



Review

Effects of heat on meat proteins – Implications on structure and quality of meat products

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Abstract

Globular and fibrous proteins are compared with regard to structural behaviour on heating, where the former expands and the latter contracts. The meat protein composition and structure is briefly described. The behaviour of the different meat proteins on heating is discussed. Most of the sarcoplasmic proteins aggregate between 40 and 60 °C, but for some of them the coagulation can extend up to 90 °C. For myofibrillar proteins in solution unfolding starts at 30–32 °C, followed by protein–protein association at 36–40 °C and subsequent gelation at 45–50 °C (conc. > 0.5% by weight). At temperatures between 53 and 63 °C the collagen denaturation occurs, followed by collagen fibre shrinkage. If the collagen fibres are not stabilised by heat-resistant intermolecular bonds, it dissolves and forms gelatine on further heating. The structural changes on cooking in whole meat and comminuted meat products, and the alterations in water-holding and texture of the meat product that it leads to, are then discussed.

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Keywords: Meat proteins; Whole meat; Meat products; Cooking; Water-holding; Texture

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1. Introduction

Cooking of meat products is essential to achieve a palatable and safe product. The meat proteins, approximately 20% of a muscle's weight, represent the main constituents that make up the structure of the meat product. They undergo substantial structural changes on heating and therefore the quality of the meat product, which is mainly governed by the meat structure, also changes drastically after cooking. This review will cover the aspects of structural changes on cooking for meat proteins per se, in whole meat and comminuted meat products, and the quality alterations of the meat product that it leads to. Earlier reviews in this area are among others Hamm (1977); Offer (1984), and Asghar, Samejima, and Yasui (1985).

2. Proteins in general

A protein is built up from a long polymer chain of amino acids, a polypeptide chain. The variable side chains give each protein chain its distinctive character. There are three general categories of side chains: non-polar, polar but uncharged and charged polar (Dickerson & Geis, 1969).

The build-up of the polypeptide chain of the proteins is called the *primary structure* (Fig. 1). This polypeptide chain forms specific conformations in solutions, the so called localised *secondary structures*, i.e., α -helix, β -pleated sheet or random coil (Fig. 1). *Tertiary and quaternary structures* of proteins are the denomination of the three-dimensional structure and the association of protein entities in solution, respectively (Fig. 1).

The stabilisation of these structures of a given protein system is dependent mostly on non-covalent forces, such as hydrogen bonding, van der Waal's forces, electrostatic and hydrophobic interactions. The formation of *globular* proteins is a typical example of how hydrophobic interactions stabilises this type of tertiary structure. There is a driving force for the non-polar, hydrophobic side chains of a protein to be removed from an aqueous to a non-polar environment. Among the meat proteins myoglobin, giving meat its colour, is a typical example of a globular protein (Fig. 2(A)).

Another structural form of proteins, which is highly prevalent among meat proteins, is the *fibrous* form of proteins. Actin, myosin and collagen in meat are typical fibrous proteins. These proteins are built up from three

main structures, namely the α -helix, the antiparallel β -pleated sheet and the triple helix (the last one only collagen). Myosin is α helical, whereas collagen uses the triple helix (Fig. 2(B)). To stabilise these structures hydrogen bonding is frequent (Dickerson & Geis, 1969).

At increased temperatures the hydrophobic side chains, for entropy reasons, can more favourably stay in the aqueous environment and in the case of the compact globular proteins that lead to an expansion and partial unfolding. However, a free energy gain can be achieved by the association of two partially unfolded proteins, thereby shielding the hydrophobic side chains from the aqueous environment (Dickerson & Geis, 1969).

If the degree of association of protein entities is too large, leading to less colloidal stability of the system, the *solubility* of the proteins is lowered, and a precipitate is formed. If, however, the three-dimensional association of the proteins occurs in such a way that the attractive and the repulsive forces are so well balanced that a three-dimensional network is formed, a gel will set. This gel binds the water in the former solution, mainly by capillary forces (Hermansson, 1986). It is also solid-like in its mechanical behaviour. The gels can in turn vary from being transparent, containing a network of strands of small cross-sections to opaque gels containing much coarser aggregated structures (Hermansson, 1986). For one of the latest review on thermally-induced gelling of globular proteins, see Clark (1998).

For the fibrous proteins the large amount of hydrogen bonds and electrostatic interactions that keep the stretched molecules in register in the large building blocks, where the fibrous proteins take part, are broken on heating. This results in the molecules having a greater freedom to form any random configuration, as driven by entropy. Since the proteins are relatively stretched in the fibrous form, fibrous proteins contract on cooking in contrast to globular proteins, which expand.

3. Meat proteins, composition and structure

The muscle consists of 75% water, 20% protein, 3% fat and 2% soluble non-protein substances. Out of the latter 2%, metals and vitamins constitute 3%, non-protein nitrogen-containing substances 45%, carbohydrates 34% and inorganic compounds 18%. The proteins can be divided into three groups; myofibrillar, sarcoplasmic and connective tissue proteins. The myofibrillar pro-

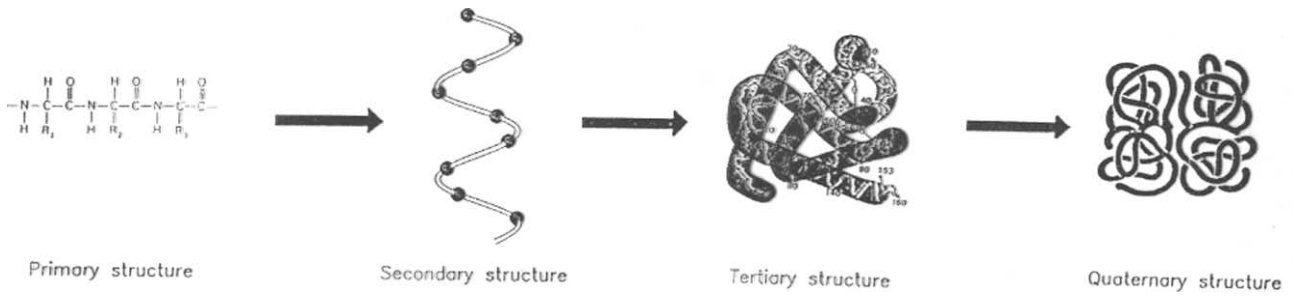


Fig. 1. The different type of structure of proteins in solution (Tornberg et al., 1990).

