

Designing an industrial protocol to develop a new fat-reduced- ice cream formulation by replacing stabilizers with microbial transglutaminase enzyme

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Abstract

In this study, the possibility of replacing stabilizers with microbial transglutaminase (MTG) enzyme in fat-reduced ice cream production was studied. In addition, the stage of adding (before or after the heat treatment) the MTG enzyme to ice cream was also investigated. Five different ice creams (A and C containing 1 unit MTG/g protein without stabilizer, B and D containing 0.5 unit MTG/g protein and 0.35 % stabilizer, which also consist of the mixture of Carrageenan (E 407), Guar gum (E 412), Xanthan gum (E 415) and Sodium alginate (E 401), and E (control) containing 0.7 % stabilizer) were manufactured. MTG has been added to samples A and B after heat treatment while it was added to C and D samples before the heat treatment. An experimental analysis related to the overrun, viscosity melting properties, pH, titratable acidity, dry matter, fat, protein, sensorial and microstructural properties of ice creams was carried out. According to the results, the amount and the adding stage of MTG significantly affected overrun, melting, viscosity, coldness, firmness, smoothness, mouth coating, color, appearance, taste, smell scores, and also microstructure of ice creams ($p<0.01$). Results also showed that MTG could be used together with other stabilizers after heat treatment in the production of ice cream. Moreover, our findings demonstrated that sample B was the closest to control in terms of sensorial properties.

Key words: fat-reduced ice cream, transglutaminase, stability, quality, microstructure

Introduction

Ice creams are complex colloidal systems which, in the frozen state, comprise of ice crystals, air bubbles, partially-coalesced fat globules and aggregates, all in discrete phases surrounded by an unfrozen continuous matrix of sugars, proteins, salts, polysaccharides, and water (Goff, 2002). Ice cream contains a high concentration of fat (Metwally, 2007), which is considered as a multifunctional ingredient and influences the creaminess

(Koxholt et al., 2001), texture, mouthfeel (Adapa et al., 2000), colour and flavour of these products (Gonzalez-Tomas et. al., 2008). Fat also contributes to the ice cream properties of during freezing and beating, especially through the formation of a three-dimensional network of partially-coalesced fat globules. Some of the fat globules surround air bubbles, which results in stabilizing the air phase and increasing levels of fat aggregation, thus improving the melting resistance (Granger et al., 2005) and ice recrystallization (Goff, 2002).

Stabilizers are other ice cream constituents that partially increase mix viscosity, improve the product aeration and body, restrict ice-crystal growth during storage, control the rate of meltdown and prevent shrinkage of product volume during storage. The increased concentration of stabilizers leads to a non-linear increase in the apparent viscosity of mix (Cottrell et al., 1980). Besides fat and stabilizer, mix proteins also affect mix viscosity and foaming ability (Cottrell et al., 1980). Milk proteins, present in ice cream formulations, emulsify the fat and contribute to partial coalescence and fat structure formation. They are adsorbed at the air interface, leading to enhanced aeration and foam stability. On the other hand, the proteins, not present at interfaces, contribute to enhancing the viscosity and textural quality of the ice cream (Vega and Goff, 2005).

The formation of ice cream structure is hindered when the fat content is reduced and certain quality attributes such as viscosity, ice crystallization, hardness, melting rate, and flavor, are also affected (El-Nagar et al., 2002). Fat components are often replaced by carbohydrates and proteins which can perform same functional properties (Benjamins et al., 2009). The enzyme transglutaminase (TG), which is Generally Recognized As Safe (GRAS) by the Food and Drug Administration (FDA, 2010), has a high affinity for dairy proteins and modifies their functional properties.

