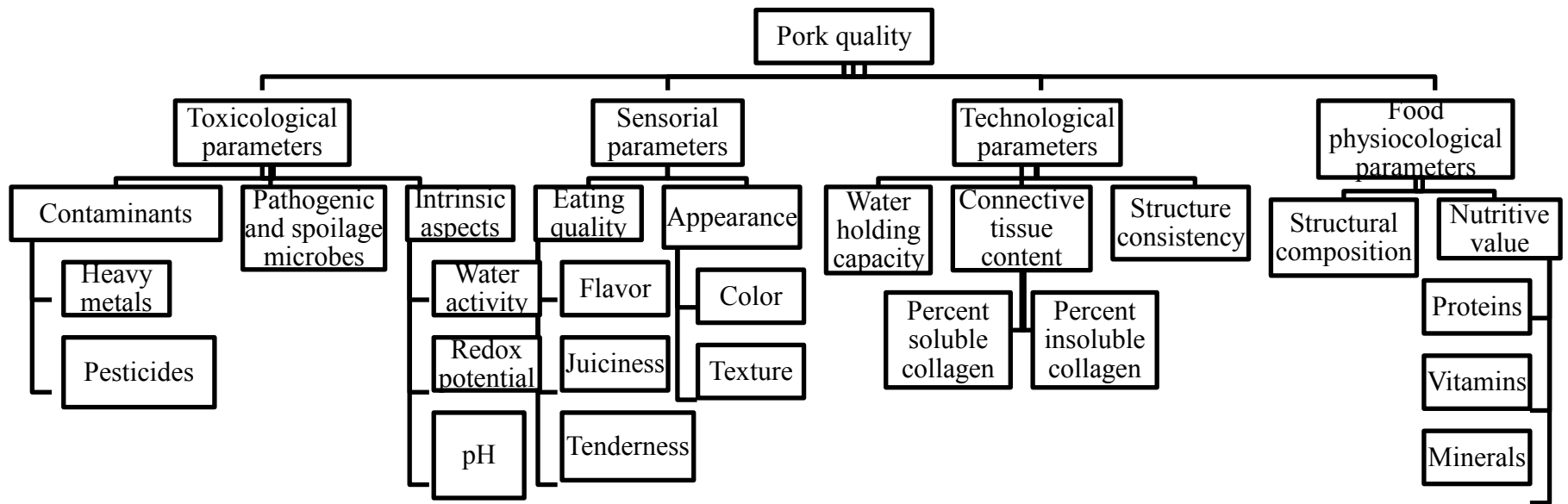
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1021 Figure 1. 3: Collagen biosynthesis and post-translational modifications. 1- hydroxylation of lysine and proline residues; 2-addition of
 1022 N- and O-linked oligosaccharides to alpha chains; 3-intrachain disulfide bonds formation in the N-terminus of the polypeptide chains;
 1023 4-interchain disulfide bond formation at the C-terminus aiding in chain alignment; 5- soluble tropocollagen (triple stranded) formation
 1024 which is subsequently transported to the Golgi vesicles.; 6- removal of C- and N-terminal propeptides after exocytosis; 7- final
 1025 processing- a lateral association of collagen molecules, crosslink formation and formation of collagen fibers. Where: Glc= glucose;
 1026 Gal= galactose; Man= mannose and GlcNAc=N-acetylglucosamine. Reproduced from Kaushal et al. (2014) with permission from
 1027 publishers (copyright 2014 Saunders Elsevier).

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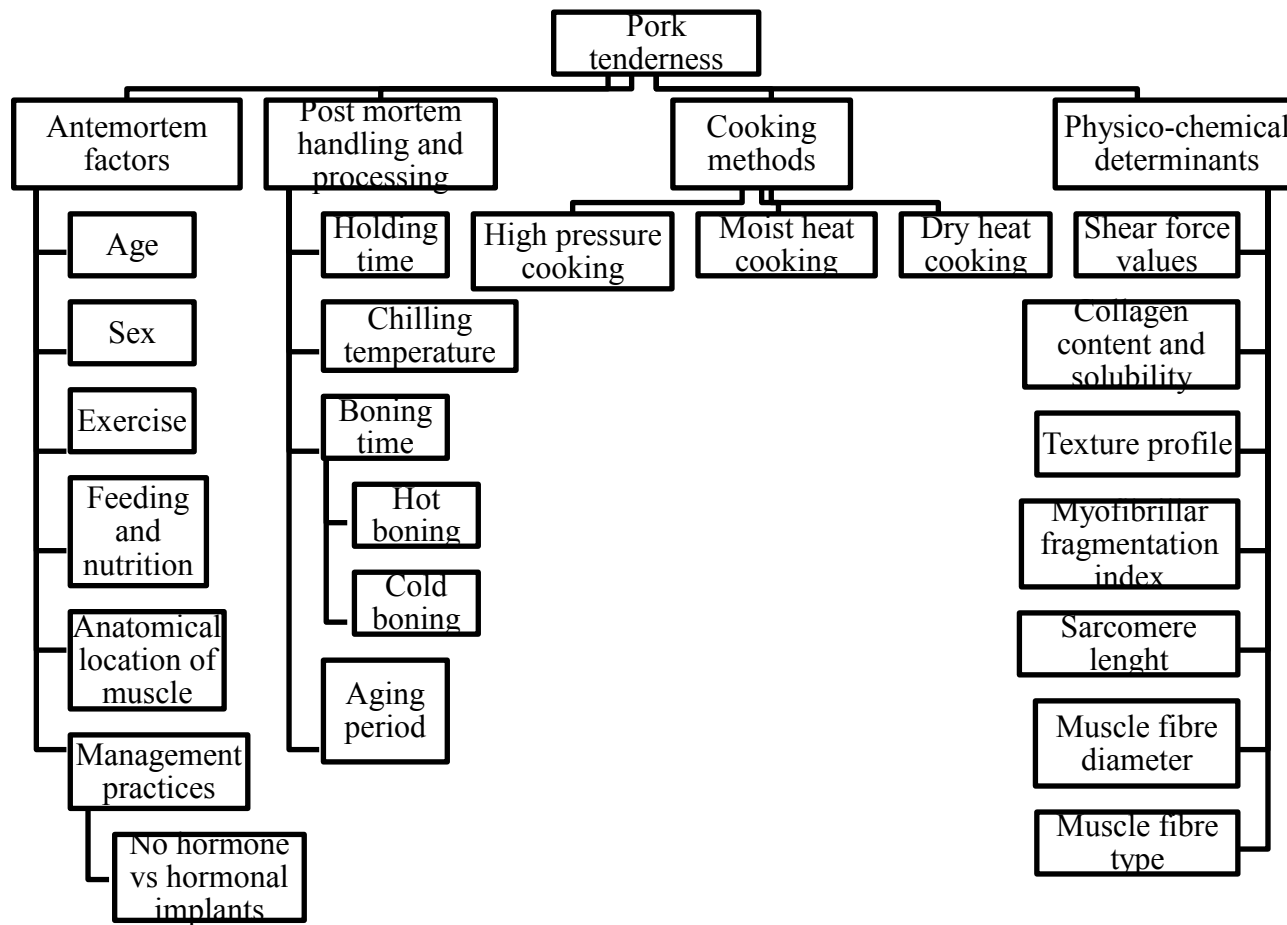
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1039 Figure 1. 5 : Major components of pork quality.

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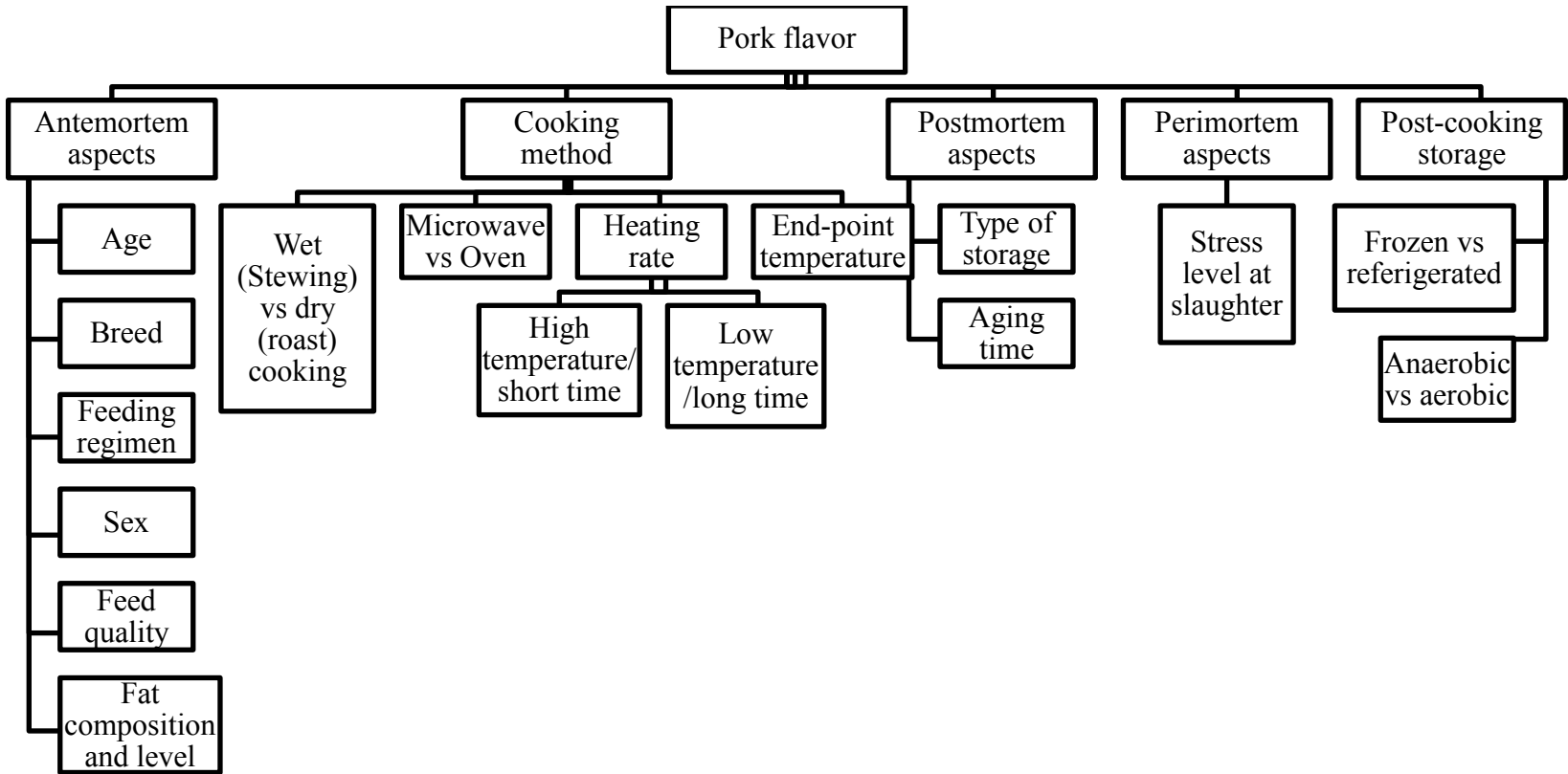
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1043 Figure 1. 6: Factors affecting pork tenderness.

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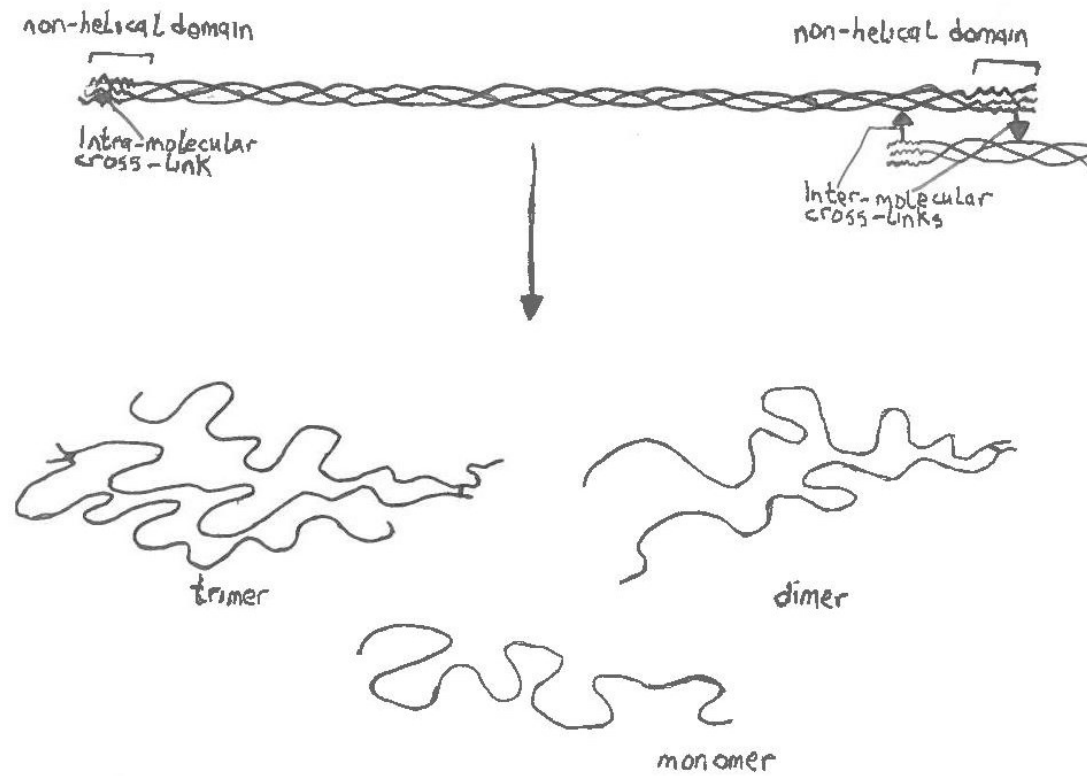


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1048 Figure 1. 7: Factors influencing perceived pork flavor

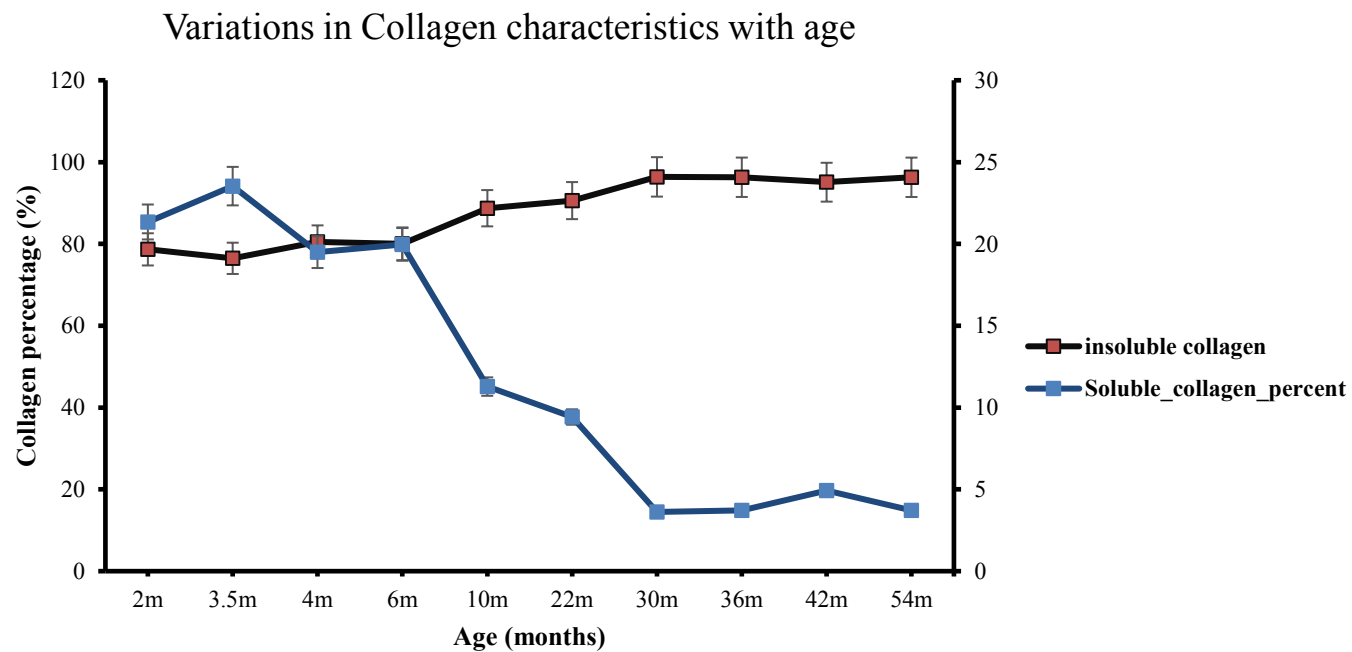
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1052 Figure 1. 8: A Schematic representation of collagen denaturation into gelatin upon heat application.



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1054 Figure 1. 9: Percent soluble and insoluble collagen after heat treatment at 77°C for 1hr in the sternomandibularis muscles of bovines of
 1055 different ages showing the decline in collagen solubility with increasing age. Adapted from Hill (1966) with modifications.

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Chapter 2

Estimates of genetic parameters for loin physical, meat quality and intramuscular collagen traits in Canadian commercial crossbred swine populations

2.1 Introduction

Since the Canadian swine industry has over the years been efficiently, reliably, and effectively producing pork to meet market demands, it is increasingly becoming imperative for the industry to progressively shift its focus towards improving meat quality. Indeed, consumer preferences and standards for quality pork have changed in recent years and for the industry to successfully meet such demands, a clear appreciation of the factors contributing to meat quality and the capacity to forecast, control and influence certain aspects of meat quality is necessary (Hocquette et al., 2012). In fact, the pork industry's ability to meet the demands and preferences of their consumers, by finding the connection between consumer benchmarks/standards for meat and eating quality, will be critical to the future survival and competitiveness of the swine industry in the overall food sector.

Profitable lean pork production has, over the years, been a major selection objective in swine breeding programs with traits like loin eye area, carcass length and reduced fat thickness being favorably weighted in indices (Enfield and Whatley, 1961; Cameron, 1990). However, research studies are showing that selection for traits of economic merit could negatively affect pork quality (Miar et al., 2014; Lonergan et al., 2001). Retail butchers and consumers alike have

1854 complained about meat quality issues associated with very lean meat (Kempster et al., 1986).
1855 They protest that such products are less juicy, flavorful, have aberrant sliceability and are
1856 unappealing (Kempster et al., 1986). This is most probably the reason why traits related to fat
1857 content are also increasingly gaining attention due to their promising potential to improve eating
1858 quality.

1859 One major aim of animal breeding in commercial livestock species is the improvement of
1860 predefined heritable traits by altering their genetic potential/abilities with the aim of shifting the
1861 population average phenotype towards a desired direction (Bourdon, 2013). Thus, determining
1862 the important traits to select in breeding programs as well as how selection for certain traits may
1863 cause responses in correlated traits is important to the success of swine breeding programs as
1864 selection on a trait may bring about changes in the phenotypes of correlated traits by causing
1865 changes in allele frequencies at loci in linkage disequilibrium or pleiotropic loci (Conner and
1866 Hartl, 2004).

1867 In recent decades, breeders are increasingly interested in the relationships between relatively
1868 easy to measure sub-primal traits and meat quality traits as well as specific biochemical traits.
1869 Intramuscular connective tissue properties and collagen characteristics to some degree influence
1870 meat structure and texture. There has indeed, been a surge in meat science research studies
1871 focusing on the inherent characteristics of intramuscular connective tissue mainly due to its
1872 potential influence on muscle texture variability and the disintegration of meat structure (Voutilla
1873 et al., 2007). In fact, pertaining to muscle disintegration, destructured muscles have lower
1874 insoluble and total collagen content compared to normal muscles (Hugenschmidt et al., 2009).
1875 Indeed, as Swatland (1990) aptly noted, the rapid growth rate of commercial meat producing
1876 animals in our time may have led to skeletal muscle fibers outgrowing their connective tissue

1877 sheaths, thus resulting in a weakening of intramuscular connective tissues and predisposing the
1878 muscle to disintegration. A possible reversal of this trend will require information on the
1879 heritabilities of inherent connective tissue characteristics. However, collagen content and heat
1880 solubility are two biochemical traits with very limited research on their heritabilities and
1881 relationships with various sub-primal and meat quality traits. Collagen indeed is the most
1882 abundant protein in mammals and a key constituent of most extracellular matrices where it offers
1883 major mechanical and structural support to tissues (Sionkowska, 2005; Wagner et al., 2009).
1884 Research in meat science has implicated the ratio of soluble to insoluble collagen as having a
1885 profound influence on meat toughness in raw meat (Hill, 1966) although a debate still exists on
1886 its role in cooked meat toughness. It is therefore imperative to examine the potential presence of
1887 genetic variability and relationships.

1888 In 2010, a series of research studies were initiated at the University of Alberta to estimate genetic
1889 and phenotypic parameters in two Canadian crossbred pig populations. Those experiments
1890 examined a total of 43 carcass and pork quality traits. However, those studies did not investigate
1891 the sub-primal physical, intramuscular fat and collagen characteristics of those swine
1892 populations. This current study seeks to fill the gap in those studies by testing the hypothesis that
1893 genetic relationships exist between collagen characteristics, sub-primal and meat quality traits.

1894 Therefore, the objectives of this study are to:

- 1895 1. Estimate the heritabilities for the loin physical, meat quality traits and collagen
1896 characteristics; and
- 1897 2. Estimate phenotypic and genetic correlations between and among the loin physical, meat
1898 quality traits and collagen characteristics.

1899 **2.2 Materials and methods**

1900 **2.2.1 Animals and study design**

1901 Experimental animals were managed in accordance with the recommendations stipulated in the
1902 Canadian Council for Animal Care guide (CCAC, 1993). The use of these animals was reviewed
1903 by an animal research ethics committee and considered Category A and so no formal approval
1904 was required. Pigs were from two Canadian breeding populations. Total number of pigs for this
1905 study amounted to 500 commercial animals of a combination of full sib and half sibs obtained
1906 from crosses between 139 Duroc boars and 429 first cross hybrid Landrace × Yorkshire sows.
1907 Pedigree information of 8 ancestral generations encompassing 8,372 animals composed of 487
1908 sires and 1105 dams was available.

1909 **2.2.2. Animal management**

1910 Management of experimental pigs in this study was as described by Miar et al. (2014) as these
1911 animals were a subset of that experimental population. Male piglets were neutered at 3 to 5 days
1912 post birth. Piglets were weaned at approximately 7.5 kg at about 21 days of age. Weanlings were
1913 subsequently housed in nursery pens for 5 weeks, then raised in pre-grower barns for a pre-
1914 growing period of 4 weeks. Following this, pigs were transitioned to commercial finishing
1915 conditions with *ad libitum* access to feed and water. The final body weight of the animals was
1916 approximately 115 kg. The pigs were transported to East 40 Packers, a provincially inspected
1917 abattoir, Brandon, MB, Canada, on a weekly basis in groups of 20-35 pigs. They were rested in
1918 lairage overnight with *ab libitum* access to water and slaughtered the following morning.
1919 Provincial regulations were followed in the slaughter of the animals. They were electrically
1920 stunned prior to exsanguination and their carcasses fabricated within 24 hours post-slaughter.

1921 They were on average 160 days old at slaughter with an average live weight of 124 kg at
1922 slaughter.

1923 **2.2.3 Loin physical measurements**

1924 A 3 and 4 rib loin section excised from the right side of the pork carcasses during fabrication
1925 were harvested, vacuum packaged and refrigerated at 4°C within 24 hours. They were then
1926 frozen (-20°C) for 96 hours and transported in coolers to the carcass and meat science laboratory
1927 at Agri-Food Discovery place, Department of Agricultural, Food and Nutritional Science,
1928 University of Alberta, and stored at -20°C until analysis. Prior to analysis, the samples were
1929 thawed at refrigerated temperature (4°C) for 61 hours.

1930 The sub-primals used in this study, that is the loin samples, were weighed and recorded as loin
1931 weight (LWT). Loin eye muscles excised from the loin sub-primals were weighed and reported
1932 as loin eye weight (LEW). Back fat trimmed from the loins were weighed and reported as thick
1933 fat weight (TFW). Meat and fat trimmings remaining after removing chops to be used for other
1934 meat quality measurements were weighed and recorded as rib trim weight (RTW) while
1935 remaining bones were reported as bone weight (BWT). The ribeye weight, thick fat weight, rib
1936 trim weight were expressed as percentages of the whole loin weight after subtracting the
1937 diaphragm weight from the whole loin weight. Diaphragm weight was comprised of nerves and
1938 diaphragm tissues of no significance to the goals of the current study.

1939 **2.2.4 Meat quality measurements**

1940 **2.2.4.1 Muscle pH**

1941 The pH measurements on frozen-thawed samples were taken at 3 different locations on the loin
1942 eye (Figure 2.1). This was done by making an incision with a knife about halfway (5cm) through

1943 the meat, waiting for some fluid to accumulate in the incision and then inserting the pH probe
1944 (Accumet® AP71 portable waterproof pH meter, Fisher Scientific, ON, Canada). The average of
1945 three measurements for each loin eye was used for statistical analysis.

1946 **2.2.4.2 Objective color measurements**

1947 A hand-held Konica Minolta CR 400 (Konica Minolta Sensing Inc., Japan) chroma meter set at
1948 illuminant D65 was used for L^* , a^* , b^* , hue and chroma determinations. These measurements
1949 were taken at three different sites on the thawed loin eye muscle after blooming for a period of 1
1950 hour at 4°C to allow for pigment oxygenation. The measuring head of the chroma meter was
1951 placed vertically on the surface of the loin chop for the readings (Figure 2.2). The colour of each
1952 loin chop was characterized using the L^* , a^* , b^* colour space, where L^* = lightness/ brightness
1953 with 0 = black and 100 = white; a^* = redness (+a) to greenness (-a) axis, and b^* = yellowness (+b)
1954 to blueness (-b) axis as recommended by the Commission Internationale de L'Eclairage (CIE).
1955 The average of 3 readings was used for statistical analysis of the color space coordinates.

1956 **2.2.4.3 Cooking loss**

1957 An approximately 200g loin roast from each animal trimmed of fat and epimysium was used for
1958 this assessment. Average weight after trimming was 192 ± 12 g. The roasts were weighed and
1959 inserted into thin walled Whirl-pak™ bags and placed in a water bath set at 73°C ensuring that
1960 the opening of the bags was above the water levels. A TinyTag™ thermistor probe (model: TV-
1961 4020, Gemini Data loggers, West Sussex, UK) was used to record temperature changes in the
1962 chops. The average cooking time was 57 ± 13 mins. When an internal temperature of 71°C was
1963 reached, chops were removed from the water bath and cooled in an ice slurry. The samples were

1964 subsequently stored at 4°C for 24 hours, gently blotted and re-weighed. Percent cooking loss was
1965 determined with the following formula:

$$1966 \frac{\text{Muscle weight precook} - \text{Muscle weight post cook}}{\text{Muscle weight precook}} \times 100$$

1967 **2.2.4.4 Instrumental measure of tenderness**

1968 Cooked pork chops were refrigerated overnight at 4°C before Warner-Bratzler shear force
1969 determinations. Six 1cm² rectangular cores were obtained longitudinal to the fiber direction of
1970 each chop. A material testing machine (Lloyd Instruments Ltd, Surrey, UK) fitted with a
1971 Warner-Bratzler-like shear head was used to shear through the cores across the muscle fibre
1972 direction at a crosshead speed of 200mm/min. NEXYGEN Plus texture analysis software was
1973 used to record resistance (peak force) values in Newtons. The average readings of six cores were
1974 used for statistical analysis.

