



## Exploring the unknowns involved in the transformation of muscle to meat

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### ABSTRACT

Meat quality development, or the transformation of muscle to meat, involves a myriad of biochemical pathways that are largely well-studied in living muscle tissue. However, these pathways are less predictable when homeostatic ranges are violated. In addition, there is far less known about how various management or environmental stimuli impact these pathways, either by substrate load or altered cellular environment. Likewise, it is largely accepted that oxygen plays little to no role in the conversion of muscle to meat, as anaerobic metabolism predominates in the muscle tissue. Even so, the oxygen tension within the tissues does not fall precipitously at exsanguination. Therefore, transition to an anaerobic environment may impact energy metabolism postmortem. Antemortem handling, on the other hand, clearly impacts meat quality development, yet the exact mechanisms remain a mystery. In this paper, we will attempt to review those factors known to affect postmortem energy metabolism in muscle and explore those areas where additional work may be fruitful.

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### 1. Introduction

Postmortem metabolism is a heavily researched topic. Many scientific and technological advances have led to improved animal welfare, feeding strategies, slaughter processes, and resulted in improved meat quality development for the consumer. Despite a myriad of advances, questions remain regarding those mechanisms controlling or impacting postmortem metabolism and how physiological and tissue-based homeostatic set points are maintained or breached by various management practices that ultimately lead to altered meat quality development. Though the rate of postmortem metabolism is quite important in driving meat quality development, it is fairly well established. Alternatively, the biochemical mechanism(s) responsible for the cessation of postmortem metabolism, or protracted carbohydrate metabolism are particularly puzzling. The role of mitochondria and curtailed oxidative metabolism play in modulating postmortem metabolism will also be explored. Finally, we will briefly review antemortem animal handling practices in an effort to understand how these management practices alter the aforementioned.

### 2. Cessation of postmortem metabolism

In order to understand those mechanisms that may control an abbreviated or protracted postmortem metabolism in muscle, one must

first reason why it stops. To date, this has not been unequivocally established. Though some would argue it is simply a function of glycogen abundance at harvest, this is not the case, especially when extreme deviates are removed from the population (Copenhaver, Richert, Schinckel, Grant, & Gerrard, 2006; Scheffler & Gerrard, 2007; Scheffler, Park, & Gerrard, 2011). Over sixty years have elapsed since meat scientists across the globe have known that some muscle, for whatever reason, is capable of breaching a final pH, which is otherwise relatively constant across myriad of animals managed and processed under a variety of conditions, yet little progress or even interest exists in this area. Essentially, there are two viable hypotheses, either there is a pH-mediated inactivation of glycolytic enzymes, which stops hydrogen accumulation at a constant endpoint, or there is loss of adenosine nucleotides preventing a glycolytic substrate to rephosphorylate (Dalrymple & Hamm, 1975; Greaser, 2001). In an attempt to stimulate or re-kindle an interest in this fascinating biochemical process, we will begin by reviewing the collective works of one of the great pioneers of postmortem metabolism, Robert K. Scopes.

Scopes and Lawrie (1963) first entered the area of postmortem metabolism by noting that an accelerated rate of postmortem metabolism resulted in denatured sarcoplasmic proteins and adulterated meat quality development. They predicted at that time that the antemortem 'state' of the animal likely dictated the pH decline in these muscle tissues (Scopes & Lawrie, 1963) and quickly extended these initial observations to show that differences in muscle temperature and pH combinations indeed altered the extractability, or solubility of sarcoplasmic, and myofibrillar muscle proteins (Scopes, 1964). These early data likely formed the foundation for a number of subsequent studies over the next 50 years based on the premise that enzymes, and other proteins, denature with time postmortem (Joo, Kauffman,