Microbial transglutaminase (MTG) (EC 2.3.2.13), which is obtained from *Streptoverticillium* S-8112 strain, catalyzes an acyl transfer reaction between c-carboxy amide groups of peptide bound glutamine residues (acyl donor) and primary amino groups in a variety of amine compounds (acyl acceptor) such as the peptide bound ϵ -(γ -glutamyl)amino groups of lysine (Kieliszek and Misiewicz, 2014; Soleymanguori et al., 2014). The enzyme catalyzes protein cross-linking, protein deamination and amine incorporation. However, cross-linking dominates the first 30 min of reaction. Caseins are good substrates for the enzyme action (de Goes-Favoni and Bueno 2014; Sanli et al., 2014), particularly after heating. Beta-lactoglobulin, on the other hand, requires heating so that the enzyme can react with it (Sharma et al., 2001). TG treatment of milk proteins modifies their functional properties such as solubility and hydration ability, rheological and emul-

sifying properties and improve protein heat stability. It has been suggested that various dairy products like yoghurt with reduced fat and protein could be produced by TG having the texture similar to the full fat and high total solid products (Faergemand et al., 1999; Kasprzyk et al., 2016; Romeih and Walker, 2017). In other words, the action of TG could simulate fat in fermented milk products and could replace fat substitutes in low-fat products (Lorenzen et al., 2002; Hinz et al., 2012; Jooyandeh et al., 2017). Therefore, considering the fact that the demand for low-fat dairy products is increasing (Romeih and Walker, 2017; Gharibzahedi et al., 2018a, b; Garibzahedi and Chronakis, 2018), TG is regarded to be of great importance in food engineering applications.

On the other hand, ice cream is one of the high-fat dairy products (approximately 10 % fat) and its fat content needs to be reduced; however, the reduction of ice-cream fat contents would adversely affect the product's physical and sensory properties (El Nagar et al., 2002). Stabilizers are one of the ingredients that control ice-cream's physical structure. But, they are sometimes expensive and their percentages in ice cream are difficult to be reduced (Lal et al., 2006).

Studies have shown that MTG enzyme improves the textural properties of low-fat ice cream and increases melting resistancy and stability of it (Metwally, 2007; Rossa et al., 2011; 2012; Kafshgari et al., 2014; 2017; Kasprzyk et al., 2016). However, there is no study on the use of MTG enzyme as a stabilizer in ice cream production.

In these contexts, the aim of this study is to reduce the amount of stabilizer in low-fat ice cream and to produce ice-cream with same properties by replacing regular level stabilizer by adding MTG enzyme. For this purpose, five different ice cream samples containing MTG enzyme with or without stabilizer were manufactured (A and C containing 1 unit MTG/g protein without stabilizer, as suggested by producer, B and D containing 0.5 unit MTG /g protein and 0.35 % stabilizer, as determined in pre-trials, and E (control) containing 0.7 % stabilizer, as determined in pre-trials. MTG enzyme was added to these ice cream mixes before and after heat treatment (A and B samples were treated with MTG enzyme after heat treatment while C and D samples were treated with MTG enzyme before

heat treatment). Then, the effects of the added amount and the stage of adding MTG enzyme on the physical, microstructural and sensory properties of ice creams were investigated.

Materials and methods

Materials

Nonfat milk powder (96 % total solids) (Pinar Dairy, Turkey) and cream (35 % fat) (Tat Can Industry A.S. SEK Plant, Turkey) were used for ice cream production. Carrageenan (E 407), Guar gum (E 412), Xanthan gum (E 415) and Sodium alginate (E 401) (Sosa Ingredients, S.L. Ctra de Granera, Spain) were used as stabilizers. Lecithin was used as an emulsifier and obtained from Sosa Ingredients (S.L. Ctra de Granera, Spain). Sucrose (100 % TS) was used as a sweetener, and vanilla was added for giving aroma. MTG was received as a gift from Activa, Ajinomoto Co. Inc. (Japan). The declared activity of the preparation was approximately 1000 U g⁻¹.

Ice cream production

The ice cream was formulated with the following composition (percentage by weight): 32-33 % of total solids for a total batch of 5 kg, 10 % non-fat total milk solids, 18 % sugar, 4 % fat, and 0.3 % emulsifier. The amount of stabilizers and MTG are listed in Table 1.