1975

1976 **2.2.4.5 Ether extract/intramuscular fat**

1977 This procedure was performed using the Soxhlet method 960.39 (AOAC, 2000). For each loin,
1978 duplicate determinations were made. A Soxtec™ 2050 Automatic Extraction apparatus (FOSS
1979 Analytical AB, Höganäs, Sweden) consisting mainly of an extraction unit, control unit and a
1980 drive unit was used for intramuscular fat determinations in the loin samples.

1981 **2.2.4.5.1 Sample preparation**

1982 Aluminum extraction cups to be used for the extraction were oven dried at 103°C for 2hours,
1983 cooled in a desiccator for 45min and weighed. Approximately 2.0g ($\pm 0.003g$) of lyophilized,
1984 ground *Longissimus dorsi* muscle were weighed into tared assembled cellulose extraction

1985 thimbles (33mm × 80mm). De-fatted cotton was subsequently plugged into the thimbles to
1986 prevent sample loss during the extraction process. With the aid of a thimble handler, the samples
1987 were transferred to a thimble stand. Sample thimbles were then inserted into the extraction unit
1988 followed by the insertion of the pre-weighed aluminum extraction cups. Petroleum ether (40-
1989 60°C) was used as the solvent for the extraction process. Approximately 80mL of petroleum
1990 ether were dispensed into each of the aluminum extraction cups by a dispenser connected to the
1991 extraction unit.

1992 **2.2.4.5.2 Ether extraction procedure**

1993 Samples were immersed in the boiling petroleum ether (solvent) for 30min, followed by 45min
1994 rinsing/refluxing which involved rinsing the thimble contents in cold solvent to ensure removal
1995 of any remaining soluble material. The thimbles were then dried for 15min.

1996 After completion of the extraction, the aluminum extraction cups containing the extracted fat
1997 were air dried in an oven for 20min at 105°C and subsequently cooled to room temperature for
1998 15min in a desiccator. After cooling, the cups were weighed again and intramuscular fat contents
1999 were calculated with the following formula: intramuscular fat

$$2000 (\%) = \frac{\text{fat recovered}}{\text{weight of the sample used for fat extraction (g)}} \times 100$$

2001 where fat recovered = weight of aluminum extraction cup after extraction - weight of aluminum
2002 extraction cup before extraction. Percent intramuscular fat was expressed on a moisture free
2003 basis.

2004 **2.2.4.6 Crude protein, fat and moisture content**

2005 Traces of meat remaining on the rib bones and vertebrae after excision of the loin eye muscle
2006 were ground into a homogenized state and analyzed for crude protein, crude fat and crude
2007 moisture content using a FOSS Food Scan™ Near-Infrared Spectrophotometer (FOSS, DK-3400,
2008 Hillerød, Denmark) calibrated with FOSS Artificial Neural Network calibration. The Artificial
2009 Neural Network uses a correlative approach to predict sample composition. Thus, it assesses the
2010 relationship between sample constituent values and established spectral characteristics to
2011 decipher the test spectra and relay results for the constituents (crude protein, fat and moisture).

2012 Each ground meat sample was uniformly spread on a glass-bottom sample cup and was
2013 subsequently inserted in the sample chamber of the FOSS Food Scan™. The product profile of
2014 the sample was then selected, and the scanning procedure initiated with the press of the start
2015 button. The analysis relies on near infra-red transmittance technology for the determination of
2016 protein, fat and moisture content simultaneously in the samples. A detector measures the amount
2017 of unabsorbed light after transmission of the light through the sample. The displayed results were
2018 in grams per 100g of sample for crude protein, fat and moisture. Duplicate readings were
2019 obtained for each sample, and the mean of the duplicates used in subsequent statistical analyses.

2020 **2.2.4 Collagen characterization**

2021 **2.2.4.1 Initial sample preparation**

2022 An approximately 2.5cm thick loin chop was excised from the loin of each animal. The steaks
2023 were weighed on a Navigator™ portable balance (Model: N08110, OHAUS, Ontario, Canada)
2024 and appropriately labelled. The epimysium around the steaks was trimmed and the chops were
2025 cut into 1cm³ cubes. The cubes were weighed again, placed in aluminum drip pans and covered

2026 with perforated aluminum foil to enhance water vapor escape during freeze drying. The samples
2027 were then frozen at -20°C for 24 hours and lyophilized for 5 days. After freeze drying, the
2028 samples were reweighed and a Waring blender (Model 7011C, Waring Commercial, Torrington,
2029 CT, USA) together with dry ice was used to crush the lyophilized samples into a coarse powder
2030 which was subsequently kept at -20°C until analysis.

2031 The moisture content in each sample was determined with the following formula:

$$2032 \text{ Moisture content} = \frac{\text{weight prefreeze drying} - \text{weight post freeze drying}}{\text{weight prefreeze dring}} \times 100$$

2033 **2.2.4.2 Heat solubility of collagen**

2034 The soluble and insoluble fractions of collagen were obtained using the method described by Hill
2035 (1966) (Figure 2.3). This method estimates the amount or proportion of collagen that is soluble
2036 or dissolves at 77°C (Zayas, 1997). Freeze dried sample powder ($1.000 \pm 0.003g$) was weighed
2037 into labelled 50mL test tubes on a Denver Analytical Balance (Model: SI-234, Denver
2038 Instrument Company, CL, USA). After weighing, 20mL of $1/4$ strength Ringer's solution (4.5g
2039 sodium chloride, 0.21g potassium chloride, 0.318g calcium chloride dehydrate, 2000mL
2040 deionized water, pH 7.4) were added to the sample tubes. The tubes were then placed in a digital
2041 controlled water bath (Fisher Scientific Isotemp™ 228, NJ, USA) set at 77°C for 60mins and
2042 vortexed every 10mins.

2043 After incubation, the tubes were cooled in ice slurry for 15mins and their contents were
2044 transferred into pre-weighed centrifuge tubes. The centrifuge tubes were balanced on a Harvard
2045 trip mechanical balance (Model 1450-SD, OHAUS® Corporation, NJ, USA). The tubes were
2046 then centrifuged (AVANTI® J-E centrifuge, Beckmann Coulter Inc., CA, USA) for 10mins at

2047 3500×g at 4°C. The resulting supernatant was decanted into pre-labelled vials. The centrifuge
2048 tubes containing the pellets were drained at a 45°C angle for 1hour (Figure 2.4) and the wet
2049 residue was subsequently scraped into separate vials for future analysis.

2050 **2.2.4.3 Acid hydrolysis and neutralization**

2051 Two 1 mL aliquots of soluble supernatant per sample were pipetted into 20mL test tubes. For
2052 insoluble fractions, 0.3000g±0.003g was weighed into the test tubes (Figure 2.5). For the
2053 soluble fraction test tubes, 1mL 12M HCL and 4mL 6M HCL were added to each tube and the
2054 head space of each tube was purged of oxygen with nitrogen gas for about 10 seconds and
2055 immediately capped. The tubes were then placed in a test tube heating block (Accublock™
2056 Digital dry bath, LABNET International Inc., NJ, USA) (Figure 2.6) and hydrolyzed for 20hours
2057 at 110°C (Suzuki et al., 1995; Colgrave et al., 2012). For the residue fraction, 6mL 6M HCL
2058 were added to each tube and hydrolyzed as described above. Such conditions ensure that the
2059 peptide bonds are hydrolyzed quantitatively (Aitken and Learmonth, 1996; Hsiang and Cole,
2060 1974).

2061 After hydrolysis, the tubes were cooled in ice slurry for 10mins. The cooled hydrolysates were
2062 then filtered through pleated Whatman™ No. 4 qualitative filter papers (GE Healthcare Ltd,
2063 Buckinghamshire, UK) wetted with deionized water into round-bottom evaporating flasks.

2064 The acid solution of each sample was evaporated using a rotary evaporator (Heidolph Collegiate
2065 LABORTA 4000WB ECO ROTOVAP, Schwabach, Germany) with a rotation speed of about
2066 30rpm equipped with a self-cleaning dry vacuum system (Model: 2026, WELCH-ILMVAC, IL,
2067 USA), heating bath temperature of 42°C, a vacuum pressure of 680-700mmHg and a circulating
2068 water bath (Model: OB-13L, Fisher Scientific, PA,USA) for maintaining condenser temperature

2069 between 7°C to 10°C; as well as a dry ice trap (Model: UST 800, KNF Neuberger Inc., NJ, USA)
2070 to improve pump efficiency and prevent condensed liquids from getting to the pump.

2071 After the solutions had been evaporated to dryness, they were reconstituted with 2mL deionized
2072 water. Reconstituted hydrolysis solution pH was tested using cut litmus paper and reconstituted
2073 solutions were neutralized as required with either 2M NaOH or 0.5 NaOH. After neutralization,
2074 samples were evaporated again as described above. After the final evaporation, dried
2075 hydrolysates from the soluble supernatant fraction were finally reconstituted with 5mL deionized
2076 water while those of the insoluble fraction were reconstituted with 8mL deionized (DI) water.
2077 Solutions were then transferred into vials and stored at -20°C for hydroxyproline determinations.

2078 **2.2.4.4 Hydroxyproline determination**

2079 The procedure for hydroxyproline quantification in experimental samples was undertaken as
2080 described by Neuman and Logan (1960) (figure 2.7). The Neuman and Logan method relies on
2081 the oxidative dehydrogenation of hydroxyproline by hydrogen peroxide in an alkaline
2082 environment (NaOH) with copper ions present to produce a chromophore which upon acid
2083 (sulphuric acid) and heat treatment, produces an orange-lilac color with p-
2084 dimethylaminobenzaldehyde (Baker et al., 1953).

2085 **2.2.4.4.1 Hydroxyproline standards preparation**

2086 One milliliter of hydroxyproline standard stock solution ([0.100g trans-4-hydroxy-L-proline
2087 SIGMA-ALDRICH, MO, USA], 25µL 12M HCL, 250mL deionized water) was added to 9mL
2088 deionized water in a screw capped glass vial to yield 10mL of 40µg/mL hydroxyproline assay
2089 standard. This solution was gently mixed by vortex and 5mL of the solution were pipetted into a
2090 different vial containing 5mL deionized water to yield 10mL of 20µg/mL hydroxyproline. This

2091 serial dilution was continued until a 10mL solution of 2.5µg/mL hydroxyproline assay standard
2092 was obtained. This process was repeated whenever hydroxyproline determinations on
2093 experimental samples were made.

2094 **2.2.4.4.2 Hydroxyproline determination in experimental samples**

2095 Two aliquots of 1mL per sample were pipetted into sealable glass test tubes. Afterwards, 1mL
2096 0.01M copper (II) sulfate solution (0.250g copper (II) sulfate pentahydrate, 100mL DI water),
2097 1mL 2.5N sodium hydroxide solution (100g NaOH pellets, 1000mL DI water) and 1mL 6%
2098 hydrogen peroxide solution (10mL 30% hydrogen peroxide, 100mL DI water) were added in
2099 succession. A yellowish-brown color forms after addition of these solutions. The tubes were then
2100 allowed to stand at room temperature for 5 minutes with occasional shaking. Tubes were then
2101 transferred to a water bath (Isotemp™ 228 Digital Control water bath, Fisher Scientific, NH,
2102 USA) set at 80°C and vigorously shaken with tube contents fully immersed in the water. This
2103 was done to ensure destruction of excess peroxide (Neuman and Logan, 1960). Tubes were then
2104 cooled in an ice slurry. After cooling, 4mL 3.0 N sulfuric acid solution and 2mL 5% p-
2105 dimethylaminobenzaldehyde (5g p-dimethylaminobenzaldehyde powder dissolved in 100mL 2-
2106 propanol) were added to each tube. The tubes were then capped, gently vortexed and incubated
2107 in a water bath set at 70°C for 16mins.

2108 After incubation, the tubes were cooled with tap water for 5mins and 1mL aliquots of tube
2109 contents were transferred into 1.5mL plastic cuvettes for measurements of the extinctions
2110 (optical densities) of the colors generated with a UV-Vis spectrophotometer (GENESYS™ 20
2111 visible spectrophotometer, Thermo Scientific, ON, Canada) at a 550nm wavelength against a
2112 blank solution (no hydroxyproline) and a set of hydroxyproline standard solutions with
2113 concentrations 2.5, 5, 10, 20 and 40µg hydroxyproline per mL. The absorbance readings of these

2114 standards were used to plot standard curves which were subsequently used to determine the
2115 hydroxyproline content of the experimental samples. That is, a linear regression model of $y=mx+$
2116 b fitted for the standards, where y = relative absorbance, x = analyte concentration, m = slope of
2117 the model and b =intercept was used to determine the hydroxyproline concentrations of the
2118 experimental samples (Colgrave et al., 2012). The average absorbance reading for duplicate
2119 solutions per sample was used for subsequent calculations. Hydroxyproline content in both the
2120 soluble and insoluble fractions of the experimental samples were multiplied by 7.14 (Dransfield,
2121 1977) This assumes that collagen comprises about 14% 4-hydroxyproline (Patiyal and Katoch,
2122 2006; Jackson et al., 1996; Dransfield et al 1983; Naveena et al., 2011; Clark et al., 2005; Ando
2123 et al., 2006).

2124 **2.2.4.5 Calculating collagen content in the raw meat**

2125 **2.2.4.5.1 Soluble collagen**

2126 Firstly, the total hydroxyproline (μg) in 1mL of reconstituted samples was determined by
2127 inserting the averages of their spectrophotometric readings into standard curve regression
2128 equations. The resultant value was multiplied by the total/ final reconstituted solution which was
2129 5mL to obtain total hydroxyproline in the 5mL reconstituted volume.

2130 Since 20mL Ringer's solution was used to incubate the freeze-dried muscle tissues, the total
2131 hydroxyproline in 5mL was multiplied by a factor of 20 to obtain total hydroxyproline (μg) in
2132 20mL Ringer's solution. This hydroxyproline amount (μg) *ceteris paribus*, is assumed to be
2133 same in the freeze-dried meat ($1\text{g}\pm 0.003$).

2134 Total soluble collagen (μg) in the freeze-dried meat used for collagen characterization was
2135 determined by multiplying the total hydroxyproline in the tissue by the conversion factor 7.14.
2136 The values obtained were subsequently converted to milligrams (mg).

2137 Total soluble collagen in total freeze-dried meat was determined by multiplying total soluble
2138 collagen per gram freeze dried meat by total freeze-dried meat (g). Total soluble collagen (mg) in
2139 total raw meat was assumed to be the same as the amount in the total freeze-dried meat. Total
2140 soluble collagen per gram raw meat was determined by dividing total soluble collagen in total
2141 raw meat by the total raw meat.

2142 **2.2.4.5.2 Insoluble collagen**

2143 In the case of the residue (insoluble fraction), after determining the total hydroxyproline in the
2144 total reconstituted volume (8mL), total hydroxyproline per gram wet residue was determined by
2145 dividing the total hydroxyproline in the used wet residue by the actual weight (g) of the wet
2146 residue used for hydrolysis which was $0.3\text{g} \pm 0.003$. Total hydroxyproline in the total wet
2147 residue that remained after the heat solubility of collagen phase was determined by multiplying
2148 the total hydroxyproline in the wet residue used for hydrolysis by the weight (g) of the wet
2149 residue that remained after heat solubility. All further determinations were as described for
2150 soluble collagen. Total collagen per gram raw meat was determined by summing the total soluble
2151 and insoluble collagens per gram raw meat.

2152 **2.2.5 Statistical analysis**

2153 Lilliefors (Kolmogorov-Smirnov) test of normality (Lilliefors 1967; Gross and Ligges, 2015)
2154 was implemented in an R statistical computing environment to test the hypothesis that the
2155 samples were drawn from normally distributed populations. Significance of fixed effects to be

2156 included in the various models were tested using the glm2 (Marschner, 2011) package which fits
 2157 generalized linear models and was further confirmed by fitted univariate models in AsReml
 2158 version 4.1 (Butler, 2009; Gilmour et al., 2015). Fixed and random effects of all fitted models are
 2159 presented in Tables 2.1 and 2.2. Means and standard deviations were obtained using the base
 2160 package (R Core Team, 2018), whilst the coefficients of variation was determined with the
 2161 FinCal package (Fan, 2016).

2162 In order to estimate phenotypic and genetic correlations among the traits under consideration,
 2163 phenotypic and genetic (co) variances were estimated with bivariate animal models using the
 2164 AsReml 4.1 software (Gilmour et al., 2015). Random additive polygenic effects were included in
 2165 animal models for all traits under consideration. Kill date (slaughter batch), sex and company of
 2166 origin were fitted as fixed effects (Table 2.1 and 2.2). Cooking time was used a linear covariate
 2167 for cooking loss (%). The model formula is as follows:

$$2168 \quad \begin{bmatrix} y_1 \\ y_2 \end{bmatrix} = \begin{bmatrix} X_1 & 0 \\ 0 & X_2 \end{bmatrix} \begin{bmatrix} b_1 \\ b_2 \end{bmatrix} + \begin{bmatrix} Z_1 & 0 \\ 0 & Z_2 \end{bmatrix} \begin{bmatrix} a_1 \\ a_2 \end{bmatrix} + \begin{bmatrix} e_1 \\ e_2 \end{bmatrix}$$

2169 Where y_1 and y_2 are vectors of phenotypic value for 2 paired traits under consideration, b_1 and
 2170 b_2 are vectors of fixed effects for the 2 paired traits and covariate, a_1 and a_2 are vectors of
 2171 random additive genetic effects of traits 1 and 2. e_1 and e_2 are vectors of random residual effects,
 2172 X_i and Z_i are known design matrices for fixed effects and random additive genetic effects
 2173 respectively.

$$2174 \quad \text{The variance-covariance matrix is as follows: } \text{var} \begin{bmatrix} a_1 \\ a_2 \\ e_1 \\ e_2 \end{bmatrix} = \begin{bmatrix} A_{\sigma_{a_1}^2} & A_{\sigma_{a_{12}}} & 0 & 0 \\ A_{\sigma_{a_{21}}} & A_{\sigma_{a_2}^2} & 0 & 0 \\ 0 & 0 & I_{\sigma_{e_1}^2} & I_{\sigma_{e_1 e_2}} \\ 0 & 0 & I_{\sigma_{e_2 e_1}} & I_{\sigma_{e_2}^2} \end{bmatrix}$$

2175 Where A is the numerator relationship matrix (Meuwissen and Luo, 1992) built from a pedigree
2176 composed of 8,372 animals. I is the identity matrix whose order is equivalent to the number of
2177 animals. Genetic (r_G) and phenotypic correlations (r_P) were estimated as: $r_G = \frac{Cov_{1,2}}{\sqrt{\sigma_{G1}^2 \sigma_{G2}^2}}$ where:

2178 $Cov_{1,2}$ = additive genetic covariance of trait 1 and 2

2179 σ_1^2 = additive genetic variance for trait 1

2180 σ_2^2 = additive genetic variance for trait 2

2181 $r_P = \frac{Cov_{p1,p2}}{\sqrt{\sigma_{p1}^2 \sigma_{p2}^2}}$ where $Cov_{p1,p2}$ = additive phenotypic covariance for trait 1 and 2

2182 σ_{p1}^2 = phenotypic variance for trait 1

2183 σ_{p2}^2 = phenotypic variance for trait 2

2184 $\sigma_p^2 = \sigma_a^2 + \sigma_e^2$

2185 $Cov_{p1p2} = Cov_{a1a2} + Cov_{e1e2}$

2186

2187 A univariate animal model: $y = Xb + Z_a a + e$; was fitted in AsReml (Gilmour et al. 2015) to
2188 estimate genetic and phenotypic variance and residual components, where:

2189 y = vector of phenotypic value

2190 b = vector of fixed effects

2191 a = vector of additive genetic effects

2192 e= vector of residual effects

2193 X= incidence matrix for the fixed effects

2194 Z= incidence matrix linking observation to random effect of animal

2195 Heritabilities were then estimated using these variance components as follows:

$$2196 \quad h^2 = \frac{\sigma_A^2}{\sigma_A^2 + \sigma_E^2} = \frac{\sigma_A^2}{\sigma_p^2},$$

2197 where σ_A^2 = direct additive genetic variance

2198 σ_E^2 =residual variance

2199 σ_p^2 = phenotypic variance

2200

2201

2202

2203 **2.3 Results**

2204 **2.3.1 Loin physical, meat quality and collagen characteristics**

2205 Phenotypic means, standard deviations, minimum and maximum values and coefficient of
2206 variation for loin physical traits are shown in Table 2.3, while those for meat quality and
2207 collagen characteristics are given in Table 2.4. A total of 11 loin physical and 13 meat quality
2208 traits were examined in this study. The loin physical traits had 500 observations except for Meat
2209 Trim Weight (MTW) and Fat Trim Weight (FTW) which had 455 and 456 respectively due to

2210 unavailability of phenotypic records for some individuals. Five hundred records are reported for
2211 meat quality traits except for intramuscular fat which had 464 records. Collagen solubility
2212 percent and total collagen had 499 observations while collagen insolubility percent had 492.
2213 Significance of fixed effects and random effect used for fitting mixed models for analysis are
2214 presented in Table 2.1 and 2.2.

2215 Loin Eye Weight (LEW) was on average 39% of the loin weight (1.4 ± 0.2 kg) (after adjusting for
2216 diaphragm weight) with a population mean of 0.5 ± 0.1 kg while thick fat weight (TFW) was 24%
2217 with a population mean of 0.3 ± 0.1 kg. Bone weight (BWT) had a mean of 0.2 ± 0.0 kg and
2218 accounted for 13% of the loin weight (LWT). The mean pH of the muscles after thawing was
2219 5.6 ± 0.1 while mean losses (CL) upon cooking to an internal temperature of 71°C was 25% and
2220 ranged from 19.0% to 36.8% while average Warner-Braztler shear force (WBSF) was
2221 $47.2\text{N} \pm 10.1$. Mean Moisture content (MC) measured after oven drying was 72% while crude
2222 moisture (FOM) measured with a FOSS Food Scan was 57%. Mean FOSS crude protein (FOP)
2223 was $16.8\% \pm 1.2$ while mean FOSS crude fat (FOF) and intramuscular fat content (IMF) were
2224 $24.1\% \pm 4.2$ and $8.1\% \pm 3.0$ respectively. The average combined-component percentage of loins in
2225 this study was 98.29. With respect to collagen characteristics, a mean of $18.0\% \pm 6.6$ was
2226 obtained for percent collagen solubility (SOLC) in raw meat while $82.2\% \pm 6.4$ was reported for
2227 collagen insolubility percent (ISOLC). Total collagen (TCOL) (mean= 3.1 ± 1.0 mg/g raw meat)
2228 ranged from 1.5 to 6.3 mg/g raw meat. The $L^* a^* b^*$ color space measured with a Minolta
2229 chroma meter produced mean values of 45.5 ± 3.0 for lightness (LOINL), 7.7 ± 1.3 for a^*
2230 (LOINA) and 3.5 ± 1.1 for b^* (LOINB) with hue (LOINH) and chroma (LOINC) means being
2231 24.6 ± 8.7 and 8.4 ± 1.2 , respectively. Of the color traits, hue had the highest coefficient of
2232 variation (35%) while L^* had the lowest 8%. As the ratio of the standard deviation to the mean,

2233 the coefficient of variation depicts the level of variability in relation to a population mean, the
2234 estimates obtained reveal that dispersion of MC was about 5 times less than for FOM while
2235 variability in insoluble collagen percent was approximately 3 times lesser than WBSF.

2236 **2.3.2 Heritabilities**

2237 Additive genetic variances, residual variances, and heritabilities with their standard errors are
2238 presented in Tables 2.5 and 2.6 for sub-primal and meat quality traits respectively. Estimates for
2239 collagen characteristics are also found in Table 2.6.

2240 Moderate heritabilities were obtained for sub-primal traits, ranging from 0.22 ± 0.10 for BWT to
2241 0.44 ± 0.11 for LEW although heritability estimates for bone percent (BP) (0.09 ± 0.08), LWT
2242 (0.19 ± 0.08) and TFW (0.16 ± 0.10) were low. Loin eye percent (LEP) and thick fat percent
2243 (TFP) were moderately heritable with estimates of 0.28 ± 0.12 and 0.42 ± 0.13 respectively. Meat
2244 trim weight (0.30 ± 0.12) and RTW (0.31 ± 0.11) were also moderately heritable while FTW
2245 (0.13 ± 0.10) was lowly heritable.

2246 Heritability estimates for meat quality traits were low to moderately heritable. Estimates ranged
2247 from 0.12 ± 0.09 for pH to 0.42 ± 0.12 for LOINA. Minolta color traits were moderately heritable
2248 ranging from 0.21 ± 0.10 for LOINB to the estimate stated earlier for LOINA. Heritability
2249 estimate for LOINL was 0.25 ± 0.11 .

2250 With respect to juiciness traits, moisture content (0.36 ± 0.11) was more heritable than crude
2251 moisture (0.21 ± 0.10). Warner-Braztler shear force value (WBSF) was moderately heritable
2252 (0.31 ± 0.12). The marbling traits, IMF and FOSS crude fat (FOF) were not heritable to the same
2253 degree, IMF (0.42 ± 0.13) had a higher heritability estimate than FOF (0.24 ± 0.10). Hue and
2254 Chroma are calculated based on LOINA and LOINB values and their heritabilities were

2255 0.21±0.10 and 0.36±0.12 which indeed gives an indication of the influence of these underlying
2256 constituent traits. Cooking loss (CL), which gives an indication of the level of shrinkage, lipid
2257 and moisture loss had a heritability estimate of 0.24±0.10.

2258 Of the traits describing collagen characteristics of the raw pork samples, collagen solubility and
2259 insolubility percent were lowly heritable with heritabilities of 0.12±0.10 and 0.15±0.10,
2260 respectively, while total collagen had a moderate heritability estimate of 0.33±0.12.

2261 **2.3.3 Correlations among traits**

2262 **2.3.3.1 Correlations among loin physical traits**

2263 The phenotypic and genetic correlations for loin physical traits are shown in Table 2.7.
2264 Generally, significant genetic correlations ($p < 0.05$) among loin physical traits were moderate to
2265 high, ranging between 0.48±0.23 to 0.93±0.23. There was a widespread negative correlation
2266 between traits related to muscle content/percent and fat content/percent.

2267 For LWT, a trait equivalent to loin sub-primal cut weight was highly correlated with LEW
2268 (0.87±0.14), MTW (0.93±0.23), FTW (0.64± 0.28), RTW (0.76±0.16) and BWT (0.77±0.26).

2269 There was a strong negative correlation between LEW and TFP (-0.60±0.16). LEW was
2270 moderately to highly correlated with BWT (0.45±0.23), MTW (0.61±0.17), RTW (0.60±0.16)
2271 and LEP (0.73±0.14); it was however not correlated with BP (-0.02±0.38). For TFW, it was
2272 highly negatively correlated (-0.76±0.14) with LEP and positively correlated with TFP
2273 (0.91±0.06). MTW had a high genetic correlation with RTW (0.92±0.08), BWT (0.64±0.22)
2274 and RTP (0.76±0.16) and a moderate negative genetic relationship with TFP (-0.50±0.22). The
2275 genetic correlations of FTW with RTW and LEP were 0.62±0.25 and -0.80±0.28 respectively.

