Effects of heating on the secondary structure of proteins in milk powders using mid-infrared spectroscopy

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ABSTRACT

Milk powder is an important source of protein for adults and children. Protein is very sensitive to heat, which may influence people's usage of nutrients in milk powder. In this study, we describe the temperature-induced secondary structure of protein in milk powders. In this study, whole milk powder containing 24% protein and infant formula containing 11% protein were heated from 25 to 100°C. Attenuated total reflectance (ATR) spectra in the mid-infrared range 400–4,000 cm⁻¹ were used to evaluate the heat effect on the secondary structure of protein in these 2 milk powders. The spectral changes as a function of temperature were maintained by difference spectra, second-derivative spectra and Gauss curve-fitted spectra. The secondary structures of protein in the whole milk powder began to change at 70°C and in the infant formula at 50°C. The β -sheet and β -turn structures in the whole milk powder both decreased in the range of 70 to 85°C, whereas α -helix structures increased. The loss of β -sheet and β -turn may contribute to the formation of α -helix in the whole milk powder. In infant formula powder, the β -sheet structure showed a decrease and then increase, whereas the β -turn structure showed an increase and then decrease in the range of 50 to 75°C, and no change was found for α -helix structures. This implies that heating may induce the transformation from β -sheet to β -turn. Overall, whole milk powder had better temperature stability than infant formula powder, probably because of the lower content of lipid in the former than in the latter. These results help us understand the thermal stability of protein in milk powder.

Key words: protein secondary structure, temperature, milk powder, mid-infrared spectroscopy

INTRODUCTION

Milk powder has high nutritional value and some useful functional properties. Milk powder contains several types of proteins, including mucins, caseins, and whey proteins (Malacarne et al., 2002; Haug et al., 2007). The nutrients in milk can be utilized exceptionally well; some of the factors may be the presence of proteins in milk. The proteins provide not only adequate amounts of essential amino acids but also biological activities from antimicrobial effects to immunostimulatory functions (Lönnerdal, 2003).

Milk powder has a long shelf life (up to 12 mo) at room temperature. However, loss of solubility occurs gradually during storage (Anema et al., 2006; Havea, 2006), which may be linked to conformational modification of protein molecules during processing and storage (Kher et al., 2007). It is a common practice to mix milk powder with hot water before feeding or drinking. The effects of hot water on the milk structure and nutrition is unclear.

Milk-based infant formulas, both liquid and powder, are very sensitive to heat damage (Puig et al., 2003). Heat treatment of human milk will influence some of its protective constituents (immunoglobulin, lactoferrin, lysozyme, and others; Ford et al., 1977). When whole milk proteins are heated (Kim and Jimenez-Flores, 1995), β -LG and other milk serum proteins interact with milk fat globule membrane proteins. As the temperature of bovine β -CN increases, the protein secondary structure turns and extended structure are stable, whereas loops and helices are unstable. The β -sheets and β -turns probably form a supporting hydrophobic core (Farrell et al., 2001). The effect of storage temperature on the solubility of milk protein concentrate showed that insolubility could be due to cross-linking of the proteins at the surface of the powder (Anema et al., 2006). The formation of heat-induced whey protein complexes in milk increases the pH of gelation and the firmness of acid milk gels (Morand et al., 2012). The increased capacity of milk proteins to bind curcumin after heat treatment can be attributed to whey pro-

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tein denaturation, as whey proteins bind to the surface of casein micelles with heating (Yazdi and Corredig, 2012).

Mid-infrared spectroscopy (from 400 to 4,000 cm^{-1}) is a vibrational spectroscopy method that can reveal a wealth of information about the constituents in molecules, including proteins (Barth and Zscherp, 2002; Etzion et al., 2004; Barth, 2007). Mid-infrared spectroscopy is an efficient method widely used to predict milk fat, protein, lactose, and more detailed milk composition traits (Soyeurt et al., 2009; Rutten et al., 2011; De Marchi et al., 2014; Pappas et al., 2015; Toffanin et al., 2015; Visentin et al., 2015), as well as energy balance and feed efficiency (McParland et al., 2014). Modern infrared spectrometers are Fourier transform infrared (FTIR) spectrometers, which are similar to the Michelson interferometer (Arrondo et al., 1993). Fourier transform infrared spectrometry is a well-established method to probe the secondary structure of protein (Surewicz et al., 1993; Pelton and McLean, 2000; Barth and Zscherp, 2002; Li-Chan, 2007). The amide I band (1,600–1,700 cm^{-1}) of protein is sensitive to changes in secondary structures (Byler and Susi, 1986; Carbonaro and Nucara, 2010; Majzner et al., 2013). The attenuated total reflectance (ATR) technique (Goormaghtigh et al., 1999) has advantages in measuring solid samples.