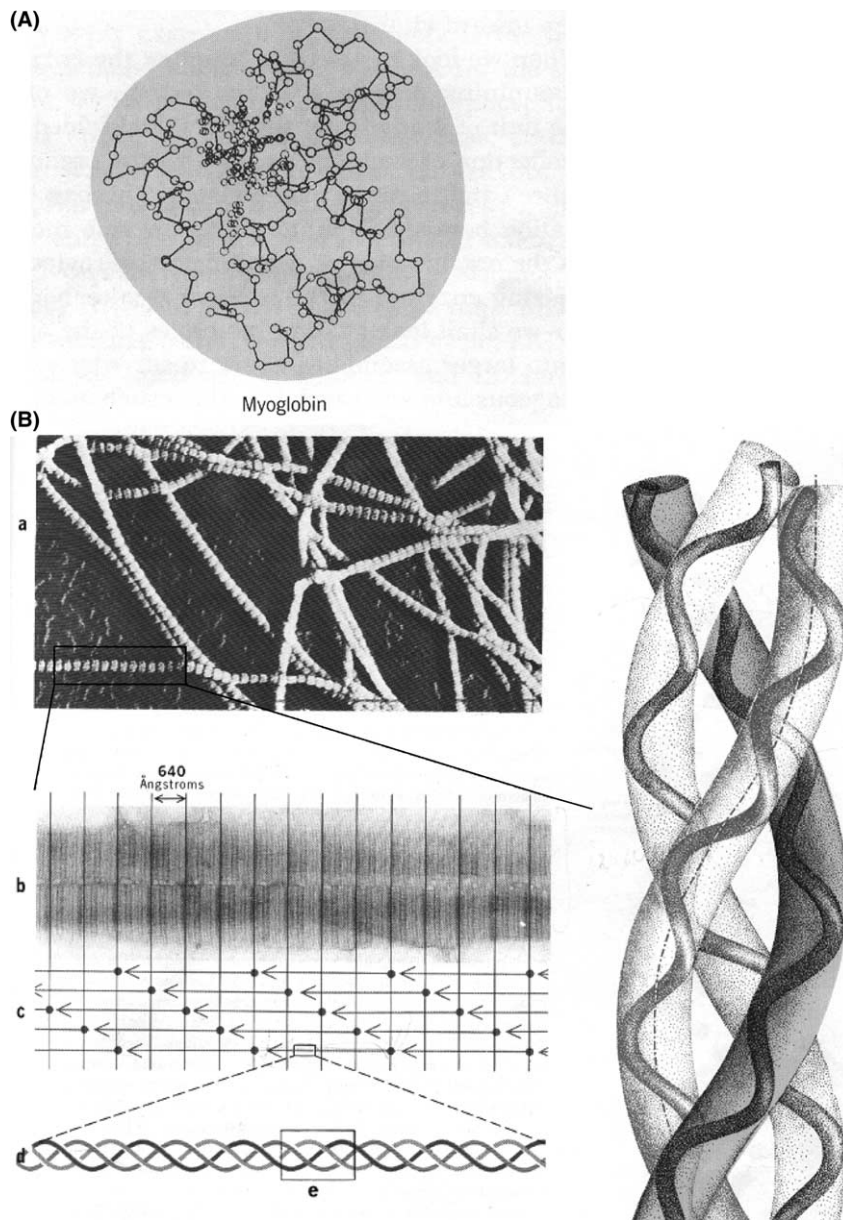


Fig. 2. Example of meat proteins being: (A) globular: myoglobin; (B) fibrous: the build-up of a collagen fibril (a) from the tropocollagen molecule (e) (Dickerson and Geis, 1969).

teins constitute between 50 and 55% of the total protein content, while the sarcoplasmic proteins account for approximately 30–34%. The remaining 10–15% of the proteins are the connective tissue proteins.

The myofibrillar proteins are further divided into three subclasses: the myofilamentous fibrous proteins myosin and actin building up the myofibrillar structure, the regulatory proteins including the tropomyosin-troponin complex, α - and β -actinin, M-protein and C-protein and ultimately the scaffold proteins, such as titin, nebulin, desmin, vimentin and synemin, supporting the whole myofibrillar structure. Titin is a massive protein with a molecular weight of around 1 million Dalton.

The sarcoplasmic proteins are the soluble proteins of the sarcoplasm, to which belong most of the enzymes of the glycolytic pathway, creatine kinase and myoglobin. About 100 different proteins are known to be present in the sarcoplasmic fraction and they are globular proteins of relatively low molecular weight ranging from 17,000 (Myoglobin) to 92,500 (Phosphorylase b).

The structures built up by the connective tissue proteins starts with an external covering sheet of connective tissue, the epimysium, around the whole muscle. This layer of connective tissue binds the individual bundles of muscle fibres into place and also binds groups of muscles together. The muscle fibre, which is the muscle cell, can vary in diameter from 10 to 100 μm and have a length of up to 30 cm. The cell has a membrane, called the sarcolemma, and is also surrounded by another type of connective tissue called the endomysium. The fibres are collected into fibre bundles, where the third type of connective tissue (perimysium) envelopes the fibre bundle (Ashgar & Pearson, 1980).

The connective tissue proteins collagen, reticulin and elastin are all fibrous proteins. Collagen, a glycoprotein, is the main structural component of the connective tissues (55–95% of the dry matter content) and is composed of tropocollagen monomers about 2800 \AA long and 14–15 \AA in diameter with a molecular weight of 300,000. The tropocollagen molecules aggregate to form either extended fibres in the epimysium and perimysium or mainly as a structural matrix in the endomysium. Collagen exists in several different genetic forms (I–V), which are present in muscle. Bailey, Restall, Sims, and Duance (1979) used immunofluorescence to show that type I is present in the epimysium, types I and III are present in the perimysium and types III, IV and V collagen are present in the endomysium.

The muscle fibres constitute 75–92% of the total muscle volume, and they hold long, thread-like structures, the myofibrils, wherein the sarcomere, the smallest contractile unit, is lined up. The structural build-up of the sarcomere can be overviewed in Fig. 3. The diameter of the myofibrils is about 1 μm and the length of a sarcomere is about 2.2 μm in a resting muscle.

According to Fig. 3 the sarcomere is built from two “building blocks”, that consist of a thick filament, extending over the A-band and a thin filament, extending from the Z-line towards the A-band in the H-zone. The thick filaments are composed of myosin. There are 200–400 molecules of myosin in each thick filament, with each being 1.5 μm long and 130 \AA in diameter (Knight & Trinick, 1987). By tryptic digestion, the myosin filament can be split into a heavy head, called heavy meromyosin (HMM), and a tail called light meromyosin (LMM). The water-insoluble LMM fraction has a molecular weight of 150,000 and is composed of either a double or triple-stranded α -helical structure. The head region consists of two globular units, each about 70 \AA in diameter with 45% α -helical content (Knight & Trinick, 1987).

The second major myofibrillar protein is actin. The fibrous actin (F-actin) is formed from longitudinal polymerization of globular actin (G-form, M_w 47,000 Da). In solution at low ionic strength actin exists in the monomeric globular form. When the ionic strength is raised, the monomers are polymerized into the fibrous structure, consisting of a double, twisted helix with a diameter of 70 \AA (Fig. 3; Ashgar & Pearson, 1980).

4. Behaviour of meat proteins on heating

Conformational changes of the proteins occurring on heating are usually called denaturation. The cooking temperature, where conformation changes occur is commonly called denaturation temperature and has been mostly investigated using differential scanning calorimetry (DSC). The unfolding of the proteins (the loss of helical structure) can also be followed by optical rotary dispersion (ORD) and circular dichroism (CD). Another way to follow the unfolding of the proteins is to measure the surface hydrophobicity of the proteins, using a fluorescent probe 8-anilino-1-naphthalene sulfonate (ANS). The next step in the structural changes to occur on heating are the protein–protein interactions, resulting in the aggregation of proteins. These processes are mainly studied by turbidity measurements and loss in protein solubility. The gel forming ability and the type of gels formed by the proteins are usually studied, using some sort of mechanical and micro-structural measurements.

4.1. Sarcoplasmic proteins

According to the review by Hamm (1977) many of the researchers have found that most sarcoplasmic proteins (i.e., those muscle proteins soluble in water or at low ionic strength) aggregate between 40 and 60 $^{\circ}\text{C}$. Davey and Gilbert (1974) found for the bull neck muscle that the

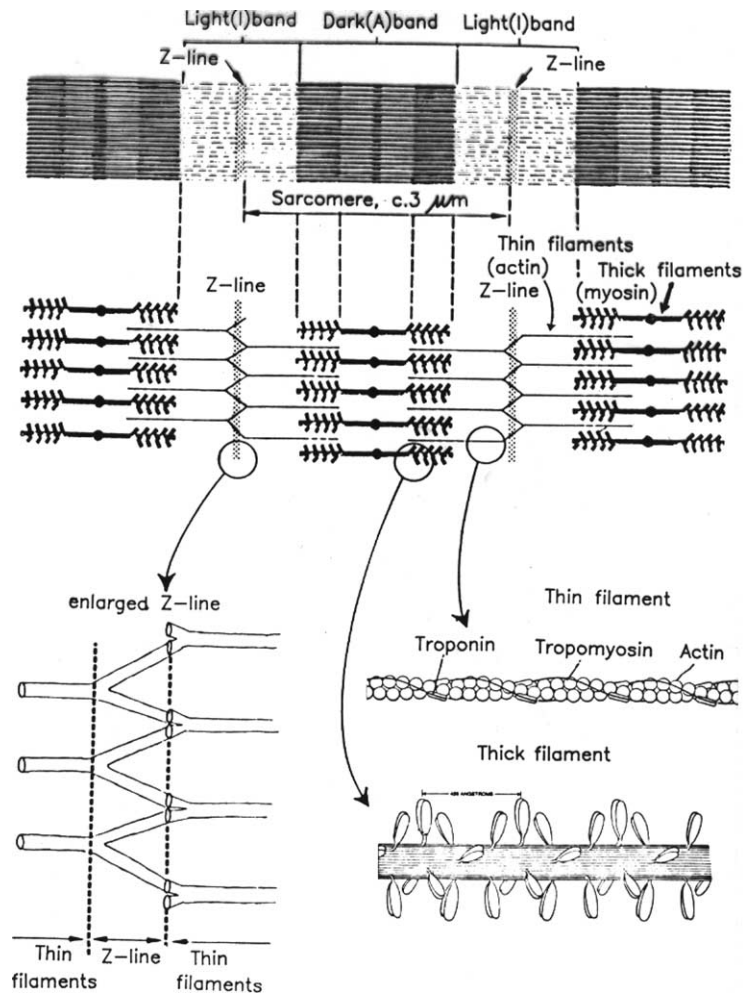


Fig. 3. The structural build-up of the sarcomere, the thin and thick filaments (Tornberg et al., 1990).

heat aggregation of the sarcoplasmic proteins could extend up to 90 °C. They were also the first to suggest that the sarcoplasmic proteins might have a role in the consistency of cooked meat in such a way that the heat-induced aggregated sarcoplasmic proteins can form a gel in between the structural meat elements and thereby link them together. In our investigations on tenderness of meat cooked to different temperatures (Tornberg, Andersson, & Josell, 1997), our measurements suggest that might well be the case. The mechanism by which this will operate will further be elaborated on later in this review.