2276 RTW was highly correlated with BWT (0.71±0.19) and RTP (0.84±0.11) but moderately

2277 correlated with TFP (-0.50 ± 0.23). Genetic correlations of BWT with LEP (-0.13 ± 0.35) and TFP
2278 (-0.25 ± 0.29) were not significant; BWT was however significantly correlated with BP
2279 (0.77 ± 0.23). A high negative correlation was obtained between LEP and TFP (-0.76 ± 0.10) and
2280 TFP was further negatively correlated with RTP (-0.66 ± 0.17).

2281 This study obtained moderately to high phenotypic correlations among the sub-primal traits.
2282 However, low phenotypic correlations were obtained between LWT with LEP (-0.20 ± 0.05), TFP
2283 (0.19 ± 0.05) and BP (-0.22 ± 0.04); LEW and FTW (0.11 ± 0.05), TFW and RTP (-0.27 ± 0.04),
2284 MTW and LEP (0.11 ± 0.05), RTW with LEP (-0.16 ± 0.05) and BP (-0.15 ± 0.05), BW and TFP ($-$
2285 0.13 ± 0.05), LEP and BP (0.12 ± 0.05).

2286 **2.3.3.2 Correlations among meat quality and collagen traits**

2287 Presented in Table 2.8 are the summaries of phenotypic and genetic correlations among meat
2288 quality and collagen traits considered in this study. In general, Minolta color traits were
2289 significantly correlated amongst themselves. Further, moisture traits were largely negatively
2290 correlated with fat traits.

2291 High negative correlations were obtained for pH with LOINL (-0.95 ± 0.35), LOINB ($-$
2292 0.87 ± 0.31), and LOINH (-0.93 ± 0.28). Muscle pH further had a strong positive correlation with
2293 MC (0.98 ± 0.27). A negative genetic correlation was obtained between muscle pH and CL ($-$
2294 0.65 ± 0.31). Cooking loss on other hand, had a strong positive genetic correlation with LOINB
2295 (0.69 ± 0.31) and moderate correlations with WBSF (0.45 ± 0.19) and LOINL (0.44 ± 0.19).

2296 With regard to CIE $L^*a^*b^*$ results, LOINL had a moderate to high genetic correlation with other
2297 color traits. It had a moderate correlation with LOINA (0.53 ± 0.20), LOINB (0.46 ± 0.22) and a
2298 high correlation with LOINH (0.85 ± 0.09). Minolta L^* was further strongly correlated with

2299 ISOLC (-0.71 ± 0.35) and SOLC (0.68 ± 0.30). For LOINA, it was highly correlated with LOINC
2300 (0.97 ± 0.01), FOM (0.62 ± 0.30), and moderately correlated with LOINB (0.49 ± 0.23). Genetic
2301 correlations of LOINB with LOINH, LOINC and IMF were 0.69 ± 0.15 , 0.64 ± 0.17 and
2302 0.61 ± 0.22 , respectively-all strong and positive. LOINB further had a strong negative correlation
2303 with MC (-0.76 ± 0.17). With respect to LOINH, a high negative correlation was with MC (-
2304 0.62 ± 0.21) although its relationship with other traits were generally moderate and not
2305 significant.

2306 Moisture content (MC) is a measure of the amount of water contained in the pork samples.
2307 Higher chemical MC was associated genetically with lower IMF content (-0.64 ± 0.14) and
2308 higher FOM percent (0.63 ± 0.22). A moderate genetic correlation was found for WBSF with
2309 SOLC (-0.38 ± 0.10) and ISOLC (0.42 ± 0.16). For FOF, it was highly genetically correlated with
2310 FOM (-0.98 ± 0.01), FOP (-0.93 ± 0.05) and IMF (0.79 ± 0.14). With respect to FOP, it was
2311 positively correlated with FOM (0.86 ± 0.09) and negatively correlated with IMF (-0.67 ± 0.18). A
2312 strong negative genetic correlation was also found between FOM and IMF (-0.83 ± 0.13). Soluble
2313 collagen percent (SOLC) was highly negatively correlated with TCOL (-0.63 ± 0.23) while
2314 ISOLC (0.66 ± 0.28) had a positive correlation with TCOL.

2315 Significant phenotypic correlations among Minolta color traits were generally moderate to high
2316 ranging from 0.58 ± 0.03 to 0.94 ± 0.02 although correlations of LOINL with LOINA (-
2317 0.24 ± 0.05), LOINA with LOINH (-0.12 ± 0.05) and LOINC with LOINH (0.15 ± 0.05) were low.
2318 Moisture content was moderately correlated with all the Minolta color traits phenotypically
2319 except with LOINA (-0.18 ± 0.05) which was low. Meat composition traits measured with the
2320 commercial FoodScan Meat analyzer were highly correlated among themselves phenotypically,

2321 ranging from 0.90 ± 0.01 between FOP and FOM, to -0.99 ± 0.00 between FOF and FOM.
2322 Phenotypic correlation found between FOM and IMF was high (-0.61 ± 0.03). TCOL had a
2323 moderate phenotypic correlation with SOLC (-0.46 ± 0.04) and ISOLC (0.48 ± 0.04).

2324 **2.3.3.3 Correlations of loin physical traits with meat quality and collagen** 2325 **traits**

2326 In Tables 2.9 and 2.10, the phenotypic and genetic correlations of loin physical traits with meat
2327 quality and collagen traits are provided. Generally, there were more significant phenotypic
2328 correlation ($p < 0.05$) than genetic correlations.

2329 Loin weight (LWT) had a moderate genetic correlation with WBSF and FOP with estimates of
2330 0.36 ± 0.12 and 0.50 ± 0.15 respectively. Loin eye weight (LEW) had moderate correlations with
2331 LOINA (-0.36 ± 0.12), FOF (-0.30 ± 0.10), FOP (0.52 ± 0.20) and TCOL (-0.32 ± 0.13). Thick fat
2332 weight (TFW) was moderately correlated with meat quality fat traits as its correlation with FOF
2333 and IMF were 0.55 ± 0.25 and 0.53 ± 0.26 respectively. Thick fat weight was further negatively
2334 correlated with CL (-0.41 ± 0.16) and FOM (-0.52 ± 0.18). With respect to MTW, it was highly
2335 correlated with CL (0.64 ± 0.27) and moderately with FOM (0.53 ± 0.26). A high positive genetic
2336 correlation was obtained between FTW and other fat traits like FOF (0.73 ± 0.24) and IMF
2337 (0.66 ± 0.31) but was negatively correlated with moisture traits like MC (-0.65 ± 0.28) and FOM
2338 (-0.70 ± 0.27). Moreover, FTW was also negative correlated with FOP (-0.71 ± 0.27). Correlations
2339 of RTW and BW were all not significant. LEP was estimated to have a negative moderately
2340 correlation with FOF (-0.41 ± 0.15), LOINA (-0.58 ± 0.19) but was positively correlated with
2341 LOINL (0.52 ± 0.26), FOP (0.47 ± 0.16) and FOM (0.38 ± 0.17). A moderate genetic correlation
2342 was obtained for FOP (-0.52 ± 0.23) and FOM (-0.53 ± 0.21) with TFP. Further, FOF and IMF as

2343 expected, were significantly correlated with TFP with estimates of 0.55 ± 0.20 and 0.43 ± 0.21
2344 respectively. RTP was highly correlated with CL (0.68 ± 0.23) and moderately correlated with
2345 IMF (-0.54 ± 0.23). Bone percent was moderately correlated with ISOLC (0.53 ± 0.10) and TCOL
2346 (0.51 ± 0.27) and lowly correlated with SOLC (-0.26 ± 0.11).

2347 In terms of their phenotypic correlations, TFW was moderately correlated with FOF
2348 (0.52 ± 0.03), FOP (-0.50 ± 0.03), FOM (-0.51 ± 0.03) and IMF (0.35 ± 0.04). loin eye percent on
2349 the other hand had moderate phenotypic correlations with FOF (-0.49 ± 0.04), FOP (0.47 ± 0.04),
2350 FOM (0.47 ± 0.04) and IMF (-0.38 ± 0.05), however, its correlation with Minolta color traits were
2351 low, ranging between -0.11 ± 0.05 with LOINH to 0.24 ± 0.05 with LOINA and LOINB.

2352 **2.4 Discussion**

2353 **2.4.1 Genetics of collagen characteristics**

2354 The mean total collagen (mg/g raw meat) obtained in this study fell within the range of the those
2355 in the literature reviewed, that is 2.6 to 5.0mg/g raw muscle (Correa et al., 2006; Therkildsen et
2356 al., 2002; Wheeler et al., 2000; Boutten et al., 2000). Correa et al. (2006) in their study of
2357 Canadian crossbred pigs with similar genetic makeup (Duroc \times (Landrace \times Large White)) as
2358 this study obtained mean total collagen slightly higher (4.0 mg/g wet tissue) than this study's
2359 mean. This difference could most probably be due to the fact they used a factor of 8 to convert
2360 hydroxyproline to collagen while this study used 7.14 as well as factors like study design and
2361 potential differences in breeding values. They further reported heat soluble collagen (%) ranging
2362 from 11.1% to 13.4%, a range less variable than the range obtained in this study. Heat soluble
2363 collagen percent gives an indication of the amount of heat labile cross-links in a sample and
2364 therefore a useful indicator of tenderness. Wheeler et al. (2000) obtained variable total collagen

2365 across different Large White composite barrow muscles ranging from 4.1mg/g raw muscle in the
2366 longissimus muscle to 7.1 mg/g raw muscle in the bicep femoris. These variations are most
2367 probably a function of anatomical location as highly exercised muscles tend to have more total
2368 collagen (Thakur et al., 2002). With respect to species-species comparison, total collagen in this
2369 study was generally lower than those reported in bovines (Cross et al., 1973; Hill, 1966; Raes et
2370 al., 2003), turkeys (Fernandez et al., 2001) and chickens (Liu et al., 1996; Sakakibara et al.,
2371 2000; Roy et al., 2007); although some studies in chickens reported relatively lower values
2372 (Nakamura et al., 2003; Nakamura et al., 2004). The ratio of soluble to insoluble collagen gives
2373 an indication of how tender or tough a muscle cut will likely be. Lebret et al. (1998) obtained
2374 similar (17.4%) collagen solubility percent from the loins of crossbred (Large White × (Large
2375 White × Landrace)) pigs slaughtered at 5.5 months. Correa et al. (2006) obtained lower (12.3%)
2376 collagen soluble percent, however, the approximate ages of the animal were not provided and as
2377 age influences solubility percent (Fang et al., 1999), interpretation of this result is difficult. Fang
2378 et al. (1999) in their study of the relationship between connective tissue development and
2379 Landrace growth and development reported heat soluble collagen percent of 24 at 6 months of
2380 age. Heat soluble collagen percent of this study was higher than those generally reported in beef
2381 studies (Nakano and Thompson, 1980; Dikeman et al., 1986) and lower than those reported for
2382 chickens and turkeys (Fernandez et al., 2001, Liu et al., 1996). Nishimura et al. (1999) in their
2383 study of the Japanese Black cattle, known for its high marbling content, obtained solubility
2384 percent 28% in 9-month olds and 16% solubility percent for 24-month olds.

2385 To the author's knowledge, this is the first study to report on the heritabilities of collagen
2386 solubility and insolubility percentages in pork. The low heritabilities of SOLC and ISOLC
2387 suggest that, of the observed variations in the population, only a small fraction is due to variation

2388 in genetics (Visscher et al., 2008) and this could limit genetic improvement of these traits. The
2389 moderate heritability estimates of TCOL (0.33) obtained in this study indicates a potential for
2390 making genetic progress in this trait. This estimate was, however, higher than the estimate
2391 reported by Suzuki et al. (2005) (0.23).

2392 The high positive genetic correlation of LOINL with SOLC coupled with its high negative
2393 genetic correlation with ISOLC suggest that selecting for LOINL with genomic tools over time
2394 will increase solubility percent and decrease insoluble collagen thereby potentially improving
2395 tenderness. However, LOINL has an antagonistic relationship with pH and therefore selecting for
2396 LOINL will lead to a decrease in pH over time, which may potentially lead to the denaturation of
2397 myofibrillar proteins and subsequently lead to a pale, soft and, if the correlation between pH and
2398 CL is taken into account, exudative meat. Indeed, Marchi et al. (2011) reported a 25 % increase
2399 in soluble collagen in PSE meat compared to normal meat. The results of this study brings to
2400 bear the genetic relationships among ISOLC, pH, LOINL and SOLC. Thus, the high correlation
2401 of LOINL with SOLC and ISOLC presents a potential to predict SOLC and ISOLC with LOINL
2402 in pigs although appropriate selection indexes may need to be established. Secondly, collagen
2403 fibers are fluorescent (Biela et al. 2013) when excited with ultra-violet light. Indeed, methods
2404 have been developed to measure biological age in humans and rats through collagen fluorescence
2405 (Odetti et al., 1992). With respect to livestock, Swatland and Findlay (1997) developed an on-
2406 line UV fibre optic fluorescence probe to predict beef toughness based on detection of collagen-
2407 linked fluorescence. They found that narrow fluorescence peaks frequencies with smaller peak
2408 widths were higher in tougher parts and peak width was significantly correlated with tenderness
2409 ($r=0.57$). It is therefore interesting that genetic relationships were found between luminosity and
2410 collagen characteristics. It is plausible to suggest that genes influencing muscle luminosity also

2411 to a degree influence collagen characteristic, although further studies may be needed to establish
2412 this relationship.

2413 Also, to the author's knowledge, this is the first study to report a genetic relationship of WBSF
2414 with SOLC and ISOLC. The moderate genetic correlation of WBSF with SOLC and ISOLC
2415 indicate a possible influence of similar gene networks on these traits. Indeed, the relationship
2416 suggests that selecting for WBSF may in time increase collagen solubility percent and decrease
2417 collagen insolubility percent ultimately resulting in a tender product. As expected, TCOL was
2418 highly correlated with SOLC and ISOLC as TCOL was calculated based on soluble and
2419 insoluble collagen mg/g raw meat.

2420

2421 **2.4.2 Loin physical and meat quality traits**

2422 Loin sub-primals are obtained through further fabrication of primals into manageable cuts. They
2423 are easier to handle and more cost effective to transport (Underly, 2012). Pork is mostly sold as
2424 packaged sub-primals on a fresh or processed basis at the retail level (Hodgson et al., 1991) and
2425 therefore taking measurements at this level of fabrication is very important as breeding
2426 objectives in pigs are driven by feedback from these cuts (Van Wilk et al., 2005).

2427 Selection for rapid growth rate has led to an increase in carcass lean percent (Newcom et al.,
2428 2002). Even as early as the 1960s, Buck et al. (1962) noted that during pig carcass assessments,
2429 lean meat percent was considered of utmost importance. It was therefore not surprising that LEP
2430 was almost twice that of TFP in this study. This agrees with the results obtained by Van Wijk et
2431 al. (2005) in their study of commercial crossbred pigs as they reported 59.3mm mean loin eye
2432 depth vs 25.1 mm mean back fat depth. Newcom et al. (2002) also reported similar observation
2433 in their study of lean efficiency pigs, in which the ratio of loin muscle area to carcass back fat

2434 was approximately 3:1. Furthermore, when compared to this study, Correa et al. (2006) reported
2435 higher loin eye and fat percent but lower bone percent in crossbred Canadian pigs (Duroc ×
2436 (Landrace × Yorkshire)) slaughtered at approximately the same slaughter weight as this study.
2437 This difference could be a function of the fact that they took their measurement on the whole loin
2438 primal while this study worked with a sub-primal. Devol et al. (1988) obtained only a 19mm
2439 difference between loin eye area and back fat while Van Wijk et al. (2005) recorded 34mm. This
2440 difference provides perspective on the level of improvement that has occurred over the years
2441 although breed differences may need to be considered when making such comparisons. Devol et
2442 al. (1988) however did not report on breed composition of their population.

2443 Intramuscular pH is a measure of muscle acidity and is crucial in the prediction of major meat
2444 quality defects. Average pH obtained was similar to those reported by Therkildsen et al. (2002),
2445 Van Wijk et al. (2005), Hodgson et al. (1991) and De Vol et al. (1987) even though pH
2446 measurements for this study were taken on frozen-thawed samples while those from the
2447 aforementioned studies were on unfrozen samples. Monin and Sellier (1985) also reported
2448 similar results in the longissimus dorsi of Large Whites, however, their values for halothane
2449 negative and positive Pietrains and Hampshires were lower: 5.45, 5.41 and 5.40 respectively. In
2450 a review by Bendall and Swatland (1988), the mean ultimate pH (pHu) of 9,600 Alberta
2451 crossbred pigs in 1978 was 5.75 with 45% of the animals having pHu fall within ranges of 5.7
2452 and 6.5. This is slightly higher than the value obtained in this study. However, this difference
2453 could be due to factors such as breed composition, sampling and measurement approaches.

2454 Juiciness is of critical significance to eating quality and is influenced mainly by the quality of the
2455 raw meat and cooking procedure (end-point temperature, heating method) (Aaslyng et al., 2003).

2456 This study used an end-point temperature of 71°C. Indeed, increasing end-point temperature is
2457 known to decrease juiciness (Heyman et al., 1990). Cooking loss is a trait intimately linked to
2458 juiciness and with tremendous influence on juiciness as it accounted for approximately 75% of
2459 the variation in juiciness scores according to a study by Bouton et al. (1975). Hodgson et al.
2460 (1991) obtained a slightly higher cooking loss percent (28%) which is mostly likely due to the
2461 final meat sample internal temperature used in their study being 75°C. Therkildsen et al. (2002)
2462 and Kristensen et al. (2002) reported higher cooking loss percent ranging between 29.9 to 33.4%.
2463 It is however worth noting that in their study, chops were cooked for 60min at 80°C. Further,
2464 their treatment effects (ageing period and feeding strategy) may have influenced these averages.
2465 Indeed, the influence of endpoint temperature in cooking loss (%) cannot be overstated. In a
2466 study by Heymann et al. (1990), changing endpoint temperature from 65.6 °C to 82.2°C resulted
2467 in a 35% increase in cooking loss. This necessitates a standard endpoint temperature in research
2468 studies to ensure ease of comparison such as those recommended by AMSA (1995).

2469 A trait closely linked to cooking loss is Warner-Bratzler shear force (WBSF) (Barbanti and
2470 Pasquini, 2005; Møller, 1981). Warner-Bratzler shear force values are widely accepted as
2471 important indicators of pork tenderness (Cai et al., 2011). Shackelford et al. (2004) reported a
2472 significantly high correlation ($r = -0.66$, $p < 0.001$) between trained sensory panel tenderness
2473 scores and WBSF of meat for porcine longissimus muscle. Warner-Bratzler shear force values
2474 reported by Hodgson et al. (1991) of 34N, De Vol et al. (1988) of 37N, Shackelford et al. (1999)
2475 of 38N, Heymann et al. (1990) of 25N, and Therkildsen et al. (2002) of 34N were lower than the
2476 mean obtained in this study. However, Kristensen et al. (2002) reported values ranging between
2477 57-70N while Oksbjerg et al. (2000) obtained a value of 50N. Oksbjerg et al. (2000) indeed
2478 compared meat quality traits between fast growing and slow growing Danish Landrace pigs and

2479 found the former to have slightly higher shear force values and cooking loss (%), giving a hint of
2480 the effect of selection for fast growth on meat quality traits. Huffman et al. (1996) suggested a
2481 tenderness threshold of 40.2N for beef steaks, as tenderness ratings corresponding to values less
2482 than or equal to 40.2N were rated as having acceptable tenderness by consumers in both
2483 restaurant and home settings. If this threshold is to be used as a standard in pork then, on
2484 average, loin chops in the present study were tough. In general, comparing WBSF values
2485 reported by studies in the 1990s (Heymann et al., 1991; Hogson et al., 1990) with those of the
2486 2000s (Oksbjerg et al., (2000); Kristensen et al., 2002), one can only get the impression that pork
2487 is getting tougher. Shear force values reported for wild boars by Cifuni et al. (2014) ranged
2488 between 44-53N and thus falls within the range of values reported for domestic pigs.

2489 Shackelford et al. (2004) employed similar WBSF methodologies as this study but reported
2490 lower WBSF values compared to this study. Their research study looked, in part, at the effect of
2491 cooking method on WBSF and found belt grill cooking to have significantly ($p<0.05$) lower
2492 mean WBSF value (3.9 kg) than open hearth broiling (4.1 kg). The WBSF coefficient of
2493 variation (CV) of this study (24) falls within the range obtained by Shackelford et al. (1999)
2494 (20.8-34.4). Jones et al. (1980) indeed reported animal to animal variability to be the major
2495 source of variation in 14 out of 19 pork quality traits they studied including WBSF. These
2496 variations are most probably the results of differences in a number of known and unexplained
2497 factors of either genetic and/or environmental origin that interact and dictate the effects of
2498 metabolic processes at the time of slaughter and after slaughter (Brocks et al., 2000). The
2499 relatively high SD and CV for some traits in this study are likely due to the wide variation in
2500 phenotypes in the populations studied. Similar high variation was reported by Hodgson et al.
2501 (1991) and DeVol et al. (1988) on meat quality traits of pork longissimus dorsi. The CV reported

2502 for ether extractable material in this study was lower than the CVs of the fat percentages reported
2503 by Hodgson et al. (1991). Indeed, DeVol et al. (1988) in their evaluation of random pork
2504 carcasses in a commercial packing plant in the US concluded that there is substantial variability
2505 in carcass composition and meat quality traits of pork moving into the market. Variability in this
2506 circumstance is not advantageous as consumers usually prefer consistency in the food products
2507 they consume.

2508 Mean chemical moisture percent and its low level of variability was similar to the results of
2509 DeVol et al. (1988), Hodgson et al. (1991), Ramsey et al. (1989) and Novakofski et al. (1989).
2510 Crude moisture content obtained with the FOSS FoodScan was lower (57%) than those obtained
2511 after freeze drying (72%). Woolley et al. (2016) made similar observations in their comparison
2512 of chemical analysis and FOSS FoodScan for proximate composition of bovine inside round lean
2513 and fat trimmings. However, they found a high correlation between chemical moisture and FOSS
2514 moisture ($r= 0.993$). The combined-component percentages reported by other researchers who
2515 have studied the proximate composition of porcine longissimus muscle (Hodgson et al. 1990;
2516 Bacon et al., 1989) were 98.54 and 98.94 respectively, which agrees with this study. The large
2517 difference between the mean ether extractable material and mean FOF is most likely due to the
2518 fact that loin trimmings were used FOF measurements while for IMF, whole loin chops cut into
2519 1cm^3 pieces and oven dried were used. Compared to this current study, Bacon et al. (1989)
2520 reported lower protein % (15.7) and moisture % (50.9) but higher fat % (32.5) using a FOSS
2521 Superscan complete meat analyzer. Crude protein % obtained by the Kjeldahl method in the
2522 studies of Ramsey et al. (1989) and Novakofski et al. (1989) 22.4 and 27.2, respectively, was
2523 higher than the mean obtained in this study with the FOSS meat analyzer. Intramuscular fat is
2524 found between muscle fibers and its greatest accumulation occurs during the later phases of

2525 physiological maturity. This trait is important due to its influence on fresh meat marketing and
2526 for meeting export standards especially in loin cuts (Lonergan et al., 2019). Ramsey et al. (1989)
2527 reported a high ether extract % (8.5) for their high marbling groups while their moderate groups
2528 had a mean of 5.5 which suggests that the mean ether extract of this study (8.1) falls within the
2529 range of moderate to abundant. The moderate-abundant marbling level of this study could be due
2530 to the Duroc sire line included in the mating system, as high levels of marbling have been
2531 reported for Durocs (Lonergan et al., 2019). With the exception of this study, all other research
2532 studies reviewed reported low ether extract % (DeVol et al., 1988; Novakosfki et al., 1989;
2533 Heymann et al., 1990).

2534 **2.4.3 Freeze-thawing affects meat colour**

2535 Meat products on retail shelves are generally selected or rejected by consumers based on their
2536 appearance and color (Brewer et al., 2006). Therefore, the ability of meat suppliers to create and
2537 maintain desirable color characteristics is essential (AMSA, 2012). Color assessment, either
2538 through instrumental or visual appraisals, is critical for tracing and correcting product
2539 development problems as well as detecting quality defects in meat research (Yancey and Kropf,
2540 2008). The state of the primary red meat pigment, myoglobin, determines muscle color (AMSA,
2541 2012; Brewer et al., 2006). However, factors such as nutrition, genetic background, species, post
2542 slaughter storage, muscle conversion to meat, lighting and display all play a role in determining
2543 perceived meat color (AMSA, 2012). Kristensen et al. (2002) and Friese et al. (2005) reported
2544 higher values for the Minolta color parameters regardless of their treatment groups. This is most
2545 probably due to the fact that they took their measurements of fresh pork chops as well as
2546 differences in blooming (myoglobin oxygenation) times. Kristensen et al. (2002) bloomed for
2547 1hr at 3°C while Freise et al. (2005) bloomed overnight at about 2°C, however, in this current

2548 study, chops were bloomed for about 1hr at 4°C. Brewer et al. (2001) in fact reported a
2549 significant effect of bloom time on the Minolta color parameters. Typical PSE pork Minolta L*
2550 is 61.0 (Friese et al., 2005), thus the LOINL range of this study (37.2-55.2) suggest that none of
2551 the loin samples exhibited PSE characteristics. Hue angle is a measurement of the level of
2552 deviation off the true red axis (a^*) in CIE $L^*a^*b^*$ color space (Brewer et al., 2006). It is
2553 computed from a^* and b^* values while chroma indicates how dull or vivid a color is (AMSA,
2554 2012). Wilkinson et al. (2006) studied the influence of modified atmosphere packaging on pork
2555 longissimus muscle and reported slightly similar chroma (8.5) and hue (22.7) as this study under
2556 their 100% CO₂ treatment group before blooming. This similarity could be due to a number of
2557 factors, including the frozen thawed state of this study's samples. Wilkinson et al. (2006) further
2558 reported increased hue and chroma values after blooming although their bloom time was not
2559 reported. Since increasing hue angle indicates that samples become less red (Brewer et al., 2006),
2560 the samples of this study were on average less red than those of the aforementioned study. This is
2561 most likely due to the fact that in fresh muscles cuts, metmyoglobin reducing activity (MRA) is
2562 high and thus formed metmyoglobin is immediately reduced to deoxymyoglobin and
2563 subsequently oxygenated during blooming. However, during aging or freeze-thaw treatment,
2564 metmyoglobin activity reduces leading to an accumulation of metmyoglobin on the meat surface.
2565 (Lanari and Cassen, 1991; Bekhit and Faustman, 2005). Also, co-factors like NADH may leach
2566 out with the exudates during thawing or be used for reactions not associated with MRA
2567 (Abdallah et al., 1999).