TABLE 1. The stabilizers and MTG enzyme amounts in ice cream samples

| Stabilizers | Sample A | Sample B | Sample C | Sample D | Control E |
|------------------------------------|----------|----------|----------|----------|-----------|
| Lecithin (%) | 0.3 | 0.3 | 0.3 | 0.3 | 0.3 |
| Guar gum (%) | - | 0.1 | - | 0.1 | 0.2 |
| Carrageenan (%) | - | 0.1 | - | 0.1 | 0.2 |
| Xanthan gum (%) | - | 0.1 | - | 0.1 | 0.2 |
| Na-Alginate (%) | - | 0.05 | - | 0.05 | 0.1 |
| MTG (unit enzyme /g protein) | 1 | 0.5 | 1 | 0.5 | - |

Five different mixes were prepared from the above listed ingredients with same calculations except for MTG addition. Ice cream mixes were treated with MTG in two different manufacturing stages. While the first treatment was prior to pasteurization of the mixture, the second treatment was after it. The first (A) and the third (C) batches contain 1 unit of MTG /g protein whereas the second (B) and the fourth (D) batches contain 0.5 unit of MTG /g protein+0.35 % stabilizer, respectively. The fifth batch was the control (E) which contains 0.7 % stabilizer mixtures. The batches A, B and E were pasteurized in a jacked boiler at 85 °C for 1 min. These mixes were homogenized by a Lab homogenizer (Microfluidics, M-110P) at 25000 rpm while being hot. Then, they were allowed to cool down to 50 °C. After homogenization, the untreated-control mix was further cooled to 4 °C and left for aging for 24 h. The batches A and B were treated with MTG. Before being added to the mixture, the calculated amount of enzyme was stirred into 50 mL of the mix. The enzyme reaction was carried out at 50 °C for 30 min and the deactivation was carried out by a short heat treatment (80 °C for 1 min) (Lorenzen et al., 2002). The batches C and D were heated to 50 °C and treated with MTG for 30 min. Then the batches were pasteurized at 85 °C for 1 min. All mixes were cooled to 4 °C for aging for 24 h. The ice-cream was produced by using a vertical freezing machine of 6 kg capacity (Uğur, Nazilli, Turkey). The partially frozen mix was packaged in 50, 100 and 200 mL cups and stored at -18 °C. The experiment was conducted in duplicate.

Physical, chemical and sensory analyses were carried out 1 day after production.

Chemical analysis

The pH of milk and ice-cream was measured using a digital pH-meter (Thermo Scientific Orion 4 Star) and titratable acidity was measured by titrating 10 g of the sample with 0.1 N NaOH using phenolphthalein indicator (T.S.E., 1994). The dry matter of ice-cream was determined by drying samples at 100±1 °C for 3.5 h using an air oven (Kendro Laboratory Products UT12, 250 °C) (T.S.E., 1994). The fat content of ice cream was determined by the Gerber method (T.S.E., 1994). The protein content of ice cream was estimated from the crude

nitrogen content of the samples determined by the Dumas method (AOAC, 2003).

Physical measurements

The overrun of the final product was determined using the formula below (Akin et al., 2007):

First dripping and complete melting times were

$$\text{Overrun} = \frac{\text{the weight of equal volume of ice-cream}}{\text{the weight of unit mix - the weight of equal volume of ice-cream}} \times 100$$

measured according to method proposed by Güven and Karaca (2002). 25 g of tempered samples were left to melt (at room temperature, 20 °C) on a 0.2 cm wire mesh screen above a beaker. First dripping and complete melting times of samples were determined as seconds.

The viscosity of ice-cream samples were determined at 4 °C using a digital Brookfield Viscometer, Model DV-II (Brookfield Engineering Laboratories, Stoughton, MA, USA). A cylindrical spindle (no.4) was used for the determination of viscosity. The rotation rate was 20 rpm and readings were recorded after 10 secs. (Akin, 2005).