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Kim, & Park, 1999; Warner, Kauffman, & Greaser, 1997), and represent events now commonly known as the conversion of muscle to meat. Most likely because of animal to animal variation, Scopes then modified his approach and used minced muscle to study postmortem metabolism. Using this approach, Newbold and Scopes (1971) showed that varying the concentrations of inorganic phosphate ( $P_i$ ) up to 50 mM reduced the ultimate pH ( $pH_u$ ) of the mince, yet concentrations greater than 50 mM did not further facilitate greater declines in the metabolizing tissue preparation. Based on these observations, they proposed that  $P_i$  may induce greater glycogen phosphorylase (GP) activity, thereby explaining the lower  $pH_u$  observed in some muscles. This hypothesis is particularly intriguing given the intimate relationship between the phosphagen system and energy metabolism in exercising muscle (Robergs, Ghiasvand, & Parker, 2004). Recall, GP is present in muscle in two forms, *a* and *b*, the former being more active than the latter (Berg, Tymoczko, & Stryer, 2007). GP is activated by AMP and inhibited by ATP and glucose 6-phosphate. Both calcium and epinephrine are capable of shifting the inactive form *b* to the active *a* form by stimulating phosphorylase kinase. Of particular significance is that though active, GP cannot generate glucose 1-phosphate without inorganic phosphate (Morgan & Parmeggiani, 1964). Given that added inorganic phosphate to a muscle mince caused a lower  $pH_u$  in glycolysing muscle preparations (Newbold & Scopes, 1971), it is possible to argue that inorganic phosphate may be rate-limiting during particular times postmortem metabolism. Addition of inorganic phosphate, on the other hand, would undoubtedly raise the pH of the mince early and ultimately could result in a higher pH of the reaction. After all, liberation of free phosphate via ATP hydrolysis, especially in heavily exercising muscle, is known to buffer muscle cells against massive hydrogen accumulation (Robergs, 2001). Regardless, defining the role of phosphate, especially free phosphate in postmortem muscle, may be quite enlightening.

Scopes further refined his *in vitro* system to include a glycolysing mixture (glycogen, ATP, NAD,  $P_i$ , etc.), organic buffers (TRIS, acetate), and purified glycolytic enzymes (Scopes, 1973). The composition of the buffer is particularly germane to the issue at hand, as buffer capacity of muscle can dramatically impact the  $pH_u$  and quality of meat (Kylä-Puhju, Ruusunen, Kivikari, & Puolanne, 2004; van Laack, Kauffman, & Greaser, 2001). Using this system, Scopes (1973) documented the extent resting muscle could rephosphorylate creatine dependent upon available inorganic phosphate, GP *a* concentration, and ATPase activity. He then showed that the rate of glycolysis is directly proportional to the amount of ATP consumed (Scopes, 1974a). Specifically, when ATPase activity was stimulated, lactate formation was increased proportionally. In addition, he noted that glycolysis stopped once adenonucleotides were metabolized. In contrast, when ATPase, or ATP consumption, was reduced or minimized, the entire system was capable of maintaining ATP concentrations in a steady-state condition, where minimal AMP is detected. As a result, a slower metabolism ensued. These results are particularly interesting as they argue that energy levels such as: phosphocreatine, ATP or the ability of ATP to be rephosphorylated (see discussion below) in the muscle tissue at harvest may shift the time at which glycolysis may begin, or even reach maximal levels. Changes in the time at which these events occur postmortem could have dramatic effects on ultimate meat quality development, as protein denaturation again is a pH-temperature phenomenon (Offer, 1991; Wismer-Pedersen, 1959). Furthermore, these data show theoretically, that removal of adenonucleotides from glycolysing muscle will arrest metabolism raising another point of control that will be discussed briefly below.

His final and arguably the most important data using this *in vitro* system, were those directly targeted at understanding the enzymes responsible for pacing both early postmortem metabolism and that responsible for extending carbohydrate metabolism in skeletal muscle (Scopes, 1974b). These data showed that regardless of enzymes present, ATPase concentration, or ATP consumption, drives the rate of metabolism. These findings formed the basis by which many understand

the role ATPase plays in controlling the rate of postmortem metabolism (Bowker, Grant, Swartz, & Gerrard, 2004; Hamm, 1977), especially where inherent differences in ATP consumption found between muscles of different fiber types change postmortem metabolism (Fernandez & Tornberg, 1991; Klont, Brocks, & Eikelenboom, 1998). Moreover, early postmortem consumption of ATP in muscle is hallmark of halothane-positive pigs containing a mutated calcium channel protein that allows cellular calcium concentrations to rise to a point where corresponding downstream ATPases force an aggressive metabolism and aberrant meat quality development (Cheah, Cheah, Crosland, Casey, & Webb, 1984; Greaser, Cassens, Briskey, & Hoekstra, 1969; Monin, Sellier, Ollivier, Gouteponge, & Girard, 1981). Results of these studies also raised the idea that GP *a*, and to a lesser extent AMP deaminase concentrations may dictate the pH at which metabolism stops (Scopes, 1974b).