2568 **2.4.4 Moderate to high heritability estimates of loin physical and meat quality**
2569 **traits presents potential for improving these traits**

2570 One key inquiry in biology is determining whether the variations observed in a trait of interest is
2571 due to biological and/or environmental factors (Hill et al., 2008). The concept of heritability is
2572 key to answering this inquiry. Heritability is a population parameter that provides an estimate of
2573 the proportion of the total phenotypic variance for a trait of interest in a population, measured at
2574 a specified age period or time that is due to additive genetic variation (Hill et al., 2008). To the
2575 authors knowledge, no literature on heritability estimates of the loin physical traits measured this
2576 study were available apart from those presented in an earlier publication from the larger study
2577 (Miar et al., 2014). The generally moderate to high heritability estimates of loin physical traits
2578 suggests a potential for improving these traits through the traditional breeding approaches.
2579 However, genomic selection, with its high accuracy on lowly heritable traits (Gorjanc et al.,
2580 2015) will be appropriate for loin physical traits like LWT, FTW, TFW and BP. Indeed, the low
2581 heritability of LWT and BP could be due to sampling errors during sub-primal excision as
2582 differences in rib numbers of pigs has been reported to range between 13.5 to 17.0 (Fredeen and
2583 Newman, 1962). Thus, this variation could cause errors in sampling sites and result in faulty
2584 representations. Generally, however, traits related to carcass composition are moderate to highly
2585 heritable (Van Wilk et al., 2005; Enfield and Whatley, 1961; Ciobanu et al., 2011).

2586 The low heritability estimate of muscle pH fell within the range (0.00-0.39) reported in the
2587 literature reviewed (Bidanel et al., 1994; Jensen et al., 1967; Knapp et al., 1997; Hovenier et al.,
2588 1992; Suzuki et al., 2005). In fact, it was in agreement with the estimates reported by Van Wijk
2589 et al. (2005) (0.11) and Hermesch et al. (2000) (0.14). Breed difference can have a tremendous
2590 influence on muscle pH estimates as Johansson, (1987) reported higher heritability estimates of

2591 pH in Landrace (0.27) and Large White (0.30). Low genetic variation, as obtained for muscle
2592 pH, limits our efforts to rank sires for their genetic endowment when using it as an indicator trait
2593 (Van Wijk et al., 2005). The low heritability of pH indeed seemed not to be species specific, as
2594 Mortimer et al. (2014) obtained an estimate of 0.08 in Sheep, Hamzah et al. (2016) reported 0.09
2595 in fish, Aass (1996) obtained 0.19 for dual purpose cattle and Bihan-Duval et al. (2003) had 0.16
2596 for turkey toms.

2597 Cooking loss (CL) is a measure of water holding capacity that quantifies the fluid released from
2598 meat samples upon heat application with or without application of external forces (Hamm,
2599 1986). Cooking loss heritability estimated in the present study (0.24) was higher than estimates
2600 reported by Lo et al. (1992) (0.06), Suzuki et al. (2005) (0.09), Scheper (1978) (0.10) and De
2601 Vries et al. (0.11) but lower than estimates of Lee et al. (2015) (0.45). These differences could
2602 most probably be due to differences in statistical modelling techniques implemented in the above
2603 studies as well as differences in population structure. Bidanel et al. (1994) however reported
2604 similar cooking loss percent heritability estimates (0.24) as this study despite their study
2605 population being Pietrain pigs. Differences in heritability estimates of cooking loss due to breed
2606 have also been reported by Malmfors and Nilson (1978) as they did not obtain additive genetic
2607 variation in Landrace ($h^2=0.00$) in Landrace but reported a high heritability estimate in Large
2608 White ($h^2 =0.51$). Heritability of a trait to a large extent depends on the originating population
2609 since variations in non-additive and additive genetic aspects as well as environmental variances
2610 are population specific. Further, genetic variances are dependent on alleles influencing a trait of
2611 interest in a population, variants effect sizes, mode of gene action and allele frequencies; and as
2612 the above variables can vary across populations, heritability of a trait in one population from a
2613 theoretical stand point does not necessarily predict heritability of that trait in a different

2614 population (Visscher et al., 2008). However, in actuality, heritability of similar traits for the most
2615 part tend to be notably similar in different populations of the same species and in some cases
2616 across species (Visscher et al., 2008).

2617 Heritability estimates of LOINL were in agreement with estimates reported by Larzul et al.
2618 (1997) (0.23) and Cherel et al. (2012) (0.20) while Van Wijk et al. (2005) obtained slightly lower
2619 estimates (0.18). Van Wijk et al. (2005) further reported lower LOINA (0.21) and LOINB (0.15)
2620 estimates. These differences could be a result of statistical modelling differences but this is
2621 unlikely as Van Wijk et al. (2005) measured tristimulus values under C illuminant while this
2622 study used D65 illuminant and Sun et al. (2017) reported a high correlation ($r>0.96$) between the
2623 aforementioned illuminants. Lee et al. (2015) and Cherel et al. (2012) also reported similar
2624 LOINA and LOINB estimates as this study. Furthermore, heritability estimates of LOINL (0.28)
2625 and LOINA (0.43) reported for Durocs by Gjerlaug-Enger et al (2010) agreed with this study's
2626 estimates although their LOINB (0.33) was higher. Cameron (1990) obtained slightly higher
2627 estimates for LOINC (0.42) although that study's LOINH (0.19) was lower than this study. The
2628 generally moderate heritabilities of the Minolta color traits in this study suggests that selection
2629 for meat color traits will lead to genetic changes in the average population means for meat color
2630 resulting in improved attractiveness of pork. It must be noted that meat color is a critical visual
2631 feature influencing the meat quality, as it influences consumer acceptability and purchasing
2632 decisions (Karumathil et al., 2016; Lonergan et al., 2019).

2633 For moisture content measurement, the estimated heritability was within range of the estimates
2634 reported by Gjerlaug-Enger et al. (2010) (0.31-0.50) while Carbling et al. (2015), Cameron
2635 (1990) and Lo et al. (1992) obtained lower estimates of 0.27, 0.26 and 0.14 respectively. Jensen
2636 et al. (1967) on the other hand, reported a high (0.81) heritability estimate for moisture content.

2637 These differences in estimates could most probably be due to differences in population structure,
2638 variations in sample size, statistical modelling, as well as differences in trait measurement
2639 procedures. For instance, Jensen et al. (1967) used hot air oven 80°C for 48hrs in the moisture
2640 content assay while the current study freeze-dried samples for 5days.

2641 Warner-Bratzler shear force is a measurement that serves as a valuable indicator of meat
2642 tenderness (Bratzler, 1932, Warner, 1952, Hovenier et al., 1993) and thus, a trait of tremendous
2643 importance in the food industry. The moderate heritability estimate of shear force conforms to
2644 estimates reported by Malmfors and Nilsson (1978) (0.34); Jensen et al. (1967) (0.25); Lee et al.
2645 (2015) (0.39); Cherel et al. (2012) (0.26) and Hovenier et al. (1993) (0.30). However, Suzuki et
2646 al. (2005) and Jung et al. (2015) obtained higher estimates of 0.45 and 0.56 respectively. Cabling
2647 et al. (2015) and Lo et al. (1992) on the other hand reported low estimates of 0.04 and 0.17
2648 respectively. These differences could be due to population differences and differences in
2649 measurement approaches. For instance, Cabling et al. (2015) used a crosshead speed of 400
2650 mm/min while this study used 200mm/min. Wheeler et al. (1997) in point of fact reported that
2651 Warner-Bratzler shear values decreased with increasing shearing cross head speed.

2652 **2.4.5 Additive genetic variance for intramuscular fat makes a substantial** 2653 **contribution to its total phenotypic variance**

2654 The moderate heritability estimate of IMF (0.42) was in conformance with estimates reported by
2655 Scheper (1978) (0.35), Suzuki et al. (2005) (0.39), De Vries et al. (1994) (0.41) and Larzul et al.
2656 (1997) (0.44). Knapp et al. (1997) obtained varying estimates for different pig breeds, with the
2657 Large White, Landrace and Pietrain having estimates of 0.38, 0.67 and 0.42 respectively. Enfield
2658 and Whatley (1961), Jensen et al. (1967) and Allen et al. (1966) reported much higher estimates

2659 of 0.63, 0.86 and 0.70 respectively. Looking at the reduction in IMF heritability estimates over
2660 time, one may be led to the opinion that selection against fat traits over time may have led to
2661 changes in allele frequencies of some genes related to IMF although studies need to be done to
2662 verify this assertion. However, the general trend of moderate to high estimate of IMF indicates
2663 that a substantial proportion of this trait is determined by additive genetic effects (Lo et al.,
2664 1992). Thus, the moderate-high estimate obtained in the present study suggests that additive
2665 genetic variance for IMF makes a substantial contribution to its total phenotypic variance and
2666 thus, there is tremendous potential to improve this trait through selective breeding. In terms of
2667 IMF heritability in other livestock species, Mortimer et al. (2014) and Mateescu et al. (2015)
2668 reported similar heritability estimates in Australian crossbred sheep (0.48) and Angus cattle
2669 (0.38) respectively, as those reported for swine breeds.

2670 The high negative correlation of FOSSF with FOSSM and FOSSP agrees with results obtained
2671 by Gjerlaug-Enger et al. (2010) for Durocs. This finding supports the established fact that an
2672 antagonistic relationship exists for fat with muscle and moisture content. Indeed, Peñagaricano et
2673 al. (2015) through a scan of the porcine genome showed that muscularity and fat deposition are
2674 influenced by the additive pleiotropic effects of a genomic region on SCC6. They also reported
2675 on 7 significant eQLT's in this region many of which play vital roles in cell differentiation and
2676 proliferation. The antagonistic relationship of moisture and muscular fat has also been reported
2677 in fish (-0.89) by García-Celrán et al. (2015) and Cattle (-0.99) by Sakuma et al. (2017).
2678 Heritabilities of meat composition traits as measured with FOSS FoodScan NIR
2679 spectrophotometer were much lower than those reported by Gjerlaug-Enger et al. (2010) as they
2680 obtained estimates of 0.54, 0.50 and 0.62 for FOP, FOM and FOF respectively. It is worth stating
2681 that these estimates were obtained from Duroc breeds as well as the fact that they used ground

2682 whole loin chop for measurements while the current study used trimmings. Even though Scheper
2683 (1978) used chemical analysis to estimate protein percent, the heritability estimate reported in
2684 that study (0.22) was similar to that found in this study.

2685 **2.4.6 Continuous selection for lean percent will negatively impact correlated** 2686 **fat traits**

2687 Carcass and meat quality characteristics in terms of its lean to fat ratio, tenderness, color and
2688 water holding capacity are determined for the most part, by structural and biochemical features
2689 of muscle and meat (Lonergan et al., 2019). However, on the fundamental level, these
2690 characteristics, for the most part, are determined by background genes. At the core of this current
2691 study was testing the hypothesis that genetic relationships exist between and among loin
2692 physical, meat quality and collagen traits. Genetic correlations provide a measure of the
2693 magnitude of association between breeding or genotypic values of individuals. Thus, it is
2694 valuable for describing how traits are connected at the genetic level and for predicting the impact
2695 of selection on a trait of interest and the changes that occurs in correlated traits (Hill, 2001). It is
2696 caused by linkage disequilibrium between genes impacting a trait and pleiotropic gene effects,
2697 although the former in most cases is transient while the latter leads to stable correlations
2698 (Goodnight, 2010). Phenotypic correlation on the other hand is a population measure that
2699 provides an estimate of the extent to which two metric traits co-vary among a population of
2700 related individuals (Conner and Hartl, 2004; Hill, 2001). Genetic and environmental correlations
2701 are two components that make up phenotypic correlation which could make phenotypic
2702 correlations of limited interpretative value as differences in environmental factors could impact
2703 its estimates. Phenotypic correlations are determined mainly by additive genetic correlation when
2704 heritability estimates are high while with lowly heritable traits, environmental correlations are

2705 paramount (Conner and Hartl, 2004). The moderate to high significant phenotypic correlations
2706 generally obtained among the loin physical traits suggest a potential for prediction among loin
2707 physical traits in the population of the present study.

2708 Even though meat quality is increasingly becoming important, meat yield is still crucial for
2709 carcass valuation as well as primal and sub-primal pricing (Van Wijk et al., 2005). Pork loin is
2710 one of the most valued cuts of a porcine carcass (Van Wijk et al., 2005) and indeed, one of the
2711 most tender parts of the carcass. The generally moderate to high genetic correlations between the
2712 sub-primal traits obtained in this study agree with the results of Newcom et al. (2002). The
2713 favorably high genetic correlation of LWT with LEW, MTW and RTW indicate a positive effect
2714 of selection for increased loin weight on muscling traits. However, LWT was also positively
2715 highly correlated with BWT, which is not a favorable correlation since selecting for increased
2716 loin weight will potentially also increase bone weight, a trait of little economic value. It must
2717 however be noted that bone weight and integrity is of vital importance from an animal welfare
2718 standpoint as it prevents broken bones from occurring during transport and handling.

2719 The high negative correlation between LEW and TFP and between TFW and LEP supports the
2720 results of other research studies. Newcom et al. (2002) also reported a negative correlation
2721 between carcass back fat and loin muscle although it was moderate (-0.43) while Johnson et al.
2722 (1999) reported a low correlation (-0.27). Furthermore, Van Wijk et al. (2005) reported a
2723 negative genetic correlation for back fat with whole loin weight (-0.41) and boneless loin weight
2724 (-0.60). Hovenier et al. (1992) also reported a high negative correlation (-0.71) between lean
2725 meat content and back fat thickness. All these studies used different pig breeds with varying
2726 population structures; however, they all obtained a negative genetic relationship between
2727 muscling traits and fat traits which are in agreement with the findings of this study and thus

2728 suggest that exercising continuous selection on lean percent could have an undesirable impact on
2729 correlated fat traits and indirectly affect eating quality. To the author's knowledge, this is the
2730 first study to report on the genetic correlations of meat trimmings. FTW, which was essentially
2731 fat trimmings from the loin was highly negatively correlated with LEP while MTW, which was
2732 essentially meat trimmings from the loin was moderately negatively correlated with TFP. These
2733 findings in a sense indicate that regardless of whether samples used were whole cuts or
2734 trimmings, plausible results can still be obtained. The results of Van Wijk et al. (2005) partially
2735 affirm this point as they obtained high genetic correlations between primals and sub-primals.

2736 **2.4.7 Linkage disequilibrium and/or pleiotropic gene effects potentially exist** 2737 **among traits related to the PSE defect**

2738 To integrate meat quality traits into swine breeding programs, relatively cheaper and objective
2739 measurements should be readily available to researchers and breeders, especially when routine
2740 data collection is necessary. The strong negative genetic correlation between pH and cooking
2741 loss obtained in this study indicates the potential of pH as an indicator trait for cooking loss (%).
2742 This result agrees with those reported by Bidanel et al. (1994) (-0.72) and De Vries et al. (1994)
2743 (-0.82), further suggesting that effective selection for higher pH may increase water holding
2744 capacity. Gjerlaug-Enger et al. (2010) and Cameron (1990) reported moderate negative genetic
2745 correlations between pH and LOINL with estimates of -0.44 and -0.38 respectively. The
2746 directions of the relationships are in congruity with this study although this current study
2747 reported higher estimates. De Vries et al. (1994) and Van Wijk et al. (2005) also reported high
2748 genetic correlations of -0.60 between pH and LOINL. A high genetic correlation of pH and L^*
2749 has also been reported in poultry (Bihan-Duval et al. 2003). Muscle pH of this study was further,
2750 highly correlated with moisture content. The relationship of pH with LOINL and MC highlights

2751 the economically important meat defects PSE and DFD, as the correlations indicate that selection
2752 for high pH will lead to high moisture content muscles and a darker color while low pH meat
2753 will tend to be exudative and pale and this supports the conclusions of Van Wijk et al. (2005). A
2754 high correlation of pH with ISOLC (0.76) and SOLC (-0.72) was obtained although associated
2755 with a high standard error. This relationship merits further investigation with a larger sample size
2756 as this may lead to a reduction in the magnitude of the standard error. Muscle pH is an easy to
2757 measure trait and thus, a good indicator trait candidate for the relatively expensive and difficult
2758 to measure traits highly correlated with it in this study.

2759 De Vries et al. (1994) obtained similar moderate genetic correlations for CL with L^* (0.47) and
2760 WBSF (0.57) as this study. Suzuki et al. (2005) on the other hand, obtained a low negative
2761 correlation (-0.22) which could be due to differences in population composition and statistical
2762 modelling. The high positive genetic relationships between LOINB and CL offer the possibility
2763 of using LOINB, a relatively easy to measure trait, to predict cooking loss in breeding programs,
2764 as the correlated response of selecting against LOINB will lead to a reductions in cooking losses
2765 and therefore an increase in water holding capacity.

2766 The strong genetic correlations of LOINB with LOINH and LOINC as well as LOINA with
2767 LOINC is expected since calculation of LOINH and LOINC is based on a^* and b^* coordinates.
2768 Van Wijk et al. (2005) and Gjerlaug-Enger et al. (2010) reported a high genetic correlation for
2769 LOINB with LOINL (0.89) and LOINA (0.79), higher than the moderate correlations obtained in
2770 this study. The correlation between MC and FOM (0.63) as well as IMF and FOF (0.79) is
2771 expected as they are just different measuring approaches for the same trait, moisture content in
2772 the case of MC and FOSSM and fat content in the case of IMF and FOSSM. It is quite
2773 interesting that these high correlations were obtained even though different parts of the loin were

2774 used for the measurements. However, the variation in correlation of these traits with other traits
2775 under consideration suggests that they must not be considered as identical traits. The high
2776 genetic correlation between MC and IMF obtained in this study were of the similar order of
2777 magnitude as those reported by Jensen et al. (1967) (-0.95), Malmfors and Nilsson (1978) (-
2778 0.72), Allen et al. (1966) (-0.78) and Cameron (1990) (-0.68) suggesting that similar background
2779 genes affect these traits.

2780 The correlation estimates obtained for TFW with CL, MC and IMF suggest that genetic selection
2781 for improved back fat is expected to result in pork with decreased cooking loss and moisture
2782 content and increased intramuscular fat content. Indeed, Jensen et al. (1967) reported a strong
2783 positive genetic relationship between IMF and pork organoleptic properties, indicating that
2784 improved IMF will improve eating quality. In a study with Duroc and Landrace pigs, the genetic
2785 correlation estimates obtained between fat and eating quality traits led Cameron (1990) to
2786 suggest that effective selection for increased lean percent would adversely affect pork juiciness,
2787 tenderness and flavor. The moderate genetic correlation of TFW and TFP with IMF suggest the
2788 deposition of back fat and intramuscular fat is brought about by similar physiological factors
2789 which disagrees with the results of Jensen et al. (1967), as in their study ether extractable lipid
2790 was not genetically correlated with back fat thickness. The negative genetic correlation of TFW
2791 with CL and FOM coupled with the correlation estimates of LEP with LOINL, LOINA and FOF
2792 supports the conclusion arrived by Jensen et al. (1967) that selecting for decreased back fat and
2793 increased lean percent would negatively affect meat quality. The moderate correlation of BP with
2794 ISOLC and TCOL suggest the influence of similar gene networks on these traits. Indeed, type 1
2795 collagen, the most abundant collagen type, can be found in both bone matrix and intramuscular
2796 connective tissues, where it provides biomechanical support.

2797 The genetic correlation estimates of LEP with LOINL, LOINA and FOSSF indicated that
2798 selection based on muscling data will result in increased LOINL, and decreased LOINA and fat
2799 content. The genetic correlation of LWT with WBSF and FOP suggests that selection for
2800 increased loin weight is expected to result in improved muscling but tougher pork over time. Lo
2801 et al. (1992) came to similar conclusions from their correlation estimates between loin muscle
2802 area and shear force.

2803 **2.5 Conclusions**

2804 Genetic correlations between WBSF and collagen characteristics indicated that similar gene
2805 networks potentially influence these traits and thus, despite the relative youthfulness of pigs at
2806 slaughter, genetic selection for collagen solubility may decrease pork toughness. Furthermore,
2807 the genetic correlations of Minolta L^* with collagen characteristics present a potential to use
2808 Minolta L^* as an indicator trait for percentage soluble and insoluble collagen. Results from this
2809 study also hint at a possible linkage disequilibrium and/or pleiotropic gene effect in color, pH
2810 and moisture content. Results further indicated that additive genetic variance for intramuscular
2811 fat and Minolta a^* makes a substantial contribution to its total phenotypic variance indicating
2812 that selection for these traits will lead to relatively rapid genetic changes in the average
2813 population means. In summary, the genetic parameters estimated in this study will make it
2814 feasible to predict response to genetic selection of loin physical and collagen characteristics and
2815 further add to the already established research database on meat quality traits. The genetic
2816 correlations agree with published reports that continuous selection for increased lean will
2817 adversely affect fat traits.

2818 **2.6 Figures**



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2820 Figure 2. 1: intramuscular pH measurement on a loin chop.



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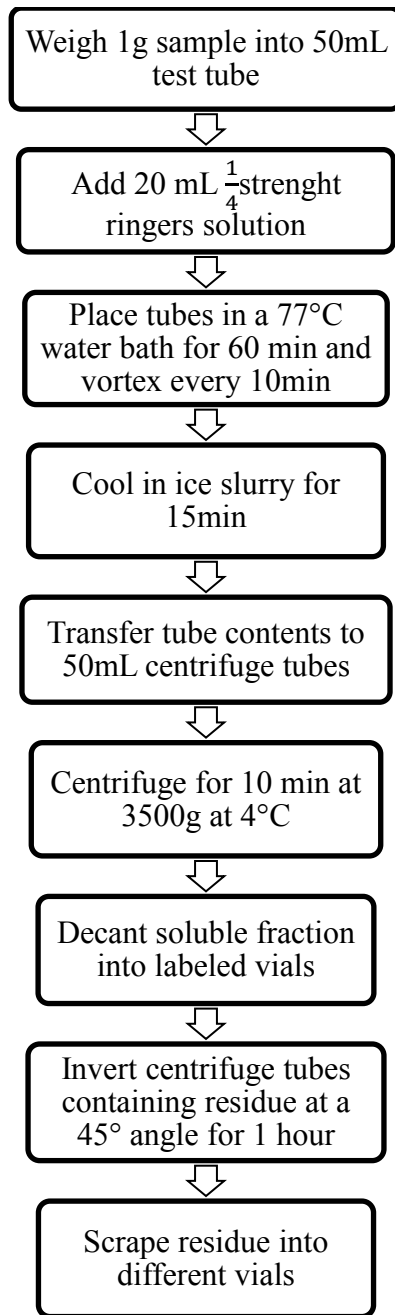
2822 Figure 2. 2: Color measurements using a Konica Minolta chroma meter.

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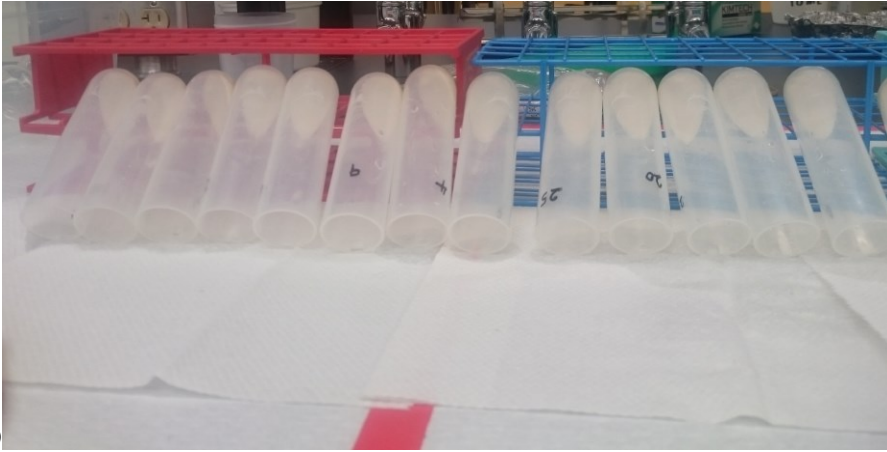
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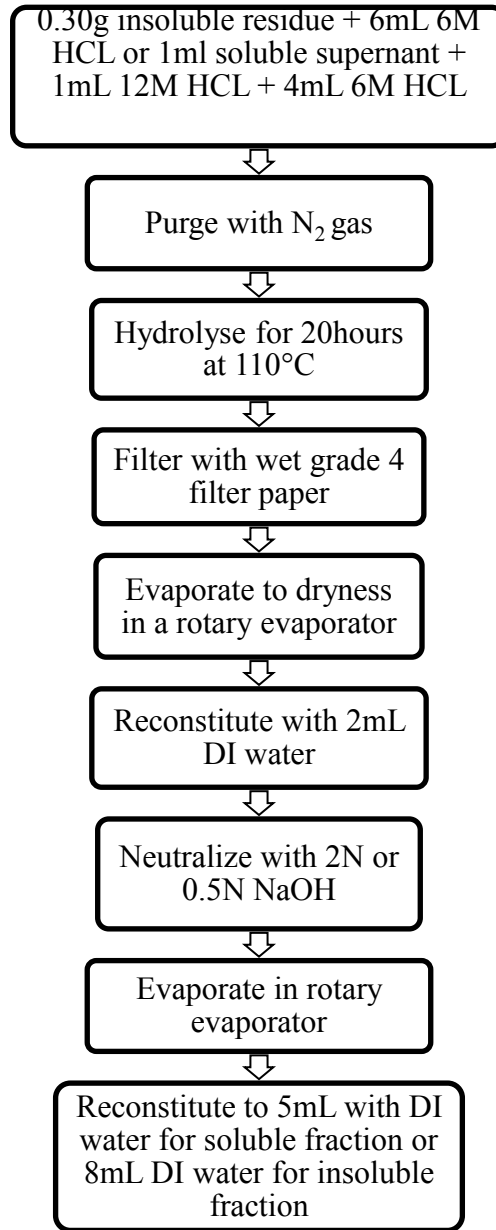
2828 Figure 2. 3: Flow chart for collagen heat solubility using the method described by Hill (1966).



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2830 Figure 2. 4: Inversion of wet residue to ensure drainage

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2833 Figure 2. 5: Flow chart for acid hydrolysis and neutralization assay used in this study.