Microstructure

Samples of ice cream (0.3 g) were excised at approximately 1 cm below the surface and mixed with 0.3 g of 3 % aqueous agar solution (Oxoid code L13) at 45 °C and the mixtures were solidified by cooling at 20 °C (Tamime et al., 1995). The gelled samples were cut into 1 mm cubes and fixed in a 2 % glutaraldehyde solution in phosphate buffer (0.1 m, pH 7.2) as primary fixative to crosslink proteins (Kalab et. al., 1995) at room temperature for 2 h, followed by 24 h at 4 °C (McManus et al., 1993). The fixed samples were washed with phosphate buffer and postfixed in a 1 % osmium tetroxide solution in phosphate buffer for 2 h for stabilizing unsaturated lipids (Kalab et al., 1995). The post-fixed samples were washed with phosphate buffer and dehydrated in a graded ethanol series (30 %, 50 %, 70 %, 80 % and 90 % respectively, 1 h in each). Afterwards, they were also dehydrated in ab-

solute ethanol for 24 h and critical point dried with a EMS 850 CriticalPoint Dryer (TEM-SEM Kim. Mad. ve Lab. Cihz., Turkey) in carbondioxide. Dry sections were fractured with a blade and fragments were mounted on aluminum SEM stubs (Modler and Kalab, 1983), and coated with vacuum gold via an EMS 550X (TEM-SEM Kim. Mad. ve Lab. Cihz., Turkey). The microstructure of the ice cream was examined with a high vacuum scanning electron microscope ZEIS EVO 50 (Carl Zeiss Microscopy, Germany) at 15-20 kV, with a magnification of 1.50K x.

Sensory assessment

The samples were subjected to sensory evaluation by untrained ten panelists using a 10 points hedonic scale in order to evaluate coldness, firmness, viscosity, smoothness, color and appearance, mouth coating, flavour and taste, and general acceptability (1 = strongly unacceptable, 10 = very good) as described by Aime et al. (2001). The panel of assessors was an external panel of non-smokers who were very familiar with dairy products and were checked on the basis of sensory acuity and consistency.

Statistical analysis

The experiment was designed according to a (3x2x2) factorial design. All statistical analyses were carried out using the SPSS statistical software program (version 5.0). Statistically different groups were determined by the Duncan's Multiple Range test (Bek and Efe, 1995).

Results and discussion

Physico-chemical properties

The gross chemical composition of ice-cream samples is given in Table 2. The effect of MTG enzyme with or without stabilizer and adding stage of it on dry matter, pH and titratable acidity was insignificant ($p>0.05$). Rossa et al. (2012), Aloglu and Oner (2013), Romeih et al. (2014) and Kafshgari et al. (2014) also reported that addition of MTG did not affect the chemical composition such as total

solids, fat, protein, lactose, and acidity of ice cream or yoghurt. However, Rahmani et al. (2014) report-

ed that the addition of MTG enzyme raised the protein content of prebiotic low fat ice cream.

TABLE 2. Gross chemical composition of ice-cream samples

| MTG treatment stage | Sample* | pH | Titratable acidity (% lactic acid) | Dry matter (%) | Fat (%) | Protein (%) |
|-----------------------|---------|--------------------------|------------------------------------|---------------------------|--------------------------|--------------------------|
| After heat treatment | A | 6.57±0.000 ^{b2} | 0.21±0.001 ^{a1} | 32.52±0.148 ^{c3} | 4.00±0.000 ^{a1} | 4.08±0.170 ^{a1} |
| | B | 6.60±0.000 ^{a1} | 0.20±0.003 ^{a1} | 32.86±0.113 ^{b2} | 4.05±0.000 ^{a1} | 4.04±0.035 ^{a1} |
| Before heat treatment | C | 6.57±0.007 ^{b2} | 0.21±0.006 ^{a1} | 32.58±0.071 ^{c3} | 4.03±0.035 ^{a1} | 4.06±0.106 ^{a1} |
| | D | 6.60±0.007 ^{a1} | 0.20±0.004 ^{a1} | 32.88±0.057 ^{b2} | 4.05±0.071 ^{a1} | 3.99±0.325 ^{a1} |
| Control | E | 6.62±0.007 ^{a1} | 0.19±0.004 ^{a2} | 33.21±0.099 ^{a1} | 4.08±0.035 ^{a1} | 3.91±0.035 ^{a1} |