Another interesting aspect of the sarcoplasmic proteins is the tenderizing effect some of these enzymes can have, using low temperature long time heating (heating rate of 0.1 °C/min) on beef muscles. Laakkonen, Sherbon, and Wellington (1970) have shown that collagenase could remain active in the meat at cooking temperatures < 60 °C, whereas at faster heating and reaching a higher end temperature of 70–80 °C they were inactivated. They also showed that a heating time of at least 6 h was needed to achieve a substantial lowering of the shear force, i.e., a tenderizing effect. Over

the same time most of the water losses between 25 and 30% (w/w) had occurred.

4.2. Myofibrillar proteins

4.2.1. Changes in secondary, tertiary and quaternary structure

With regard to changes in secondary structure on heating of myosin Morita and Yasui (1991) have for example measured the change in helix content (circular dichroism) and surface hydrophobicity (ANS) of LMM, light meromyosin, at pH 6 and 0.6 M KCl. The helix content of LMM began to decrease at about 30 °C and attained a minimum at 70 °C. Simultaneously, heating to 65 °C progressively increased the surface hydrophobicity, whereas at higher temperatures it decreased again. The decrease in hydrophobicity observed at the higher temperatures suggests that part of the hydrophobic residues take part in protein–protein interactions leading to a network formation of aggregates, a gel.

The advantage of the DSC method is that it can be used in complex mixtures and at high concentrations of

proteins, which is the situation occurring in meat. A typical curve (Fig. 4) from thermal transitions found in a muscle is composed of three major transition zones A, B and C. The first transition displays its maximum between 54 and 58 °C and has been attributed to myosin (Martens & Vold, 1976; Wright, Leach, & Wilding,

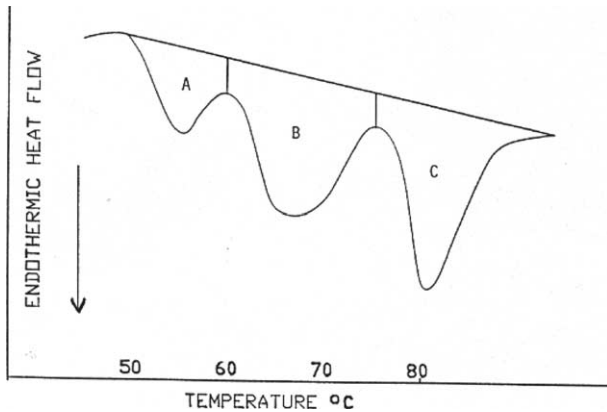


Fig. 4. A typical thermal curve of muscle is composed of three major zones: A, Myosin subunits; B, Sarcoplasmic proteins and collagen, and C, Actin (Findlay et al., 1989).

1977). The second transition, which occurs between 65 and 67 °C, was assigned to collagen (Martens & Vold, 1976; Stabursvik & Martens, 1980) and to sarcoplasmic proteins (Wright et al., 1977). The third transition has been assigned to actin and is found between 80 and 83 °C (Wright et al., 1977). For the second transition it has also been shown that both isolated actomyosin and myosin and its sub-units undergo transitions in the same temperature range (Wright & Wilding, 1984). Recently, the thermal denaturation of titin from pork and beef has been investigated, using DSC. Denaturation was characterised by a single DSC peak at 78.4 and 75.6 °C for beef and pork titin, respectively (Pospiech, Greaser, Mikolajczak, Chiang, & Krzywdzinska, 2002).

Xiong and Brekke (1990a, 1990b) have studied salt-soluble proteins (SSP, the myofibrillar proteins) from pre- and postrigor chicken on heating at 0.6 M NaCl, pH 6, using ANS, turbidity and rigidity measurements for unfolding (hydrophobicity), protein–protein association and gelation, respectively (Fig. 5(a)–(c)). Despite some differences between breast and leg muscles and between pre- and postrigor SSP samples, a general trend was seen in all samples that protein–protein association beginning at 36–40 °C was preceded by protein unfold-

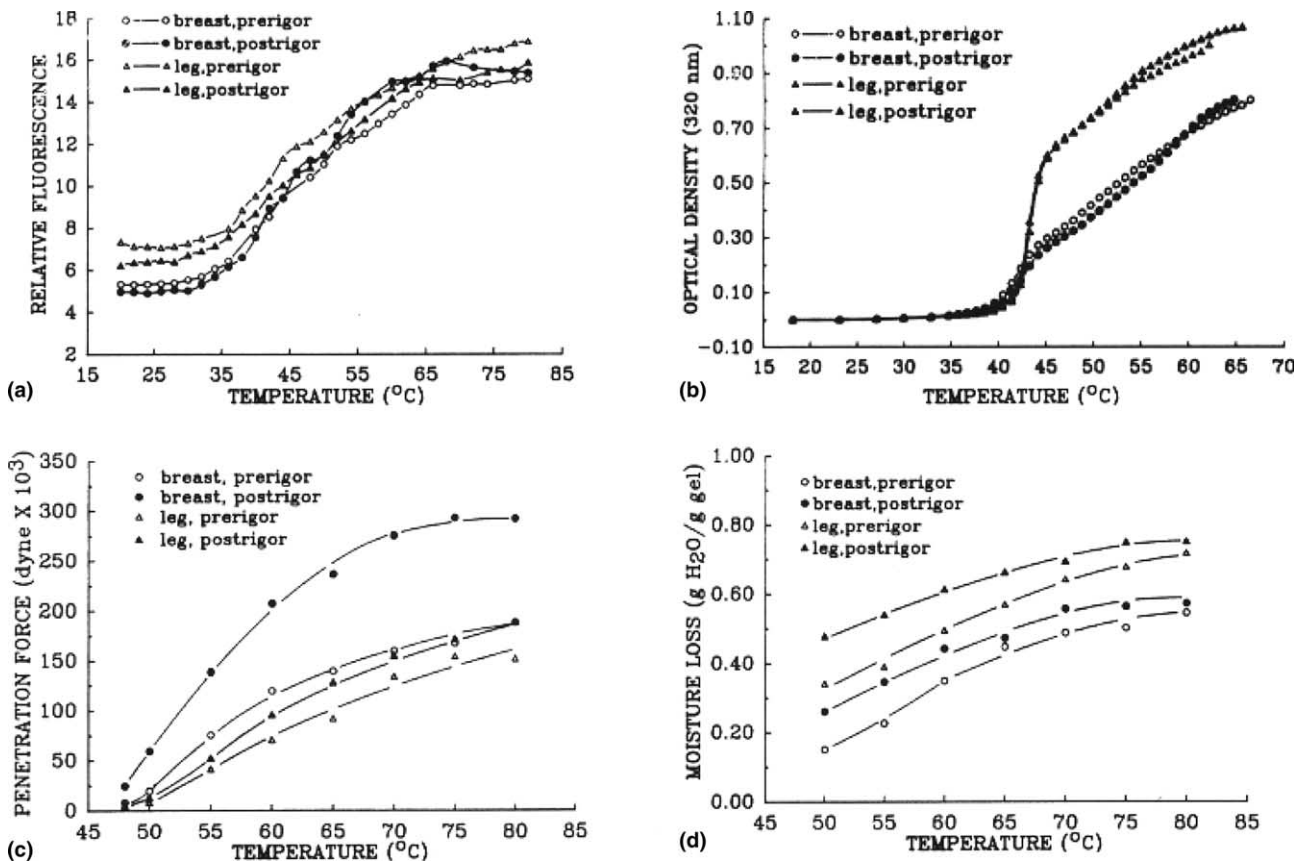


Fig. 5. Salt-soluble proteins (SSP) prepared from pre- and postrigor chicken muscles as a function of temperature. SSP were suspended in 0.6 M NaCl, pH 6.0. Heating rate = 1 °C/min. (a) ANS-protein fluorescence intensity. (b) Protein–protein interactions measured as turbidity. (c) Gel strength as penetration force. (d) Water loss (Xiong and Brekke, 1990a, 1990b).

ing at 30–32 °C. Gelation at 45–50 °C was initiated after interactions of proteins had taken place. They further stated from these investigations that the hydrophobicity was not dependent on the type of muscle (white, breast and red, leg) nor on muscle rigor state, but rather the protein–protein association varied due to these variables, thereby forming different type of gels. According to Fig. 5(b) the optical density of the SSP solution increased drastically above 40 °C and after 45 °C the increase was less. There are no differences between pre- and post-rigor meat. However, breast SSP showed a great difference from leg SSP in aggregation behaviour. But when gels are formed the penetration force is the highest for the breast post-rigor SSP (Fig. 5(c)). Comparing the water-holding of these different gels, as visualised in Fig. 5(d), the best is achieved by breast, pre-rigor. High penetration force for gels usually reflect highly aggregated junction points in the gel, as Hermansson (1982) has shown for blood plasma gels, and good water-holding gels suggest a fine-stranded gel with a higher amount of junction points per volume unit (see Fig. 6 for clarification). This suggests that breast pre-rigor SSP form more the last type of gel, as revealed by Fig. 5(b) and (d), having a lower degree of aggregation and better water-holding, whereas the leg SSP forms the former type of gel, having a high degree of aggregation and a higher penetration force.