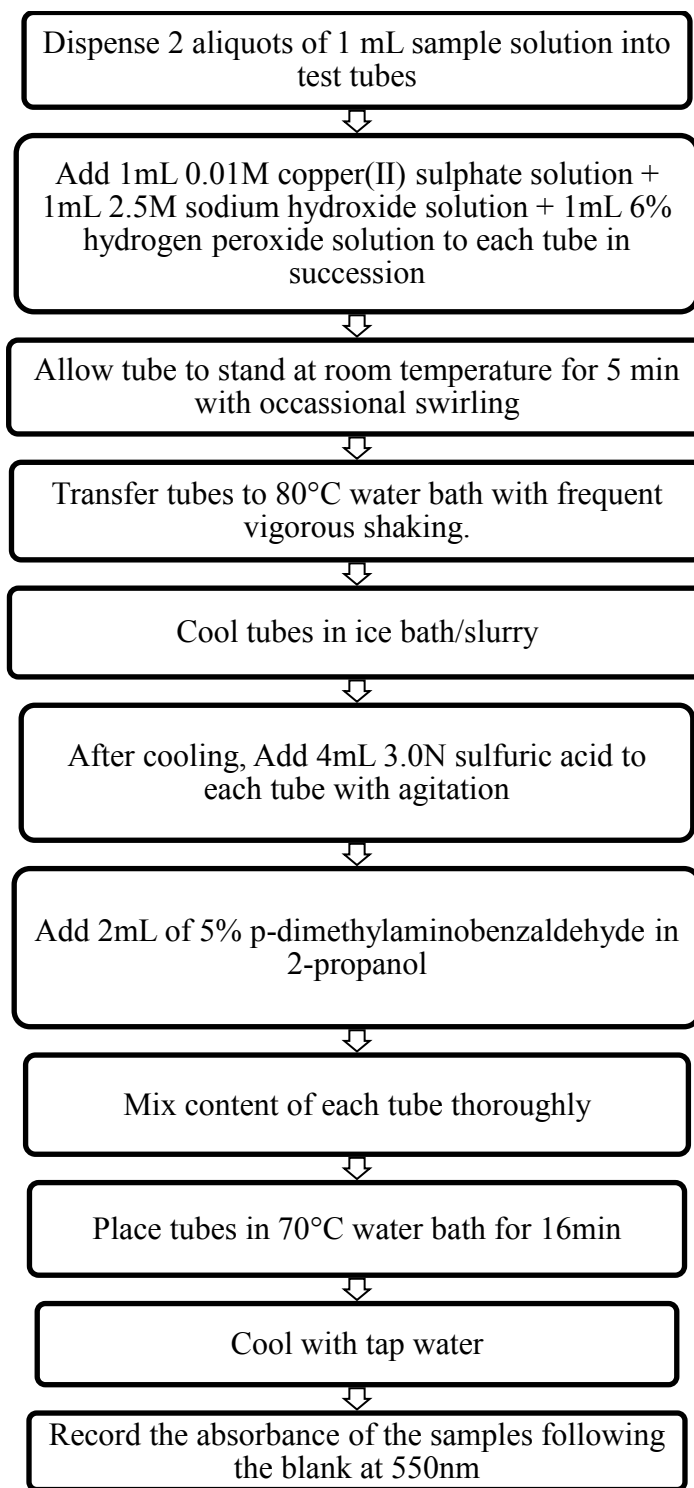
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2836 Figure 2. 6: Acid hydrolysis on a heating block

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2839 Figure 2. 7: Flow chart for assay for hydroxyproline determination.

2840 **2.7 Tables**

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2842 Table 2. 1: Significance of fixed and random effects included in the models for the analysis of
 2843 loin physical traits.

Trait	Fixed effects				Random effect
	Slaughter batch	Company	Sex	TCT ²	Animal
Loin Weight (kg)	*** ¹	***	ns	-	✓
Loin Eye Weight (kg)	***	***	ns	-	✓
Thick Fat (kg)	***	***	ns	-	✓
Meat Trim Weight (kg)	***	***	ns	-	✓
Fat Trim Weight (kg)	***	**	ns	-	✓
Rib Trim Weight (kg)	***	**	ns	-	✓
Bone/Neural Weight (kg)	***	***	ns	-	✓
Loin Eye percent (%)	***	***	ns	-	✓
Thick fat percent (%)	*	***	ns	-	✓
Rib trim percent (%)	***	***	ns	-	✓
Bone percent (%)	***	ns	ns	-	✓

2844 ¹ ns, *, **, ***, represent not significant, p<0.10, p<0.05 and p<0.001, respectively; ² Total
 2845 cooking time.

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2849 Table 2. 2: Significance of fixed and random effects included in the models for the analysis of
 2850 meat quality traits.

Trait	Fixed effects				Random effect
	Slaughter batch	Company	Sex	TCT ²	Animal
PH	*** ¹	***	ns	-	✓
Cooking loss (%)	***	***	ns	**	✓
<i>L</i> *	***	ns	ns	-	✓
<i>a</i> *	***	*	ns	-	✓
<i>b</i> *	***	***	ns	-	✓
Hue	***	**	ns	-	✓
Chroma	***	ns	ns	-	✓
Weight loss (%)	***	ns	ns	-	✓
Warner-Bratzler Shear force (N)	***	ns	ns	-	✓
Fat	***	*	ns	-	✓
Protein	***	ns	ns	-	✓
Moisture	***	**	ns	-	✓
Ether Extract	*	ns	ns	-	✓
Collagen solubility (%) in raw meat	***	*	ns	-	✓
Collagen insolubility (%) in raw meat	***	*	ns	-	✓
Total collagen mg/ g raw meat	***	ns	ns	-	✓

2851 ¹ ns, *, **, ***, represent not significant, p<0.10, p<0.05 and p<0.001 respectively; ²total
 2852 cooking time

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 2855

2856 Table 2. 3: Descriptive statistics of loin physical traits measured on frozen-thawed loin:
 2857 abbreviations, number of animals (n) per measure trait, mean, standard deviation (SD), Minimum
 2858 value (Min), Maximum value (Max), Coefficient of variation (CV).

Trait	Abbreviation	Pigs (n)	Mean	SD	Min	Max	CV (×100)
Loin Weight (kg)	LWT	500	1.42	0.17	1.11	1.81	11.82
Loin Eye Weight (kg)	LEW	500	0.54	0.07	0.40	0.68	12.53
Thick Fat Weight (kg)	TFW	500	0.32	0.08	0.15	0.49	24.45
Meat Trim Weight (kg)	MTW	455	0.23	0.05	0.10	0.49	23.74
Fat Trim Weight (kg)	FTW	456	0.12	0.04	0.04	0.35	36.15
Rib Trim Weight (kg)	RTW	500	0.32	0.05	0.21	0.47	14.95
Bone/Neural Weight (kg)	BWT	500	0.17	0.03	0.1	0.29	18.00
Loin Eye percent (%)	LEP	500	38.85	3.25	31.62	45.96	8.37
Thick fat percent (%)	TFP	500	23.59	4.18	15.95	32.98	17.71
Rib trim percent (%)	RTP	500	23.79	2.66	19.03	33.33	11.17
Bone percent (%)	BP	500	12.52	1.45	9.52	15.91	11.58

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2862 Table 2. 4: Descriptive statistics of meat quality and collagen traits measured on frozen-thawed
 2863 loin: (standard deviation (SD), minimum value (Min), maximum value (Max), coefficient of
 2864 variation (CV)).

Trait	Abbrev- iation	Pigs (n)	Mean	SD	Min	Max	CV (×100)
pH	-	500	5.56	0.14	5.20	5.92	-
Cooking loss (%)	CL	500	25.17	2.99	18.99	36.76	11.89
<i>L</i> *	LOINL	500	45.46	3.02	37.23	55.18	7.55
<i>a</i> *	LOINA	500	7.65	1.31	5.09	11.17	17.12
<i>b</i> *	LOINB	500	3.50	1.14	1.07	7.48	32.50
Hue	LOINH	500	24.64	8.66	6.15	44.22	35.13
Chroma	LOINC	500	8.39	1.16	5.83	11.18	13.82
Chemical Moisture (%)	MC	500	72.21	0.73	70.81	73.96	1.01
Warner-Bratzler Shear force (N)	WBSF	500	47.16	10.05	21.62	84.68	24.00
FOSS fat	FOF	500	24.08	4.20	14.57	32.64	17.44
FOSS protein	FOP	500	16.79	1.19	14.33	19.98	7.08
FOSS moisture	FOM	500	57.42	3.23	50.63	64.87	5.62
Intramuscular fat	IMF	464	8.11	2.95	2.42	22.08	36.43
Collagen solubility (%) in raw meat	SOLC	499	18.03	6.59	1.98	34.28	36.52
Collagen insolubility (%) in raw meat	ISOLC	492	82.18	6.38	68.51	98.02	7.76
Total collagen (mg/ g raw meat)	TCOL	499	3.09	0.99	1.53	6.29	32.00

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2872 Table 2. 5: Heritabilities (h^2) among loin physical traits on frozen thawed three-rib pork loin
 2873 sections with standard errors

Trait	$h^2 \pm se^{(1)}$	$\sigma_e^2^{(2)} \pm se$	$\sigma_a^2^{(3)} \pm se$
Loin Weight (kg)	0.19±0.08	0.019±0.002	0.005±0.002
Loin Eye Weight (kg)	0.44±0.11	0.002±0.000	0.002±0.000
Thick Fat (kg)	0.16±0.10	0.004±0.000	0.001±0.000
Meat Trim Weight (kg)	0.30±0.11	0.001±0.000	0.000±0.000
Fat Trim Weight (kg)	0.13±0.09	0.001±0.000	0.000±0.000
Rib Trim Weight (kg)	0.31±0.11	0.001±0.000	0.001±0.000
Bone Weight (kg)	0.22±0.10	0.002±0.000	0.001±0.000
Loin Eye percent (%)	0.28±0.12	6.745 ±0.966	2.658±1.231
Thick fat percent (%)	0.42±0.13	9.445±1.752	6.83±2.495
Rib trim percent (%)	0.39±0.12	2.350±0.376	1.430±0.514
Bone percent (%)	0.09±0.08	1.645±0.161	0.165±0.153

2874 (1)-Standard error; (2)-Residual variance; (3)-Additive genetic variance.
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2878 Table 2. 6: Heritabilities (h^2) of meat quality and collagen traits measured on frozen-thawed
 2879 pork longissimus muscle with standard errors

Trait	$h^2 \pm se^{(1)}$	$\sigma_e^{2(2)} \pm se$	$\sigma_a^{2(3)} \pm se$
PH	0.12±0.08	0.116±0.01	0.002±0.001
Cooking loss (%)	0.24±0.10	4.226±0.503	1.242±0.589
L^*	0.25±0.11	3.525±0.452	1.176±0.552
a^*	0.42±0.11	0.559±0.092	0.394±0.130
b^*	0.21±0.10	0.646±0.076	0.176±0.088
Hue	0.25±0.10	20.847±2.574	6.981±3.102
Chroma	0.36±0.11	0.689±0.106	0.384±0.144
Moisture content (%)	0.36±0.11	0.293±0.044	0.167±0.060
Warner-Bratzler shear force (N)	0.31±0.11	56.414±8.188	24.917±10.603
Crude Fat	0.24±0.10	11.904±1.462	3.721±1.747
Crude Protein	0.17±0.09	0.994±0.108	0.196±0.117
Crude Moisture	0.21±0.09	7.171±0.836	1.865±0.961
Collagen solubility (%) in raw meat	0.12±0.09	32.241±3.497	4.413±3.677
Collagen insolubility (%) in raw meat	0.15±0.09	29.409±3.289	5.084±3.555
Total collagen mg/ g raw meat	0.33±0.12	0.617±0.095	0.306±0.127
Intramuscular fat	0.42±0.13	4.851±0.913	3.541±1.297

2880 (1)-Standard error; (2)-Residual variance; (3)-Additive genetic variance

2881 Table 2. 7: Genetic (below diagonal) and phenotypic (above diagonal) correlation estimates and their standard errors among loin
 2882 physical traits

Trait ¹	LWT	LEW	TFW	MTW	FTW	RTW	BWT	LEP	TFP	RTP	BP
LWT		0.65 ±.02	0.56 ±.03	0.48 ±.04	0.50 ±.03	0.71 ±.02	0.49 ±.03	-0.20 ±.05	0.19 ±.05	-0.04 ±.05	-0.22 ±.04
LEW	0.87 ±.13 ²		-0.04 ±.05	0.51 ±.04	0.11 ±.05	0.51 ±.04	0.41 ±.04	0.47 ±.04	-0.37 ±.05	0.01 ±.05	-0.05 ±.05
TFW	0.32 ±.36	-0.37 ±.29		-0.08 ±.05	0.45 ±.04	0.30 ±.05	0.12 ±.04	-0.71 ±.02	0.88 ±.01	-0.27 ±.04	-0.34 ±.04
MTW	0.93 ±.22	0.61 ±.17	-0.31 ±.36		-0.10 ±.05	0.73 ±.02	0.44 ±.04	0.11 ±.05	-0.30 ±.05	0.52 ±.04	0.09 ±.05
FTW	0.64 ±.28	0.06 ±.36	0.35 ±.42	0.30 ±.43		0.61 ±.03	0.09 ±.05	-0.35 ±.04	0.28 ±.05	0.28 ±.05	-0.33 ±.04
RTW	0.76 ±.16	0.60 ±.16	-0.14 ±.35	0.92 ±.08	0.62 ±.25		0.44 ±.04	-0.16 ±.05	-0.05 ±.05	0.61 ±.03	-0.15 ±.05
BWT	0.77 ±.26	0.48 ±.23	0.32 ±.39	0.64 ±.22	0.50 ±.44	0.71 ±.19		-0.07 ±.05	-0.13 ±.05	0.09 ±.05	0.64 ±.02
LEP	0.28 ±.39	0.73 ±.14	-0.76 ±.14	0.18 ±.30	-0.80 ±.28	0.10 ±.29	-0.13 ±.35		-0.78 ±.02	-0.01 ±.05	0.12 ±.05
TFP	-0.09 ±.38	-0.60 ±.16	0.91 ±.06	-0.50 ±.22	0.32 ±.35	-0.50 ±.23	-0.25 ±.29	-0.76 ±.10		-0.37 ±.04	-0.35 ±.04
RTP	-0.08 ±.40	0.21 ±.23	-0.48 ±.27	0.76 ±.15	0.44 ±.35	0.84 ±.11	0.43 ±.28	0.01 ±.29	-0.66 ±.17		0.04 ±.05
BP	0.35 ±.66	-0.02 ±.38	0.19 ±.67	0.12 ±.44	0.44 ±.82	0.32 ±.47	0.77 ±.23	-0.15 ±.47	-0.21 ±.39	0.36 ±.39	

2883 ¹LWT=Loin Eye Weight; LEW= Loin Eye Weight; TFW= Thick Fat weight; MTW= Meat Trim Weight; FTW= Fat Trim Weight;
 2884 RTW= Rib Trim Weight; BWT= Bone Weight; LEP= Loin Eye percent; TFP= Thick fat percent; RTP= Rib trim percent; BP= Bone
 2885 percent. ²Significant correlations (P < 0.05) are highlighted in bold.
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2889 Table 2. 8: Genetic (below diagonal) and phenotypic (above diagonal) correlation estimates among meat quality traits

Trait 1	pH	CL	LOI NL	LOI NA	LOI NB	LOI NH	LOI NC	MC	WBSF 1	FOF	FOP	FOM	IMF	SOLC	ISOLC	TCOL
PH		-0.13 ±.05	-0.40 ±.04	0.04 ±.05	-0.32 ±.04	-0.37 ±.04	-0.06 ±.05	0.34 ±.04	0.01 ±.05	0.05 ±.05	-0.10 ±.05	-0.02 ±.05	0.05 ±.05	-0.08 ±.04	0.09 ±.05	-0.10 ±.05
CL	-0.65² ±.31		0.07 ±.05	0.00 ±.05	0.05 ±.05	0.03 ±.05	0.01 ±.05	0.07 ±.05	0.26 ±.04	-0.14 ±.05	0.14 ±.05	0.12 ±.05	-0.14 ±.05	0.07 ±.05	-0.07 ±.05	0.05 ±.05
LOI	-0.95 ±.35	0.44 ±.19		-0.24 ±.05	0.58 ±.03	0.74 ±.02	-0.03 ±.05	-0.39 ±.04	-0.16 ±.05	0.16 ±.05	-0.14 ±.05	-0.18 ±.05	0.27 ±.05	0.14 ±.05	-0.14 ±.05	0.01 ±.05
NL	±.35	±.19			±.03	±.02	±.05	±.04	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05
LOI	0.20	0.32	-0.53		0.41	-0.12	0.94	-0.18	-0.10	0.08	-0.08	-0.07	0.21	-0.00	0.02	-0.03
NA	±.36	±.27	±.20		±.04	±.05	±.00	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05
LOI	-0.87 ±.31	0.69 ±.31	0.46 ±.22	0.49 ±.23		0.77 ±.02	0.65 ±.03	-0.54 ±.03	-0.13 ±.05	0.16 ±.05	-0.15 ±.05	-0.20 ±.04	0.40 ±.04	0.06 ±.05	-0.06 ±.05	0.01 ±.05
NB	±.31	±.31	±.22	±.23		±.02	±.03	±.03	±.05	±.05	±.05	±.04	±.04	±.05	±.05	±.05
LOI	-0.93 ±.28	0.48 ±.32	0.85 ±.09	-0.30 ±.25	0.69 ±.15		0.15 ±.05	-0.33 ±.05	-0.13 ±.05	0.12 ±.05	-0.12 ±.05	-0.15 ±.05	0.29 ±.05	0.06 ±.05	-0.08 ±.05	0.02 ±.05
NH	±.28	±.32	±.09	±.25	±.15		±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05
LOI	0.06	0.30	-0.43	0.97	0.64	-0.15		-0.33	-0.14	0.11	-0.13	-0.11	0.30	0.01	-0.01	-0.04
NC	±.37	±.28	±.25	±.02	±.17	±.29		±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05	±.05
MC	0.98 ±.27	0.32 ±.34	-0.39 ±.28	-0.10 ±.24	-0.76 ±.17	-0.62 ±.21	-0.25 ±.23		0.06 ±.05	-0.27 ±.05	0.20 ±.05	0.35 ±.04	-0.61 ±.03	-0.02 ±.05	0.04 ±.05	0.03 ±.05
WB	0.01	0.45	-0.22	0.08	0.04	-0.23	0.01	-0.21		-0.22	0.20	0.17	-0.13	-0.05	0.06	0.04
SF	±.42	±.19	±.29	±.28	±.34	±.31	±.30	±.28		±.04	±.04	±.05	±.05	±.05	±.05	±.05
FOF	-0.20 ±.42	-0.43 ±.31	0.35 ±.27	-0.55 ±.28	0.06 ±.34	0.32 ±.30	-0.54 ±.28	-0.39 ±.26	-0.20 ±.32		-0.93 ±.00	-0.99 ±.00	0.45 ±.04	-0.01 ±.05	0.01 ±.05	-0.00 ±.05
FOP	0.04 ±.47	0.33 ±.36	-0.08 ±.34	0.40 ±.34	0.16 ±.40	-0.25 ±.35	0.40 ±.35	0.05 ±.32	0.33 ±.36	-0.93 ±.05		0.90 ±.01	-0.45 ±.04	0.01 ±.05	-0.01 ±.05	-0.00 ±.05
FO	0.40 ±.44	0.32 ±.34	-0.37 ±.32	0.62 ±.30	-0.15 ±.35	-0.40 ±.31	0.56 ±.30	0.63 ±.22	-0.00 ±.35	-0.98 ±.01	0.86 ±.09		-0.47 ±.04	-0.02 ±.05	0.02 ±.05	0.03 ±.05
M	±.44	±.34	±.32	±.30	±.35	±.31	±.30	±.22	±.35	±.01	±.09		±.04	±.05	±.05	±.05
IMF	0.15 ±.36	-0.20 ±.28	0.12 ±.26	0.17 ±.23	0.61 ±.22	0.40 ±.25	0.25 ±.24	-0.64 ±.14	-0.02 ±.30	0.79 ±.14	-0.67 ±.18	-0.83 ±.13		-0.05 ±.05	0.05 ±.05	0.01 ±.05
SOL	-0.72 ±.64	-0.10 ±.43	0.68 ±.30	-0.50 ±.39	0.00 ±.45	0.66 ±.50	-0.40 ±.41	-0.60 ±.43	-0.38 ±.10	0.17 ±.42	-0.33 ±.45	-0.17 ±.45	0.37 ±.41		N.E*	-0.46 ±.04
C	±.64	±.43	±.30	±.39	±.45	±.50	±.41	±.43	±.10	±.42	±.45	±.45	±.41			

ISO	0.76	0.11	-0.71	0.45	-0.12	-0.57	0.34	0.60	0.42	-0.12	0.16	0.12	-0.35	N.E		0.48
LC	±.56	±.40	± .35	±.33	±.42	±.41	±.36	±.38	± .16	±.40	±.44	±.42	±.37			± .04
TCO	0.03	0.07	-0.08	0.12	0.31	0.06	0.06	0.17	0.10	0.19	-0.08	-0.17	-0.21	-0.60	0.66	
L	±.41	±.32	±.28	±.26	±.32	±.30	±.28	±.27	±.31	±.32	±.37	±.34	±.26	± .23	± .28	

2890 1-CL= Cooking loss (%); LOINL=Minolta *L**; LOINA=Minolta *a**; LOINB= Minolta *b**; LOINH= Minolta hue; LOINC= Minolta
2891 chroma; MC= Moisture content (%); WBSF= Warner-Bratzler shear force (N); FOF= FOSS crude fat; FOP= FOSS crude protein;
2892 FOM= FOSS crude moisture; IMF= Intramuscular fat; SOLC= Soluble collagen percent (%) raw meat; ISOLC= Insoluble collagen
2893 percent (%) raw meat; TCOL= Total collagen mg/g raw meat. 2-Significant correlations (P < 0.05) are highlighted in bold.

2894 *Not estimable-ASReml assumes the 2 traits are not perfectly correlated, but here, the 2 traits are perfectly correlated, one is a simple
2895 transformation of the other, soluble collagen percent =100-insoluble collagen percent; consequently, the two US matrices are singular
2896 and cannot be inverted. Thus, variance components for the two traits are the same.

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Table 2. 9: Phenotypic correlation between sub primal and meat quality

Trait ¹	LWT	LEW	TFW	MTW	FTW	RTW	BWT	LEP	TFP	RTP	BP
PH	0.13 ²	0.08	0.09	0.08	0.05	0.10	0.09	0.01	0.10	-0.04	-0.06
	\pm .04	\pm .05	\pm .04	\pm .05	\pm .05	\pm .05	\pm .04	\pm .05	\pm .05	\pm .03	\pm .04
CL	-0.02	0.12	-0.12	0.24	-0.03	0.09	0.03	0.09	-0.17	0.12	0.01
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .01	\pm .05	\pm .05	\pm .05	\pm .00
LOINL	-0.02	-0.11	0.04	-0.08	0.00	-0.06	-0.01	-0.08	0.04	-0.03	0.01
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
LOINA	0.04	-0.15	0.15	-0.05	0.13	0.03	0.02	-0.24	0.18	-0.03	-0.02
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
LOINB	0.02	-0.16	0.18	-0.09	0.08	-0.02	0.01	-0.24	0.17	-0.05	-0.04
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
LOINH	-0.00	-0.09	0.08	-0.06	0.03	-0.03	0.04	-0.11	0.05	-0.03	0.02
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
LOINC	0.06	-0.14	0.19	-0.05	0.14	0.03	0.05	-0.26	0.20	-0.05	-0.02
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
MC	0.07	0.24	-0.13	0.20	-0.10	0.10	0.05	0.25	-0.20	0.08	0.08
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
WBSF	-0.09	0.08	-0.23	0.11	-0.18	-0.10	-0.04	0.21	-0.24	-0.01	0.05
	\pm .04	\pm .05	\pm .04	\pm .05	\pm .05	\pm .05	\pm .04	\pm .05	\pm .05	\pm .05	\pm .04
FOF	0.18	-0.16	0.52	-0.41	0.56	0.06	-0.07	-0.49	0.58	-0.19	-0.24
	\pm .04	\pm .05	\pm .03	\pm .04	\pm .03	\pm .05	\pm .05	\pm .04	\pm .03	\pm .05	\pm .04
FOP	-0.16	0.17	-0.50	0.33	-0.60	-0.10	0.02	0.47	-0.53	0.13	0.18
	\pm .04	\pm .05	\pm .03	\pm .04	\pm .03	\pm .05	\pm .05	\pm .04	\pm .03	\pm .05	\pm .04
FOM	-0.13	0.16	-0.51	0.42	-0.55	-0.05	0.07	0.47	-0.57	0.18	0.26
	\pm .04	\pm .05	\pm .03	\pm .04	\pm .03	\pm .05	\pm .05	\pm .04	\pm .03	\pm .05	\pm .04
SOLC	-0.04	0.04	-0.09	-0.03	0.00	-0.02	0.02	0.09	-0.08	0.02	0.06
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .04	\pm .05	\pm .05	\pm .05	\pm .04
ISOLC	0.03	-0.07	0.10	0.01	-0.01	-0.00	-0.01	-0.12	0.10	-0.03	-0.05
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05
TCOL	-0.01	-0.09	0.03	-0.03	0.06	0.02	-0.03	-0.09	0.00 \pm	0.06	-0.02
	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	\pm .05	.05	\pm .05	\pm .05
IMF	0.07	-0.26	0.35	-0.21	0.23	-0.03	-0.06	-0.38	0.44	-0.18	-0.15

±.05 ±.05 ±.04 ±.05 ±.05 ±.05 ±.05 ±.05 ±.05 ±.04 ±.05 ±.05

2910 ¹LWT=Loin Eye Weight; LEW= Loin Eye Weight; TFW= Thick Fat weight; MTW= Meat Trim Weight; FTW= Fat
2911 Trim Weight; RTW= Rib Trim Weight; BWT=Bone Weight; LEP= Loin Eye percent; TFP= Thick fat percent; RTP=
2912 Rib trim percent; BP= Bone percent; CL= Cooking loss (%); LOINL=Minolta *L**; LOINA=Minolta *a**; LOINB=
2913 Minolta *b**; LOINH= Minolta hue; LOINC= Minolta chroma; MC= Moisture content (%); WBSF= Warner-Bratzler
2914 shear force (N); FOF= FOSS crude fat; FOP= FOSS crude protein; FOM= FOSS crude moisture; IMF= Intramuscular
2915 fat; SOLC= Soluble collagen percent (%) raw meat; ISOLC= Insoluble collagen percent (%) raw meat; TCOL= Total
2916 collagen mg/g raw meat. ²Significant correlations (P < 0.05) are highlighted in bold.
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2931 Table 2. 10: Genetic correlation between loin physical and meat quality traits

Trait ¹	LWT	LEW	TFW	MTW	FTW	RTW	BWT	LEP	TFP	RTP	BP
PH	-0.30 ±.54	-0.24 ±.35	0.14 ±.49	0.26 ±.38	-0.32 ±.56	0.06 ±.39	0.24 ±.42	-0.58 ±.40	0.30 ±.35	0.08 ±.38	0.56 ±.62
CL	-0.42 ±.42	-0.33 ±.26	-0.41 ±.16²	0.64 ±.27	-0.06 ±.45	0.20 ±.30	0.20 ±.34	-0.33 ±.31	-0.10 ±.29	0.68 ±.23	0.34 ±.42
LOINL	0.22 ±.40	0.20 ±.24	-0.07 ±.35	0.10 ±.30	0.38 ±.41	-0.08 ±.28	-0.24 ±.32	0.53 ±.26	-0.36 ±.24	-0.04 ±.26	-0.52 ±.44
LOINA	-0.19 ±.35	-0.37 ±.12	-0.21 ±.35	0.26 ±.30	-0.01 ±.38	0.08 ±.26	0.35 ±.32	-0.58 ±.19	0.17 ±.23	0.13 ±.25	0.20 ±.41
LOINB	0.61 ±.51	0.25 ±.32	-0.18 ±.43	0.40 ±.36	0.76 ±.39	0.56 ±.35	0.39 ±.38	-0.05 ±.32	-0.39 ±.31	0.15 ±.31	0.01 ±.48
LOINH	0.23 ±.44	0.31 ±.27	-0.27 ±.39	0.19 ±.33	0.60 ±.39	0.11 ±.31	0.31 ±.35	0.29 ±.31	-0.50 ±.25	0.02 ±.28	0.25 ±.45
LOINC	-0.11 ±.38	-0.25 ±.22	-0.29 ±.36	0.34 ±.31	-0.04 ±.39	0.05 ±.27	0.56 ±.33	-0.35 ±.25	-0.09 ±.27	0.03 ±.26	0.47 ±.45
MC	0.01 ±.40	-0.10 ±.25	0.20 ±.35	0.17 ±.27	-0.65 ±.28	-0.09 ±.28	-0.15 ±.31	-0.19 ±.29	0.23 ±.25	0.08 ±.25	0.06 ±.40
WBSF	0.36 ±.12	0.23 ±.26	0.28 ±.38	0.45 ±.29	-0.41 ±.42	0.23 ±.30	0.21 ±.35	-0.30 ±.32	0.05 ±.29	-0.02 ±.29	-0.27 ±.47
FOF	-0.06 ±.45	-0.30 ±.10	0.55 ±.25	-0.47 ±.26	0.73 ±.24	-0.19 ±.31	-0.12 ±.34	-0.41 ±.15	0.55 ±.20	-0.41 ±.27	-0.10 ±.46
FOP	0.50 ±.15	0.52 ±.20	-0.26 ±.37	0.57 ±.29	-0.71 ±.27	0.50 ±.34	-0.07 ±.39	0.47 ±.16	-0.52 ±.23	0.51 ±.29	-0.79 ±.76
FOM	0.15 ±.48	0.26 ±.27	-0.52 ±.18	0.53 ±.26	-0.70 ±.27	0.23 ±.32	0.07 ±.36	0.38 ±.17	-0.53 ±.21	0.44 ±.29	0.07 ±.49
SOLC	0.02 ±.58	0.00 ±.36	-0.06 ±.51	-0.13 ±.42	0.36 ±.53	-0.10 ±.42	-0.04 ±.46	0.06 ±.43	0.23 ±.39	-0.17 ±.40	-0.25 ±.11
ISOLC	-0.19 ±.53	-0.15 ±.33	0.20 ±.47	0.02 ±.39	-0.42 ±.49	-0.09 ±.38	0.12 ±.43	-0.21 ±.40	-0.04 ±.36	-0.00 ±.36	0.53 ±.10

TCOL	-0.18	-0.32	0.43	-0.24	0.20	-0.03	0.03	-0.40	0.03	0.03	0.51
	±.40	±.13	±.37	±.29	±.41	±.29	±.32	±.30	±.28	±.27	±.27
IMF	0.29	-0.04	0.53	-0.29	0.66	-0.05	0.22	-0.34	0.43	-0.54	0.16
	±.39	±.25	±.26	±.26	±.31	±.27	±.34	±.25	±.21	±.22	±.48

2932 ¹LWT=Loin Weight; LEW= Loin Eye Weight; TFW= Thick Fat weight; MTW= Meat Trim Weight; FTW= Fat Trim Weight; RTW=
2933 Rib Trim Weight; BWT= Bone Weight; LEP= Loin Eye percent; TFP= Thick fat percent; RTP= Rib trim percent; BP= Bone percent;
2934 CL= Cooking loss (%); LOINL=Minolta *L**; LOINA=Minolta *a**; LOINB= Minolta *b**; LOINH= Minolta hue; LOINC= Minolta
2935 chroma; MC= Moisture content (%); WBSF= Warner-Bratzler shear force (N); FOF= FOSS crude fat; FOP= FOSS crude protein;
2936 FOM= FOSS crude moisture; IMF= Intramuscular fat; SOLC= Soluble collagen percent (%) raw meat; ISOLC= Insoluble collagen
2937 percent (%) raw meat; TCOL= Total collagen (mg/g raw meat)

2938 ²Significant correlations (P < 0.05) are highlighted in bold.