Understanding how Scopes settled on GP *a* as a driver of  $pH_u$  in porcine skeletal muscle is logical though difficult to explain in living, or dying muscle. In his studies (Scopes, 1974b), GP *a* concentrations were included at sub- and supra-physiological levels. While the increased levels of GP *a* resulted in a reduced  $pH_u$  *in vitro*, it also accelerated the rate of pH decline, begging the question of whether phosphorylase differs between pigs with altered muscle metabolism. GP *a* activity in resting halothane positive pig muscle is lower compared to wild-type pigs (Fernandez, Neyraud, Astruc, & Sante, 2002), yet increases above that of wild-type animals during postmortem metabolism (Monin, Talmant, Laborde, Zabari, & Sellier, 1986). This increase in phosphorylase *a* content is likely due to the calcium-mediated activation of phosphorylase kinase, the enzyme responsible for the conversion of GP *b* to *a* form (Meyer, Fischer, & Krebs, 1964). Alternatively, it could be due to increased catecholamine release and action at harvest (Althen, Ono, & Topel, 1977). However, classically halothane pigs do not differ in  $pH_u$  (Copenhafer et al., 2006; De Smet et al., 1996; Fernandez et al., 2002; Klont, Lambooy, & van Logtestijn, 1993; Kocwin-Podsiadla, Przybylski, Kuryl, Talmant, & Monin, 1995) which argues against GP *a* content driving the  $pH_u$  of meat. Even so, a number of investigators have shown that meat of halothane positive pigs results in a lower  $pH_u$  (Fisher, Mellett, & Hoffman, 2000; Hamilton, Ellis, Miller, McKeith, & Parrett, 2000; Klont & Lambooy, 1995; Klont, Lambooy, & van Logtestijn, 1994; Monin et al., 1981). The latter issue makes it difficult to study cessation of postmortem metabolism in halothane-sensitive pigs. In comparison, no differences have been noted in GP *a* activity between wild-type and Rendement Napole (RN) pigs (Estrade, Ayoub, Talmant, & Monin, 1994). Recall, RN pigs possess a gene mutation that somehow allows for a breach in the normal postmortem set points and results in lower  $pH_u$  and a type of 'acid meat'. Regardless, the aforementioned data strongly support the notion that GP may be involved in controlling the extent of postmortem metabolism and should be closer scrutinized.

Though not specifically addressed by Scopes' work, the enzyme most frequently implicated as responsible for the cessation of postmortem metabolism is phosphofructokinase (PFK) (Bendall, 1973; Hamm, 1977), due in part, to its rate-limiting status in glycolysis and its complex control in living tissues (Berg et al., 2007). Moreover, glucose 6-phosphate increases late postmortem (Copenhafer et al., 2006; Kastenschmidt, Hoekstra, & Briskey, 1968) suggesting a loss in PFK activity sometime earlier. Data to directly support this hypothesis are scant. However, a recent study comparing normal and RN pig muscle showed a number of sarcoplasmic and myofibrillar proteins experience phosphorylation events postmortem (Lametsch et al., 2011). To that end, the increased phosphorylation of PFK in RN pig muscle may increase its stability and subsequent pH inhibition during the postmortem period (Sola-Penna, Da Silva, Coelho, Marinho-Carvalho, & Zancan, 2010). This may help explain the lower  $pH_u$  of fresh pork derived from this genotype.

Schwäglele and Honikel (1988) quantified a host of glycolytic enzyme activities over a wide range of pH values (5.3 to 6.8) found in postmortem tissue. Results showed the activities of PFK, glyceraldehyde 3-phosphate dehydrogenase, phosphoglycerate kinase, pyruvate kinase,

and adenylate kinase varied with pH conditions studied; however no enzyme lost activity completely. Enzymes were extracted in a phosphate buffer (pH 7.0) suggesting assays were conducted at a pH close to the physiological norm, rather than those conditions found postmortem. If this was indeed the case, then the activity of these enzymes at normal postmortem conditions remains unknown, though Scopes argues that PFK remains active at pH 5.35 (Scopes, 1974b). Regardless, these findings provide some evidence that glycolytic-based enzyme inactivity may be subtle. These data do not, however, rule-out enzyme inactivation as a means of arresting postmortem metabolism.

