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## Abstract

Milk proteins are known to possess a wide range of functional properties, such as emulsification, thickening, gelling and foaming. Milk proteins facilitate the formation and stabilisation of oil droplets in emulsions or of air bubbles in foams in formulated foods. These functional properties of milk proteins are exploited in the manufacture of dairy and other products, such as recombined milk, cream, butter, yoghurt, ice cream, cream liqueurs, dressings, mayonnaise, sauces and desserts. This chapter provides an overview of the emulsifying and foaming properties of milk proteins, focusing on the adsorption of milk proteins at oil–water and air–water interfaces with emphasis on the preferential adsorption among milk proteins and the stability of milk-protein-based emulsions and foams. Highlights on the behaviour of milk-protein-stabilised emulsions after consumption that have recently attracted a great deal of research interest are discussed briefly.

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## Keywords

Milk-protein-stabilised emulsions • Foam stability • Emulsion stability • Emulsion digestion • Gastrointestinal tract

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## 5.1 Introduction

Milk proteins are generally classified into caseins and whey proteins. Caseins are flexible proteins that have no rigid  $\alpha$ -helix and  $\beta$ -pleated sheet structure and comprise four distinct proteins,  $\alpha_{s1}$ -,  $\alpha_{s2}$ -,  $\beta$ - and  $\kappa$ -casein, all of which are phosphoproteins (Fox, 2009; Singh, 2011). In contrast, whey proteins are globular in nature and

possess high levels of secondary, tertiary and, in most cases, quaternary structure. Whey proteins can be fractionated into  $\beta$ -lactoglobulin ( $\beta$ -lg),  $\alpha$ -lactalbumin ( $\alpha$ -la), bovine serum albumin, lactoferrin, immunoglobulins and several minor proteins. Most commercially available milk protein ingredients are mixtures of various caseins or whey proteins, e.g., caseinates, whey protein concentrates (WPCs), whey protein isolates (WPIs) and milk protein concentrates (MPCs). These ingredients are widely used in the preparation of a broad range of food emulsions and foams.

Emulsions (milk, cream, butter, mayonnaise, coffee whiteners, whipped toppings, cream liqueurs and low fat spreads) and foams (whipped cream and ice cream) are dispersed oil–water and air–water systems, respectively, and represent a major proportion of processed food formulations. During the emulsification or foaming process, both caseins and whey proteins adsorb rapidly at oil–water or air–water interfaces, forming a film around the oil droplets or air bubbles (Damodaran, 1997; Dickinson and Patino, 1999). This adsorbed layer or film protects the oil droplets or air bubbles against various physicochemical processes of instability. Knowledge of protein structures at the interfaces and their mechanical and rheological properties is essential for controlling the stability of these dispersed systems. Protein interfacial structures and properties are affected by changes in pH, ionic strength, temperature, shear and pressure, which in turn alter the stability of these dispersed systems. The aim of this chapter is to provide readers with an overview of the formation and properties of emulsions and foams stabilised by various forms of milk proteins, focusing mainly on recent studies.

In the case of emulsions, major advances have been made in understanding the adsorption process, the composition and structure of adsorbed layers of proteins and how these proteins influence their physical and chemical properties (Dickinson and Stainsby, 1988; Dickinson, 1998, 1999a; Damodaran, 2005; McClements, 2005). Interestingly, in recent years, the physical and biochemical stability of emulsions after consumption has generated a great deal of research interest (McClements *et al.*, 2009; Singh *et al.*,

2009; Golding and Wooster, 2010; Le Révérend *et al.*, 2010; Singh, 2011; Singh and Sarkar, 2011; Singh and Ye, 2013). Some progress has been made on understanding how the adsorbed layers and the physical structures of food emulsions influence the rates of lipid digestion. Knowledge of these complex interactions between the emulsion droplets and the physiological components, such as mucin, gastric and intestinal enzymes (e.g., pepsin, trypsin and lipases) and bile salts, is key to understanding the physiological behaviour of emulsions during their transit through the gastrointestinal tract. Hence, in this chapter, we discuss current advances in our understanding of the physiological behaviour of emulsions, particularly those stabilised by milk proteins, from a physicochemical viewpoint.

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## 5.2 Formation and Stability of Protein-Stabilised Emulsions

Generally, emulsions can be prepared using a wide range of high shear apparatus, such as colloid mills, high speed blenders, high pressure valve homogenisers and ultrasonic equipment, that mix an oil phase and an aqueous phase together in the presence of a surfactant (McClements, 2005). During high pressure valve homogenisation a coarse mixture of the oil and aqueous phases is forced through a narrow slit under the action of high pressure, resulting in cavitation, intense laminar shear flow and turbulence. Consequently, the structurally amphiphilic emulsifier molecules, such as proteins, are adsorbed at the interface, creating a stabilising interfacial layer at the droplet surface and leading to the generation of fine, uniformly dispersed droplets (Dickinson, 2003).

The physicochemical properties and the stability of emulsions depend on a number of factors such as the types and concentrations of the dispersed phase and the continuous phase, the nature of the stabilising layer, temperature, pH, the viscosity of both phases, the homogenisation conditions and other processing parameters employed, such as heat treatment, high pressure

processing and enzymatic hydrolysis (McClements, 2005). The stability of an emulsion therefore refers to its ability to resist any alteration in its properties and structure over the time scale of observation. Interfacial layers of different structures, compositions and charges can be carefully designed using specific proteins to meet the physicochemical demands and the required stability of food emulsions. Interestingly, as long as sufficient surfactant to cover the newly-created interface is present during homogenisation, emulsions are generally very stable to coalescence over prolonged storage periods. However, these emulsions are susceptible to different types of instability as a result of various types of physical and chemical processes, which in turn lead to enhanced creaming or serum separation. Generally, physical instability refers to modifications in the spatial arrangement or size distribution of the emulsion droplets, such as creaming, flocculation and coalescence, whereas chemical instability includes changes in the composition of the emulsion droplets themselves, such as oxidation and hydrolysis (McClements, 2005).

Stokes' Law can be used to describe creaming, which involves the movement of oil droplets under gravity or an applied centrifugal force to form a concentrated cream layer at the top of the emulsion without any change in the droplet size distribution. The rate of creaming can be calculated using the following mathematical expression (Hunter, 1989; McClements, 2005; Singh *et al.*, 2009):

$$v_{stokes} = \frac{2r^2(\rho_1 - \rho_2)}{9\eta} \quad (5.1)$$

where  $v_{stokes}$  = velocity of creaming,  $r$  = radius of the emulsion droplets,  $\rho_1$  and  $\rho_2$  = densities of the continuous and dispersed phases, respectively, and  $\eta$  = shear viscosity of the continuous phase.

Hence, the kinetic stability of an emulsion can be increased or the creaming rate can be decreased by lowering the radius of the droplets, by increasing the viscosity of the continuous phase or by decreasing the difference in density between the two phases.

However, instabilities other than creaming, such as flocculation or coalescence, cannot be described by this law. Emulsion flocculation is an aggregation process that arises when droplets associate because of unbalanced inter-atomic attractive and repulsive forces (Dalglish, 1997). Commonly, there are two types of droplet–droplet interaction, i.e., depletion flocculation and bridging flocculation. Generally, depletion flocculation occurs because of the presence of a non-adsorbing biopolymer in the continuous phase of the emulsion, which can promote the association of emulsion droplets by inducing an osmotic pressure gradient within the continuous phase surrounding the droplets. In contrast, bridging flocculation occurs when a high molecular weight biopolymer at a sufficiently low concentration adsorbs on to two or more emulsion droplets, resulting in bridges (McClements, 2005).