As shown above the denaturation of myofibrillar proteins in solution usually results in gel formation, because especially myosin is unique in the sense that it form gels at very low concentration of 0.5% by weight (Hermansson & Langton, 1988). For comparison sarco-plasmic proteins need about 3% by weight to gel

(Hamm & Grabowska, 1978). When purified myosin is heated, the firmness of the gel reaches its maximum at 45 °C at pH 5.5 or at 60 °C at pH 6 (Sharp & Offer, 1992). If actin is present in the solution, a firmer gel is obtained (Yasui, Ishioroshi, & Samejima, 1980). Ionic strength (mostly NaCl) and pH are important factors since they determine if the myosin exists in monomeric form or as filaments. At ionic strengths >0.3 and at neutral pH, the myosin molecules are dispersed as monomers, forming a coarse network with large pores. At lower ionic strength the myosin molecules are assembled in filaments, resembling the natural thick filaments in the muscle. During heating a firmer gel is formed, especially if the filaments are very long. Such a gel consists of a finer and more uniform network, with smaller pores (Sharp & Offer, 1992).

The gel formation of myosin occurs in two steps, in two separate temperature regions on heating. The first part of the reaction occurs between 30 and 50 °C and the second step at temperatures above 50 °C. The first step involves aggregation of the globular heads of myosin. Sharp and Offer (1992) have studied heating of purified myosin at different constant temperature for 30 min in an electron microscope. They found that after heating at 30 °C in 30 min the appearance of the myosin molecule had not changed. After heating at 35 °C, the presence of native myosin molecules with two heads was still dominating, but other types of new structure had been formed, such as two myosin molecules, that had aggregated by dimerisation of their heads. After heating at 40 °C there were no native myosin molecules left and the only monomers present had coalesced heads. Heating to 50 °C resulted in further aggregation. At this temperature it was hard to distinguish between the individual tails. Heating at 50–60 °C lead to the formation of large globular aggregates. No tails were seen after these temperatures. The second stage involved structural changes in the helix-structure of the myosin tail, leading to network formation, where hydrophobic groups interact with each other.

4.3. Connective tissue proteins

At temperatures between 53 and 63 °C the collagen denaturation occurs according to DSC measurements (Martens, Stabursvik, & Martens, 1982), which probably involves first the breakage of hydrogen bonds loosening up the fibrillar structure and then the contraction of the collagen molecule. If unrestrained, collagen fibres shrink to one-quarter of its resting length on heating to temperatures between 60 and 70 °C. If the collagen fibres then are not stabilised by heat-resistant intermolecular bonds, it dissolves and forms gelatine on further heating. The presence of heat-stable bonds means that intermolecular linkages are retained at these temperatures and

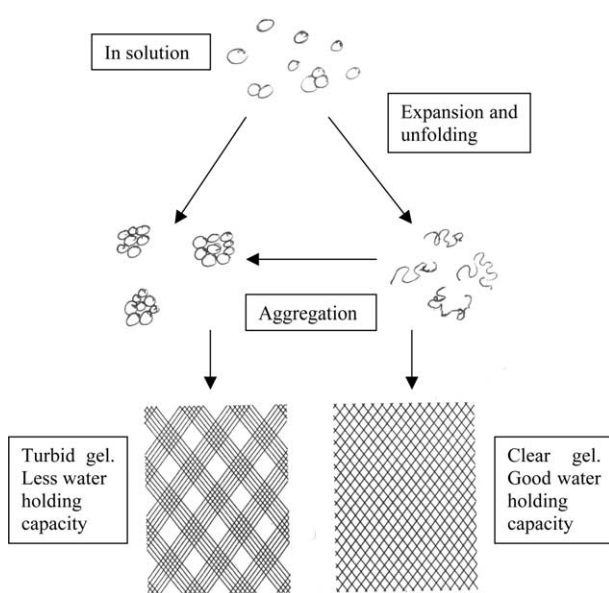


Fig. 6. Different type of gel formation on heating of globular proteins with varying degree of aggregation. Freely after Hermansson (1982).

a proportion of the fibre matrix does not dissolve (Light, Champion, Voyle, & Bailey, 1985). In young animals the epimysium contains primarily of thermally labile cross-links, the perimysium a mixture of thermally labile and stable cross-links and the endomysium of thermally stable cross-links (Sims & Bailey, 1981). As the animal age increases the thermally labile increasingly converts into thermally stable cross-links (Shimokomaki, Eisdien, & Bailey, 1972). Higher levels of heat-stable cross-links lead to the development of greater tension in the connective tissue during cooking (Sims & Bailey, 1981).

Wu, Dutson, and Smith (1985) have followed the structural alterations, using scanning electron microscopy, of the epimysium, perimysium and endomysium from bovine *sternomandibularis* caused by heating to 60 and 80 °C for 1 h. The epimysium did not show large alterations after cooking, whereas the perimysial collagen and endomysial collagen became granular at 60 °C and started to gelatinise at 80 °C. There are also differences in solubilization for the different type of collagens, where type I is more easily solubilized on heating than type III (Burson & Hunt, 1986).

5. Structure and quality of meat products on cooking

5.1. Different quality aspects of meat products

Important quality aspects of meat can embrace quality factors, such as fat- and water-holding, appearance, and the eating quality of meat products. The concept of the eating quality implies texture/tenderness, juiciness, flavour and aroma. In this review we will confine ourselves to quality aspects such as water-holding and texture of meat products.

Besides whole meat, meat products, such as hamburgers and emulsion sausages, will be referred to in this review. The procedure of manufacturing hamburger pat-

ties normally includes mincing, blending and forming, after which the patties are either frozen or fried and then frozen. Frankfurters and similar sausages are made by chopping meat, with the addition of water and NaCl, in a bowl chopper into a fine meat homogenate, in which pork fat is further dispersed and emulsified. Heat treatment of the stuffed sausage batter is performed in a smoke-chamber. The microstructure of the different meat products, such as a transverse cut of whole meat (a), a beef burger (b) and an emulsion sausage (c) can be compared in Fig. 7. A hamburger is composed of more or less intact meat fibres and fibre bundles up to 50–70%, but they are randomly distributed compared to the well-defined anisotropic structure of the whole muscle. In the so called emulsion-type sausages, however, a meat protein network formation constitutes the major part of the structure, as shown by Andersson, Andersson, and Tornberg (2000).

5.2. Structural changes of meat products during cooking

5.2.1. Whole meat

The structural events during cooking of the different proteins in their structural environment of the meat have been investigated by among others by Cheng and Parrish (1976); Jones, Carroll, and Cavanaugh (1977) and Bendall and Restall (1983). During heating, the different meat proteins denature as described above and they cause meat structural changes, such as the destruction of cell membranes (Rowe, 1989), transversal and longitudinal shrinkage of meat fibres, the aggregation and gel formation of sarcoplasmic proteins and the shrinkage and the solubilisation of the connective tissue.