*A: 1 U/g protein, B:0.5 U/g protein, C: 1U/g protein, D:0.5 U/g protein, E: control

a, b, c Means in the same column followed by different letters and numbers were significantly different according to MTG level and treatment stage, respectively (p<0.01)

TABLE 3. Physical properties of ice-cream samples

| MTG treatment stage | Sample* | Viscosity (cP) | Overrun (%) | First dripping time (secs) | Complete melting time (secs) |
|-----------------------|---------|-----------------------------|---------------------------|------------------------------|------------------------------|
| After heat treatment | A | 890±4.243 ^{d4} | 23.12±0.332 ^{d4} | 765.50±14.849 ^{c4} | 2297±32.527 ^{c4} |
| | B | 4060±28.284 ^{b2} | 32.88±0.269 ^{a1} | 1229.50±54.447 ^{a1} | 3413.50±61.518 ^{a1} |
| Before heat treatment | C | 573.50±37.477 ^{e5} | 21.95±0.332 ^{e5} | 715.50±30.406 ^{c5} | 2126±53.740 ^{d5} |
| | D | 2823±24.042 ^{c5} | 26.83±0.573 ^{c5} | 1059±12.728 ^{b5} | 3072±33.941 ^{b5} |
| Control | E | 5840±123.037 ^{a1} | 30.83±0.544 ^{b2} | 1107.50±45.962 ^{b2} | 3166.50±54.447 ^{b2} |

*A: 1U/g protein, B:0.5U/g protein, C: 1U/g protein, D:0.5U/g protein, E: control

a, b, c Means in the same column followed by different letters and numbers were significantly different according to MTG level and treatment stage, respectively (p<0.01)

Physical properties of ice-cream samples are shown in Table 3. Sample B has the highest overrun value. Overrun values of ice creams without stabilizers are lower in comparison to the other samples. For a fact, stabilizers partially improve the product aeration. Using MTG enzyme alone as a stabilizer caused lower overrun values in the samples, but when MTG enzyme was used with stabilizers, overrun values increased from 21.95 to 32.88 % (p<0.01). The enzyme cross-linking of milk serum proteins and proteins of fat globules would have enhanced the cohesion properties of air cells lamella and the adherence of the adsorbed film of fat globules (Metwally et al., 2007). This action resulted in a high overrun with stable foam in low

stabilizer samples B and D. Increasing overrun of ice cream treated by MTG enzyme was also reported by several researchers (Metwally et al., 2007; Rossa et al., 2012; Rahmani et al., 2014; Kafshgari et al., 2014). Similarly, the addition stage of the MTG enzyme had a significant effect on overrun values of the ice cream samples (p<0.01). The overrun values of samples treated with MTG enzymes after pasteurization were higher compared to the samples treated with MTG enzymes before pasteurization. This can be related to the stabilization of the foam emulsion system by MTG enzyme which makes crosslinks of inter-or intra-molecular ϵ -(γ -glutamine)-lysine isopeptide bonds and leads to keep the air cells into the three-dimensional net-

work structure (Gharibzahedi et al., 2018a). Thus, the formation of casein polymers involving air bubbles increases volume and air bubble stabilization as in the sample B. Similar results were found in low-fat ice cream by Rossa et al. (2012) and Metwally et al. (2007).