In contrast to flocculation, coalescence refers to a completely irreversible increase in droplet size by the accretion of two or more primary emulsion droplets, gradually leading to the separation of the oil phase and the aqueous phase. Coalescence generally occurs when the stabilising film surrounding the emulsion droplets is thinned to a certain critical thickness, resulting in film breakage, thus joining emulsion droplets (van Aken *et al.*, 2003; van Aken, 2004). Generally, emulsions are stable to coalescence as the proteins or other biopolymer molecules adsorb at the droplet surfaces, forming a dense viscoelastic interfacial layer (Dickinson and Stainsby, 1988). However, any extreme processing conditions, such as high shear or enzymatic hydrolysis, that lead to significant attrition of the interfacial film can give rise to gradual agglomeration of bare emulsion droplets, resulting in coalescence and oiling-off. For instance, coalescence has been widely reported in emulsions stabilised by whey protein hydrolysates because of the formation of a thinner interfacial film and the reduced surface viscosity of an interface formed with predominantly short peptides as opposed to intact proteins (Agboola *et al.*, 1998 Singh and Dalglish, 1998).

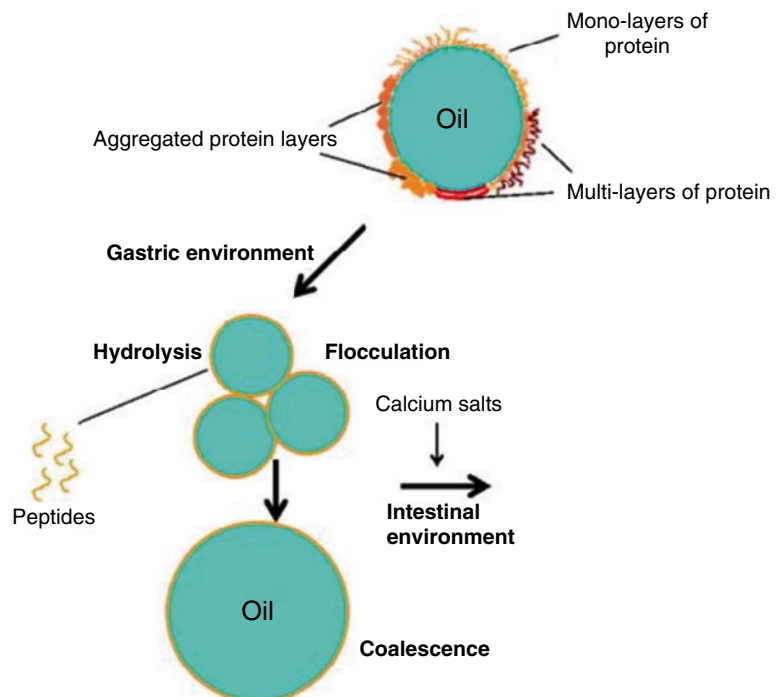
### 5.2.1 Aspects of Emulsions Stabilised by Milk Proteins

Milk proteins such as caseins, caseinates, WPIs,  $\beta$ -lg and bovine serum albumins are known to be excellent emulsifiers because of their amphiphilic nature (Morr, 1982; Mulvihill and Fox, 1989). In most food emulsions, the oil droplets are coated by a continuous film of adsorbed material, such as caseins and/or whey proteins. They reduce the interfacial tension between the oil and aqueous phases, form films with different rheological properties and, thus, stabilise the emulsion droplets. The structure and the composition of the adsorbed layers can be quite complicated because foods in general contain a variety of surface-active agents; all are possibly adsorbed at the interface either individually as monolayers or aggregates or in combination, resulting in complex multi-layered interfacial layers (Singh *et al.*, 2009), as illustrated in Fig. 5.1. The nature of the interfacial layers formed depends largely on the type, concentration, charge and conformation of the adsorbed milk protein, and on the types of

interaction and competition that occur between the adsorbed species (Dickinson, 2003; McClements, 2005).

The role of the caseins and whey proteins in stabilising emulsions has been thoroughly investigated (see reviews by Kinsella, 1984; Morr and Ha, 1993; Dalgleish, 1995, 2006; Wong *et al.*, 1996; Dickinson, 1999b, 2001; Singh, 2005). The most commonly used forms of milk protein in food emulsions are sodium caseinate and whey proteins (WPIs or WPCs). Because of their highly surface-active properties, it is possible to make stable emulsions at a relatively low ratio of milk protein to oil (about 1:60). In these emulsions, the surface protein coverage is a function of increasing protein concentration until it reaches a plateau value of about 2.0–3.0 mg/m<sup>2</sup> (Euston and Hirst, 1999; Srinivasan *et al.*, 2001). Because of the flexible structures of caseins, they adsorb rapidly at the interface, forming extended adsorbed layers up to about 10 nm thick (Holt and Sawyer, 1988; Dalgleish, 1990, 1995, 1996a; Mackie *et al.*, 1993; Dickinson and McClements, 1995; Fang and Dalgleish, 1998). In contrast, globular

**Fig. 5.1** Schematic illustration of the possible changes in milk protein-stabilised emulsions as they pass through the *in vitro* physiological model (Singh and Ye, 2013: reproduced with the permission of Elsevier Inc.)



whey proteins such as  $\beta$ -lg unfold partially, somewhere intermediate between the native state and the fully denatured conformation, resulting in compact adsorbed layers that are only about 2 nm thick (Dickinson, 1998). The sequence of surface activity reported for milk proteins is  $\beta$ -casein > monodispersed casein micelle > serum albumin >  $\alpha$ -1a >  $\alpha$ <sub>s</sub>-casein =  $\kappa$ -casein >  $\beta$ -lg > euglobulins (Ennis and Mulvihill, 2000).

An overview of milk protein layers adsorbed at oil–water interfaces and their relationship to the physicochemical stabilisation of emulsions is given in the following subsections.

### 5.2.1.1 Emulsions Stabilised by Caseins and Caseinates

Caseins, because of their surface activity, are known to adsorb strongly at an oil–water interface during emulsification, thus protecting the emulsion droplets against physicochemical instability (Dickinson, 1999b). The long-term stability of emulsions against coalescence can be attributed to both electrostatic and steric stabilisation effects (Dickinson, 2006). The relative absence of tertiary and secondary structure (Holt and Sawyer, 1988) and the presence of distinct hydrophobic and hydrophilic domains in the primary structure (Swaisgood, 1992) contribute to the relatively high surface activity of the caseins. However, as there is a lack of clarity of the native structure of caseins, the conformational changes upon adsorption at the oil droplet surface are not completely understood (Dalglish, 2004).

Generally,  $\alpha$ <sub>s1</sub>-casein and  $\beta$ -casein, which contribute almost 75 % of the total casein of milk, provide similar emulsifying properties based on their amino acid sequences (Swaisgood, 1982). Both  $\alpha$ <sub>s1</sub>-casein and  $\beta$ -casein carry a net negative charge at neutral pH, are distinctly amphiphilic, have similarities in terms of linear disordered chains of around 200 residues with phosphoserine chains and have strong abilities to adsorb at oil–water interfaces. However,  $\beta$ -casein is reported to have higher surface activity and is more flexible in nature, because of its numerous proline residues, little ordered structure and negligible intermolecular cross-links, than  $\alpha$ <sub>s1</sub>-casein (Swaisgood, 1982; Dickinson, 1994).

Experimental analysis has shown that  $\beta$ -casein adsorbs to the droplet surface with its hydrophobic region strongly anchored to the oil phase and its hydrophilic region (4–50 residues at the N-terminal) protruding into the aqueous phase (Dalglish, 1996a; Dickinson, 1999b). In contrast to the tail-like anchoring phenomenon of  $\beta$ -casein,  $\alpha$ <sub>s1</sub>-casein has a loop-like conformation that binds to the droplet surface *via* peptides towards the middle of the sequence (compared with the end of the sequence in the case of  $\beta$ -casein).