The idea that postmortem metabolism stops in response to adenonucleotides loss is intriguing and rational. After all, the enzymes thought to be rate-limiting in glycogenolysis and glycolysis (PFK, GP, pyruvate kinase) are all allosterically activated/inactivated by ATP or its metabolites (Berg et al., 2007). Under normal circumstances, ATP is hydrolyzed to ADP, which then can be rephosphorylated to ATP by either phosphoglycerate kinase, pyruvate kinase or adenylate kinase. Adenylate kinase converts 2 ADP molecules into one ATP and one AMP molecule ( $ADP + ADP \rightarrow ATP + AMP$ ) (Pearson, 1971). Once formed, AMP is quickly converted by AMP deaminase ( $AMP \rightarrow IMP$ ) and unavailable for use by glycolysis (Pearson & Young, 1989), thus reducing the concentration of the adenonucleotide pool and eliminating a substrate for glycolysis to phosphorylate. The adenylate kinase and AMP deaminase reactions are coupled so an increase in activity of one results in an increase of the other. Though subtle differences exist between species (Fishbein, Davis, & Foellmer, 1993), AMP deaminase or adenylate kinase activity do not differ greatly, mitigating excitement for either in controlling differences in the extent of postmortem metabolism postmortem. Even so, these enzymes likely deserve some additional consideration regarding our quest to understand fully postmortem metabolism.

One means of testing whether adenonucleotides dictate the cessation of glycolysis would be to eliminate them experimentally from the tissue. A complete loss of ATP would necessarily stimulate adenylate kinase and AMP deaminase and speed up the arrest of glycolysis. The only approach readily available is electrical stimulation, which stimulates glycolysis (Hwang, Devine, & Hopkins, 2003). Theoretically, it may be possible to hydrolyze enough ATP that sufficient energy would not be available to break rigor bonds, and thereby arrest glycolysis. Then, is it possible to stimulate carcasses with electricity enough to stop postmortem metabolism? If so, would sustained electrical stimulation create a dark, firm, and dry (DFD)-like meat condition if the loss of adenosine nucleotides is solely responsible for the cessation of glycolysis and establishment of the  $pH_u$  of meat? Across numerous electrical stimulation studies reviewed, we have yet to read of any form of electrical stimulation that ever resulted in higher  $pH_u$  values or DFD-like meat, at least in pigs (Hallund & Bendall, 1965; Hammelman et al., 2003; Maribo, Ertbjerg, Andersson, Barton-Gade, & Møller, 1999; Taylor, Nute, & Warkup, 1995). In fact, electrical stimulation is a reliable tool to produce pale, soft, and exudative (PSE) meat in pigs (Bowker, Grant, Forrest, & Gerrard, 2000). Taken together, these data indicate that despite a rapid loss of ATP caused by electric stimulation, sometimes repeated, enough nucleotides remain in the muscle to facilitate glycolysis to a normal  $pH_u$ . Even so, until we find a way to eliminate ATP experimentally, we cannot rule out this hypothesis.

### 3. Mitochondria and meat quality

At harvest, muscle must adapt to new physiological circumstances in order to maintain ATP and homeostasis. Delivery of oxygen to muscle is eliminated, which impedes mitochondria from contributing to ATP production. Thus, the conversion of muscle to meat is traditionally viewed as an anaerobic process that is largely governed by degradation of glycogen to lactate and  $H^+$ . While fundamentally true, this has propagated a dubious conclusion that because mitochondria can no longer generate ATP, they are irrelevant to pH decline and postmortem

metabolism. In fact, mitochondria are functionally complex and operate quite differently as the cellular milieu changes.

Although typically thought of as individual organelles, mitochondria are dynamic and plastic. Fusion and fission events allow mixing of mitochondrial contents and also contribute to a range of mitochondrial morphology, from small, round-oval organelles to a branched tubular network (reviewed by Chan, 2006). Structure of mitochondria is well-suited to their function. Mitochondria consist of an outer membrane and a highly folded inner membrane, resulting in two aqueous compartments: the intermembrane space and the matrix. The outer membrane is permeable to most small molecules and ions, while the inner membrane is completely impermeable to most small molecules. The inner membrane also contains the proteins of the electron transport chain, and the transfer of electrons through these protein complexes leads to protons being pumped out of the mitochondrial matrix into the intermembrane space. This results in an uneven distribution of charge or mitochondrial membrane potential ( $\Psi_{mt}$ ), with the intermembrane space being more positive. Under normal physiological circumstances, the flow of protons through  $F_1F_0$  ATP synthase (complex V) back to the mitochondrial matrix powers ATP synthesis. The rate of oxidative phosphorylation is primarily determined by the need for ATP. Thus, high ADP concentrations stimulate maximal oxygen consumption or respiration, referred to as state 3 respiration (Chance & Williams, 1955). Additionally, greater workloads contribute to increases in intracellular calcium; in turn,  $Ca^{2+}$  may be taken up by mitochondria, primarily via the  $Ca^{2+}$  uniporter in the inner membrane. Subsequently, increases in matrix  $Ca^{2+}$  enhance activity of enzymes such as pyruvate dehydrogenase, and this helps generate  $\Psi_{mt}$  (reviewed by McCormack, Halestrap, & Denton, 1990). Not all protons flow back through ATP synthase; proton leak across the inner membrane represents inefficiency because oxygen consumption is not coupled to ATP synthesis. Experimentally, an ATP synthase inhibitor can be used to eliminate the contribution of proton flow through the ATP synthase to determine basal or state 4 respiration. The ratio of oxygen consumption rate in state 3 relative to state 4 (respiratory control ratio, RCR) indicates the tightness of coupling between respiration and phosphorylation, and also serves as an index of mitochondrial integrity (reviewed by Brand & Nicholls, 2011).