In terms of the viscosity, the control sample was the highest ($p<0.01$). On the contrary, Metwally et al. (2007) and Rossa et al. (2012) reported that the samples containing TG enzyme had higher viscosity compared to control samples (without TG enzyme). This could be explained by a lack of stabilizer or the half amount of stabilizer in MTG enzyme treated to ice cream when compared to the control. Stabilizers increase the water binding capacity, hence do not allow the water to flow. Samples A and C without added stabilizer had the lowest viscosity value. Also the samples B and D contained about half the amount of stabilizer in comparison to the control (E) sample. The samples contained a regular stabilizer as mentioned in the studies above. The addition of MTG enzyme with stabilizer caused a significant increase in the viscosity ($p<0.01$), probably due to the ability of MTG enzyme to form high-molecular-weight polymers from monomers of proteins, conferring greater resistance to flow and the effects of the stabilizer. Garibzahedi et al. (2018a) reported that the polymerization of milk proteins through crosslinking reactions caused a notable increase of the water binding capacity by the formed high-molecular weight matrix, and so this bioengineered protein structure can greatly allow more resistance to flow. Similarly, in this study, viscosity values in the samples (B and D) treated with MTG enzyme after pasteurization were higher

than the samples treated with MTG enzyme before pasteurization ($p<0.01$). This can be explained by the formation of cross-linking with denatured whey protein and caseins (O'Sullivan et al., 2002).

First dripping and complete melting times of ice creams without stabilizers have shorter than other samples. The first dripping times and complete melting times prolonged when MTG enzymes were used with stabilizers ($p<0.01$). This could be related to the polymerization of milk proteins by the action of TG enzyme (Rossa et al., 2011) which led to an increase in the stability of ice cream, especially when the amount of fat in the formulation was reduced (Rossa et al., 2012). Furthermore, stabilizers provide greater resistance to the flow of the liquid phase resulting in slower melting. Our results indicated that the addition stage of the MTG enzyme had also a significant effect on the firstdripping and complete melting times ($p<0.01$). Sample B had the longest first dripping and complete melting times. Sharma et al. (2001) reported that a heating step can cause a noticeable increase in the crosslinking of proteins. This fact could be related to the kinetic improvement of protein reorganization induced by MTG enzyme at high temperatures. In heated milk, the MTG enzyme keeps more whey proteins into the matrix with reduced syneresis by changing the physical features of protein gels with low porosity (Gharibzahedi et al., 2018a). According to Cruz et al. (2009), the melting time of ice cream is related to its stability after overrun. Likewise, in the present study, stability of the samples (A and B) treated with MTG enzyme after pasteurization was better than in the samples (C and D) treated with MTG enzyme before pasteurization.

TABLE 4. Sensory characteristics of ice cream samples

| MTG treatment stage | Sample* | Coldness | Firmness | Viscosity | Smoothness | Mouth coating | Color and appearance | Taste and smell | General acceptability |
|-----------------------|---------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|
| After heat treatment | A | 6.94±0.177 ^{b2} | 4.63±0.276 ^{c3} | 4.88±0.233 ^{c5} | 4.20±0.247 ^{c5} | 4.00±0.163 ^{c5} | 7.15±0.113 ^{c5} | 5.87±0.219 ^{c5} | 5.12±0.247 ^{c5} |
| | B | 5.19±0.219 ^{d4} | 8.07±0.156 ^{a1} | 8.45±0.085 ^{a1} | 8.44±0.134 ^{a1} | 8.24±0.014 ^{a1} | 9.13±0.099 ^{a1} | 8.70±0.297 ^{a1} | 8.26±0.099 ^{a1} |
| Before heat treatment | C | 7.60±0.170 ^{a1} | 3.89±0.255 ^{d4} | 4.55±0.191 ^{c5} | 3.84±0.184 ^{c4} | 3.82±0.148 ^{c5} | 7.08±0.191 ^{c5} | 4.82±0.226 ^{d4} | 4.55±0.106 ^{d4} |
| | D | 6.16±0.191 ^{c5} | 6.94±0.240 ^{b2} | 7.19±0.290 ^{b2} | 7.51±0.170 ^{b2} | 6.83±0.219 ^{b2} | 8.63±0.269 ^{b2} | 7.55±0.191 ^{b2} | 7.20±0.191 ^{b2} |
| Control | E | 5.07±0.226 ^{d4} | 8.14±0.198 ^{a1} | 8.57±0.134 ^{a1} | 8.24±0.276 ^{a1} | 8.29±0.177 ^{a1} | 9.09±0.205 ^{a1} | 8.69±0.205 ^{a1} | 8.23±0.170 ^{a1} |