Both  $\beta$ -casein and  $\alpha$ <sub>s1</sub>-casein have been used to prepare stable oil-in-water emulsions (Dickinson, 1989, 1999b; Swaisgood, 1992). Both caseins lower interfacial tension at the oil–water interface but the rate of lowering of the interfacial tension is greater for  $\beta$ -casein than for  $\alpha$ <sub>s1</sub>-casein. Moreover,  $\alpha$ <sub>s1</sub>-casein-stabilised emulsion droplets have higher negative charge at neutral pH and are relatively more susceptible to flocculation at high ionic strengths than  $\beta$ -casein-stabilised emulsion droplets (Dickinson *et al.*, 1988). Based on a few experimental studies, it can be inferred that  $\alpha$ <sub>s1</sub>-casein accounts for less surface coverage, resulting in thinner interfacial films, than  $\beta$ -casein (Brooksbank *et al.*, 1993; Dalglish, 1993, 1996b).  $\beta$ -Casein, because of its relatively higher surface activity, also adsorbs preferentially at the oil–water interface compared to  $\alpha$ <sub>s1</sub>-casein, and appears to displace  $\alpha$ <sub>s1</sub>-casein from the droplet surface (Dickinson and Stainsby, 1988).

In the food industry, individual caseins are generally not used to prepare emulsions because of the cost of pure fractions and their sparse availability. Instead, various types of caseinate, such as sodium caseinate, are widely used in the preparation of emulsion-type products. Caseinates are produced from skim milk by lowering the pH to 4.6, by adding either lactic or hydrochloric acid or microbial cultures to precipitate the casein, then resolubilising it with alkali or alkaline salts of sodium, potassium or calcium at neutral pH followed by spray drying (Mulvihill, 1989). Sodium caseinate comprises not only  $\alpha$ <sub>s1</sub>- and  $\beta$ -caseins but also  $\kappa$ - and  $\alpha$ <sub>s2</sub>-caseins and small quantities of lipids and

inorganic salts. The stability of an oil-in-water emulsion made with sodium caseinate largely depends on the composition of the adsorbed layer, the quantities of proteins and the conformation of the casein in the continuous phase. In sodium caseinate-stabilised oil-in-water emulsions, all forms of caseins ( $\alpha_{s1}$ -,  $\alpha_{s2}$ -,  $\beta$ - and  $\kappa$ -caseins) are adsorbed at the emulsion droplet surface (Robson and Dalgleish, 1987; Hunt and Dalgleish, 1994a; Srinivasan *et al.*, 1996), providing stability against coalescence and flocculation.

In contrast to pure caseins, the competitive adsorption of  $\beta$ -casein rather than the other casein fractions in sodium caseinate appears to be driven by the total protein content or the volume ratio of caseinate to oil in the emulsion. It has been shown that  $\beta$ -casein adsorbs preferentially at the interface only at lower caseinate concentrations (<2.0 %) and/or when the ratio of caseinate to oil is low, i.e., when the caseins are predicted to exist as monomers (Srinivasan *et al.*, 1996, 1999). However, at higher total caseinate concentrations (caseinate:oil ratio of >1:60),  $\alpha_{s1}$ -casein adsorbs preferentially and  $\beta$ -casein loses its competitive adsorption ability; this has been attributed to the self-aggregating tendency of  $\beta$ -casein to form micelles or to complex with other casein fractions, such as  $\alpha_{s1}$ -casein, via hydrophobic interactions (Lucey *et al.*, 2000). Furthermore, these aggregated complexes appear to have less emulsifying capability, as the hydrophobic areas are mutually blocked in the process of complex formation (Lorient *et al.*, 1989). Irrespective of the caseinate concentration,  $\kappa$ -casein from sodium caseinate has been found to be least adsorbed at droplet surfaces.

The creaming stability of sodium caseinate emulsions (20–30 % w/w oil) shows a complex dependence on the caseinate content. At lower caseinate concentrations, the emulsion is destabilised by bridging flocculation because there is insufficient protein to fully cover all the droplets in the emulsion. At an intermediate caseinate concentration of about 2.0 % w/w, the emulsion is stabilised against flocculation, coalescence and creaming for several weeks as the protein content is sufficient to cover the droplet surface. However,

when the caseinate concentration is increased to above 3.0 % w/w, unadsorbed caseinate gives rise to depletion flocculation (Dickinson and Golding, 1997; Srinivasan *et al.*, 2001). Further increasing the protein concentration above 6.0 % w/w results in a very high degree of depletion flocculation, leading to a strong emulsion droplet network, which is stable to creaming.

Interestingly, concentration-dependent depletion flocculation is not common in whey-protein-stabilised emulsions. It appears that depletion flocculation in sodium caseinate-stabilised emulsions is caused by the presence of casein aggregates (sub-micelles) formed from the self-assembly of casein molecules in the aqueous phase of the emulsion at concentrations above 2 % w/w. The addition of moderate amounts of calcium chloride to emulsions containing excess sodium caseinate has been shown to eliminate depletion flocculation and to improve the creaming stability (Ye and Singh, 2001). This effect can be attributed to an increase in the average size of the casein aggregates in the aqueous phase, resulting in a large increase in the molecular mass of the caseins (Dickinson *et al.*, 2001). In addition, there is a reduction in the concentration of unadsorbed caseinate.

### 5.2.1.2 Emulsions Stabilised by Whey Proteins

Whey proteins ( $\beta$ -lg,  $\alpha$ -la, bovine serum albumin, lactoferrin and immunoglobulins) are characterised by three-dimensional structures that are held together by disulphide bridges (Kinsella, 1984). They are soluble over a wide pH range. Whey proteins in general are highly susceptible to thermal denaturation above 70 °C because of their globular nature (Kinsella and Whitehead, 1989; Hunt and Dalgleish, 1995; Singh, 2005). The most important whey protein fractions include  $\beta$ -lg and  $\alpha$ -la, which account for ~70–80 % of the total whey protein and possess excellent emulsifying properties. These proteins adsorb on to oil-water interfaces and form stable emulsions, although the emulsions formed are slightly less stable than casein-stabilised emulsions under the same conditions (Hunt and Dalgleish, 1994a; Dalgleish, 1995).

Structurally,  $\beta$ -lg is a compact, folded, globular protein, containing 162 amino acids along with two disulphide bonds and one free thiol group (Swaisgood, 1982). The three-dimensional structure of  $\beta$ -lg comprises nine strands of anti-parallel  $\beta$ -sheets, joined together into a conical  $\beta$ -hydrophobic barrel unit, and a flanking three-turn  $\alpha$ -helix (Sawyer *et al.*, 1985; Papiz *et al.*, 1986; Oliveira *et al.*, 2001). Under ambient temperatures ( $\sim 25$  °C) and at neutral pH,  $\beta$ -lg exists mostly as a non-covalently linked dimer with a molecular weight of  $\sim 36$  kDa (McKenzie and Sawyer, 1967; Ziegler and Foegeding, 1990). Because of its amphiphilic nature,  $\beta$ -lg shows good emulsifying properties by adsorbing at the interfacial layer, where it partially unfolds and forms a continuous interfacial film through intermolecular  $\beta$ -pleated sheet interactions. The exposed reactive free thiol groups at the interface lead to slow polymerisation of the adsorbed protein *via* sulphhydryl–disulphide interchange mechanisms (Dickinson and Matsumura, 1991; McClements *et al.*, 1993; Lefèvre and Subirade, 2003).  $\beta$ -Lg has been the most extensively studied of all food proteins for its role in stabilising oil-in-water emulsions, because of its well-defined structure and properties (McKenzie, 1971; Kinsella and Whitehead, 1989).