Transverse shrinkage to the fibre axis occurs mainly at 40–60 °C, which widens the gap already present at rigor between the fibres and their surrounding endomysium. There is, however, a controversy regarding these observations. Davey and Gilbert (1974) for exam-

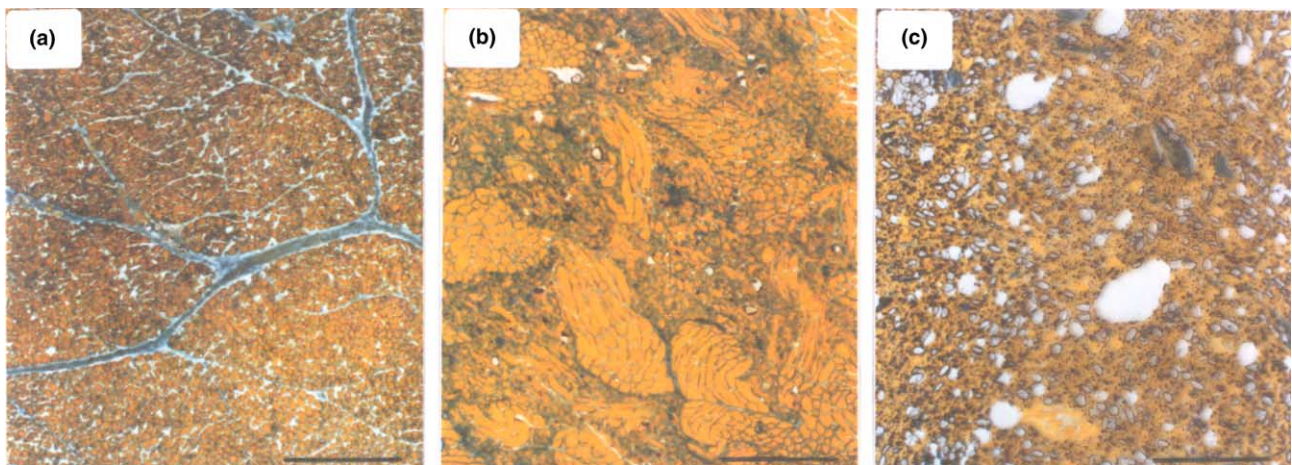


Fig. 7. Microstructure of three meat products, using light microscopy. (a) A transverse cut of a cutlet. (b) Beef burger. (c) Emulsion sausage. The bar represents 1 mm.

ple found no change in the cross-sectional area on cooking of the neck muscle, whereas Bendall and Restall (1983) found that the transverse shrinkage of both fibres and fibre bundles of *M. psoas major* starts at about 40 °C. There is also a disagreement between the results presented in the literature with regard to the temperature, where the longitudinal shrinkage of the fibre starts. Offer (1984) and Bendall and Restall (1983) have observed that fibres do not shorten until 60 °C, whereas Hostetler and Landman (1968) have reported that both sarcomere and fibre shortening usually begin at temperatures of 40–50 °C. At 60–70 °C the connective tissue network and the muscle fibres co-operatively shrink longitudinally, the extent of shrinkage increasing with temperature. This is mainly based on the fact that the intra-muscular collagen (mainly perimysial) shrinks longitudinally at 64 °C (Mohr & Bendall, 1969).

We have in one of our investigations (Tornberg et al., 1997) followed the structural changes in the different protein systems of the meat during cooking. Firstly, we separated the water phase from whole meat and studied the aggregation of the sarcoplasmic proteins, recording the change in absorbance in a spectrometer (Fig. 8(a)). From Fig. 8(a) it can be deduced that the increase in absorbance starts at about 40 °C and will be more or less terminated at 60 °C. This is in accordance with the behaviour of the sarcoplasmic proteins on heating as discussed earlier. Secondly, the shrinkage during cooking of the separated connective tissue per se and fibre shrinkage, both transversally and longitudinally, have been followed under the light microscope during cooking. The micrographs taken continuously during heating were quantified with an image analysing system (Fig. 8(b)–(d)). The different lines represent sample variation.

The transverse shrinkage of the fibre starts at 35–40 °C and then increases almost linearly as a function of temperature. The total shrinkage at 80 °C can, though, according to Fig. 8(b), vary from 3 up to 14% area. The shrinkage of the connective tissue per se starts at 60 °C, where after at around 65 °C it contracts more intensively. However, the amount of shrinkage varies substantially from about 7% area up to 19% area. The discrepancy in results could, among other things, be due to large biological variation within a muscle and between different muscles. These observations are similar to those of Bendall and Restall (1983).

5.2.2. Comminuted meat products

Comminution in combination with salt addition and subsequent heating, which are processes used in the making of hamburgers and emulsion sausages, drastically alter the structure of the meat system. On cooking hamburgers the higher occurrence of whole fibres and pieces of fibres causes more shrinkage as compared to emulsion sausages. In the latter product, higher amounts

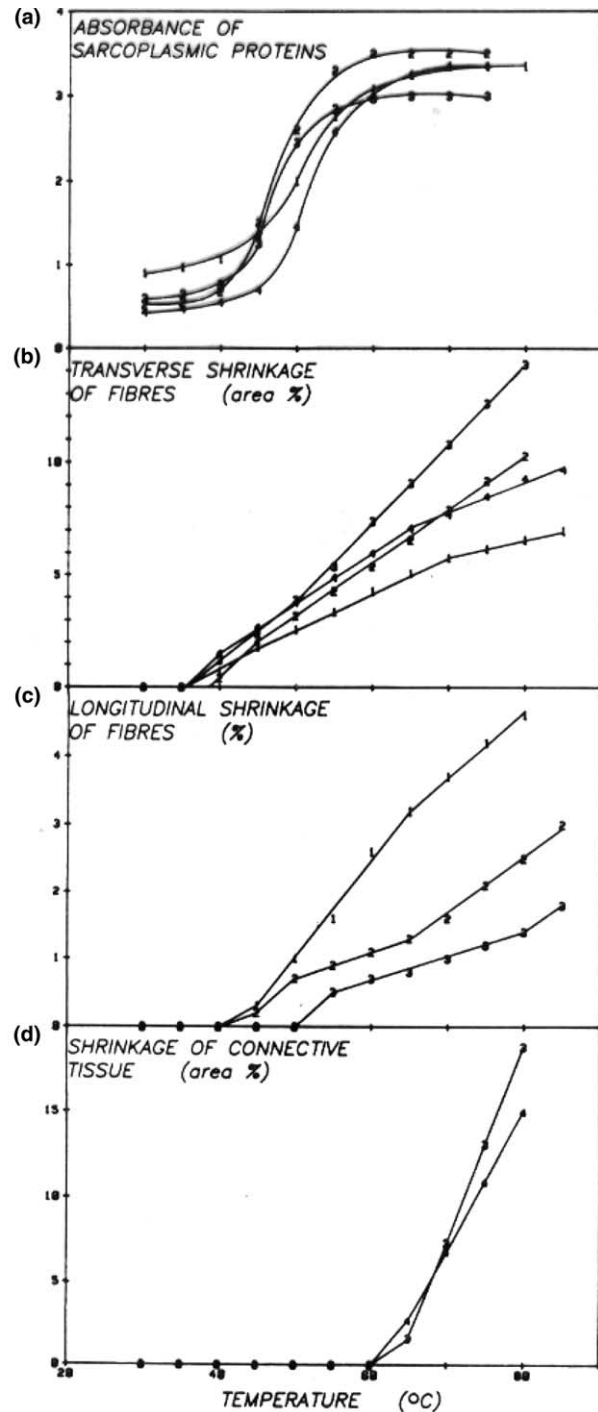


Fig. 8. Effect of cooking temperature on the absorbance of the sarcoplasmic fraction (a), the transverse (b) and longitudinal (c) shrinkage of fibres and the shrinkage of connective tissue (d) prepared from four different animals, using beef *M. biceps femoris* (Tornberg et al., 1997).

of myofibrillar proteins are extracted, which on heating create a dense protein network, a gel, that holds water efficiently by capillary forces.

The quality of that protein network is influenced by a number of factors interacting in a complicated way, as

visualised in Fig. 9. The amount of myofibrillar proteins that are extracted into the water-phase during comminution and blending is generally considered to be the most important factor for the quality of the meat network. Moreover, the type of gel matrix formed is related to the dispersed or aggregated state of the protein prior to gelation. The complex meat system consists of, not only dissolved proteins, but also insoluble components like meat fibres, connective tissue and fat. The amount and state of these components have a large impact on the gelation properties. The properties of the gel are also influenced by the heating process.

5.3. Water-holding of meat products on cooking

5.3.1. Whole meat

Considering the structural basis of water-holding in whole meat, the work of Offer and co-workers cannot be overlooked (Offer, 1984; Offer & Knight, 1988; Offer et al., 1989). It is important to point out that the structural origin of water-holding in whole meat and highly comminuted products is different. In the former it is the shrinkage and swelling of myofibrils that is the crucial factor (Offer & Knight, 1988), whereas the ability of meat proteins to form different types of gel comes more into play in comminuted meat products (Hermansson, 1986).