*A: 1U/g protein, B:0.5U/g protein, C: 1U/g protein, D:0.5U/g protein, E: control

a, b, c Means in the same column followed by different letters and numbers were significantly different according to MTG level and treatment stage, respectively ($p<0.01$)

Sensory evaluations

Adding MTG enzyme with or without the stabilizer and the addition stage significantly ($p<0.01$) affected the tested sensory characteristics (Table 4) of ice cream samples, as well. For instance, Sample C had the highest coldness value and was followed by samples A, D, B, and E, respectively. This result could be related to MTG enzyme and stabilizer amount of ice cream samples. Whilst sample A and C had MTG enzyme without stabilizer, sample B and D had MTG enzyme with stabilizer, which was half of the stabilizer of the control sample (E). Coldness is related to large ice particles. When the water content increases, larger ice particles are formed and the sensation of coldness is intensified (Aime et al., 2001). MTG enzyme causes an increase in the water content of food. In addition, stabilizers can bind water and thus cause small ice crystals to occur in ice cream. On the other hand, after control, sample B and D had the highest firmness and viscosity scores. Because of crosslinking between milk serum proteins and fat globules in ice cream, which is formed by MTG enzyme, a higher cohesiveness effect was obtained (Sharma et al., 2001). Therefore, these ice cream samples had a firmer and more viscous structure than other samples. Rossa et al. (2012) reported that the addition of the MTG enzyme increased the hardness of ice creams. In our study, Sample B had the highest smoothness and mouth coating scores. The use of MTG with stabilizer resulted in a higher cohesion effect, more air entrapment in the ice cream, a smoother structure and slower melting of ice creams. The use of MTG enzymes with stabilizers also affected color and appearance and taste and smell scores of ice cream samples in a positive way. Kirmaci (2005) reported that the MTG enzyme improved physical and rheological properties of the product without affecting the taste and odor characteristics. But, in our study, the highest general acceptability score was recorded for sample B. Thus, the addition of MTG enzyme after heat treatment improved sensory characteristics of ice cream samples, due to an increase in the crosslinking of proteins. Gharibzahedi et al. (2018a) reported that high overall acceptability of ice cream formulations, modified via MTG enzyme crosslinking, could be resulted from improving the structural stability and rheolog-

ical behavior of the frozen ice cream foams with a reduction in ice crystals formation. As a result, samples A and B had lower coldness and higher firmness, viscosity, smoothness, mouth coating, color and appearance, taste and smell and general acceptability scores than Samples C and D.