$\alpha$ -La, another major whey protein, is a globular, calcium metallo-protein, which is stabilised by four intra-chain disulphide bonds (Swaisgood, 1982). In contrast to  $\beta$ -lg,  $\alpha$ -la does not contain a free thiol group. These two whey proteins also differ in their amino acid composition, with  $\beta$ -lg having more proline residues than  $\alpha$ -la (eight and two, respectively), resulting in higher hydrophobicity, and  $\alpha$ -la having more cysteine residues than  $\beta$ -lg (eight and five, respectively), resulting in more internal disulphide bridges (Ng-Kwai-Hang, 2003).  $\alpha$ -La denatures at a relatively low temperature ( $\sim 66$  °C) compared with  $\beta$ -lg ( $\sim 73$  °C) but does not aggregate because of the absence of a free thiol group (Dalgleish *et al.*, 1997; Schokker *et al.*, 2000; Considine *et al.*, 2007). Native  $\alpha$ -la has good emulsifying capabilities, but has poor gelation properties.

Dickinson *et al.* (1989) studied competitive adsorption at the oil–water interface in emulsions

stabilised by  $\beta$ -lg and  $\alpha$ -la model systems using classical exchange measurements. They showed that competitive displacement between  $\beta$ -lg and  $\alpha$ -la was rather slow and very limited, in contrast to that between caseins, which was shown to be much faster and essentially reversible in character (Dickinson *et al.*, 1988). For instance,  $\beta$ -lg was not displaced from the interface in  $\beta$ -lg-stabilised emulsions when  $\alpha$ -la was added at a level of 1:1 % w/w until a prolonged time period of 32 h. On increasing the proportion of  $\alpha$ -la to  $\beta$ -lg (to 10:1), only 15 % of the  $\beta$ -lg was displaced to the serum phase. In contrast, on increasing the proportion of  $\beta$ -lg in an  $\alpha$ -la-stabilised emulsion (to 10:1), nearly 30 % of the  $\alpha$ -la was displaced. This suggests that, for  $\beta$ -lg and  $\alpha$ -la model systems, the interfacial adsorption is relatively irreversible (in comparison with casein systems) and the protein that is initially introduced to the interface will probably dominate at the interface, irrespective of its relative surface activity (Dickinson *et al.*, 1988, 1989). Both these whey proteins are highly structured globular proteins and undergo conformational changes upon adsorption at an interface (Fang and Dalgleish, 1997, 1998). However, of the two globular proteins,  $\beta$ -lg is even more difficult to displace than  $\alpha$ -la. This can be attributed to the sulphhydryl–disulphide interchange reactions that occur in  $\beta$ -lg-stabilised emulsions, but not in pure  $\alpha$ -la-stabilised emulsions because of the absence of a free thiol group in  $\alpha$ -la (Dickinson and Matsumura, 1991; Monahan *et al.*, 1995; Damodaran and Anand, 1997). Upon adsorption,  $\beta$ -lg undergoes partial unfolding, stretches and becomes densely packed at the interface, enabling the free thiol group on each molecule of  $\beta$ -lg to link *via* intermolecular covalent disulphide bridges at the droplet interface. As the extent of polymerisation increases during storage, the interfacial film continues to strengthen irreversibly with time, resulting in high surface rheology (Dickinson, 1989; Damodaran and Anand, 1997; Dalgleish, 2004). Thus, its displacement from the interface by  $\alpha$ -la becomes highly unlikely.

Whey proteins also contain low levels of lactoferrin, a glycoprotein of molecular weight  $\sim 80$  kDa, which has about 700 amino acid

residues and is well known for its iron-binding capacity (Baker and Baker, 2005). Unlike most milk proteins, which have isoelectric points ( $pI$ s) ranging from 4.5 to 5.5, lactoferrin has a relatively high  $pI$  of  $\sim 8.0$  and thus has the unique property of possessing a high positive surface charge at neutral pH (almost +50 mV) (Ye and Singh, 2006a). This high positive charge density of lactoferrin has been predicted to allow the formation of cationic emulsion droplets over wide pH ranges. The adsorption behaviour of lactoferrin in oil-in-water emulsions was explored by Ye and Singh (2006a). Similar to other milk proteins such as caseinates and  $\beta$ -lg, lactoferrin adsorbed to the oil-water interface, producing stable emulsion droplets with a net positive charge.

In contrast to caseinates and  $\beta$ -lg, emulsions stabilised by lactoferrin were stable over a wide range of pH from 7.0 to 3.0. The droplet sizes of lactoferrin emulsions were reported to be very similar to those of  $\beta$ -lg emulsions prepared under the same conditions of pH, oil:protein ratio and homogenisation pressure. However, lactoferrin-stabilised emulsions had a comparatively higher surface coverage because of the higher molecular weight of lactoferrin. As lactoferrin in solution is highly positively charged, lactoferrin has been shown to exhibit electrostatic complexation with anionic  $\beta$ -lg at neutral pH (Wahlgren *et al.*, 1993). Using this theory, multi-layered oil-in-water emulsions were produced from the electrostatic interactions of oppositely charged milk proteins, i.e., lactoferrin and  $\beta$ -lg, at neutral pH at the droplet surface, resulting in stable emulsion droplets with thick multi-layered interfacial layers and greater amounts of protein adsorbed at the oil-water interface (Ye and Singh, 2007). The primary emulsion, containing either cationic (lactoferrin-coated) or anionic ( $\beta$ -lg-coated) droplets, was produced initially. The secondary emulsion was then formed by adding either lactoferrin or  $\beta$ -lg solution to the primary emulsion based on opposite charges. Interestingly, the overall charge of emulsion droplets stabilised using the binary protein mixtures was close to zero at some concentrations. However, the multi-layered emulsions were protected against floccu-

lation because of the strong steric effects of the dense interfacial film at the droplet surface.

In addition to pure proteins, the commercially available forms of whey protein that are widely used as emulsifiers in food industries are WPCs (comprising 25–80 % protein) and WPIs (comprising >90 % protein). These concentrated forms of whey protein are produced by ultrafiltration, diafiltration or ion exchange followed by drying steps to obtain protein levels of  $\sim 80$ –95 % (Morr and Ha, 1993; Mulvihill and Ennis, 2003). Both WPC and WPI are widely used in processed food applications because of their water-binding, gelling, foaming and surface-active properties (Mulvihill and Ennis, 2003; Singh, 2005). Processing treatments during the manufacture of WPC and WPI tend to denature some of the whey proteins because globular proteins are highly susceptible to conformational changes (denaturation) and aggregation when the pH, ionic strength or temperature is changed and this generally affects their functional properties.

It is interesting to note here that, when WPC or WPI is used to stabilise emulsions, similar in quantities to pure fractions of the individual proteins, there is little preferential adsorption of  $\beta$ -lg over  $\alpha$ -la or *vice versa* at the droplet surface regardless of the proportion of protein to oil (Euston *et al.*, 1996; Ye and Singh, 2000, 2006b). Ye and Singh (2000) showed that in WPC-stabilised emulsions (30 % w/w oil, 0.5 % w/w WPC), the proportions of adsorbed  $\alpha$ -la and  $\beta$ -lg were  $\sim 18$  % or  $\sim 82$  %, respectively, compared with those in the original WPC solution ( $\alpha$ -la  $\sim 25$  %;  $\beta$ -lg  $\sim 75$  %), suggesting that  $\beta$ -lg was adsorbed slightly in preference to  $\alpha$ -la under these conditions.

Preferential adsorption of  $\alpha$ -la and  $\beta$ -lg was more clearly demonstrated when WPI-stabilised emulsions were subjected to pH changes. Shimizu *et al.* (1981) showed that the total protein adsorption was highest at pH 5 in whey-protein-stabilised emulsions, possibly because of the dense network formed at a pH close to the  $pI$ . Interestingly, they also observed that the adsorption of  $\beta$ -lg decreased as a function of decreasing pH from 9 to 3, whereas the adsorption of  $\alpha$ -la



increased in the same pH range. This decrease in the adsorption of  $\beta$ -lg was attributed to pH-dependent conformational changes of tertiary and quaternary structures (Shimizu *et al.*, 1985; Hunt and Dalgleish, 1994b).