Water loss from whole raw meat can be obtained by evaporation from the surface, and as exudate, when a muscle is cut. This exudate, a solution of sarcoplasmic proteins and called drip loss, is drained from the cut surface of the meat by gravity, if the viscosity of the exudate is low enough and the capillary forces do not

retain it. Offer et al. (1989) have confirmed that this drip loss arises predominantly from the longitudinal channels through the meat between the fibre bundles. In order to be able to understand and control changes in water-holding, the main question that then arises is how water is accumulated and lost in those channels.

Most of the water in the living muscle is held within the myofibrils ($\approx 80\%$), in the spaces between the thick and thin filaments (Offer et al., 1989). Any large changes in the distribution of water within the meat structure, by necessity, originate from changes in this spacing. Lateral shrinkage of the filament lattice is brought about by a pH-fall closer to the isoelectric point, rigor contraction and myosin denaturation (Offer & Knight, 1988). There will only be changes in the water distribution, if the myofibrils change in volume. The fact that fibre and fibre bundles shrink when their constituent myofibrils shrink, has been shown by Offer and Cousins (1992), thereby giving rise to the two extracellular compartments around fibres and fibre bundles.

Cooking induces structural changes, which decrease the water-holding capacity of the meat. The review by Offer (1984) summarises the structural changes occurring on cooking as follows: When the transverse shrinkage to the fibre axis occurs mainly at 40–60 °C this widens the gap already present at rigor between the fibres and their surrounding endomysium. At 60–70 °C the connective tissue network and the muscle fibres cooperatively shrink longitudinally, the extent of shrinkage increasing with temperature. This shrinkage causes the great water loss that is obtained on cooking. It is then presumed that water is expelled by the pressure exerted by the shrinking connective tissue on the aqueous solution in the extracellular void.

A convenient way of studying water-holding in meat is to investigate the multi-exponential decay of the relaxation time, T_2 , of water protons in the muscle, using the ^1H pulse-NMR (nuclear magnetic resonance). Two dominating, discernible relaxation processes have mostly been observed, where the major fraction ($\approx 80\%$) of the muscle water has a T_2 (called T_{23}) between 35 and 50 ms, while the rest of the water relaxes in the range of 100–150 ms (called T_{22}). The percentage of water relaxing with the shortest relaxation time can be considered as mainly held by the myofibrils, since a very high fraction of the water is occupied by the myofibrils. Tornberg and Larsson (1986) have further shown, by comparing the percentage of water having a $T_2 = 100\text{--}150$ ms with the percentage of area around the fibre bundles (evaluated by light microscopy), that the latter could be predicted with an 80% probability using the pulse-NMR method.

Studies of the water distribution on cooking of meat, using ^1H pulse-NMR, have been done by Fjelkner-Modig and Tornberg (1986), Tornberg and Larsson (1986), Borisova and Oreshkin (1992), Tornberg,

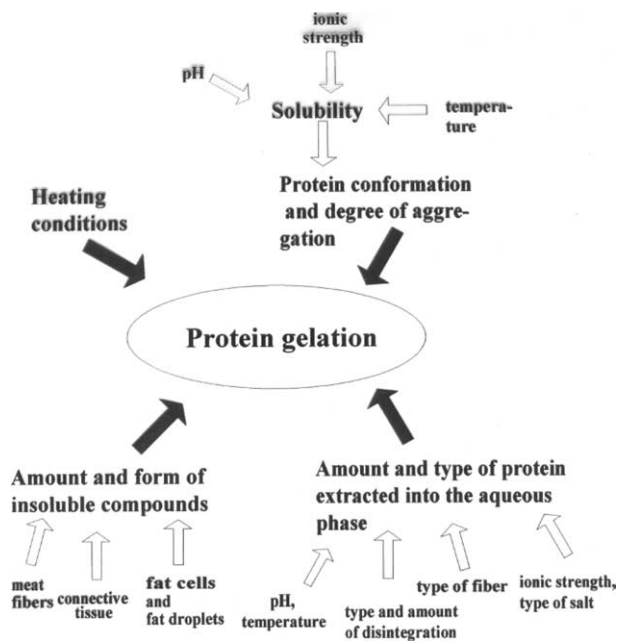


Fig. 9. Factors affecting protein gelation, when cooking a meat batter (Andersson et al., 1997).

Andersson, Göransson, and von Seth (1993), Micklander, Peshlov, Purslow, and Engelsen (2002) and Bertram, Balling Engelsen, Busk, Karlsson, and Andersen (2004). It was noted both by Fjelkner-Modig and Tornberg (1986), Tornberg and Larsson (1986) and Tornberg et al. (1993) that the T_2 for the major fraction of the water (T_{23}) in both beef and pork was lowered on cooking from around 40 to around 30 ms. This suggests a more aggregated, dense protein structure in the myofibrils on heating, which is in accordance with the denaturation behaviour of the proteins as discussed earlier. For the cooked meat samples, the percentage water of T_{22} and extracellular space around fibre bundles, as evaluated by light microscopy, was compared at the different cooking temperatures (Fig. 10). The amount of water around fibre bundles increases up to 50 °C, in comparison with the raw meat, which seems to be in accordance with the transverse shrinkage of fibres and fibre bundles. Above 50 °C this widened gap diminishes, again up to 70 °C, probably mainly due to the shrinkage of the connective tissue. The increase in extracellular space from 70 to 90 °C is more difficult to comprehend but, according to the light micrographs a swelling of the perimysium seems to occur at these cooking temperatures.

5.3.2. Comminuted meat products

On comminution of meat and salt addition, leading to the extraction of meat proteins, which occurs when producing hamburger patties and/or emulsion type sausages, more of the structure consists of a protein gel network after heat treatment. This is especially the case for the highly comminuted emulsion-type sausages (4% salt on meat basis), which is reflected in the water-holding of this meat product, being superior to both hamburgers (0.4% salt) and whole meat, reflected in Fig. 11. In Fig. 11 the cooking loss of the whole meat, the ham-

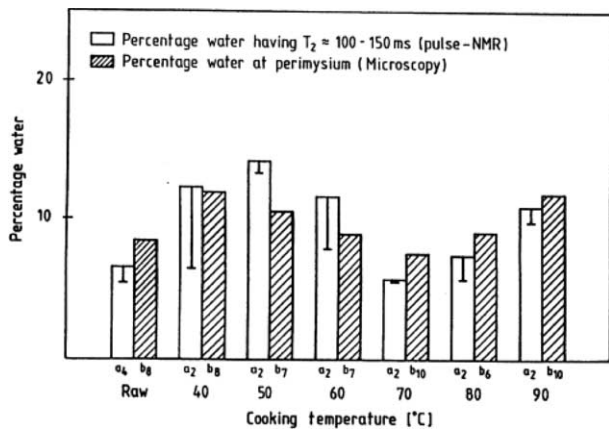


Fig. 10. Percentage water having a T_{22} of 100–150 ms, using proton pulse NMR, and percentage water around fibre bundles (at perimysium), as determined with light microscopy, as a function of cooking temperature (a_n , n = number of samples; b_n , n = number of photos).

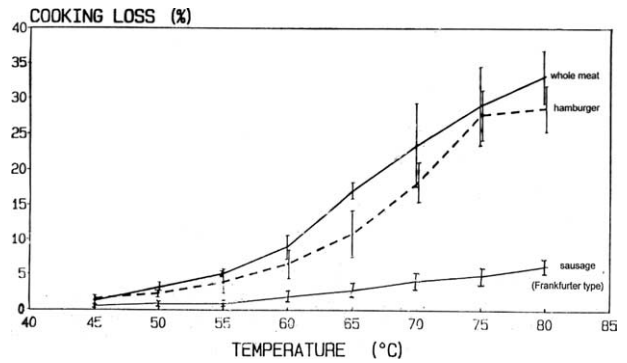


Fig. 11. Cooking losses (%) as a function of cooking temperature for whole meat, hamburger and an emulsion sausage. The meat raw material used in all cases was *M. biceps femoris*.

burger and the emulsion sausage of beef *M. biceps femoris* can be compared at different cooking temperatures (The heating gradient was 1.5 °C/min up to 60 °C, thereafter 0.7 °C/min was used until the final temperature was reached in the centre of the meat sample). The temperature dependence on cooking loss for the whole meat, the highest increase in water loss is achieved in the temperature region of 60–80 °C, which is similar to the results of other investigators (Davey & Gilbert, 1974; Bouton, Harris, & Shorthouse, 1976; Bendall & Restall, 1983). Furthermore, it is interesting to note that although the hamburgers have been comminuted, the cooking losses are almost as large as for the whole meat. This is probably due to the more prevalent shrinkage of whole fibres and pieces of fibres, causing larger water losses in the hamburgers as compared to emulsion sausages. Because in the latter product, the higher amounts of myofibrillar proteins extracted create a dense protein network that holds more water (in combination with 4% added potato starch, which is usually added in Swedish emulsion-type sausages).