While limited oxygen dictates that mitochondria have limited ability to generate ATP postmortem, they appear to remain functionally competent for some time. In fact, intact mitochondria could be isolated from ox neck muscle at 144 h postmortem (Cheah & Cheah, 1971). Furthermore, mitochondria isolated from dark-cutting beef are reasonably well-coupled at 5 d postmortem and perform better than mitochondria obtained from control animals at 24 h postmortem (Ashmore, Parker, & Doerr, 1972). Moreover, there are clear improvements in functional parameters of mitochondria when the pH of isolation and assay medium is increased to physiological pH (Ashmore et al., 1972; Tang et al., 2005). Thus, the main factor affecting function of mitochondria isolated in the postmortem period is development of low muscle pH. Similarly, mitochondria isolated from “normal” muscle immediately postmortem resemble those found in intact muscle, whereas some mitochondria from PSE muscle are already swollen and show decreased matrix density (Greaser et al., 1969). In permeabilized fibers from halothane positive muscle, mitochondrial state 3 respiration declines to a greater extent from 0 to 12 h postmortem compared to control (Werner, Natter, Schellander, & Wicke, 2010). Together, this suggests that acidic cellular environment and rapid pH decline are major factors influencing mitochondrial morphology and function postmortem.

Although mitochondria are still capable of oxidative phosphorylation postmortem, the oxygen deficient environment limits respiration. So, how do mitochondria contribute to postmortem metabolism and pH decline? It is important to consider that, at harvest, muscle is living and all metabolic systems are functionally competent; the subsequent struggle to “survive” or maintain homeostasis may parallel that of living

muscle exposed to anoxic or ischemic conditions. When oxygen is lacking, electron transport in mitochondria ceases, and responsibility for ATP production shifts to anaerobic glycolysis. The mismatch between ATP production and utilization creates several challenges for cellular homeostasis. First, cellular pH decreases, and this “deenergizes” or gradually decreases the  $\Psi_{mt}$ . Concurrently, decreased efficiency of sarcoplasmic reticulum  $Ca^{2+}$ -ATPase contributes to intracellular  $Ca^{2+}$  overload. In turn, mitochondria accumulate substantial amounts of  $Ca^{2+}$ , and sustained increases in matrix  $Ca^{2+}$  contribute to permeability transition whereby the inner membrane becomes permeable to solutes (Bernardi et al., 2006). This, in turn, can trigger mitochondrial matrix swelling and collapse of  $\Psi_{mt}$ . Under these extreme circumstances, the mitochondrial ATP synthase may actually run in reverse, hydrolyzing glycolytically-produced ATP in order to maintain a membrane potential (Scott & Nicholls, 1980). Net consumption of ATP by mitochondria further compromises cellular energy balance and viability.

As Hudson (2012) has recently suggested, this “cellular treason” by mitochondria may be applicable to postmortem muscle and meat quality development. ATP hydrolysis by the  $F_1F_0$  ATPase would directly influence cellular ATP consumption, and could further promote glycolysis and pH decline postmortem. In turn, rate of ATP consumption would be expected to be positively correlated with mitochondrial content. Therefore, as Hudson (2012) proposed,  $F_1F_0$  ATPase driven ATP hydrolysis could partly explain the paradoxical observation that red, oxidative muscles sometimes exhibit faster pH decline rates than whiter, more glycolytic muscles.