We used ATR-FTIR to detect the heat-induced changes of 2 milk powders: whole milk powder and infant formula. To avoid the interference of water (Etzion et al., 2004), we used milk powders directly as our experimental samples without mixing with water.

MATERIALS AND METHODS

Apparatus and Materials

The measurements were carried out using an FTIR spectrometer (Tensor 27, Bruker, Bremen, Germany), equipped with global source, KBr beamsplitter, and deuterated triglycine sulfate detector. The crystal in the ATR attachment for the FTIR (Pike Technologies, Madison, WI) was germanium, and the angle of incidence was 45°. A press was used so that the milk powders could contact the crystal, avoiding trapped air. Infrared spectra were recorded with 16 scans in 400–4,000 cm^{-1} range with a resolution of 4 cm^{-1} . A vacuum blast-drying oven (DGG-9030A, Shanghai Senxin, Shanghai, China) was used to dry and heat the samples. The whole milk powder and infant formula (0–12 mo) were manufactured by the Inner Mongolia Yili Industrial Group Co. Ltd. (Hohhot, China), a well-known corporation in the Chinese milk powder market and were purchased in a local supermarket.

Methods

Sixteen tubes in parallel, each with a volume of 10 mL and with 20 g of sample each, were put into the vacuum blast-drying oven and then heated at a rate of 5°C/min. The temperature was increased from 25 to 100°C at 5°C intervals (16 samples). One tube was taken out after the temperature had reached each preset cut-off temperature, capped immediately, and stored in a desiccator with silica gel. To guarantee the samples were completely heated, heating was maintained for 5 min. About 2 to 3 g of milk powder from each capped tube was placed on the ATR crystal for measurement.

A linear baseline was subtracted from each spectrum to give a straight baseline in the spectral region of 1,800–2,000 cm^{-1} , and the spectra were smoothed with a 4-point Savitzky-Golay function to remove the possible white noise (Dong et al., 1990). For the secondary structure analysis, deconvolution of each spectrum was performed according to the methods of Fourier self-deconvolution (Kauppinen et al., 1981) and the finite impulse response operator using the software Opus 6.5 (Bruker, Bremen, Germany). A spectrum of a single band that is characteristic of a secondary structure is broadened in the liquid or solid states. The bands overlapped and could not be distinguished from each other. Second derivatives of the amide I (1,600–1,700 cm^{-1}) bands were used to indicate the position of individual component peaks of secondary structure within the amide I envelope. A curve-fitting procedure was used to estimate the area of each component representing secondary structures (Kumosinski and Farrell, 1993). The second derivatization and Gaussian curve fitting in the amide I region were analyzed using the software Origin 8.0 (OriginLab, Northampton, MA).

RESULTS AND DISCUSSION

FTIR

The mid-infrared spectra (400–4,000 cm^{-1}) of whole milk and infant formula powders at 25°C are shown in Figure 1. The characteristic infrared spectral bands of the 2 milk powders were very similar. The absorption bands of 1,630 to 1,680 cm^{-1} and 1,510 to 1,570 cm^{-1} were from protein (Carbonaro and Nucara, 2010) assigned to C=O stretching vibration absorption of amide I and N–H and C–H bending vibration absorption of amide II, respectively. The characteristic peaks of 2,920 cm^{-1} , 2,850 cm^{-1} , and 1,743 cm^{-1} were from lipids in the milk powders, which can be assigned to the bands of antisymmetric CH_2 stretching, symmetric CH_2 stretching, and C=O double-bond stretching, respec-

tively (Mizutani et al., 2004; Zhou et al., 2006). The absorption bands located at 3,200 to 3,800 cm^{-1} , 1,030 to 1,200 cm^{-1} , 900 to 930 cm^{-1} , and 755 to 785 cm^{-1} were attributed mainly to carbohydrate (Barth, 2007; Wenstrup et al., 2014). The NH stretching vibration gives rise to the amide A band of protein in the 3,200 to 3,800 cm^{-1} range (Barth and Zscherp, 2002).