In emulsions formed with both caseinate and whey protein, Hunt and Dalgleish (1994b) reported that the preferential adsorption depended on the protein concentration. This was further validated by a study by Ye (2008), which suggested that in emulsions made with mixtures of sodium caseinate and WPC, caseins adsorb preferentially at the oil–water interface at high protein concentrations whereas whey proteins adsorb preferentially at low protein concentrations (<3 %).

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### 5.3 Formation and Stability of Protein-Stabilised Foams

Because of their surface-active properties, proteins are known to contribute to the formation of foams and to the physical stability of foam-based food formulations such as whipped cream, mousse and ice cream. Proteins are less effective than low molecular weight surfactants in reducing the air–water interfacial tension but they form an interfacial film that exhibits viscoelastic properties and that enables the foam to resist destabilisation (Murray and Ettelaie, 2004).

In general, foams can be generated by two mechanical methods, i.e., bubbling and stirring (Bikerman, 1973; Prud'homme and Khan, 1996; Exerowa and Kruglyakov, 1998; Weaire and Hutzler, 1999). In the bubbling method, the foam is produced by bubbling gas or air through the aqueous phase containing the foaming agent (protein, surfactants, etc.) using a single capillary, a set of capillaries or a porous plate. The size of the foam bubbles thus generated depends on the pore size of the capillaries or the porous plate, the properties of the surfactant solution, such as dynamic surface tension, surface elasticity and bulk viscosity, and the conditions of foam formation, i.e., rate of gas flow, temperature, pressure, etc. In contrast, the stirring method involves

mixing the gaseous phase and the aqueous phase, which contains the foaming agent, mechanically using a stirrer or shaker or allowing simultaneous flow of the gas and the liquid in a tube.

The foaming capacity (or foamability) is defined either as the volume of foam generated under fixed conditions of temperature and intensity of mechanical agitation or by the time needed to generate a certain volume of foam. Both the foam formation process and foamability depend largely on the physicochemical properties of the stabilising substances. From a protein perspective, foam formation depends on the rate at which the proteins can transfer to the air–liquid interface and the stability of the foam generated depends on the ability of the adsorbed proteins to form a cohesive viscoelastic film *via* intermolecular bonds (Damodaran, 1997).

Similar to emulsions, foams also require high energy and subsequent thermodynamic instability makes them liable to separate into their two original phases over time. Thus, foams are also kinetically stable colloidal dispersions and undergo destabilisation over different time scales mainly by three mechanisms, i.e., liquid drainage, bubble coalescence and disproportionation of individual bubbles (Ivanov, 1988; Prud'homme and Khan, 1996; Exerowa and Kruglyakov, 1998; Weaire and Hutzler, 1999; Pereira *et al.*, 2003; Denkov, 2004; Murray and Ettelaie, 2004; Saint-Jalmes *et al.*, 2005; Denkov and Marinova, 2006). Drainage is driven mainly by gravity and involves a gradual rise of bubbles through the foam mass, while the aqueous phase drains through the lamellae and the plateau borders between the foam bubbles. In contrast, bubble coalescence involves thinning and rupturing of the isolated liquid interfacial films separating two neighbouring bubbles. Foam bubbles are stabilised against coalescence by the generation of strong colloidal forces that act between the film surfaces and the adsorption of surface-active molecules such as proteins to form a dense film. The third type of foam destabilisation is disproportionation, which involves bubble coarsening because of the diffusion of gas through the foam films, from the smaller bubbles to the larger bubbles.

### 5.3.1 Aspects of Foams Stabilised by Milk Proteins

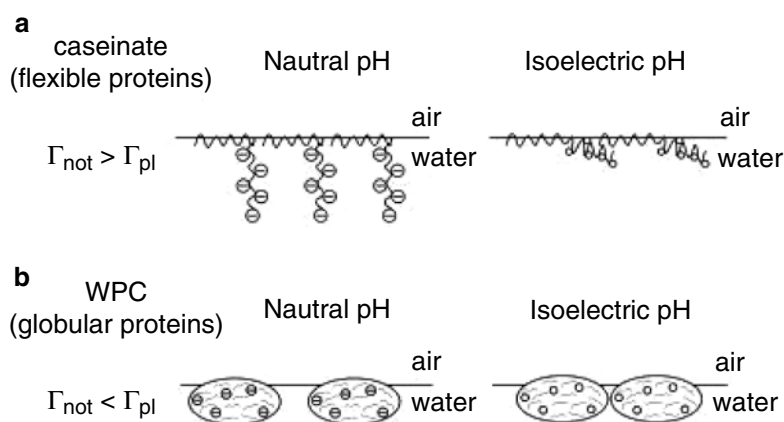
Milk proteins are widely known for their foam-forming and foam-stabilising properties (Anderson and Brooker, 1988). Despite their high molecular weight and (in the case of whey proteins) their complex secondary and tertiary structures, milk proteins are able to diffuse from the aqueous phase and adsorb at the air–water interface during foam formation because of the compatibility of the hydrophobic regions of their structure with the hydrophobic nature of the gaseous phase. The different molecular structures of flexible and globular milk proteins generally lead to different structures of the adsorbed layers at the air–water interface. Milk proteins are well known to alter their charge and surface activity with pH and accordingly their foamability is also affected.

Marinova *et al.* (2009) compared the foaming behaviours of caseins and WPC as a function of pH and over a range of ionic strengths. As expected, both caseins and whey proteins showed a rapid increase in foam volume as a function of protein concentration until a plateau value was reached; there was a corresponding increase in protein adsorption at the air–water interface and a decrease in dynamic surface tension with increasing protein concentration. Both types of protein led to the stabilisation of foams against bubble coalescence. However, there were significant differences between foams stabilised with WPC and

foams stabilised with caseins. For example, a relatively lower concentration of sodium caseinate (>0.3 % w/w) than of WPC (>1 % w/w) was needed to generate stable foam. Also, the volume of foam generated by WPC (10–11 mL) was nearly half of that generated by sodium caseinate (22–23 mL). When the pH was varied, WPC had maximum foamability near the *pI* whereas sodium caseinate had minimum foamability near the *pI*.

Marinova *et al.* (2009) explained these differences based on the molecular structures of the adsorbed milk protein layers and the different aggregation behaviours, as schematically presented in Fig. 5.2. At the natural pH (pH 6.5–6.8), i.e., far from the *pI* (pH 4.6), flexible casein molecules allowed the formation of denser adsorbed layers comprising hydrophobic amino acid residues in a “loop”-like configuration and the hydrophilic chain extending farther away as a “tail” within the serum phase, thus ensuring better foam stabilisation (Dickinson *et al.*, 1993) (Fig. 5.2). In contrast, WPC could not anchor strongly at the surface of the foam bubbles because of its intact globular conformation, at least initially (Gurkov *et al.*, 2003; Freer *et al.*, 2004). The minimum foamability of sodium caseinate near the *pI* can be explained on the basis of the unavailability of sufficient quantities of casein for adsorption at the surfaces because of the precipitation of casein at the *pI*. However, in the case of whey proteins at pH 4.5, slightly positively charged  $\beta$ -lg (*pI* 5.1), and slightly

**Fig. 5.2** Schematic presentation of sodium caseinate (a) and WPC (b) molecules adsorbed at the water–air interface (Marinova *et al.*, 2009; reproduced with the permission of Elsevier Inc.)



negatively charged  $\alpha$ -la ( $pI$  4.5), may have interacted and strengthened the compaction of the molecules at the air–water interface *via* electrostatic interactions and thus allowed stronger interfacial films, ensuring better bubble coverage and thus high foamability. This is in line with the results obtained by other authors; it was found that, even in mixed protein systems, globular whey proteins, such as  $\beta$ -lg, and flexible random coil caseins, such as  $\beta$ -caseins, typically have different adsorption rates and also different foaming abilities because of their different abilities to adapt their conformation at the air–serum interface (Martin *et al.*, 2002; Ridout *et al.*, 2004).