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3386

Chapter 3

Contributions of company of origin, sex and intrinsic pork characteristics to measures of loin physical, meat and collagen characteristics in a Canadian context

3.1 Introduction

Consumer acceptability of pork is a factor of significant importance to the success of pork products in the market. It is driven mainly by product appearance at retail or point of purchase and the subsequent satisfaction experienced upon consumption (Jeremiah et al., 1999). Yet producing pork at reduced cost by aiming for increased animal growth rates, using cheaper feed ingredients and increasing lean percent (Ngapo and Garipey, 2008) with minor emphasis on product quality has been the focus of swine production over the past decades (Gjerlaug-Enger et al., 2010). Recently, shifts in consumer preferences suggest that pork market standing could be bolstered by putting additional weight on quality to enhance consumer product acceptance.

Indeed, pork quality in the last decade has received appreciable attention from industry players like retailers, producers, processors, packers and researchers (Newcom et al., 2004) due to the potential for consumer discontent with eating quality of pork from current genotypes with higher lean-to-fat ratios (Kempster et al., 1986). Based upon this, producing pork products tailored to meet demands of certain market niches and export markets may be critical to the long-term competitiveness of the Canadian pork industry on the export market.

3407 Several factors come into play to determine pork quality. These can be of genetic origin like
3408 breed and sex or environmental factors like pre-slaughter handling, post-slaughter management
3409 and contemporary groups (Ngapo and Garipey, 2008). Due to the fact that Canadian pork for the
3410 most part is obtained from crosses between Large White sows and Duroc boars, minimum
3411 differentiation among different populations in terms of loin physical and pork characteristics
3412 would be expected owing to the fact that selection is applied only to a defined number of swine
3413 breeds and is directed by grading systems that place premiums on lean percent (Zhang et al.,
3414 2016). However, that is usually not the case, as inconsistencies in pig meat products are a major
3415 challenge confronting the pig industry in many countries (Ellis et al., 1996). A stratification
3416 system generated from compositional or relatively easy to measure pork quality measurements
3417 can be useful to retailers as a greater degree of information could be provided to consumers with
3418 respect to the physical characteristic of a cooked product, thus allowing individuals to buy cuts
3419 with tailored levels of protein, fat, moisture, and predicted yields and losses.

3420 Although several studies have been undertaken over the years, debates still exist on the effect of
3421 sex and location/origin/slaughter plant on pork quality. Indeed, significant differences in
3422 tenderness, juiciness and overall quality have been reported for pork from different locations in
3423 Australia under their present production and product processing schemes (Channon et al., 2000).
3424 Further, differences in intramuscular fat, color and tenderness attributes were found to be due to
3425 sex (Channon et al 2004) while Jeremiah et al (1999) reported the opposite. In general, a perusal
3426 of research on the influence of sex on quality attributes reveals differences in its effect on
3427 quality; while some studies report influence of sex on quality (Channon et al., 2000; Ellis et al.,
3428 1995), others do not (Jeremiah et al., 1999; Ellis et al., 1996). It is therefore necessary to explore
3429 the influence of these factors on Canadian pork quality attributes.

3430 Tenderness is the most important palatability trait affecting pork acceptability (Suman, 2012)
3431 and a positive relationship between tenderness and collagen characteristics has been proposed in
3432 beef (Cross et al., 1973). Further, drip loss has been showed to be significantly associated with
3433 meat protein oxidation (Traore et al., 2011) and myofibrillar protein oxidation has been found to
3434 decrease beef tenderness (Rowe et al., 2004) hinting at a potential relationship between drip loss
3435 and tenderness. However, few studies have looked at the potential presence of these relationships
3436 in pork especially in the Canadian context.

3437 Further, meat quality traits like tenderness tend to be of low to moderate heritability (Ciobanu et
3438 al., 2011) and are mostly difficult to measure on live animals, thus, there is increasing interest in
3439 the level of breeding effort to be allocated to these quality traits relative to the importance of
3440 crucial factors of pork production like sex, feeding regimen, slaughter plant, terminal sire
3441 genotype and other management practices which ultimately, also influence product quality (Ellis
3442 et al., 1996). Therefore, it is imperative to investigate the factors that predict and/or influence
3443 these traits and the degree of their effect. Thus, this study sought to investigate the relationship
3444 between company, sex, loin physical, meat quality and collagen characteristics of cross-bred pigs
3445 subjected to commercial finishing conditions. Specific hypotheses tested included: Drip loss
3446 influences Warner-Bratzler shear force values; tenderness of cooked pork is influenced more by
3447 myofibrillar proteins than by intramuscular fat content; collagen characteristics influence
3448 cooking loss percent and tenderness decreases with increased muscling. In addition, this study
3449 further assesses the relationship between sub-primal, meat quality and collagen characteristics
3450 when stratified into 4 subclasses based on compositional and quality measures (intramuscular fat
3451 (%), shear force values, pH and L^*) to test the hypothesis that varying levels of intramuscular
3452 (%), shear force, pH and L^* values can be used to predict pork quality.

3453 **3.2 Materials and methods**

3454 **3.2.1 Animals and husbandry**

3455 Data were from 500 pigs derived from 3-way crosses between Duroc sires and hybrid Large
3456 White × Landrace sows. They were collateral relatives (full and half sibs) obtained from 2
3457 breeding populations. All animals were raised on farms located in 2 Canadian provinces under
3458 commercial growing conditions. Male piglets were neutered at 3 to 5 days post birth. Piglets
3459 were weaned at approximately 7.5 kg which was on average 21 days. The weanlings were
3460 subsequently housed in nursery pens for 5 weeks, subsequently raised in pre-grower barns for a
3461 pre-growing period of 4 weeks and then subjected to commercial finishing conditions in
3462 randomly allocated finishing units with *ad libitum* access to feed and water. The final body
3463 weight of the animals was approximately 115 kg. The pigs were transported to East 40 Packers
3464 (provincially inspected abattoir), Brandon, MB, Canada, on a weekly basis in groups of 20-25
3465 pigs for animals from company B and 20-35 pigs for those from company A. They were rested in
3466 lairage overnight with *ab libitum* access to water and slaughtered the following morning.
3467 Guidelines recommended by the Government of Canada were adhered to in the slaughter of the
3468 animals. They were electrically stunned prior to exsanguination and their carcasses fabricated
3469 within 24 hours post-slaughter. They were on average 160 days old at slaughter with an average
3470 live weight of 124 kg at slaughter. Further descriptive information on the animals can be found
3471 in section 2.2.1.

3472 **3.2.2 Loin physical measurements**

3473 For a complete description of loin physical measurements see section 2.2.2. In brief, a 3 and 4 rib
3474 loin section from the right side of each carcass was weighed and recorded as the loin weight

3475 (LWT). The loin eye muscle (longissimus thoracis) was removed and recorded as (LEW). The
3476 back fat was removed, weighed and recorded as thick fat weight (TFW). Bone weight (BWT)
3477 was obtained by weighing the loin bones.

3478 **3.2.3 Meat quality measurements**

3479 A complete description of meat quality and collagen characteristics measurements is found in
3480 section 2.2.3 and 2.2.4. Briefly, pH of the thawed muscles was measured using a glass electrode
3481 equipped pH meter after a 2-point calibration with 4.01 and 7.01 buffer solutions. The probe was
3482 inserted at 3 different sites for a total of 3 readings per loin. The average of the readings was
3483 used for statistical analysis. Drip loss was determined following the procedures elaborated by
3484 Honikel (1998). The loin chops to be used were cut across the muscle fiber direction. They were
3485 subsequently deboned and trimmed of excess fat and epimysium. Average weight after trimming
3486 was $81\text{g}\pm 9$. Intramuscular pH of the chops was on average, 5.54 ± 0.19 . The chops were
3487 weighed, and this weight was recorded as the initial weight. They were then suspended by hooks
3488 in inflated plastic bags ensuring that no other external forces are exerted on the samples except
3489 for gravity. Samples were stored at 4°C for 24 hours and afterwards gently blotted with soaker
3490 pads and re-weighed and this weight recorded as the final weight. Percent drip loss was
3491 subsequently calculated with the following formula: $\frac{\text{Initial weight} - \text{Final weight}}{\text{Initial weight}} \times 100$.

3492 Cooking loss (CL) was determined by cooking approximately $192\text{g}\pm 12$ of loin roast in a 73°C
3493 water bath until an internal roast temperature of 71°C was reach. Cooking loss was calculated as
3494 a percentage of the roast weight before cooking.

3495 Minolta L^* , a^* , and b^* measurements were taken with a Konica Minolta CR 400 (Konica
3496 Minolta Sensing Inc., Japan) set at CIE standard illuminant D65. The average of 3 readings was

3497 used for statistical analysis. Shear force measurements were taken on six 1cm² rectangular cores
3498 using a Lloyd material testing machine (Lloyd Instrument Ltd, Surrey, UK) fitted with a Warner-
3499 Bratzler-like shear blade. Cores were cut at a 200mm/min shear speed and force exerted was
3500 recorded with NEXYGEN software (NEXYGEN Corp. USA). Moisture content (MC) was
3501 determined by expressing weight loss after freeze drying as a percentage of initial weight.

3502 A Soxtec™ 2050 Automatic Extraction apparatus (FOSS Analytical AB, Höganäs, Sweden) was
3503 used for intramuscular fat content determinations. Two grams of ground lyophilized loin muscle
3504 were weighed into pre-weighed cellulose extraction thimbles for the analysis. Petroleum ether
3505 was used as the extraction solvent for the procedure. All determinations were made in duplicate
3506 and the average used for statistical analysis.

3507 **3.2.4 Collagen characteristics**

3508 A complete description of collagen solubility analysis, acid hydrolysis and hydroxyproline
3509 content determination is found in Section 2.2.4. Sample preparation and solubility analysis were
3510 undertaken as described by Hill (1966) while hydroxyproline content in the soluble and insoluble
3511 fractions were determined with the colorimetric assay of Neuman and Logan (1960). Regression
3512 equations obtained from trans-4-hydroxy-L-proline (Sigma Aldrich, MO, USA) standards were
3513 used to predict hydroxyproline content of the samples based on their absorbance readings at
3514 550nm. Hydroxyproline content was subsequently converted to collagen by multiplying by the
3515 conversion factor 7.14.

3516 **3.2.5 Stratification design**

3517 Loin physical, meat quality and collagen characteristics data were stratified into 4 subclasses
3518 based on their intramuscular fat content, shear force, intramuscular pH, and Minolta *L** values

3519 with group 1 having the lowest and group 4 having the highest values (De Vol et al. 1987). For
3520 IMF groupings, group 1 contained individuals with levels ranging between 2.4-4.99% which
3521 were the low IMF individuals (De Vol et al., 1987), groups 2 and 3 (5.01-7.99% and 8.01-
3522 10.98%, respectively) were allocated the next two 2.98% increases in IMF, and group 4 whose
3523 range was 11.00-22.09% were the greater than 11% IMF animals. For the shear force group,
3524 group 1 encompassed individuals with relatively tender pork (Hodgson et al., 1991) with values
3525 ranging between 21.62-39.89N, and groups 2 and 3 had the next two 10 N ranges of WBSF
3526 values of 40.01-49.01N and 50.01-60.99N, respectively, while group 4 contained individuals
3527 with tough pork (Oksbjerg et al., 2001) with WBSF values ranging from 61.49-84.66 N. With
3528 respect to intramuscular pH groupings, group 1 consisted of individuals with potentially
3529 exudative pork with very low pH values ranging from pH 5.20-5.39, group 2 contained the loins
3530 considered to have normal pH values ranging from 5.40-5.68, group 3 contained the high normal
3531 pH loins (pH 5.69-5.89) and group 4 had the loins with pH values equal to and greater than 5.90
3532 (pH 5.90-6.23) that were potentially dark, firm and dry (Honkavaara, 1988). Lastly, for the L^*
3533 groups, group 1 accommodated individuals with pork color that would be considered dark
3534 purplish red to dark reddish pink (Friese et al., 2005) with an L^* range of 33.74-41.97, group 2
3535 had pork considered dark reddish pink with an L^* the range of 41.99-44.99, group 3 had a color
3536 considered reddish pink with L^* values ranging from 45.01-50.89, and group 4 contained pork
3537 considered pink to grayish pink with L^* values ranging from 51.00-56.85 (Friese et al., 2005).

3538 **3.2.6 Statistical analysis**

3539 Data were analyzed as a 2×2 factorial design in R version 3.5.2 software (R Core Team, 2017)
3540 with company (company A vs company B), sex (barrow vs gilt) and their 2-way interactions as
3541 the sources of variation. Least-squares means for specified factors and factor combinations were

3542 obtained with the lsmeans package version 2.27-62 (Lenth, 2016). Mean separation analysis was
3543 undertaken using Tukey multiple comparisons to compare treatment means at an alpha level of
3544 0.05 with the Agricolae package version 1.2-8 (Mendiburu, 2017)

3545 Pearson product-moment correlations were used to explore the linear relationships between and
3546 among the loin physical, meat quality traits and collagen characteristics using the rcorr function
3547 in the Harrell Miscellaneous (Hmisc) package version 4.2-0 (Harrell et al., 2019) in R to
3548 compute matrices of Pearson's r and p-values for all possible pair of traits, based on the
3549 hypothesis that no correlation (zero) exists between any 2 traits under consideration. Pearson
3550 correlations coefficients are calculated using the following formula:

3551 $r_{trait1,trait2} = \frac{covariance(trait1,trait2)}{standard\ deviation\ trait1 \times standard\ deviation\ trait2}$ and standard error (S.E) for the

3552 correlation coefficient is: $S.E = \sqrt{\frac{1-r^2}{n-2}}$ where n= sample size, r= Pearson correlation coefficient.

3553 The t-test for Pearson's $r = \frac{r-\rho}{S.E}$ and since the null hypothesis is that the correlation between the
3554 variables is zero, the ρ term will be dropped out and thus, the t-statistic is essentially Pearson's r
3555 divided by its standard error. Bonferroni multiple-comparison correlation was used to minimize
3556 probability of type 1 error (false positives) results. Thus, because 31 comparisons were made at
3557 an alpha level of 0.05, significant correlations were determined as follows: $0.05/31=0.002$,
3558 therefore correlation coefficients with $p < 0.002$ were reported as significant.

3559 In order to ascertain statistical relationships among the variables in this study, multiple
3560 regression analyses were performed to fit models to predict/estimate loin weight, thick fat, loin
3561 eye weight, bone weight, pH, drip loss, cooking loss, L^*, a^*, b^* , moisture content, Warner-
3562 Bratzler shear force, intramuscular fat content and insoluble collagen using other loin physical,
3563 meat quality traits and collagen characteristics as predictors/independent variables (X) to

3564 generate regression models with the best regression coefficient (R^2) for the regressands (Y) in R
3565 statistical computing environment.

3566 The fitted models were of the form: $\hat{Y}_i = \hat{\beta}_0 + \hat{\beta}_1 X_{1i} + \hat{\beta}_2 X_{2i} + \dots + \hat{\beta}_k X_{ki} + e_i$

3567 Where: \hat{Y}_i = response variable

3568 β_0 =population intercept/mean

3569 $\hat{\beta}_1, \hat{\beta}_2, \dots, \hat{\beta}_k$ = regression coefficients

3570 $X_{1i}, X_{2i} \dots X_{ki}$ =predictors

3571 e_i = error term with expectation $E(\varepsilon|X_{1i}, X_{2i} \dots X_{ki})=0$

3572 Correlation among independent variables was tested using the VIF function in car package
3573 version 3.0-2 (Fox and Weisberg, 2011), which computes variance inflation factor (VIF) for
3574 linear models and variables with VIF's greater than 5 were removed. Results from the Pearson
3575 correlations were further used to check for significant relationships among the variable.

3576 The step function in the stats package (R Core Team, 2018) was used for model selection based
3577 on their Akaike information criterion (AIC) values. Backward elimination method was used for
3578 variable selection in the regression models. The method starts with a model including all the
3579 predictors (full model). Predictors with the least contribution to decline in error sum of squares
3580 or increase in R^2 were deleted. This process continued until deletion of any predictor resulted in
3581 substantial decline in R^2 , thus achieving a parsimonious but best model. Pearson correlation
3582 coefficients of the independent variables were examined to ensure that variables in the various
3583 models were not correlated to each other.

3584 **3.3 Results**

3585 **3.3.1 Effect of company of origin and sex on loin physical, meat quality traits**
3586 **and collagen characteristic**

3587 Results for treatments (company and sex) on the traits considered are presented in Tables 3.1-
3588 3.3. The results obtained showed that company had a significant effect ($p < 0.01$) on the loin
3589 weight (LWT), loin eye weight (LEW) and thick fat weight (TFW) of the populations with the
3590 mean values of company A being significantly ($p < 0.01$) greater than company B. There was
3591 however no significant difference ($p > 0.05$) between barrows and gilts for loin physical traits.
3592 Pertaining to intramuscular collagen characteristics, soluble collagen percent was significantly
3593 ($p < 0.01$) higher in company B compared to company A while for insoluble collagen percent,
3594 individuals from company A had significantly ($p < 0.01$) higher mean values relative to company
3595 B. The effect of sex on collagen characteristics was however not significant ($p > 0.05$). Total
3596 collagen was also statistically not significant ($p > 0.05$) regardless of company of origin.
3597 Intramuscular pH and IMF were significantly influenced by company with animals from
3598 company A having significantly ($p < 0.01$) higher pH values. Sex's influence on pH and IMF were
3599 however not significant ($p > 0.05$). Cooking loss (%) was significantly ($p < 0.01$) higher in
3600 company B compared to company A but was not significant ($p > 0.05$) with respect to sex. The
3601 Minolta color traits (L^* , a^* , b^*) were not affected by company of origin, sex or their interaction
3602 ($p > 0.05$), nor were moisture content and shear force.

3603 **3.3.2 Correlation analysis**

3604 Pearson product-moment correlation coefficient results for loin physical traits, meat quality traits
3605 and collagen characteristics are presented in Tables 3.3-3.6.

3606 Loin weight (LWT) was highly significantly related with loin eye weight (LEW) ($r = 0.73$,
3607 $p < 0.0001$), meat trim weight (MTW) ($r = 0.60$, $p < 0.0001$), rib trim weight (RTW) ($r = 0.72$,
3608 $p < 0.0001$) and moderately correlated fat trim weight (FTW) ($r = 0.40$, $p < 0.0001$), and loin eye
3609 percent (LEP) ($r = -0.35$, $p < 0.0001$). Loin eye weight on the other hand was not highly correlated
3610 with any of the loin physical traits but had low-moderate correlations. It was moderately
3611 correlated with MTW ($r = 0.56$, $p < 0.0001$), RTW ($r = 0.50$, $p < 0.0001$), bone weight (BWT) ($r =$
3612 0.43 , $p < 0.0001$), and LEP ($r = 0.37$, $p < 0.0001$) and lowly correlated with bone percent (BP) ($r =$
3613 -0.20 , $p < 0.001$) and TFW ($r = 0.26$, $p < 0.001$).

3614 Pertaining to TFW, it was highly negatively correlated with LEP ($r = -0.70$, $p < 0.0001$) but
3615 positively correlated with thick fat percent (TFP) ($r = 0.90$, $p < 0.0001$) as anticipated. It was
3616 further moderately associated with FTW ($r = 0.39$, $p < 0.0001$), RTW ($r = 0.39$, $p < 0.0001$), RTP (r
3617 $= -0.32$, $p < 0.0001$) and BP ($r = -0.37$, $p < 0.0001$) and lowly related with MTW ($r = 0.18$, $p < 0.01$)
3618 and BWT ($r = 0.28$, $p < 0.0001$). Rib trim weight, BWT and RTP were significantly correlated
3619 with MTW with correlation coefficients of 0.77, 0.50 and 0.42 respectively. Fat trim weight
3620 (FTW) was positively correlated with RTW ($r = 0.63$, $p < 0.0001$), TFP ($r = 0.28$, $p < 0.0001$) and
3621 RTP ($r = 0.43$, $p < 0.0001$) but negatively correlated with LEP ($r = -0.41$, $p < 0.0001$) and BP ($r = -$
3622 0.21 , $p < 0.0001$). Rib trim weight had low to moderate correlations with LEP ($r = -0.28$,
3623 $p < 0.0001$), BW ($r = 0.48$, $p < 0.0001$) and RTP ($r = 0.59$, $p < 0.0001$). An expected high but
3624 negative correlation was obtained between TFP and LEP ($r = -0.77$, $p < 0.0001$). Thick fat percent
3625 (TFP) was further negatively correlated with RTP ($r = -0.36$, $p < 0.0001$) and BP ($r = -0.34$,
3626 $p < 0.0001$).

3627 Correlation coefficients for meat quality traits and collagen characteristics are shown in Table
3628 3.5. Muscle pH was negatively correlated with L^* ($r = -0.29$, $p < 0.0001$), b^* ($r = -0.36$, $p < 0.0001$)

3629 and Hue ($r = -0.39$, $p < 0.0001$) but positively correlated with moisture content (MC) ($r = 0.36$,
3630 $p < 0.0001$). Drip loss (DL) was positively associated with CL ($r = 0.23$, $p < 0.0001$) and WBSF (r
3631 $= 0.39$, $p < 0.0001$) but was not significantly related to any other meat quality trait. Total cooking
3632 time (TCT) had low to moderate correlation with the Minolta color traits ranging from $r = 0.20$,
3633 $p < 0.0001$ for a^* to $r = -0.36$, $p < 0.0001$ for Hue. Total cooking time was further positively
3634 correlated with cooking loss (%) ($r = 0.43$, $p < 0.0001$). Cooking loss percent (CL) on the other
3635 hand, was negative related with all the Minolta color traits except with LOINA ($r = 0.31$,
3636 $p < 0.0001$) and its correlations with LOINL, LOINLB and LOINH were -0.34 , -0.27 and -0.41
3637 respectively. Cooking loss percent further had a low but positive correlation with WBSF ($r =$
3638 0.27 , $p < 0.001$).

3639 Minolta color traits were generally moderately to highly collinearly related amongst themselves
3640 with correlation coefficients ranging from 0.40 between LOINB and LOINC to 0.91 between
3641 LOINA and LOINC, although LOINL was lowly correlated with LOINC (-0.26). Moisture
3642 content as anticipated, was moderately but negatively correlated with fat traits; its correlation
3643 with CF and IMF were $r = -0.36$, $p < 0.0001$ and $r = -0.59$, $p < 0.0001$, respectively. Furthermore,
3644 its relationship with FOSS crude protein (FOP) (0.30) and FOSS crude moisture (FOM) (0.38)
3645 were also moderate. Warner-Bratzler shear force (WBSF) had a significant but low positive
3646 correlation with FOM ($r = 0.25$, $p < 0.001$). There was strong correlation among the traits
3647 measured with the commercial meat analyzer namely Foss crude fat (FOF), FOP and FOM
3648 ranging from 0.80 between FOP and FOM to -0.98 between FOF and FOM. FOSS crude fat was
3649 positively correlated with intramuscular fat (IMF) ($r = 0.45$, $p < 0.0001$) while FOM ($r = -0.42$,
3650 $p < 0.0001$) and FOP ($r = -0.41$, $p < 0.0001$) had negative correlations with IMF. With regard to
3651 collagen characteristics, soluble collagen percent (SOLC) had a low negative correlation ($r = -$

3652 0.30) with total collagen (TSOL) while insoluble collagen percent's (ISOLC) correlation with
3653 TSOLC was positive (0.30).