The absorbance difference between whole milk powder and infant formula powder can be seen by focusing on the spectral window from 1,500 to 1,800 cm^{-1} , as shown in Figure 2. In the spectrum of whole milk powder, the absorbance at 1,740 cm^{-1} was almost the same as that at the 1,540 cm^{-1} and 1,650 cm^{-1} bands. The absorbance ratios of 1,740/1,540 cm^{-1} and 1,740/1,650 cm^{-1} were 0.81 and 1.08, respectively. In the spectrum of infant formula powder, the absorbance at 1,743 cm^{-1} was much higher than that at 1,540 cm^{-1} and 1,650 cm^{-1} . The absorbance ratios of 1,740/1,540 cm^{-1} and 1,740/1,650 cm^{-1} were 1.50 and 2.20, respectively. Table 1 shows the ratio of the 3 absorbances for clarity, which indicates that the whole milk powder contained similar amounts of lipid and protein, whereas infant formula contained more lipid than protein, which is consistent with the contents indicated on the respective packages. As indicated on the label, 100 g of the whole milk powder contained 28 g of lipid and 24 g of protein, whereas 100 g of the infant formula contained 27 g of lipid and 11 g of protein. The percentages of lipid and protein are given in Table 1.

The presence of lipid in the milk powder does not make the spectroscopy of protein more complicated. Instead, lipid enhances the changes in protein because strong interactions occur between lipid and protein in a lipid-protein powder system. Mizutani et al. (2004) showed more drastic changes in protein secondary structures in a lipid-zein mixture than in zein only when powders were heated.

Difference Spectra

The mid-infrared absorption spectra of the whole milk and infant formula powders changed when the temperature was increased from 25°C to 100°C. The underlying spectral changes as a function of temperature were easier to discern when these data were pre-

Figure 1. Fourier transform infrared (FTIR) spectra of whole milk and infant formula powders at 25°C.

sented as difference spectra, generated by subtracting the spectrum at 25°C from those obtained at higher temperatures. To avoid interference from noise and mechanical drifting, baseline correction and smoothing were performed on the spectra. As shown in Figure 3a, the FTIR difference spectra of whole milk powder exhibited a positive feature in absorbance centered around 1,650 cm^{-1} at 45°C, arising primarily from formation of α -helical conformations with increasing temperature. This band disappeared and reappeared with increasing temperature. The band centered around 1,540 cm^{-1} exhibited a similar feature, whereas the band around 1,740 cm^{-1} showed a negative feature in absorbance and a new band around 1,750 cm^{-1} appeared as tem-

Figure 2. Fourier transform infrared (FTIR) spectra of whole milk and infant formula powders from 1,500 to 1,800 cm^{-1} at 25°C.

Table 1. The absorbance ratio and percentage of protein and lipid in the whole milk and infant formula powders

Absorbance ratio	Whole milk	Infant formula
1,740/1,540 (cm^{-1})	0.81	1.50
1,740/1,650 (cm^{-1})	1.08	2.20
Protein (%)	0.24	0.11
Lipid (%)	0.28	0.27

perature increased. Temperature-induced variations in backbone (lipid C=O double-bond stretching) may shift the band of lipid toward a higher frequency (Yu and Damiran, 2011). Different features were found in infant formula as indicated in Figure 3b. The FTIR difference spectra of infant formula powder displayed a negative feature centered around $1,650\text{ cm}^{-1}$ at 45°C , resulting primarily from loss of α -helical conformations with increasing temperature. The α -helical structure partially reformed at higher temperatures. The band around $1,740\text{ cm}^{-1}$ showed the negative feature as that of whole milk powder. No higher frequency band of $1,740\text{ cm}^{-1}$ was observed for infant formula.

Second-Derivative Spectra

To elucidate the spectral differences, second-derivative spectra were used to find the number of secondary structure components and the approximate positions of these peaks (Kumosinski and Farrell, 1993). The second-derivative analysis can enhance the resolution for the spectra (Susi and Michael Byler, 1983). The second-derivative spectra were calculated using the Savitzky–Golay derivative algorithm. The frequencies at which amide I and amide II bands appeared were highly dependent on the secondary structure of the protein.