While pH decline and ATP consumption may be related directly to mitochondrial content, the relationship is likely much more convoluted. Specifically, the properties of mitochondria may vary not only by fiber type, but by their location within the muscle cell. Conventionally, energetic differences between red and white muscles are attributed primarily to greater mitochondrial density of slow fibers (Schwartzmann, Hoppeler, Kayar, & Weibel, 1989). Yet, others recently suggested that mitochondria exhibit a certain phenotype based on fiber type (Picard, Hepple, & Burelle, 2012). Mitochondrial phenotype encompasses not only ATP producing capacity, but also reactive oxygen species production, and sensitivity to  $Ca^{2+}$  and apoptosis. The ability of mitochondria to buffer  $Ca^{2+}$  influences the amplitude and frequency of  $Ca^{2+}$  oscillation within the cytosol. The nature of cytosolic  $Ca^{2+}$  oscillations varies by fiber type. Thus,  $Ca^{2+}$  handling properties of mitochondria and their tolerance for aberrant  $Ca^{2+}$  levels may differ according to fiber type. With net ATP consumption postmortem and reduced sarcoplasmic reticulum  $Ca^{2+}$ -ATPase function, mitochondria may play a significant role in accumulating  $Ca^{2+}$  and restraining glycolysis. Subcellular localization of mitochondria in muscle cells may also impact function and response to cellular stress. Muscle mitochondria are distributed primarily either beneath the cell membrane (subsarcolemmal) or between the myofibrils (intermyofibrillar) (Müller, 1976). Although slow and fast fibers have a similar volume of intermyofibrillar mitochondria, slow fibers have a much higher volume of subsarcolemmal mitochondria (Philippi & Sillau, 1994). Subsarcolemmal and intermyofibrillar mitochondrial likely form different networks that are not connected, which contributes to distinct functional and biochemical properties.

Another point of consideration, however, is that there are inhibitory mechanisms that prevent the ATPase from wasting ATP during cellular stress. For example, certain species, such as frogs, are exposed to extended periods of anoxia which would be expected to threaten their livelihood. However, in this situation, ATP hydrolysis is greatly reduced via inhibition of the  $F_1F_0$  ATPase (St-Pierre, Brand, & Boutilier, 2000). In turn, this restricts cellular ATP depletion during ischemia, and delays cell damage and death. Moreover, different species possess different capacities for ATPase inhibition. This is based on content and affinity of  $F_1$ -ATPase inhibitory subunit (IF<sub>1</sub>). The IF<sub>1</sub> subunit binds to the ATPase under nonenergizing conditions, when the pH of the mitochondrial matrix is low (Rouslin, Frank, & Broge, 1995). During anoxia, species possessing greater content of high affinity

IF<sub>1</sub> demonstrate marked inhibition of ATPase, whereas species with low amounts of high affinity IF<sub>1</sub> exhibit little inhibition of ATPase. Additionally, IF<sub>1</sub> content varies by tissue, and importance of IF<sub>1</sub> may depend on conditions (i.e. presence of uncoupler) (Rouslin & Broge, 1996; Rouslin et al., 1995).

Despite their complex role in cellular function and metabolism, the role of mitochondria in postmortem muscle has remained largely unexplored. While the loss of oxygen impedes “normal” mitochondrial function, it does not preclude mitochondria from contributing to meat quality development. Mitochondrial content, as well as functional differences between distinct mitochondrial populations may influence the course of postmortem metabolism. Furthermore, mitochondrial properties may be intrinsically beneficial or detrimental to the course of postmortem metabolism and pH decline. For example, pro-survival or anti-apoptotic proteins may associate with mitochondrial membrane proteins and modify response to energetic depression. Understanding these aspects of mitochondria in postmortem muscle may improve our ability to predict pH decline and meat quality.

#### 4. Antemortem handling

Feed withdrawal or fasting prior to slaughter is commonly used to improve meat quality by elevating  $pH_u$  through a reduction in liver (Warriss, 1982) and muscle glycogen (Fernandez & Tornberg, 1991; Sterten, Oksbjerg, Frøystein, Ekker, & Kjos, 2010; Wittmann, Ecolan, Levasseur, & Fernandez, 1994). A Bayesian meta-analysis using 16 studies showed fasting duration increases the  $pH_u$  of the *longissimus* muscle of pigs (Salmi et al., 2012) yet, the correlation is weak likely due to the unique relationship between glycogen and the  $pH_u$  of meat. In fact, this relationship is curvilinear where the muscle glycogen content is primarily responsible for dictating  $pH_u$  between a  $pH_u$  of 7.2 and 5.7 (Bendall, 1973; Henckel, Karlsson, Jensen, Oksbjerg, & Petersen, 2002), but below 5.7, other as yet unknown mechanisms must be responsible for the variation in this range. Some of the possible mechanisms have already been discussed in earlier sections, yet others likely remain.