Interestingly, the foaming properties of  $\beta$ -lg and WPI can be improved by increasing the ionic strength in the range 0–0.1 M NaCl at pH levels above or below their  $pI$ , as evidenced by dynamic surface tension measurements (Davis *et al.*, 2004; Zhang *et al.*, 2004). This is because of the charge screening effect of the salt, which allows the proteins to be compacted at the interface, resulting in increased protein adsorption, and the formation of a strong viscoelastic network, which provides stabilisation of the foam against bubble coalescence.

Furthermore, certain chemical (Enomoto *et al.*, 2007; Wooster and Augustin, 2007), physical (Phillips *et al.*, 1990; Yang *et al.*, 2001) or enzymatic (Davis *et al.*, 2005) treatments are also known to improve the foaming properties of whey proteins. These treatments generally influence the conformation of whey proteins, especially  $\beta$ -lg, by modifying the protein structure, and thus alter the kinetics of protein adsorption at the interface, the time needed for the whey protein to rearrange upon adsorption at the interface and the ability to interact with other interfacial proteins, in the case of mixed systems.

The effects of heat treatment on the foaming properties of  $\beta$ -lg have been studied because heat treatment is commonly used in food processing to make many food foams. When native  $\beta$ -lg was compared with thermally treated  $\beta$ -lg, it was demonstrated that heated  $\beta$ -lg had higher surface hydrophobicity, adsorbed at a much faster rate at the air–water interface and had better foaming properties with respect to the initial rheology of

the interfacial film (Phillips *et al.*, 1995; Kim *et al.*, 2005; Croguennec *et al.*, 2006). It has been shown that  $\beta$ -lg aggregates produced by heat treatment of  $\beta$ -lg solution at 85 °C for 15 min can significantly improve the foaming properties (Schmitt *et al.*, 2007; Unterhaslberger *et al.*, 2007; Rullier *et al.*, 2008). Moro *et al.*, (2011) showed that preheating a 5.5 % (w/v)  $\beta$ -lg solution at 85 °C for 3 min generated a considerable change in its aggregation profile, producing non-native monomers (51 %) and dimers (33 %) and trimers (16 %). Because of the formation of these polymeric aggregates, the surface hydrophobicity was increased dramatically, which in turn improved the foamability; the foamability was of the order of ~800 % higher than that of the corresponding foam formed with unheated  $\beta$ -lg. This greater foam stability against disproportionation or collapse was attributed to an increase in the viscosity of the protein solution because of the presence of aggregates, which slowed the rate of liquid drainage because of compaction of the interfacial film.

As well as the individual proteins, the interactions between  $\beta$ -lg and  $\alpha$ -la during the thermal treatment of WPI have been found to play a significant role in the foaming properties and the stability of foams. Zhu and Damodaran (1994) showed that the heat treatment of WPI at neutral pH generated aggregates that improved foamability (at a monomeric:polymeric ratio of 60:40) or foam stability (at a monomeric:polymeric ratio of 40:60). Davis and Foegeding (2004) further confirmed these findings by showing similar foam stability improvements even when native whey protein was mixed with whey protein polymers generated by heat treatment at a similar monomeric:polymeric ratio of 40:60. This can be explained on the basis of rapid movement of monomeric whey protein at the interface upon foaming to decrease the surface tension followed by the formation of a viscoelastic interfacial network (mainly driven by disulphide bond formation and hydrophobic interactions) at the interface by the soluble aggregates. Nicorescu *et al.* (2009) showed that an optimal heat treatment of 2 % w/v WPI at 80 °C at ionic strength of 50 mM NaCl and at neutral pH was effective in obtaining an

improved firmness of the interfacial films because of the simultaneous formation of a cohesive network of protein aggregates (the generation of approximately 10 % soluble whey protein aggregates) at the interface and throughout the foam lamellae, which enhanced stability against liquid drainage. However, at temperatures of 80–100 °C, the generation of more than 50 % soluble aggregates caused weakening of the interfacial network because the presence of a large number of heavy clusters of aggregates, which behaved as a “solid”, led to more rapid drainage.

Our understanding of the role of the molecular structure and the processing of milk proteins on the formation and stability of foams has advanced significantly over recent years. For the future, it is crucial to understand how factors such as protein–protein interactions, interactions with other ingredients such as sugar, the emulsifiers in food foams and the interfacial composition of mixed interfaces contribute to foam stability holistically and the generation of “novel” foam properties.

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#### 5.4 Interactions of Milk-Protein-Stabilised Emulsions Under Physiological Conditions

For the last decade or so, the effects of processing (e.g., heat, high pressure and shear) on the properties of food emulsions (e.g., viscosity, droplet size distribution and phase stability) have been studied extensively. Interestingly, much of this work has involved the use of milk proteins and has focused on understanding the functionality of milk proteins in stabilising emulsions and exploiting their unique properties to produce novel structures and sensory perceptions. In contrast, efforts to elucidate the fate of milk-protein-stabilised emulsions following consumption during *in vitro* gastrointestinal digestion are relatively recent and are now generating a great deal of interest.

When a complex food emulsion is consumed, the properties of each of its components, together with its interactions with physiological factors, including mucin, pepsin, lipase, gastric mucins and phospholipids, should be considered. These

biochemical agents may interact with the emulsion and result in modification of the adsorbed protein layers and the droplet characteristics, affecting the stability of the emulsion and the digestibility of its components (Fig. 5.1). Although emulsions can be carefully manipulated using various physical or chemical processes before they are consumed, understanding of the interactions during transit through the gastrointestinal tract is of great importance to gain insights into the post-consumption structural and physicochemical changes in these emulsions. As this chapter is concerned with milk proteins, we briefly describe the behaviour of milk-protein-stabilised emulsions in physiological environments and discuss how the milk protein-based interfacial layer influences the various steps involved in the digestibility of emulsified lipids.

When a milk-protein-stabilised food emulsion is consumed, it resides for a short period in the mouth and is exposed to a wide range of biochemical conditions, such as dilution effects, because of mixing with saliva, and access to salivary enzymes such as amylases, biopolymers such as mucins and different electrolytes in the saliva, as well as physicochemical conditions, such as moderate changes in pH and temperature (to around 37 °C) and shear forces between the tongue and the oral palate (Malone *et al.*, 2003a, b; de Wijk *et al.*, 2004; de Wijk and Prinz, 2005; Vingerhoeds *et al.*, 2005). Interestingly, there is some evidence to show that the behaviour of milk-protein-stabilised emulsions in the mouth is largely driven by the non-covalent interactions of salivary components with the adsorbed milk protein layer at the oil droplet surface (van Aken *et al.*, 2005; Sarkar *et al.*, 2009a; Vingerhoeds *et al.*, 2009). Emulsions formed with WPI, sodium caseinate and lactoferrin showed flocculation of the droplets when mixed with human saliva. This flocculation was predominantly driven by the highly glycosylated negatively-charged mucin present in human saliva. The emulsion flocculation in the presence of saliva was considered to be regulated by depletion forces, van der Waals' forces and/or electrostatic interactions between emulsion droplets and salivary proteins, and was largely dependent on the

initial charge of the milk-protein-stabilised emulsion droplets (Silletti *et al.*, 2007; Sarkar *et al.*, 2009a).