The peaks were assigned to different protein secondary structures. Peak assignment of amide I bands, which was shown in Table 2, was done according to the references (Pelton and McLean, 2000; Carbonaro and Nucara, 2010). Peaks between $1,620$ and $1,640\text{ cm}^{-1}$ were assigned to β -sheets, $1,651\text{ cm}^{-1}$ to α -helix, and between $1,660$ and $1,695\text{ cm}^{-1}$ were assigned to β -turns. The peak at $1,697\text{ cm}^{-1}$ was assigned to unordered structures.

Gauss Curve-Fitted Spectra

According to the positions of negative peaks observed in Figure 4, we performed Gauss curve fitting on spectra after subtracting baseline from $1,600$ to $1,700\text{ cm}^{-1}$. Figure 5 shows the results of Gauss curve fitting.

Table 2. Assignment of observed negative peaks to secondary structure¹

Observed negative peaks (cm^{-1})	Corresponding secondary structure
1,624, 1,627, 1,639	β -Sheet
1,651	α -Helix, loop
1,662, 1,666, 1,678, 1,681, 1,693	β -Turn
1,697	Unordered

¹Pelton and McLean (2000); Carbonaro and Nucara (2010).

Figure 3. Fourier transform infrared (FTIR) difference spectra of (a) whole milk powder, and (b) infant formula powder generated by subtracting the spectra collected at 25°C from those collected at higher temperatures, as indicated.

The results demonstrate that despite differences in the contents of main ingredients between whole milk powder and infant formula powder as described above, the principal secondary structure contained in the 2 milk powders was very similar.

The relative contents of the secondary structures of protein were calculated from the areas of the individual bands of the Gauss curve-fitting results. Changes in the secondary structure at different temperatures for the 2 milk powders are illustrated in Figure 6. Relatively, the contents of β -sheet and β -turn for both milk powders were higher compared with α -helix contents. Heating the 2 milk powders had different effects. In the case of whole milk powder, the contents of β -sheet, β -turn, and

Figure 4. Second-derivative spectra of original attenuated total reflectance (ATR)-Fourier transform infrared (FTIR) spectra of whole milk and infant formula powders from 1,600 to 1,700 cm^{-1} at 25°C. Negative peaks with small positive lobes represent different secondary structure components.

α -helix showed substantial changes at 70°C, whereas structures in infant formula changed at 50°C. The β -sheet and β -turn structures in whole milk powder displayed decrease and recovery at 70 to 85°C, whereas the α -helix structure displayed increase and recovery at this temperature range. The loss of β -sheet and β -turn may contribute to the formation of α -helix. In infant formula powder, the β -sheet structure showed a decrease and then increase, whereas the β -turn structure showed an increase and then decrease with the temperature increasing from 50 to 75°C, which implies that the β -sheet and β -turn may transform to each other with heating. The relative content of α -helix changed much less in infant formula than in whole milk powder, which may be due to the lower content of protein in infant formula. Proteins are stable in their low-energy native (folded) state and aggregation via protein-protein or lipid-protein interactions will lead to destabilization (Arrondo and Goñi, 1999). Time-resolved mid-infrared spectroscopy is needed for future research to determine how quickly protein secondary structures change with heating.

CONCLUSIONS

Mid-infrared spectra technology was used to monitor the absorbance changes of whole milk and infant formula powders at different temperatures. The infant formula powder contained more lipid relative to protein, whereas the whole milk powder had similar lipid and protein contents. Both milk powders had higher

proportions of β -sheet and β -turn structures than α -helix structures. In whole milk powder, the β -sheet, β -turn, and α -helix structures began to change at 70°C, whereas changes in secondary structure in the infant formula began at 50°C. Compared with infant formula powder, whole milk powder had better stability against heating. The lipid may help to maintain the protein structure when heated.

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Figure 5. Gauss curve-fitted spectra of whole milk and infant formula powders. Gauss curve fitting was performed on spectra of whole milk and infant formula powders after subtracting the baseline from 1,600 to 1,700 cm^{-1} .

Figure 6. Relative areas of the bands fitted to the Fourier-deconvoluted spectra of whole milk and infant formula powders at different temperatures from 1,600 to 1,700 cm^{-1} . Each band was assigned to the component of the secondary structure.

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