3654 Pearson correlation coefficients of sub-primal traits with meat quality traits and collagen
3655 characteristics are shown in Table 3.6. Significant correlation coefficients of sub-primal
3656 variables with the meat quality traits and collagen characteristics were generally low to moderate
3657 except for correlations for TFP with FOM ($r = -0.60$, $p < 0.0001$) and FOF ($r = 0.62$, $p < 0.0001$)
3658 which were high. Loin weight was moderately correlated with pH (0.30) but lowly related to
3659 LOINL (-0.22), LOINH (-0.21), FOF (0.24), FOP (-0.27), FOM (-0.21). Thick fat weight was
3660 moderately correlated with FOF (0.54), FOP (-0.50) and FOM (-0.51) and IMF (0.36). Bone
3661 weight had a low correlation with MC (0.23). Loin eye percent was moderately correlated with
3662 FOF (-0.55), FOP (0.42), FOM (0.57) and IMF (-0.41). Thick fat percent on the other hand had
3663 moderate relations with FOP (-0.54) and IMF (0.46). Significant correlations of RTP and BP
3664 with meat quality traits and collagen characteristics were all low. No significant associations
3665 were obtained between sub-primal variables and the collagen characteristics.

3666 **3.3.3 Regression analysis**

3667 Presented in Table 3.7 and 3.8 are the significant multiple linear regression analyses between
3668 loin physical traits, meat quality trait and collagen characteristics.

3669 The multiple regression analysis for LWT was performed with, LEW ($p < 0.001$), DL ($p = < 0.05$),
3670 a^* ($p < 0.001$) and MC. Fifty three percent of the variation in loin weight was explained by
3671 variation in loin eye weight. Sixty one percent of the variation in TFW was explained by the
3672 following variables: LWT ($p < 0.001$), DL ($p < 0.05$), a^* ($p < 0.05$) and ISOLC ($p < 0.001$). Model
3673 for LEW included, LWT ($p < 0.001$), DL ($p < 0.001$), and a^* ($p < 0.01$) and they explained 56% of

3674 the variation in loin eye weight. The statistical significance of the p-values showed that DL and
3675 a^* play vital roles in prediction of the regressands above.

3676 Regression equations for loin physical regressands were:

3677 $LWT = 0.25 + 1.94 \times (\text{Loin eye weight}) - 0.01 \times (\text{Drip loss}) + 0.02 \times (\text{Minolta } a^*)$

3678 $TFW = -0.35 + \text{Loin weight} \times (0.37) - 0.00 \times (\text{Drip loss}) - 0.01 \times (\text{Minolta } a^*) + 0.00 \times (\text{Insoluble}$
3679 $\text{collagen percent})$

3680 $LEW = 0.15 + 0.28 \times (\text{Loin weight}) + 0.00 \times (\text{drip loss}) - 0.01 \times (\text{Minolta } a^*)$

3681 Summaries of regressions for meat quality regressands are presented in Table 3.8. For
3682 intramuscular pH, 12% of its variation were explained by moisture content while L^* and Thick
3683 fat weight explained 9% and 8% respectively. Loin eye percent explained only 4 % of the
3684 variations in drip loss. For CL, the regression analysis retained total cooking time (TCT)
3685 ($p < 0.001$), Minolta chroma ($p < 0.001$), and SOLC ($p < 0.001$) and intramuscular fat content (%)
3686 and they described 25% of the variation in cooking loss. Twenty-six percent of the variation in
3687 WBSF was model by DL ($p < 0.001$), FOF ($p < 0.001$) and Minolta chroma ($p < 0.001$) with DL
3688 explaining 15% of the variation in WBSF. Minolta L^* was best modeled with pH ($p < 0.01$), and
3689 a^* ($p < 0.001$), and they explained 37% of the variation in L^* . The backward selection retained
3690 TFW ($p < 0.01$), WBSF ($p < 0.001$), and L^* ($p < 0.001$) in the multiple regression model of Minolta
3691 a^* . Minolta b^* was best modelled by pH and chroma as they explained 13 and 16% of its
3692 variation respectively. Stepwise regression to assess the contribution various independent
3693 variables to variations in insoluble collagen percent selected intramuscular pH, total collagen
3694 (mg/g raw meat) and intramuscular fat and these variables predicted 12% of its variation.

3695 With reference to MC, the stepwise approach retained pH ($p<0.001$) and IMF ($p<0.001$) and
3696 described 50% of the variation in moisture content. P-values for TFW ($p<0.001$), MC ($p<0.001$),
3697 and ISOLC ($p<0.05$) partial regression coefficients were significant versus IMF and explained
3698 44% of the variation in intramuscular fat content. Cooking time explained 18% of the variance in
3699 CL while a^* explained the most variation in L^* (28%). IMF on the other hand, explained most of
3700 the variation in MC, explaining 35% of the variation.

3701 Regression equations for cooking loss (%), pH and shear force (N) were:

3702 $CL = 16.23 + 0.09 \times (\text{cooking time}) + 0.48 \times (\text{Minolta chroma}) + 0.08 \times (\text{soluble collagen } (\%))$

3703 $pH = 0.95 + 0.56 \times (\text{Thick fat weight}) - 0.01 \times (\text{Minolta } L^*) + 0.07 \times (\text{Moisture content})$

3704 $WBSF = 58.80 + 1.46 \times (\text{Drip loss}) - 0.30 \times (\text{FOSS crude fat}) - 1.61 \times (\text{Minolta chroma})$

3705 **3.3.4 Stratification analysis**

3706 Data obtained from the 500 loins were stratified into 4 groups based on their intramuscular fat
3707 content, shear force, pH and Minolta L^* values. Mean separation analysis for selected physical
3708 and meat quality measurements stratified according to their ether extractable intramuscular fat
3709 content are presented in Table 3.9, those stratified by Warner-Bratzler shear force are in Table
3710 3.10 while Tables 3.10 and 3.11 represent those of pH and L^* respectively. The stratification
3711 system effectively partitioned the loins into groups that differed in physical, meat quality and
3712 collagen characteristics. Stratification using IMF level or shear force could not successfully
3713 explain variation in pH, stratification by L^* however achieved significant differences in pH.
3714 Stratification based on IMF content successfully explained variations in back fat weight, loin eye
3715 percent, back fat percent, moisture content, shear force and FOSS protein. Stratification based on

3716 WBSF successfully explained variations in loin eye percent, back fat percent, drip loss, cooking
3717 loss and FOM and FOM, but could not explain variation in back fat weight, moisture content,
3718 and soluble collagen percent. Stratification by pH on the other hand explained differences in
3719 back fat weight, loin eye weight, a^* , L^* , cooking loss, moisture content and soluble collagen
3720 percent. Groupings based on L^* explained dissimilarities in pH, loin eye weight, a^* , IMF, CL,
3721 FOSS protein and total collagen.

3722 Specifically, intramuscular fat groups 1 and 2 had significantly ($p<0.01$) higher loin eye percent
3723 and moisture content but lower back fat weight and back fat percentages. Intramuscular fat group
3724 4 (which contained the high IMF individuals) on the other hand, had the lowest ($p<0.01$)
3725 moisture content of the 4 IMF groups as well as the highest backfat percent. Shear force group 1
3726 had significantly ($p<0.01$) lower drip loss and cooking loss (%). Samples in shear force group 4
3727 had the highest ($p<0.01$) loin eye percent, FOSS moisture content and lowest back fat percent
3728 although not significantly different from group 2 and 3. For the pH groups, groups 1 and 2 had
3729 significantly ($p<0.01$) lower backfat weight and loin eye weight but higher a^* . Group 1 further
3730 had the lowest ($p<0.01$) moisture content of the 4 pH groups. For pH group 4, it had the lowest
3731 a^* , L^* , cooking loss, although not significantly different from group 3. Intramuscular pH group 4
3732 further had the lowest soluble collagen percent although not significantly different from groups 2
3733 and 3. Animals in L^* group 1 had significantly ($p<0.01$) lower IMF but higher loin eye weight,
3734 pH and moisture content of the L^* groups while those in group 4 had the lowest ($p<0.01$) a^* and
3735 cooking loss. Total collagen was significantly lower ($p<0.01$) for group 1 although not
3736 significantly different from group 2.

3737 **3.4 Discussion**

3738 The regression coefficient of the LWT model indicated that LEW is the most important variable
3739 to variations in loin weight. Regression analysis by Lisiak et al. (2015) and Wilson et al. (2016)
3740 for loin primals and loin respectively however both retained back fat in the final models. The
3741 present study did not include backfat (TFW) in the LWT models due to issues with
3742 multicollinearity.

3743 Cooking time being the most important factor predicting cooking loss in the present study is not
3744 surprising as Bouton et al. (1976) indeed obtained higher cooking loss with increasing cooking
3745 time. Products with higher fluid losses upon cooking are also known to have lower juiciness
3746 ratings (Channon et al., 2016) thus, making cooking loss important to eating quality. A key
3747 variable influencing variations in pork quality is pH. The pH accounted for variation in L^* (10%)
3748 and MC (12%) in this study and it was also among the traits that predicted variations in insoluble
3749 collagen percent. This is expected as the quality defects PSE and DFD are intimately related to
3750 these traits. Muscle pH contributions to variations in L^* reported by Boler et al. (2010) were
3751 28%, evidently higher than the estimate obtained in this study. These differences could be due to
3752 differences in postmortem treatment of the samples. For instance, Boler et al. (2010) stored their
3753 samples at 4°C for 21 days before taking quality measurements while this current study's
3754 samples were stored at -20°C and thawed for 61hrs at 4°C before measurements. Dransfield et al.
3755 (1995) reported that pH accounted for as high as 53% of the variations in L^* . However, it must
3756 be noted that their L^* measurements were taken with a Hunter D25 color difference meter with a
3757 sample size of 64 pigs while this current study employed a Minolta chromameter with a sample
3758 size of 500 pigs. The contribution of pH to variations in the meat quality traits under
3759 consideration in this study indicate that for frozen-thawed samples, other traits in addition to

3760 ultimate pH is needed to accurately predict pork quality. The fact that intramuscular fat content
3761 (lipid content) explained 35% of the variation in moisture content while moisture content also
3762 explained 35% of the variation in intramuscular fat content confirms the already established
3763 relationship between those two traits for predicting variations in each other. Indeed, a strong
3764 genetic relationship was obtained between those two traits in study 2.

3765 A prediction equation for Warner Bratzler shear force by DeVol et al (1988) also retained fat
3766 percentage as a variable contributing to variability of shear force, which agrees with the finding
3767 of this study. Intramuscular fat may have contributed to variation in insoluble collagen percent as
3768 explained by Nishimura (2010), who suggested that deposition of fat in the perimysium results in
3769 remodeling of the extracellular matrix and reduction in the mechanical strength of the
3770 intramuscular connective tissue through disruption of the collagen fibers' ability to form
3771 longterm crosslinks.

3772 Freezing is a technology extensively used in the meat industry to preserve meat and to ensure
3773 long term usability (Estévez et al., 2011). However, in the course of freezing pork for storage, its
3774 myofibrillar proteins undergo oxidative reactions resulting in the formation of inter and
3775 intramolecular crosslinks, carbonylation of specific amino acids, modification of side chains,
3776 peptide scission and loss of functional groups (Xiong, 2000; Estévez et al., 2011). Indeed, studies
3777 by Estévez et al. (2011) on the effect of frozen-thawed storage on protein carbonylation
3778 confirmed that myofibrillar proteins of porcine *m. longissimus dorsi* undergo oxidative reactions
3779 during postmortem frozen storage, leading to the formation of α -amino adipic and γ -glutamic
3780 semialdehydes from certain amino acids namely: arginine, proline and lysine. This oxidation of
3781 myofibrillar proteins has been found to significantly decrease water holding capacity (Lu et al.,
3782 2017; Bao and Ertbjerg, 2018) and porcine longissimus muscle with higher drip loss also tend to

3783 have higher protein oxidation as shown in the results of Traore et al. (2012). Rowe et al. (2004)
3784 further found oxidation conditions to decrease beef tenderization through μ -calpain inactivation.
3785 Indeed, m and μ -calpain possess oxidizable cysteine residues at their active sites and thus
3786 oxidizing conditions inhibit their proteolytic activity (Guttmann, 1997). It is therefore not
3787 surprising that drip loss predicted most of the variation in tenderness in the present study which
3788 goes on to suggests that pork samples in the present study may have gone extensive myofibrillar
3789 protein oxidation during storage.

3790 Pork producers are always looking for ways to improve production efficiency and profitability
3791 through the use of the best animals for breeding and the fine tuning of production practices. The
3792 finding that sex had no influence on loin weight and loin eye weight agree with the results
3793 obtained by Bertol et al. (2015) and Cisneros et al. (1996). Ramirez and Cava (2007) however
3794 obtained statistically higher loin weights in barrows compared to gilts. Their measurements were
3795 however on whole loin primals while the present study took measurements on sub-primals.
3796 Cisneros et al. (1996) further reported no significant differences in backfat thickness (cm)
3797 measured at the 1st rib between barrows and gilts, however for the measurement taken at the 10th
3798 rib barrows had a significantly higher backfat thickness (cm). Lattore et al. (2004), Ramirez et al.
3799 (2007) and Kim et al. (2018) also reported significantly higher backfat (mm) in barrows
3800 compared to gilts which disagrees with results of this study as no significant differences were
3801 obtained among the sexes although it must to stated that in the current study, sub-primal backfat
3802 weight was measured rather than backfat depth.

3803 The results from the stratification analysis suggest that samples with lower IMF levels tend to be
3804 tougher and have higher moisture content, lean percent, and cooking losses while vice versa can
3805 be said for high IMF samples. These results agree with the results obtained by Hodgson et al.

3806 (1991) after that study stratified samples into 10 marbling groups. Hodgson et al. (1991) further
3807 reported an increase in protein percent and moisture percent but a decrease in fat percent for
3808 samples in their high shear force groups which indeed agrees with the results reported in the
3809 present study.

3810 Moisture content in meat plays a crucial role in product shelf life, palatability and functionality
3811 (Branden, 2013). Indeed, approximately 75% of muscle composition is water (Voyle, 1979).
3812 However, after slaughter, alterations in homeostatic processes causes an increase in lactic acid
3813 concentration leading to a decline in meat pH (Braden, 2013). This causes changes in water
3814 holding capacity and consequently, moisture content. Thus, water holding capacity and moisture
3815 content decreases as muscle pH approach its isoelectric point (minimum hydration). Indeed, the
3816 major muscle proteins, myosin and actin have their isoelectric points at 5.3 and 4.8 respectively
3817 (Rusell, 2000) which suggest that myosin's ability to hold on to water molecules will be minimal
3818 at the aforementioned pH. This decline in pH can lead to muscle defects like PSE condition
3819 resulting in economic losses. It is therefore not surprising to observe that, animals that fell into
3820 the high pH group (group 4) had significantly higher ($p < 0.01$) moisture content while those in
3821 group 1 had the lowest means. The stratification based on pH further suggested that animals with
3822 higher pH tend to also have significantly higher backfat but lower L^* , drip loss, cooking loss,
3823 and soluble collagen percent, this result sheds further light on the potential of stratification by pH
3824 to predict certain quality attributes. Results from the L^* stratification suggests that animal with
3825 lower luminosity values also tend to have higher pH, lower IMF and total collagen, but higher
3826 loin eye muscle weight and moisture content.

3827 Meat appearance on retail shelves has tremendous influences on initial consumer perceptions of
3828 product quality. Latorre et al. (2004), Kim et al. (2018) and Zemra et al. (2015) reported no

3829 significant differences in Minolta color traits regardless of sex. These results are in agreement
3830 with those of this study and thus suggests that pork color traits are not significantly influenced by
3831 animal sex. One important indicator of pork quality is its ability to retain moisture (Huff-
3832 Lonergan and Lonergan, 2005). Muscle pH and drip loss are indirect indicators of moisture
3833 retention potential. The finding that irrespective of sex, muscle pH and drip loss were similar
3834 between barrows and gilts agrees with reports by Ramirez and Cava (2007), Cisneros et al.
3835 (1996) and Franco et al. (2014). Indeed Barton-Gade (1987) in an investigation of meat quality
3836 in castrates and gilts from 3 breed combination: Large White×(Large White× Landrace),
3837 Duroc×(Large White× Landrace) and Hampshire×(Large White× Landrace) concluded that sex
3838 had no significant effect on the traits that reflect the PSE/DFD defect in their study's carcasses.

3839 Castrated males in the studies of Franco et al. (2014) and Alonso et al. (2009) had higher
3840 intramuscular fat contents compared to entire females. Also, studies in Quebec by Correa et al.
3841 (2006) of pigs from similar breed combination as this study (Duroc× (Landrace× Large White)
3842 revealed significantly higher IMF in barrows. These differences with the present study could be
3843 due to population variations and differences in experimental design. It is, however, worth noting
3844 that barrows in this study on average, had higher IMF although not statistically significant
3845 ($p>0.05$). Ramirez and Cava (2007), Kim et al. (2018) and Sundrum et al. (2011) however found
3846 no effect of sex on intramuscular fat content, which agrees with the results of the present study.
3847 Although, it is also worth noting that in all those studies, castrated males had numerically higher
3848 intramuscular fat content that was not significant statistically.

3849 Meat tenderness is a highly ranked quality attribute in meat products (Williams et al., 1986).
3850 Warner Bratzler shear force test is a method established as an objective measure of meat
3851 tenderness. Shear force was not significant ($p>0.05$) between sex in the study of Magowan et al.

3852 (2011) and Kim et al. (2018), which agrees with the findings of this study. Barton-Gade (1987)
3853 however reported a 3N difference between gilts and barrows with gilts having higher estimates
3854 on average though not statistically significant ($p>0.05$). Sex, however, had a significant effect on
3855 cooking loss as reported by Kim et al. (2018), Magowan et al. (2011) and Latorre et al. (2004)
3856 which contradicts the findings of the present study. These differences could, however, be due to
3857 several factors such as statistical modeling and differences in data collection. As an instance,
3858 Kim et al. (2018) cooked samples at 80°C for 30min while Magowan et al. (2011) cooked
3859 samples at 70°C for 50min. However, in the present study, samples were cooked to an internal
3860 temperature of 71°C as monitored by a probe and therefore, resident time in the water bath
3861 varied. Other investigators who have studied the effect of sex on cooking loss, however, reported
3862 no significant ($p>0.05$) influence of sex on cooking loss (Cisneros et al., 1996; Beattie et al.,
3863 1999 and Ramirez and Cava, 2007). With respect to collagen characteristics, sex had no
3864 influence on intramuscular collagen characteristics in the study of Correa et al. (2006) which
3865 agrees with the results of the present study. The results of the present study generally agree with
3866 the conclusions drawn by Correa et al. (2006) that sex has no significant effect on pork quality
3867 traits.

3868 Pearson correlation is used to give an indication of the strength and direction of the linear
3869 relationship between 2 variables. The high correlation of LWT with LEW and TFW indicate that
3870 samples with heavier loins were likely to also have heavier loin eye's and backfat. The negative
3871 association of LEP with TFW suggests that heavily muscled loins tend to however have lower
3872 backfat contents. Furthermore, the moderate-high correlation of loin percent with crude fat,
3873 crude protein, and crude moisture indicates that highly muscled samples tend to have lower fat
3874 content but higher protein and moisture contents.

3875 Brewer et al. (2001) and Van Laack and Kauffman (1999) obtained a high negative correlation
3876 between pH and L^* while the degree of association in this study was low. Hambrecht et al.
3877 (2003) however obtained similar associations of pH and L^* (-0.33) as this study even though that
3878 study measured pH at 18hrs. Correlation between shear force and drip loss was low and not
3879 significant in the study of Cannata et al. (2010). However, the present study obtained a
3880 significant moderate correlation between those two traits. These differences could be due to
3881 several factors such as sample size and trait measurement differences as Cannata used a sample
3882 size of 53 and measure slice shear force with a crosshead speed of 500 mm/min. In fact,
3883 crosshead speed has been found to influence shear force values (Wheeler et al., 1997).

3884 Cooking loss is a trait of appreciable importance in fresh as well as processed meat products as
3885 the ability of a product to retain its moisture during cooking enhances juiciness for the consumer
3886 and profitability for the processor (Channon et al., 2016). Cooking rate (Boles and Swan, 2002),
3887 cooking temperature (Shin et al., 1992) and end-point temperature (Fjelkner-Modig, 1986) are
3888 some factors affecting cooking loss. The moderate positive correlation of cooking time with
3889 cooking loss indicates that steaks that take longer to reach an internal temperature of 71°C also
3890 tend to have more fluid losses associated with it. The low positive correlation of drip loss with
3891 cooking loss obtained in this study agrees with the results of Watanabe et al. (2018). The positive
3892 correlation of shear force with cooking loss agrees with the results of Bouton et al. (1975) and Li
3893 et al. (2013). Thus, samples that loose more fluids during cooking tend to be tougher. Since a
3894 significant proportion of muscle water can be found between and within myofibrils, and between
3895 muscle bundles, heat application will cause a release of loosely bound water molecules resulting
3896 in an increase in fiber density and ultimately a tougher product.

3897 The high association of crude fat with crude protein and moisture reported in this study agrees
3898 with the results of Hodgson et al. (1991). Hodgson et al. (1991) however reported a moderate
3899 correlation of moisture percent and protein percentage while correlation obtained in this study
3900 was high. Further, Watanabe et al. (2018) also obtained a high negative correlation of IMF with
3901 moisture content but not with cooking loss or drip loss in their study with commercial Japanese
3902 hybrid (Large White × Landrace) × Duroc pigs which agrees with results obtained in the
3903 present study. Matured adipose cells tend to have very sparse cytoplasm with approximately 85%
3904 of its volume being filled with triglyceride (Swatland, 1994) and as a consequence, increased
3905 deposition of intramuscular fat results in a reduction in moisture content.

3906 **3.5 Conclusion**

3907 Although the debate on the role of collagen fiber state on fluid losses from cooked meat may
3908 persist in the meat science community, results from the present study indicated that collagen
3909 characteristics do indeed influence variations in cooking loss. This study further indicated that
3910 Warner-Bratzler shear force values may be driven more by denaturation of myofibrillar protein
3911 native structure than by intramuscular fat content. Mean separation analysis also showed WBSF
3912 to increase with increased loin eye muscle weight and reduced backfat which suggests that
3913 selection for rapid growth rate with its associated reduction in calpain activity may increase pork
3914 toughness. This study further implicated pH and intramuscular fat content as variables
3915 influencing insoluble collagen percent, indicating that deposition of fat in the perimysium may
3916 be affect the level and/or amount of crosslink formation. The stepwise regression further
3917 suggested that pH is not a reliable sole predictor of meat quality. The present study also
3918 confirmed that high lean or muscle content tended to be associated with high moisture but lower

3919 fat content which suggests that lean sub-primals will also tend to have low fat content which
3920 indirectly affect eating quality.

3921 **3.6 Tables**

3922 Table 3. 1: Effect of company and sex on collagen characteristics of porcine m. longissimus dorsi

Traits	Factors				P-value		
	Company		Sex		Company	Sex	Company× Sex
	Company A	Company B	Barrow	Gilt			
Soluble collagen (%)	16.83 ±0.44	19.10 ±0.57	17.83 ±0.48	18.15 ±0.53	<0.01	0.43	0.18
Insoluble collagen (%)	83.21 ±0.44	80.89 ±0.57	82.65 ±0.48	81.94 ±0.53	<0.01	0.43	0.18
Total collagen (mg/g raw meat)	3.00 ±0.07	3.17 ±0.09	3.06 ±0.07	3.07 ±0.08	0.14	0.94	0.52

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Table 3. 2: Effect of company and sex on loin physical traits of porcine m. longissimus dorsi

Traits	Factors				P-value		
	Company		Sex		Company	Sex	Company× Sex
	Company A	Company B	Barrow	Gilt			
Loin weight (kg)	1.46 ±0.01	1.30 ±0.01	1.40 ±0.01	1.39 ±0.01	<0.01	0.85	0.61
Loin eye weight (kg)	0.54 ±0.01	0.50 ±0.01	0.53 ±0.01	0.52 ±0.01	<0.01	0.52	0.89
Thick fat weight (kg)	0.36 ±0.01	0.26 ±0.01	0.32 ±0.01	0.32 ±0.01	<0.01	0.55	0.39

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3934 Table 3. 3: Effect of company and sex on meat quality traits of porcine m. longissimus dorsi

Traits	Factors				P-value		
	Company		Sex		Company	Sex	Company × Sex
	Company A	Company B	Barrow	Gilt			
PH	5.61±0.01	5.50±0.01	5.56±0.01	5.57±0.01	<0.01	0.41	0.56
Drip loss (%)	6.48±0.20	7.26±0.26	6.80±0.22	6.74±0.24	0.02	0.72	0.46
Cooking loss (%)	24.21±0.24	26.45±0.31	25.10±0.25	25.00±0.28	<0.01	0.50	0.92
Minolta <i>L</i> *	45.26±0.24	45.39±0.31	45.25±0.26	45.38±0.28	0.75	0.75	0.65
Minolta <i>a</i> *	7.59±0.10	7.64±0.12	7.53±0.10	7.70±0.11	0.75	0.28	0.46
Minolta <i>b</i> *	3.36±0.08	3.39±0.10	3.32±0.09	3.43±0.09	0.79	0.39	0.94
Moisture content (%)	72.26±0.06	72.19±0.08	72.21±0.07	72.25±0.07	0.49	0.64	0.28
Warner-Bratzler Shear force (N)	47.32±0.83	47.93±1.07	48.47±0.90	46.40±0.99	0.64	0.10	0.15
Intramuscular fat	8.35±0.20	7.51±0.26	8.10±0.22	7.95±0.24	0.01	0.78	0.90

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Table 3. 4: Pearson correlation coefficients between loin physical traits of porcine m. longissimus dorsi

Trait	LEW	TFW	MTW	FTW	RTW	BWT	LEP	TFP	RTP	BP
LWT ¹	.73*	.76*	.60*	.40*	.72*	.53*	-.35*	.42*	-.11	-.30*
LEW ²	1.00	.26*	.56*	.10	.50*	.43*	.37*	-.12	-.14	-.20*
TFW ³		1.00	.18*	.39*	.39*	.28*	-.70*	.90*	-.32*	-.37*
MTW ⁴			1.00	-.01	.77*	.50*	-.02	-.12	.42*	.02
FTW ⁵				1.00	.63*	.14	-.41*	.28*	.42*	-.21*
RTW ⁶					1.00	.48*	-.28*	.09	.59*	-.12
BWT ⁷						1.00	-.16	.04	.03	.63*
REP ⁸							1.00	-.77*	0.00	.12
TFP ⁹								1.00	-.36*	-.34*
RTP ¹⁰									1.00	.14
BP ¹¹										1.00

3947 1-Whole loin Weight; 2- Loin eye Weight; 3- Thick fat weight; 4- Meat trim weight; 5- Fat trim
 3948 weight; 6- Rib trim weight; 7-Bone weight; 8- Loin eye percent; 9- Thick fat percent; 10- Rib
 3949 trim percent; 11- Bone percent *P<0.002.