During normal postmortem metabolism, the *longissimus* muscle (white, glycolytic muscle) utilizes 35–40  $\mu\text{mol}$  glycogen/g tissue to achieve a  $pH_u$  near 5.5 (Copenhafer et al., 2006). Oftentimes, this muscle contains greater than necessary amounts of glycogen (60  $\mu\text{mol}/\text{g}$ ) resulting in 2–30% of the total glycogen remaining at 24 h (Pearson & Young, 1989). This residual glycogen content may help explain the weak relationship between fasting and  $pH_u$ . Fasting reduces glycogen content of muscle by as much as 20% in the *longissimus* muscle of pigs over a 24 h period (Wittmann et al., 1994), yet this reduction is insufficient to reduce glycogen below the aforementioned threshold that affects  $pH_u$ . Henckel et al. (2002) suggests that increases in  $pH_u$  will not develop unless muscle glycogen is reduced below 53  $\mu\text{mol}/\text{g}$  tissue supporting this argument. Furthermore, red, oxidative muscles contain less glycogen and produce meat with higher  $pH_u$  (Monin & Talmant, 1987). Curiously, a 24 h fast reduces the glycogen content of the more oxidative *semispinalis* muscle by 50% and results in a higher  $pH_u$  (Wittmann et al., 1994). Thus, the percentage decrease in glycogen between the two muscles in response to a 24 h fast is quite different, but the absolute decrease of each is quite similar (~15  $\mu\text{mol}/\text{g}$ ) (Wittmann et al., 1994). Therefore, fasting may reduce the stored carbohydrate content of muscle below this ‘critical’ threshold and thus, fasting may be more effective on red muscles due to their lower resting glycogen content. This is especially apparent with RN pigs. Again, RN pig muscle contains extremely high levels of glycogen and produce meat with low  $pH_u$ . When fasted for 36 h, the  $pH_u$  of meat from RN mutant pigs still remained lower than controls (Bidner, Ellis, Witte, Carr, & McKeith, 2004), arguing that a 36 h fast is not sufficient to lower glycogen levels below the critical threshold where it can impact  $pH_u$  and recover a normal phenotype. Regardless of whether fasting impacts postmortem metabolism and meat quality characteristics, the practice of fasting should not be eliminated from the standard slaughter

procedure because of the microbiological ramifications mitigated by reducing gastrointestinal tract content (Miller, Carr, Bawcom, Ramsey, & Thompson, 1997).

Of particular significance to the issue at hand is the idea that different sources of dietary energy impact antemortem muscle glycogen concentrations (reviewed by Andersen, Oksbjerg, Young, & Therkildsen, 2005). Most intriguing is the idea that easily digestible carbohydrates may alter postmortem metabolism to prevent DFD meat. Muscle of pigs fed a diet of sucrose resulted in an accelerated pH decline, lower  $pH_u$  (Briskey, Bray, Hoekstra, Phillips, & Grummer, 1959) and elevated glycogen (Briskey, Bray, Hoekstra, Phillips, & Grummer, 1960). To explain the accelerated metabolism, a follow-up study compared the GP activity between control fed and sucrose pigs, but found no difference between the two treatments were detected (Sayre, Briskey, & Hoekstra, 1963). High-sugar diets encourage the incorporation of carbohydrate into the muscle as glycogen and alters glucose metabolism (reviewed by Fernandez & Tornberg, 1991) which may alter the enzymatic activity within the muscle. Though GP activity was not altered in sucrose-fed pigs (Sayre et al., 1963), other enzymes may have been altered to facilitate an accelerated metabolism. Alternatively, a lower  $pH_u$  in meat may have been directly a result of increased glycogen deposition. However, very high inclusion rates may be necessary to achieve this endpoint and this alone may have detrimental effects on tissue energy metabolism as up to 15% sucrose incorporation to grower and finishing pig diets has no effect on glycogen content or postmortem metabolism (Camp, Southern, & Bidner, 2003).