Sarkar *et al.* (2009a) investigated the behaviour of negatively and positively charged oil-in-water emulsions stabilised by milk proteins in the presence of artificial human saliva. At neutral pH, negatively-charged  $\beta$ -lg-stabilised emulsions underwent some degree of depletion flocculation because of strong repulsive forces with anionic mucin. In contrast, positively charged lactoferrin-stabilised emulsions interacted with mucin *via* electrostatic interactions, which led to bridging-type flocculation under certain conditions. These kinds of emulsion–saliva electrostatic interactions might occur when emulsions are consumed in real situations and could result in different sensory and textural perceptions *in vivo*.

After oral transit, emulsions are swallowed, are subjected to shear effects in the oesophagus and are finally exposed to a highly acidic pH (typically between 1 and 3) and shear forces because of peristaltic movements of the stomach (Weisbrodt, 2001; Kalantzi *et al.*, 2006). During the gastric phase, emulsions are exposed to digestive juices, containing proteolytic (pepsin) and lipolytic (gastric lipase) enzymes, mucins and salts. It is obvious that milk-protein-stabilised emulsions would undergo major changes in the stomach because of the possible action of pepsin on the protein layer at the interface, the effects of low pH and ionic strength on the droplet charge and the interactions of gastric mucin with interfacial protein. For example, flexible caseins are highly susceptible to hydrolysis by pepsin in aqueous solutions (Guo *et al.*, 1995). However, globular whey proteins, particularly  $\beta$ -lg, are known to be largely resistant to peptic hydrolysis in their native state (Schmidt and Poll, 1991).

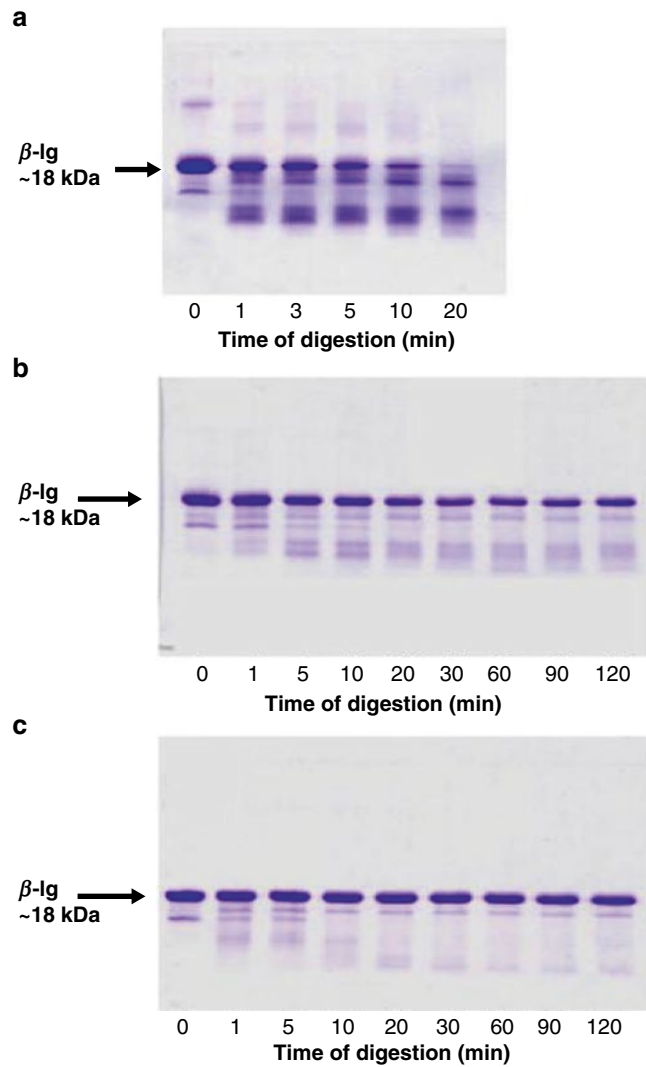
Studies by Macierzanka *et al.* (2009) and Sarkar *et al.* (2009b), which are in agreement with other reports, suggest that  $\beta$ -lg becomes highly prone to hydrolysis by pepsin when present as the interfacial layer in an emulsion. This is due to a conformational change of the  $\beta$ -lg molecules upon adsorption at the droplet surface, which opens up the peptic cleavage sites for enzymatic attack by pepsin. Interestingly, in

milk-protein-stabilised emulsions, generally not all the protein is present at the droplet surface, i.e., in the adsorbed state; a considerable proportion remains in the aqueous phase. As the adsorbed and unadsorbed proteins are likely to exist in different conformational states, their susceptibilities to pepsin could be different under gastric conditions; the rate of hydrolysis of  $\beta$ -lg, from sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS-PAGE) gels, is shown in Fig. 5.3. Sarkar *et al.* (2009b) showed that adsorbed  $\beta$ -lg was more susceptible to pepsin hydrolysis (85 % decrease in the intact protein) than unadsorbed protein (about 50 % decrease in the intact protein). In contrast,  $\beta$ -lg in solution in its native state was largely resistant to pepsin digestion (about 20 % decrease in the intact protein) under the same simulated gastric conditions. This suggested that the conformation of unadsorbed  $\beta$ -lg is significantly altered during emulsion formation, possibly as a result of the high turbulence during homogenisation, as mentioned in the previous section.

The hydrolysis of the adsorbed layers by pepsin also results in a loss of positive charge on the droplet surface as well as a reduction in the thickness of the adsorbed layer (Sarkar *et al.*, 2009b). The peptides that remain at the interface are unable to provide sufficient electrostatic and steric stabilisation effects. As a result, these emulsions with hydrolysed interfaces are highly susceptible to flocculation and coalescence (Fig. 5.1). Such a phenomenon has been demonstrated for lactoferrin- and  $\beta$ -lg-stabilised emulsions, which undergo flocculation followed by some degree of coalescence on exposure to simulated gastric conditions (Sarkar *et al.*, 2009b; Sarkar, 2010).

In addition to the important effect of gastric enzymes, highly glycosylated mucin, which forms a self-associated gel-like structure at gastric pH and protects the stomach from digesting itself, also has an important role in its interaction with milk-protein-stabilised emulsions (Bansil and Turner, 2006). The work of Sarkar *et al.* (2010a) suggests that the addition of a low level of soluble mucin promotes the flocculation of

**Fig. 5.3** Sodium dodecyl sulphate polyacrylamide gel electrophoretograms of  $\beta$ -lg emulsions: (a) cream phase; (b) continuous phase; (c) native  $\beta$ -lg solutions [containing 0.36 %  $\beta$ -lg (the same as the concentration of  $\beta$ -lg in the continuous phase of the emulsion)] after mixing with simulated gastric fluid as a function of incubation time (Sarkar *et al.*, 2009b: reproduced with the permission of Elsevier Inc.)



$\beta$ -lg-stabilised emulsions, possibly through a bridging mechanism, but does not significantly affect the action of pepsin on the adsorbed  $\beta$ -lg layer. Overall, when they reach the gastric tract, emulsions stabilised by milk proteins show a significant level of instability, which is predominantly driven by the proteolytic effect of pepsin.

Following transit through the gastric tract, emulsions enter the human intestinal tract, which is a very complex environment, as it contains various salts, pancreatic enzymes, coenzymes, bile salts, phospholipids, remnants of oral and gastric digestion and various microbial species at neutral to alkaline pH (6.0–7.5) (McClements

*et al.*, 2009; Singh *et al.*, 2009; Singh and Ye, 2013). As milk protein might be susceptible to the actions of pancreatic enzymes such as trypsin and chymotrypsin, both the adsorbed proteins/peptides and the unadsorbed proteins/peptides might be further hydrolysed into smaller peptides and amino acids. Over the last few decades, several studies have reported the effects of the hydrolysis of milk-protein-stabilised oil-in-water emulsions by trypsin on their physical stability (Kaminogawa *et al.*, 1987; Leaver and Dalglish, 1990). As expected, extensive hydrolysis of adsorbed protein layers results in rupturing of the interfacial layer, leading to the coalescence of

emulsion droplets and subsequent oiling-off (Agboola and Dalgleish, 1996). However, the behaviour of milk protein-based emulsions in a more complex intestinal environment, simulating intestinal conditions, has not been reported until very recently, by Sarkar *et al.* (2010b, c).