In one of our latest studies, by Kovascne (2004), on beef-burgers patties, fried on a double-sided pan fryer to a core temperature of 72 °C and having different pan temperatures (100, 150 and 175 °C), we found that the water losses, sometimes were reaching up to 60% of the initial water content. The pressure driven water loss, due to meat protein contraction, never went below 80% of the total water losses. This means that the pressure driven water loss is a substantially more important mechanism governing the water loss on frying of beef-burgers than the evaporation losses occurring at the surface crust formed at the higher frying temperatures of 150 and 175 °C.

5.4. Textural properties of meat products on cooking

5.4.1. Whole meat

Meat texture can be evaluated by both sensory and instrumental methods. Numerous studies have been

conducted in the area of meat cookery, but especially attempts to relate sensory evaluated tenderness with mechanical and structural changes in the meat during cooking are scarce (Josell & Tornberg, 1994; Martens et al., 1982; Tornberg et al., 1997). Textural properties of meat with emphasis on the instrumental methods have been reviewed by amongst others Purslow (1991), Lepetit and Culioli (1994) and Harris and Shorthose (1988). In this review only the cooking temperature dependence of texture properties of meat will be dealt with.

During the mastication of meat, deformation and fracture of the samples takes place. The mechanical forces acting on meat can include shear, compressive and tensile forces and they should be defined in the mechanical test in use. As meat is a composite, it is important to study in which structural elements failure takes place, and where cracks propagate, to be able to understand its mechanical properties. Tensile tests on cooked strips of both beef (Bouton & Harris, 1972; Purslow, 1985) and pork (Christensen, Purslow, & Larsen, 2000; Josell & Tornberg, 1994; Mutungi, Purslow, & Warkup, 1995) have been used.

For the assessment of the texture of whole meat, the empirical method of the Warner–Bratzler shear device is the most widely used. The most commonly used configuration is the one in which the shearing plane is perpendicular to the muscle fibres. Tensile, shear and compression forces operate in this type of test. The W–B-technique is, however, the instrumental technique that usually yields the best correlation with sensory panel scores for meat toughness.

When relating the sensory properties of the meat with the mechanical, structural traits, Josell and Tornberg (1994) used *M. longissimus* from pork and Tornberg et al. (1997) whole and minced meat from beef (*M. biceps femoris*). From the former investigation the sensory evaluated tenderness of pork loin increases substantially from 50 to 65 °C, where after it decreases again to 80 °C (Fig. 12). In this study also the temperature dependence of longitudinal and transversal fracture stress and strain were recorded, using tensile measurements (Fig. 13(a) and (b)). As shown by Purslow (1985) for beef the longitudinal rupture of the fibres require higher stresses than transversal rupture, which was also the case for pork. The increase in tenderness in the temperature interval 50–65 °C was observed as a decrease in the transversal fracture stress and in the transversal and longitudinal fracture strain (Fig. 13(a) and (b)). The elastic modulus of the meat registered both at small and large deformations increased with cooking temperature from 50 up to 80 °C (Fig. 13(d)).

Bouton and Harris (1981) showed, using W–B shear peak values, that there is a decrease between 50 and 65 °C, which they attributed to a weakening of the collagenous connective tissue, due to denaturation. This

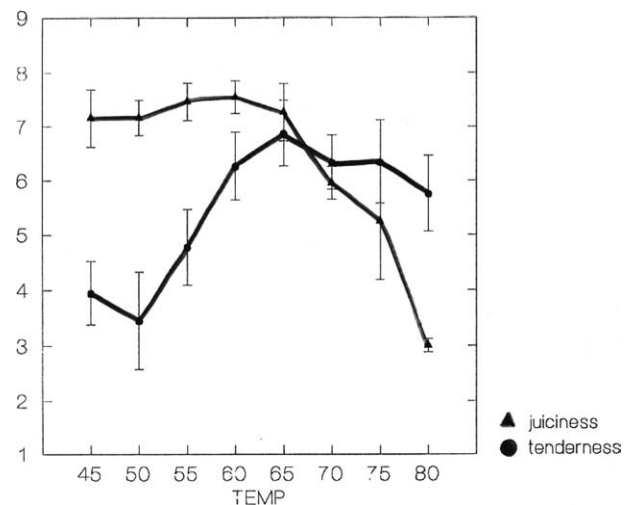


Fig. 12. Sensory evaluated tenderness and juiciness for pork *M. longissimus dorsi* versus cooking temperature (Josell and Tornberg, 1994).

phenomenon could also be the reason for the fracture stress to diminish from 50 to 65 °C in our investigation. However, we are more inclined to suggest another hypothesis to explain the observed phenomena. In the tensile measurements the strain at fracture tells us how much the meat yields on tearing and this yielding would increase between 50 and 65 °C, if the weakening of the perimysium connective tissue would be the dominating mechanism in the tensile measurements. However, the opposite is observed, i.e., from 50 to 65 °C a decrease in fracture strain from 70% to 30% is achieved (Fig. 13(b)). Concomitantly, the elastic modulus of the whole meat system increases (Fig. 13(c) and (d)), which means that the meat converts from a viscoelastic to a more or less elastic material.

We suggest that the reason for the toughness observed for the raw meat up to about 50 °C is based on the fact that the applied stress during mastication is reduced by viscous flow in the fluid-filled channels in between fibres and fibre bundles. This situation is, however, improved from 50 to 65 °C by the formation of a gel of aggregated sarcoplasmic proteins gluing the fibres and fibre bundles together. The viscous flow becomes then lower as the elasticity of the meat increases in that temperature region. That means there is a higher probability of the applied stress being transferred within the material to the crack without any viscous dissipation of energy and there propagating it. The piece of meat is then more easily fractured in the mouth and mastication is facilitated and tenderness improved. However, above 65 °C elasticity acts adversely and impairs the tenderness. For a fully brittle fracture in a linearly elastic material (which we suppose cooked meat above 65 °C behaves like) higher elastic modulus gives rise to larger tensile stresses to extend a crack (Jowitt, 1979), i.e., a tougher meat. We suggests that the contraction of the