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3957 Table 3. 5: Pearson correlation coefficients between and meat quality traits and collagen characteristics of porcine m. longissimus
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Trait	DL	TCT	CL	<i>L</i> [*]	<i>a</i> [*]	<i>b</i> [*]	hue	chr	MC	WBSF	FOF	FOP	FOM	SOLC	ISOLC	TCOL	IMF
PH	.07	.12	-.06	-.29*	.00	-.36*	-.34*	-.10	.36*	.10	.10	-.11	-.05	-.09	.09	-.14	.07
DL ¹	1.00	-.11	.23*	.12	-.12	.09	.12	-.08	.12	.39*	-.14	.21*	.14	.08	-.08	.05	-.11
TCT ²		1.00	.43*	-.30*	.20*	-.29*	-.36*	.07	.05	.00	.02	-.17	.04	.00	.00	-.12	-.05
CL ³			1.00	-.34*	.31*	-.27*	-.41*	.15	.00	.27*	-.06	-.12	.12	.13	-.13	-.02	-.12
<i>L</i> ⁴				1.00	-.53*	.68*	.89*	-.26*	-.11	-.07	-.06	.26*	-.05	.04	-.04	.18	.08
<i>a</i> ⁵					1.00	.03	-.45*	.91*	-.28*	-.18	.26*	-.34*	-.21*	.04	-.04	-.01	.26*
<i>b</i> ⁶						1.00	.86*	.40*	-.38*	-.21*	.13	.10	-.24*	.03	-.03	.23*	.30*
Hue ⁷							1.00	-.10	-.19	-.11	-.03	.27*	-.09	.00	.00	.22*	.13
Chr ⁸								1.00	-.38*	-.25*	.28*	-.26*	-.28*	.04	-.04	.07	.34*
MC ⁹									1.00	.17	-.36*	.30*	.38*	-.08	.08	-.05	-.59*
WBSF ¹⁰										1.00	-.22*	.17	.25*	-.05	.05	-.09	-.12
FOF ¹¹											1.00	-.88*	-.98*	-.02	.02	.02	.45*
FOP ¹²												1.00	.80*	.01	-.01	.05	-.41*
FOM ¹³													1.00	.03	-.03	-.06	-.42*
SOLC ¹⁴														1.00	-1.00*	-.30*	.09
ISOC ¹⁵															1.00	.30*	-.09
TCOL ¹⁶																1.00	.02
IMF ¹⁷																	1.00

3959 1- Drip loss; 2- Total cooking time; 3- Cooking loss (%); 4-Minolta *L*^{*} ; 5-Minolta *a*^{*} ; 6-Minolta *b*^{*} ; 7-Minolta hue ; 8-Minolta
 3960 chroma ; 9-Moisture content ; 10- Warner-Bratzler shear force; 11-FOSS crude fat ; 12-FOSS crude protein ; 13-FOSS crude moisture
 3961 ; 14-soluble collagen percent ; 15-Insoluble collagen percent ; 16-Total collagen (mg/g raw meat); 17- Intramuscular fat; *P<0.002

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Table 3. 6: Pearson correlation coefficients between loin physical traits, meat quality traits and collagen characteristics of porcine *m. longissimus dorsi*

Trait	LWT ¹	LEW ²	TFW ³	MTW ⁴	FTW ⁵	RTW ⁶	BWT ⁷	LEP ⁸	TFP ⁹	RTP ¹⁰	BP ¹¹
PH	.30*	.28*	.29*	.13	.10	.16	.11	-.01	.20*	-.10	-.14
DL ¹²	-.01	.13	-.12	.03	.04	.04	.06	.19	-.17	.04	.05
TCT ¹³	.16	.25*	.06	.10	-.04	.05	.01	.15	-.02	-.07	-.12
CL ¹⁴	.12	.22*	-.07	.16	-.11	.06	-.04	.19*	-.16	.00	-.12
<i>L</i> ¹⁵	-.22*	-.29*	-.13	-.09	.08	-.02	.00	-.13	-.05	.18*	.17
<i>a</i> ¹⁶	.14	-.04	.23*	-.05	.02	-.03	-.12	-.21*	.26*	-.17	-.23*
<i>b</i> ¹⁷	-.14	-.35*	.01	-.16	.17	-.02	-.02	-.32*	.12	.10	.09
Hue ¹⁸	-.21*	-.31*	-.11	-.12	.14	-.01	.04	-.18*	-.03	.18*	.21*
Chr ¹⁹	.09	-.14	.23*	-.11	.08	-.03	-.11	-.31*	.29*	-.13	-.18*
MC ²⁰	.11	.35*	-.14	.30*	-.02	.22*	.23*	.32*	-.28*	.15	.14
WBSF ²¹	-.04	.16	-.13	.08	-.21*	-.07	.06	.26*	-.19*	-.07	.08
FOF ²²	.24*	-.17	.54*	-.34*	.55*	.09	-.09	-.55*	.62*	-.12	-.31*
FOP ²³	-.27*	.05	-.50*	.24*	-.37*	-.05	.04	.42*	-.54*	.19*	.28*
FOM ²⁴	-.21*	.21*	-.51*	.35*	-.57*	-.09	.10	.57*	-.60*	.08	.30*
SOLC ²⁵	-.08	-.03	-.16	.02	.06	.05	.00	.07	-.15	.16	.05

ISOLC ²⁶	.08	.03	.16	-.02	-.06	-.05	.00	-.07	.15	-.16	-.05
TCOL ²⁷	-.05	-.14	-.02	-.03	.16	.08	-.06	-.12	-.01	.14	-.01
IMF ²⁸	.11	-.20*	.36*	-.16	.20*	.00	-.09	-.41*	.46*	-.10	-.19*

3967 1-Whole loin weight; 2- Loin eye weight; 3- Thick fat weight 4-Meat trim weight; 5- Fat trim weight; 6- Rib trim weight; 7-
3968 Bone weight; 8- Loin eye percent; 9-Thick fat percent total cooking time; 10- Rib trim percent Cooking loss; 11- Bone percent
3969 ; 12- Drip loss; 13- Total cooking time; 14- Cooking loss (%); 15-Minolta *L** ; 16-Minolta *a** ; 17-Minolta *b** ; 18-Minolta hue
3970 ; 19-Minolta chroma ; 20-Moisture content ; 21- Warner-Bratzler shear force; 22-FOSS crude fat ; 23-FOSS crude protein ; 24-
3971 FOSS crude moisture ; 25-soluble collagen percent ; 26-Insoluble collagen percent ; 27-Total collagen (mg/g raw meat); 28-
3972 Intramuscular fat; *P<0.002.

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3990 Table 3. 7: Multiple linear regression analysis of loin weight, thick fat and bone weight using other loin physical, meat and collagen
 3991 characteristics.

Dependent Variable	R ²	S. E	Independent variables	Regression coefficient	Partial R ²	P-value
Loin Weight	0.56	0.11	Constant	0.25	-	0.0001
			Loin eye weight	1.94	0.5314	<0.0001
			Drip loss	-0.01	0.0002	0.0128
			Minolta <i>a</i> *	0.02	0.0198	<0.0001
Thick fat	0.61	0.06	Constant	-0.35	-	<0.0001
			Loin Weight	0.37	0.5731	<0.0001
			Drip loss	-0.00	0.0134	0.0184
			Minolta <i>a</i> *	-0.01	0.0507	0.0013
			Insoluble collagen (%)	0.00	0.0240	0.0069
Loin Eye Weight	0.56	0.04	Constant	0.15	-	<0.0001
			Loin weight	0.28	0.5314	<0.0001
			Drip loss	0.00	0.0174	0.0007
			Minolta <i>a</i> *	-0.01	0.0013	0.0008

3992 R² = coefficient of determination (correlation coefficient square); SE = standard error of estimates

3993 All p-values for R² were significant p<0.05

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Table 3. 8: Multiple linear regression analysis of meat quality traits using loin physical, other meat quality traits and collagen characteristics.

Dependent Variable	R ²	S. E	Independent variables	Regression coefficient	Partial R ²	P-value
Intramuscular pH	0.29	0.14	Constant	0.95	-	0.1
			Thick fat weight	0.56	0.0821	<0.0001
			Minolta <i>L</i> *	-0.01	0.0864	<0.0001
			Moisture content	0.07	0.1269	<0.0001
Drip loss	0.04	2.86	Constant	-3.10	-	0.1
			Loin eye percent	-4.74	0.0355	<0.0001
Cooking Loss	0.25	3.05	Constant	16.23	-	<0.0001
			Cooking time	0.09	0.1888	<0.0001
			Minolta chroma	0.48	0.0235	0.001
			Soluble collagen percent	0.08	0.0178	0.004
			Intramuscular fat content (%)	-0.21	0.0136	0.001
Warner-Bratzler Shear force (N)	0.26	10.24	Constant	58.80	-	<0.0001
			Drip loss	1.46	0.1542	<0.0001
			Crude fat	-0.30	0.0494	0.02
			Minolta chroma	-1.61	0.0608	<0.0001
Minolta <i>L</i> *	0.37	2.65	Constant	88.55	-	<0.0001
			pH	-5.99	0.0864	0.0009
			Minolta <i>a</i> *	-1.28	0.2830	<0.0001
Minolta <i>a</i> *	0.35	1.09	Constant	18.01	-	<0.0001
			Thick fat weight	1.98	0.0508	0.0055
			Minolta <i>L</i> *	-0.22	0.2830	<0.0001
			Warner-Bratzler shear force	-0.22	0.0323	<0.0001
Minolta <i>b</i> *	0.27	1.01	Constant	13.46	-	<0.0001
			pH	-2.30	0.1323	<0.0001
			Minolta Chroma	0.32	0.1613	<0.0001
Moisture Content	0.50	0.64	Constant	61.56	-	<0.0001
			pH	2.19	0.1269	<0.0001
			Intramuscular fat	-0.19	0.3523	<0.0001

Intramuscular fat	0.44	2.11	Constant	134.14	-	<0.0001
			Thick fat weight	9.69	0.1314	<0.0001
			Moisture content	-1.73	0.3525	<0.0001
			Insoluble collagen percent	-0.04	0.0086	0.0269
Insoluble collagen percent	0.12	6.00	Constant	46.93	-	<0.0001
			pH	5.55	0.0088	0.01
			Total collagen (mg/g raw meat)	2.06	0.0904	<0.0001
			Intramuscular fat content	-0.23	0.0086	0.04

4004 R^2 = coefficient of determination (correlation coefficient square); SE = standard error of estimates; All p-values for R^2 were
4005 significant $p < 0.05$.

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4018 Table 3. 9: Mean separation analysis for loin physical and meat quality measurements stratified according to their ether extractable
 4019 intramuscular fat content (%)

Intramuscular fat (%) range	Group	n	Backfat weight (kg)	Loin eye (%)	Backfat (%)	pH	Drip loss (%)	Cooking loss (%)	Moisture content (%)	Shear force (N)	FOSS protein (%)
2.4-4.99	1	57	0.28 ±0.01 ^b	40.89 ±0.52 ^a	20.73 ±0.67 ^c	5.58 ±0.02	7.56 ±0.46	25.15 ±0.57	72.81 ±0.12 ^a	49.52 ±1.84 ^{ab}	17.80 ±0.20 ^a
5.01-7.99	2	193	0.30 ±0.01 ^b	39.64 ±0.27 ^a	21.87 ±0.35 ^c	5.56 ±0.01	6.79 ±0.24	25.20 ±0.30	72.57 ±0.06 ^a	49.86 ±0.96 ^a	17.15 ±0.10 ^b
8.01-10.98	3	148	0.35 ±0.01 ^a	37.75 ±0.33 ^b	24.88 ±0.42 ^b	5.55 ±0.01	6.55 ±0.29	25.12 ±0.36	71.98 ±0.07 ^b	44.44 ±1.17 ^b	16.38 ±0.12 ^c
11.00-22.09	4	66	0.39 ±0.01 ^a	36.50 ±0.49 ^b	27.35 ±0.63 ^a	5.61 ±0.02	6.60 ±0.43	23.98 ±0.54	71.19 ±0.11 ^c	45.70 ±1.74 ^{ab}	15.87 ±0.19 ^c
P-value			<0.01	<0.01	<0.01	0.22	0.31	0.24	<0.01	<0.01	<0.01

4020 ^{a, b, c} Means without a common superscript letter within a column are significantly different (P < 0.05); n: sample size.

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4030 Table 3. 10: Mean separation analysis for loin physical and meat quality measurements stratified according to their Warner-Bratzler
 4031 shear force (WBSF) value (N)

WBSF (N)	Group	n	Back-fat weight (kg)	Loin eye (%)	Back-fat (%)	pH	Drip loss (%)	Cooking loss (%)	Moisture content (%)	FOSS fat (%)	FOSS moisture (%)	FOSS protein (%)	Soluble collagen (%)
21.62-39.86	1	149	0.33 ±0.01	37.81 ±0.35 ^b	24.44 ±0.47 ^a	5.55 ±0.01	4.98 ±0.27 ^b	23.69 ±0.36 ^b	72.12 ±0.09	25.80 ±0.47 ^a	55.88 ±0.36 ^b	16.51 ±0.14	17.50 ±0.65
40.01-49.01	2	193	0.32 ±0.01	38.58 ±0.36 ^{ab}	23.17 ±0.48 ^{ab}	5.56 ±0.01	7.34 ±0.28 ^a	25.13 ±0.36 ^a	72.14 ±0.09	24.76 ±0.48 ^{ab}	56.86 ±0.36 ^{ab}	16.83 ±0.14	18.57 ±0.66
50.01-60.99	3	153	0.31 ±0.01	39.43 ±0.35 ^a	23.26 ±0.47 ^{ab}	5.59 ±0.01	7.61 ±0.27 ^a	25.47 ±0.36 ^a	72.34 ±0.09	24.08 ±0.47 ^{ab}	57.39 ±0.36 ^a	17.01 ±0.14	17.11 ±0.65
61.49-84.66	4	66	0.31 ±0.01	40.04 ±0.53 ^a	21.68 ±0.72 ^b	5.58 ±0.02	7.82 ±0.41 ^a	26.67 ±0.54 ^a	72.44 ±0.14	23.11 ±0.71 ^b	58.28 ±0.55 ^a	17.06 ±0.21	17.37 ±0.99
P-value			0.29	0.03	0.01	0.30	<0.01	<0.01	0.11	<0.01	<0.01	0.05	0.43

4032 ^{a, b} Means without a common superscript letter within a column are significantly different (P < 0.05); n: sample size.

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4048 Table 3. 11: Mean separation analysis for loin physical and meat quality measurements stratified according to their pH values

pH	Group	n	Back-fat weight (kg)	Loin eye weight (kg)	<i>a</i> *	<i>L</i> *	Drip loss (%)	Cooking loss (%)	Moisture content (%)	FOSS fat (%)	FOSS moisture (%)	FOSS protein (%)	Soluble collagen (%)
5.20-5.39	1	66	0.31 ±0.01 ^b	0.52 ±0.01 ^b	7.80 ±0.15 ^a	46.39 ±0.35 ^a	6.86 ±0.31	25.14 ±0.40 ^a	71.60 ±0.08 ^c	23.81 ±0.54	57.53 ±0.41	16.99 ±0.14	19.88 ±0.87 ^a
5.40-5.68	2	331	0.32 ±0.01 ^b	0.53 ±0.01 ^b	7.70 ±0.08 ^a	45.74 ±0.19 ^a	6.72 ±0.21	24.91 ±0.19 ^a	72.05 ±0.04 ^b	24.47 ±0.26	56.94 ±0.21	16.93 ±0.08	17.99 ±0.35 ^{ab}
5.69-5.89	3	98	0.36 ±0.02 ^a	0.57 ±0.02 ^a	7.12 ±0.16 ^b	45.10 ±0.40 ^{ab}	6.86 ±0.32	24.00 ±0.39 ^{ab}	72.51 ±0.11 ^a	24.44 ±0.51	57.11 ±0.38	16.90 ±0.16	17.26 ±0.70 ^{ab}
5.90-6.23	4	26	0.38 ±0.02 ^a	0.59 ±0.02 ^a	7.01 ±0.28 ^b	43.65 ±0.72 ^b	6.41 ±0.53	22.11 ±0.93 ^b	72.89 ±0.14 ^a	24.61 ±0.98	57.15 ±0.78	16.60 ±0.25	15.62 ±1.46 ^b
P-value			<0.01	<0.01	<0.01	<0.01	0.78	<0.01	<0.01	0.77	0.70	0.72	0.01

4049 ^{a, b, c} Means without a common superscript letter within a column are significantly different (P < 0.05); n: sample size.

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4067 Table 3. 12: Mean separation analysis for loin physical and meat quality measurements stratified according to their L^* values

L^*	Group	n	pH	Loin eye weight (kg)	a^*	IMF (%)	WBSF (N)	Cooking loss (%)	Moisture content (%)	FOSS fat (%)	FOSS moisture (%)	FOSS protein (%)	Total collagen (mg/g raw meat)	Soluble collagen (%)
33.74-41.97	1	71	5.68 ±0.01 ^a	0.57 ±0.01 ^a	8.40 ±0.15 ^a	6.96 ±0.32 ^b	48.87 ±1.36	24.89 ±0.42 ^{ab}	72.62 ±0.09 ^a	23.70 ±0.61	58.00 ±0.46	16.73 ±0.15 ^b	2.72 ±0.12 ^b	16.65 ±0.88
41.99-44.99	2	189	5.54 ±0.01 ^b	0.53 ±0.01 ^b	8.19 ±0.08 ^a	8.13 ±0.19 ^a	48.42 ±0.85	25.68 ±0.21 ^a	72.06 ±0.05 ^b	24.55 ±0.34	57.17 ±0.26	16.67 ±0.08 ^b	3.09 ±0.07 ^{ab}	17.93 ±0.45
45.01-50.89	3	220	5.57 ±0.01 ^b	0.53 ±0.01 ^b	7.12 ±0.09 ^b	8.44 ±0.24 ^a	46.43 ±0.77	24.26 ±0.25 ^b	72.03 ±0.06 ^b	24.62 ±0.31	56.79 ±0.24	16.93 ±0.09 ^b	3.13 ±0.08 ^a	18.45 ±0.47
51.00-56.85	4	51	5.54 ±0.02 ^b	0.55 ±0.01 ^{ab}	6.00 ±0.13 ^c	8.45 ±0.51 ^a	45.69 ±0.61	21.62 ±0.52 ^c	72.14 ±0.16 ^b	23.51 ±0.80	56.63 ±0.64	18.01 ±0.32 ^a	3.35 ±0.12 ^a	18.72 ±0.84
P-value			<0.01	<0.01	<0.01	<0.01	0.15	<0.01	<0.01	0.28	0.09	<0.01	<0.01	0.24

4068 ^{a, b, c} Means without a common superscript letter within a column are significantly different ($P < 0.05$); n: sample size.

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Chapter 4

Summary and general conclusion

4.1 Summary of results

The resemblance between parents and progeny serves as the foundation for selective breeding (Bourdon, 1995). Thus, the use of desirable animals as sire and dams over time, leads to improvements in the mean phenotypic value of subsequent generations for traits of interests. Now, just as some traits tend to exhibit more resemblance than others, so also some are generally more responsive to selection (Falconer and Mackay, 1996). This notion is the fulcrum on which the progress made by humans in animal improvement and breeding rests.

Global pork production has seen a rapid rise over the past half century from 25 million metric tons in the 1960's to 112 million metric tons in 2013 (Ritchie and Roser, 2019). This ability to produce pork at such high throughput has been possible through the adoption of systematic scientific approaches selection and breeding and the fine-tuning of management practices. The ability to produce pork at such a high level has come with some surmountable challenges. For one, unlike the beef industry which has been able to successfully align its grading system with their consumer choices, with quality grades ranging from Canada A to prime in a country like Canada (BCRC, 2019) with similar grading systems in other developed countries, the pork industry still lags behind with its production at the packing and producer level for the most part, being driven by pounds/kilos of lean pork. However, in the wake of the increasing consumer demand for pork of better quality, coupled with the high variability in terms of quality

characteristics of pork on retail shelves (Salas and Mingala, 2017; De Vol et al., 1988) the pork industry may stand a chance of improving its standing in the food industry by having meat quality standards and updating its grading system. For this to be achieved, various research studies may need to be undertaken to determine phenotypic and genetic relationships of pork quality traits as well as how certain extrinsic and intrinsic characteristics of pork contribute and/or predict quality may need to be examined. This thesis sought to contribute to knowledge in this respect.

In the first study, phenotypic and genetic parameters for loin physical, meat quality and intramuscular collagen characteristics (IMCC) of crossbred Canadian swine populations were examined. Data used were from 500 animals with pedigree information dating back to 8 ancestral generations. The results indicated substantial variation in loin physical attributes, meat quality and IMCC of the populations studied. A similar trend has been reported in other countries (De Vol et al., 1988). Consistency of expectation in the perceptible attributes of the products consumers purchase off-shelf is crucial in their purchasing decisions (Uri, 1995; Ha and Jang, 2012) and thus, this variability may prove to be a disadvantage for the pork industry. In general, moderate to high heritability estimates of loin physical traits were obtained in this study which confirms the potential for these traits to respond to selection. Indeed, other researchers have also found traits related to carcass characteristics to be moderate to highly heritable (Ciobanu et al., 2011; Van Wilk et al., 2005).

The proportions of additive variance obtained for meat color traits and total collagen in the present study indicated that genetic changes in the average population mean can be achieved through traditional breeding approaches. Further, additive genetic variance for intramuscular fat obtained in this study indicates that genetics make a substantial contribution to its total

phenotypic variance and this agrees with reports in literature (De Vries 1994; Larzul et al., 1997). The results from the genetic correlations indicated that continuous selection for lean percent will negatively impact correlated fat traits as the correlation between LEW and TFP (0.61 ± 0.16), TFW and LEP (-0.76 ± 0.10), FOF and FOP (-0.93 ± 0.05), FOP and IMF (-0.67 ± 0.18), LEP and FOF (-0.41 ± 0.15) were all moderate to highly negative.

Pale, soft and exudative condition in pork is a muscle defect known to cause considerable financial loss to the pig sector (Grandin, 2000). It is a quality defect caused by variables of both genetic and/or environmental origin. Over the course of studying this condition, a number of indicator traits have been used to predict its presence or absence in carcasses. These include L^* , pH, moisture content, cooking loss and collagen characteristics. This current study obtained a strong genetic relationship among these traits indicating a likely presence of linkage disequilibrium and/or pleiotropic gene effect existing among the traits related to the PSE defect. Further, the strong genetic correlation of L^* with SOLC and ISOLC indicates a potential for using it as an indicator trait for collagen characteristics.

One important specific hypothesis tested in this study was the presence of a genetic relationship between WBSF and collagen characteristics. Indeed, a moderate genetic correlation was found for WBSF with SOLC (-0.38 ± 0.10) and ISOLC (0.42 ± 0.16) indicating that similar gene networks to some degree, influence tenderness and IMCC.

The second study used a number of statistical modelling approaches to assess the contribution of company of origin, sex and certain intrinsic pork characteristics to measures of loin physical, meat and collagen characteristics. Estimation of the differences among group means in the population indicated that there was no significant effect of sex on loin physical, meat and

collagen characteristics while company significantly influenced loin physical, collagen characteristics, pH, drip loss and cooking loss but was not significant for total collagen, moisture content, WBSF and Minolta color traits.

Results from regression analysis indicated that pH predicted significant variations in L^* and moisture content which agrees with results from other studies (Boler et al., 2010; Dransfield et al., 1995). However, pH was also revealed as an unreliable sole predictor of quality characteristics and that other intrinsic variables will need to be included in order to accurately predict quality. Prediction equation for ISOLC retained IMF indicating that deposition of fat in the perimysium may influence crosslinking ability of collagen fibers.

Research studies (Estevez et al., 2011; Traore et al., 2012; Rowe et al., 2004) have found a relationship among water holding capacity, protein oxidation and tenderness after freeze-thawing. Based on these studies, a hypothesis that drip loss predicts variations in tenderness was formulated. The results indeed showed that drip loss predicted the most variation in tenderness after frozen storage indicating that the samples may have undergone myofibrillar protein oxidation during storage.

Findings from the stratification analysis indicated that roasts with lower IMF content also tend to be tougher with higher lean and cooking loss percent. Further, muscle pH stratification showed that high pH animals tend to also have higher backfat but lower drip loss, cooking loss and L^* while stratification based on L^* revealed that roast with lower L^* values also tend to have lower IMF and total collagen but higher muscling. The findings from this exercise shed light on the potential of these traits for predicting quality.

4.2 Future work and study limitations

In this research, tenderness was shown to be genetically correlated with SOLC and ISOLC. However, as these collagen traits were estimated to be lowly heritable and difficult and expensive to measure, they may be good candidates for prediction using approaches that predicts animals' genetic merit through DNA markers. Further, studies may need to be undertaken to identify candidate functional genes associated with WBSF and SOLC and ISOLC. This research found moderate to high heritability estimates for traits related to fat content and a strong negative genetic relationship between these traits and those related to muscling. However, their fatty acid profiles were not examined. It would have been beneficial to have examined these traits. Also, it would have been revealing to examine the glycolytic potential and the muscle fiber type characteristics of the population and their relationships with muscling and meat quality traits. In addition, an examination of the expression levels of matrix metalloproteinases at different growing phases of pigs and their influence on collagen turn over may be warranted.

In this thesis, textural properties (tenderness) of the samples were examined through WBSF tests only. However, as the evaluation of structural and textural properties of meat is vital in quality assessment (Ávila et al., 2014), an approach which will reveal more information about the samples may be necessary. Thus, texture profile analysis is recommended for use in future studies as data on hardness, fracturability, cohesiveness, adhesiveness, gumminess and chewiness (Novaković and Tomašević, 2017) will also then be available for interpretation.

Another area of interest that could be explored is genetic parameters for collagen characteristics in different pig breeds especially purebreds as well as estimates from comparatively tougher but important primals like the ham. Indeed, collagen characteristics have been shown to vary with

carcass part at the phenotypic level (Jeremiah and Martin, 1981) with the least exercised muscles tending to have higher ratio of soluble to insoluble collagen. It will be interesting to assess if this is just a function of environmental factors or genetics play a role. In addition, future studies should also be designed to make it feasible for data on a greater number of samples than used in this thesis to be obtained as this might reduce some of the high standard error estimates associated with a number of genetic correlations in the second study (Chapter 2).

Normally age of an animal is taken into consideration when determining how collagen affects meat quality, as age is known to influence collagen characteristics (Hill 1966). Animal age was not considered in this thesis because the pigs were very similar in age being between 5.5 and 6 months of age, but size or muscle fiber type should perhaps be considered in future studies. Muscle fiber composition not only affects muscle color but can affect the content of collagen (Kovanen et al. 1984). Relationships between collagen solubility and L^* value in this thesis appeared to indicate that as L^* value increased so too did collagen solubility. If this is the case, then glycolytic fibers, which are often associated with increased lightness of muscle color (Listrat et al., 2016), may have more soluble collagen than oxidative fibers. This warrants further investigation to test this hypothesis, as it may have implications for meat quality if pigs are slaughtered at ages greater than 6 months and at slaughter weights greater than 135 kg.

The second study (Chapter 3) found a significant relationship between drip loss and WBSF. Future studies may therefore need to examine the level of protein oxidation pre- and post-freezing and how it affects texture profile in a Canadian context.

4.3 References

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