In addition to intestinal enzymes, the interaction of surface-active bile salts with the adsorbed protein layer are important. Bile salts, which originate from the liver *via* the gall bladder, consist mainly of sodium salts of taurocholic, taurodeoxycholic, taurochenodeoxycholic, glycocholic and glycodeoxycholic acids. These surface-active compounds displace the adsorbed proteins/peptides from the surface of emulsion droplets because of their relatively higher surface activity, thus promoting the accessibility of the active site of lipase to the hydrophobic lipid core (Wickham *et al.*, 1998; Fave *et al.*, 2004; Mun *et al.*, 2007; Sarkar *et al.*, 2010b). The nature and, in particular, the charge of the adsorbed milk protein layer seem to drive the preferential adsorption of bile salts and the subsequent displacement of the protein layer (Singh *et al.*, 2009). For instance, bile salts have been shown to displace whey proteins more readily than caseinates from the interface of emulsion droplets during storage (Mun *et al.*, 2007). In emulsions stabilised by negatively charged  $\beta$ -Ig, displacement of protein was observed even at the lowest concentration of bile salts (Sarkar *et al.*, 2010b), but the bile salts did not displace positively-charged lactoferrin from the emulsion droplets. The bile salts appeared to bind to the adsorbed lactoferrin layer *via* an electrostatic mechanism.

Upon entering the intestine, pancreatic lipase adsorbs to the droplet interface, usually *via* complexation with colipase and/or bile salts (Bauer *et al.*, 2005). Colipase is a short polypeptide with a molecular weight of 10 kDa, which forms a stoichiometric complex with lipase in a 1:1 w/w ratio, enabling the water-soluble pancreatic lipase to attach firmly to the hydrophobic lipid core at the oil droplet surface (Erlanson-Albertsson, 1992). Bile salts may either facilitate or inhibit the activity of pancreatic lipase depending on their concentration (Lowe, 2002; Bauer *et al.*, 2005). At low concentrations, bile salts promote

pancreatic lipase activity, mainly by allowing the adsorption of lipase to the oil–water interface (Gargouri *et al.*, 1983; Mun *et al.*, 2007) as well as solubilising and removing the inhibitory reaction products from the oil–water interface. However, at high concentrations, bile salts generally compete with lipases for adherence to the droplet surface, thus inhibiting the point of contact between the hydrophobic lipid core and the lipase (Gargouri *et al.*, 1983) and retarding lipase activity. Pancreatic lipase cleaves triacylglycerols to form 2-monoacylglycerols and fatty acids; some of these digestion products are surface active and could potentially displace the initial adsorbed material from the droplet surface (McClements *et al.*, 2009; Singh *et al.*, 2009; Sarkar *et al.*, 2010c).

Most studies of lipid digestion in milk protein-based emulsions have used *in vitro* intestinal models containing pancreatic lipase and bile salts. The extent of lipid hydrolysis was found to be similar in caseinate- and whey protein-stabilised emulsions, although the oil droplets in the whey protein-stabilised emulsions were less stable (Mun *et al.*, 2007). Our study showed that lactoferrin- and  $\beta$ -Ig-stabilised emulsions underwent a significant degree of coalescence on the addition of physiological concentrations of pancreatin and bile salts (Sarkar *et al.*, 2010c). For both emulsions, destabilisation in simulated intestinal fluid was largely attributed to the lipolysis of the hydrophobic lipid core by the lipase fractions of the pancreatin as well as the proteolysis of the adsorbed protein layer by the trypsin or other proteolytic fractions present in pancreatin.

In addition to pure whey protein systems, studies on the *in vitro* digestion of WPI emulsions showed that they did not undergo pronounced structural changes during simulated gastric digestion although the  $\alpha$ -la and a portion of the  $\beta$ -Ig adsorbed at the interface were hydrolysed by pepsin. However, during the subsequent intestinal phase of digestion, the partially digested WPI-stabilised emulsion droplets underwent coalescence (Li *et al.*, 2013). In contrast, in the case of sodium caseinate-stabilised emulsions, the droplets underwent droplet flocculation

with some degree of coalescence during the gastric phase itself (Li *et al.*, 2012). Because of its open flexible structure, casein was easily hydrolysed by pepsin, which in turn led to coalescence of droplets under gastric conditions. Overall, at both sodium caseinate- and whey-protein-stabilised interfaces, digestion in the gastric fluid containing pepsin apparently accelerated the coalescence of the emulsion droplets during subsequent exposure to intestinal fluid containing pancreatic lipase. However, for both milk-protein-stabilised emulsions, the rate and the extent of lipid digestion in the intestinal environment were found not to be influenced by the previous structural changes that may have occurred during the gastric phase.

Recently, a number of researchers (Hur *et al.*, 2009; Li *et al.*, 2012; Kenmogne-Domguia *et al.*, 2013) have studied the effect of proteolysis of the adsorbed milk protein layer and subsequent physicochemical changes of the emulsion droplets during the entire physiological transit. Emulsions stabilised by casein and bovine serum albumin were treated under *in vitro* gastric conditions at various pH values and at various concentrations of pepsin, as a function of incubation time. The adsorbed protein was hydrolysed to different degrees by pepsin, which resulted in droplet flocculation and coalescence in the emulsions. When these gastric-treated samples were exposed to *in vitro* intestinal digestion, the results showed that gastric conditions could modify the kinetics of lipolysis, but had limited impact on the final extent of lipolysis in the intestinal step of digestion.

Studies in our laboratory on lipid droplets initially coated with lactoferrin (cationic) and  $\beta$ -lg (anionic) and sequentially treated with simulated oral, gastric and intestinal fluids in an *in vitro* physiological model further validated that milk-protein-stabilised interfaces, irrespective of their high original electrostatic charges, offer little protection to the droplets against pepsin- and pancreatin-induced destabilisation and thus cannot help, individually, in controlling the rate and the extent of lipid digestion (Sarkar, 2010; Singh and Sarkar, 2011). The mechanism of destabilisa-

tion and re-stabilisation in intestinal fluid following pre-processing in oral and gastric fluids could not be interpreted reliably because of interference from one or more of the factors in the chemically complex, simulated physiological media used. There is a clear need for further research in this area to have a better understanding of the different competitive displacement mechanisms and hydrolytic reactions occurring in the intestine and to characterise the final state of the droplets and the products of lipid hydrolysis. More complete *in vitro* digestion models to simulate various physiological processes occurring in the mouth, stomach and small intestine need to be developed and then validated by *in vivo* and clinical studies. Further research in this area is likely to lead to new knowledge that can be used in designing food matrices by manipulating milk proteins effectively at the oil–water interface during physiological transit for controlled lipid delivery applications.

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## 5.5 Conclusions

Milk proteins in both soluble and dispersed forms have excellent surface-active, foaming and emulsion-stabilising properties. Differences in structure, flexibility and state of aggregation of the different milk proteins give rise to differences in their emulsion- and foam-stabilising properties. These attributes of milk proteins have been exploited to manufacture various prepared foods. For decades, research has been performed on oil-in-water emulsions and foams using purified or simple mixtures of caseins and whey proteins to manufacture a wide range of products and there is now a great deal of understanding on the conformation of proteins at oil–water interfaces, competitive exchange reactions and factors controlling the rheology and stability of emulsions under different environmental conditions (temperature, pH and ionic conditions). However, much less is known about the further processing of emulsions after they have been consumed, i.e., during oral processing in the mouth and during the digestion processes. This area of research



needs to be developed further before the interactions between milk-protein-stabilised emulsions and physiological factors can be carefully utilised to develop novel products with sensory and/or health benefits.

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