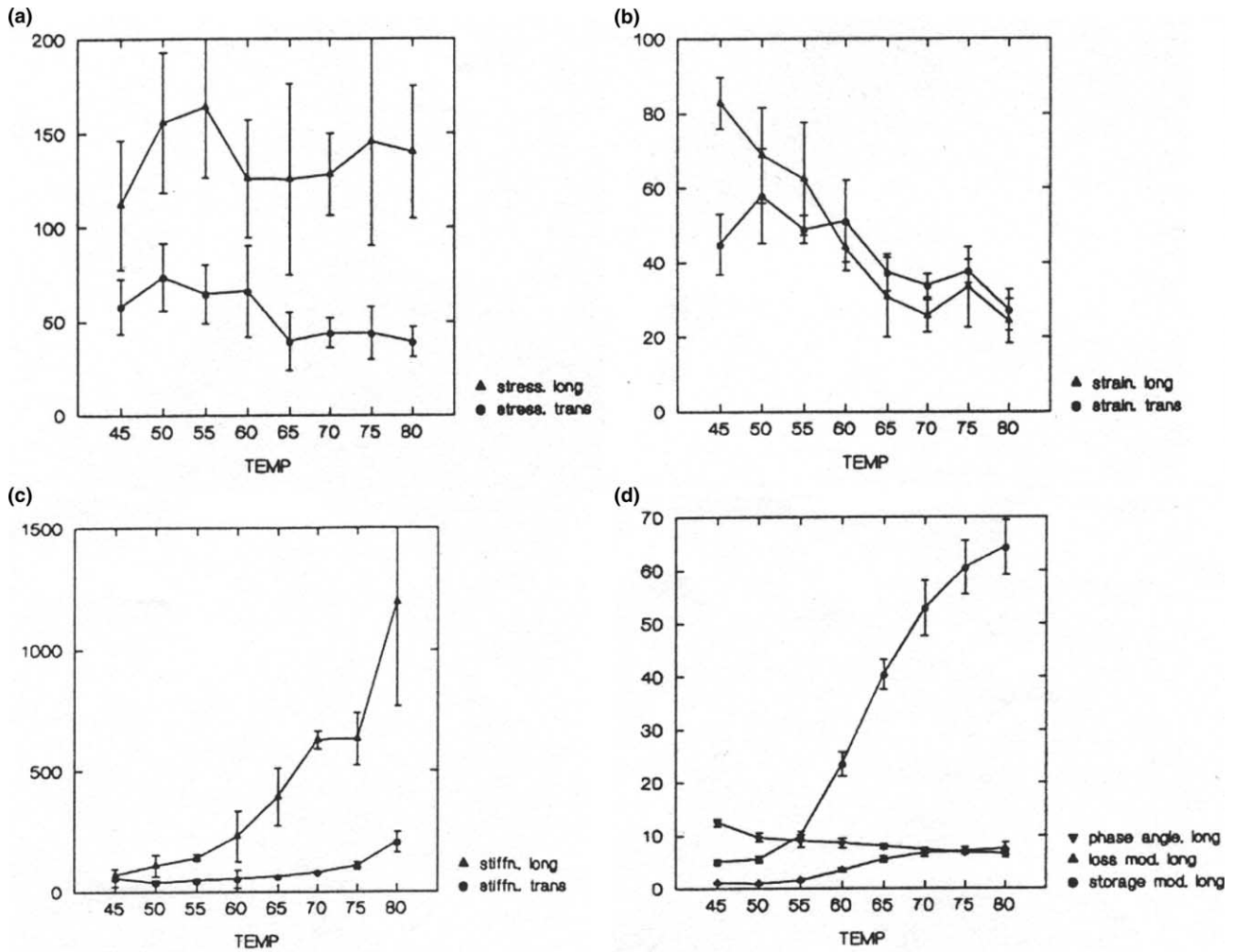


Fig. 13. Different mechanical properties for pork *M. longissimus dorsi* versus cooking temperature. (a) Longitudinal and transversal fracture stress (kPa). (b) Longitudinal and transversal fracture strain (%). (c) Longitudinal and transversal initial stiffness (kPa). (d) Longitudinal storage and loss moduli (kPa). (Josell and Tornberg, 1994).

connective tissue, mainly occurring after 65 °C as seen in Fig. 8, gives rise to an increase in the elasticity of the meat by forming a much denser material in the temperature region of 65–80 °C and thereby a tougher meat.

5.4.2. Comparison between whole and minced meat

These findings are further substantiated, when comparing the rheological properties of whole and minced meat (*M. biceps femoris*, beef) in relation to sensory and structural characteristics (Tornberg et al., 1997). The sensory evaluation (Fig. 14) showed that, for the whole meat, the toughness decreased drastically from 55 to 60 °C, thereafter increased again up to 80 °C. For the minced meat, however, the hardness increased over the whole temperature range and was significantly lower than the toughness of the whole meat at cooking temperatures below 60 °C. The rheological properties of the whole meat, cut transversally to the fibre direction, and the minced meat were also recorded, using a low deformation dynamic shear test, and the results

can be seen in Fig. 15. The storage modulus increases steeply with the temperature from 50 to 65 °C both for the whole and the minced meat, where after it levels

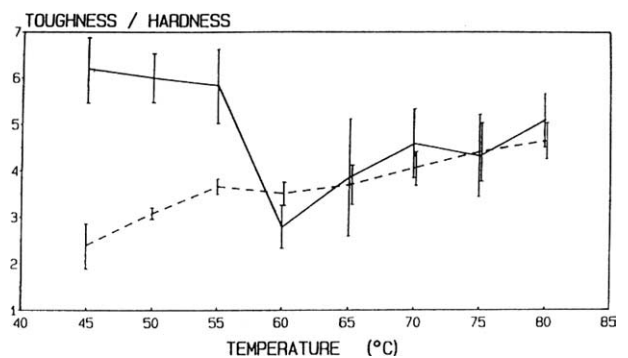


Fig. 14. Toughness of whole meat (—) and hardness of minced patties (---), made out of beef *M. biceps femoris*, as a function of cooking temperature. The bars give the standard deviation (Tornberg et al., 1997).

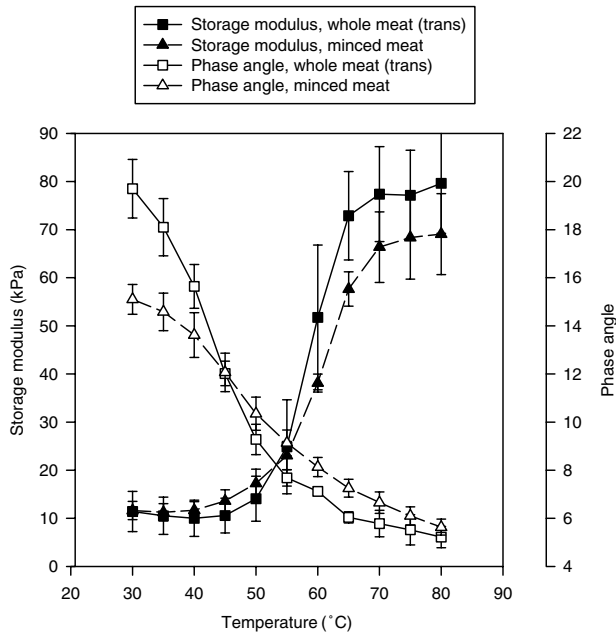


Fig. 15. Storage modulus (kPa) and phase angle (degrees) for whole meat (■, □) and for minced meat (▲, △), respectively, made out of beef *M. biceps femoris*, as a function of cooking temperature. The bars give standard deviation (Tornberg et al., 1997).

off to values for the modulus of 80 and 70 kPa for the whole and minced meat, respectively. The phase angle, which is the quotient between the loss and the storage modulus, decreases already from 35–40 °C down to a low plateau round 65–70 °C.

These results tell us that the spatial arrangement of the fibres is of utmost importance for the textural behaviour of the meat. In the whole meat the crack that fractures the meat has to pass over the fluid filled channels at temperatures below 55 °C (for the case of biceps femoris) and therefore the fracture energy will dissipate as viscous flow and it is therefore anticipated as tough. For the minced meat, however, the structure is no longer anisotropic and the more or less disintegrated muscle fibre and bundles are now randomly distributed in the batter and the crack then preferentially goes through the extracted myofibrillar mass that holds the fibre together in the batter. The fracture stress to propagate a crack in that material is much less than the stress needed to pass through a meat fibre. As the temperature increases from 45 to 65 °C in the beef burger batter the extracted proteins form a gel and after 65 °C the contraction of the connective tissue and the substantial loss of water contribute to a denser product. All these structural changes give rise to an increased elasticity of the minced meat (Fig. 15), and also an enhanced sensorial perceived toughness (Fig. 14).

If this increased toughness of the minced meat on heating was beneficial for the overall impression of it was not investigated in that study. For an emulsion sau-

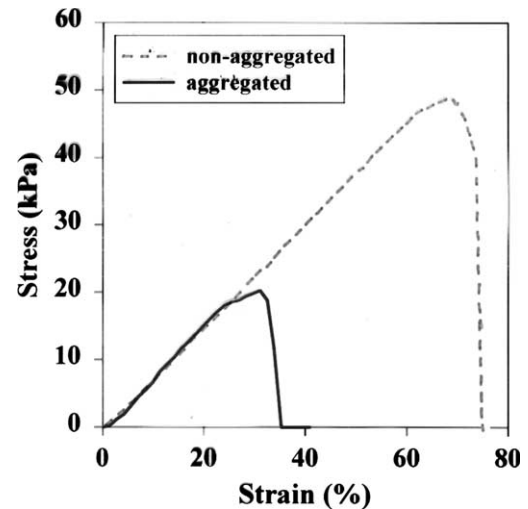


Fig. 16. A comparison of the stress–strain curves in the tensile mode for aggregated and non-aggregated cooked emulsion sausages (Andersson et al., 1997).

sage, however, where different textures were produced by using varying recipes, salt concentrations and pH (Andersson, Andersson, & Tornberg, 1997), the overall acceptability of the emulsion-type sausage was mostly governed by the properties of the protein network. This was best characterised by the fracture strain in the tensile test used. When comparing the stress–strain curves obtained for aggregated and non-aggregated sausages, which can be seen in Fig. 16, the latter were characterised by having larger fracture strains and they were also rated having the highest overall acceptability. Even the elastic modulus (G') was significantly higher for non-aggregated sausage (31.5 kPa) compared to the aggregated (24.9 kPa). Evidently, a dense, elastic meat protein gel is sensorial preferred to a more brittle, grainy protein gel in an emulsion sausage.

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