Regardless of the mechanism, diet can be used to mitigate low pH values by reducing carbohydrates from the diet to less than 5% and increasing fat to approximately 18% on a weight basis (Rosenvold et al., 2001). When fed three weeks prior to slaughter, these diets reduce the glycogen content of the *longissimus dorsi* by 11–26% (Rosenvold et al., 2001) and increase  $pH_{45}$  and  $pH_u$  (Rosenvold et al., 2002; Tikk, Tikk, Karlsson, & Andersen, 2006) and accelerate temperature decline (Tikk, Lindahl, Karlsson, & Andersen, 2008; Tikk et al., 2006). Presumably, this feeding strategy does not alter key glycolytic enzymes like PFK, GP *a*, or GP *b* activity, but may lower  $\mu$ -calpain activity (Rosenvold et al., 2002), the enzyme responsible for postmortem proteolysis of myofibrillar proteins (Huff-Lonergan & Lonergan, 2007).

Strategic feeding protocols and fasting are intriguing because both appear to reduce glycogen content, which are thought to improve meat quality, yet there is no indication of an interaction between the two. To our knowledge, only one study has been conducted to address this issue (Partanen, Honkavaara, & Ruusunen, 2007). In this study, the longer fasting time and a fibrous diet increased  $pH_u$  and lowered muscle glycogen, and indicated that the combination of the two may produce an additive effect to elevate  $pH_u$ .

Pre-slaughter stress is complicated, but clearly reduces meat quality through altered metabolism. If an animal is stressed long-term, glycogen stores may be depleted resulting in abbreviated metabolism and a high  $pH_u$  (Briskey, 1964). Adrenaline injections prior to slaughter can produce a similar result if administered appropriately (Bendall & Lawrie, 1962; Henckel, Karlsson, Oksbjerg, & S holm Petersen, 2000; Henckel et al., 2002; Rosenvold et al., 2001). Long-term stress requires a recovery period known as lairage. From a tissue standpoint, this is interpreted as sufficient time to re-establish the glycogen that was in the muscle, or eliminate entirely the stress hormones, mostly catecholamines, that may have been released in the body in response to the stress (Faucitano, 2010). For the most part, increased lairage times elevate  $pH_{45}$ ,  $pH_u$  and thereby improve meat quality measurements, especially  $L^*$  (lightness) values (Salmi et al., 2012; Zhen et al., 2013). Alternatively, short-term stress immediately prior to slaughter is known to accelerate metabolism and result in higher muscle temperatures and a lower  $pH_u$  (van der Wal, Engel, & Reimert, 1999), a condition similar to the HAL mutation. Short-term stress is difficult to remedy due to its occurrence immediately prior to slaughter during handling or movement of the animal to the abattoir. Rosenvold and Andersen

(2003) provide a much more comprehensive review about the differences between long-term and short-term stress upon postmortem metabolism.

To complicate matters further, the effects can be combined. Shen et al. (2006) induced pre-slaughter transport stress upon pigs both with and without rest. Transport resulted in decreased (glycogen + glucose + G6P) early postmortem as well as reduced muscle pH. Muscle from transported pigs, without rest, produced meat with a lower  $pH_u$  compared to control animals, exhibited an accelerated (AMP + IMP)/ATP ratio, maintained higher GP *a* activity through 4 h postmortem, and resulted in meat with a higher drip loss and a PSE condition. While muscle glycogen was indeed reduced in these animals, it was insufficient to produce DFD meat and only resulted in accelerated metabolism. Recall, muscle glycogen must be reduced below 53  $\mu\text{mol/g}$  tissue to produce DFD meat. A general lack of consensus reflects the complexity of studying the effect of transportation on ultimate energy metabolism in the muscle, especially in pigs. Even so, it argues that pre-slaughter stress is capable of producing both DFD and PSE meat, yet the exact biochemical mechanisms remain quite unclear.

## 5. Summary

When thinking about the conversion of muscle to meat, unanswered questions still remain. The mechanism responsible for the cessation of anaerobic glycolysis postmortem has eluded researchers for the past 60 years, but a pH-mediated inactivation of glycolytic enzymes or a loss of adenosine nucleotides is likely. It is also quite difficult to accurately depict postmortem metabolism without recognizing the contribution of mitochondria, despite the loss of oxygen that inhibits “normal” function. Finally, prior to slaughter, the energy charge created in the animal’s muscles dictates the rate and extent of metabolism. Fasting, diet, transport and lairage all affect postmortem metabolism and must not be overlooked when attempting to improve meat quality.

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