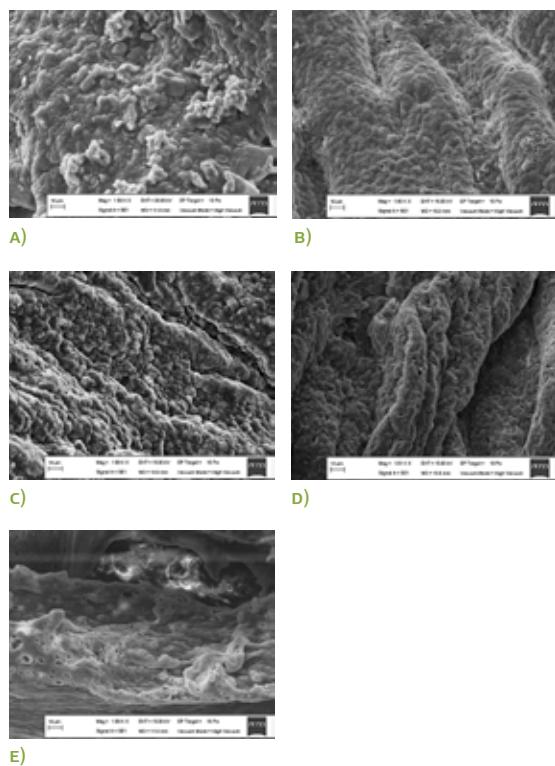


FIGURE 1. Microstructure of ice cream samples: (a) Sample A: 0.7 U/g protein MTG was added after heat treatment, b) Sample B: 0.35 U/g protein MTG was added after heat treatment, c) Sample C: 0.7 U/g protein MTG was added before heat treatment, d) Sample D: 0.35 U/g protein MTG was added before heat treatment, e) Control (MTG not added)

Microstructures of ice cream samples

Microstructure of ice cream samples are shown in Figures 1-5. Therefrom it could be observed that there were more air bubbles in the control sample and the structure was hollower compared to MTG enzyme treated ice creams. The more MTG enzyme was added, the less compact structure was obtained in the MTG treated samples due to a lack of stabilizers (Fig 1-4). Actually, it is expected that higher amount of MG would cause a more compact structure because of more intense cross-linking between the proteins. But, we determined that the addition of stabilizer with MTG enzyme improved

the structure of ice cream. This can be explained by the water binding capacity of the stabilizer. Gharibzahedi et al. (2018a) reported that the viscosity could further be intensified via the interaction of MTG enzyme with other carbohydrates such as inulin contained in the ice cream matrix. On the other hand, MTG treatment after heating resulted in a more compact structure than MTG treatment before heating. This could be related to an increase in denatured proteins caused by heating. Accordingly, the kinetic improvement of proteins was reorganized and MTG enzyme kept more whey proteins into the matrix (Gharibzahedi et al., 2018a). As a result, syneresis reduced and structure became more compact (Fig 1-2 and 3-4).

Conclusions

According to the results, the stabilizer amount can be reduced halved by using MTG enzyme in the fat-reduced ice cream. Results also showed

that the application of MTG enzyme with stabilizer obviously improved the sensory and physical properties of ice cream samples. However, application of MTG enzyme without stabilizer was not suitable in the production of fat-reduced ice cream. Also, it has been determined that the addition of MTG after heat treatment was more appropriate in terms of physical and sensory properties of ice cream samples. Consequently, the MTG enzyme can be used to reduce the amount of stabilizer in the fat-reduced ice cream production after heat treatment.

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Dizajn industrijskog protokola razvoja nove recepture sladoleda smanjenog udjela masti uz uporabu mikrobne transglutaminaze kao zamjene za stabilizatore

Sažetak

U ovoj studiji je istražena mogućnost zamjene stabilizatora mikrobnom transglutaminazom (MTG) u proizvodnji sladoleda sa smanjenim udjelom masti. Pri tom je također ispitivan utjecaj vremena dodavanja MTG (prije ili nakon toplinske obrade) na svojstva sladoleda. Proizvedeno je 5 različitih uzoraka sladoleda: A i C s 1 jedinicom MTG/g proteina, bez stabilizatora; B i D s 0,5 jedinica MTG/g proteina i 0,35 % stabilizatora, (smjesa karagenana (E 407), guar gume (E 412), ksantan gume (E 415) i natrijevog alginata (E 401)) i E (kontrola) s 0,7 % stabilizatora. MTG je dodavan uzorcima A i B nakon, a uzrocima C i D prije toplinske obrade. Svim uzorcima su određivani porast volumena, viskoznost, svojstva topljenja, pH, titracijska kiselost, suha tvar, udio masti i proteina, senzorska svojstva i mikrostruktura. Sukladno dobivenim rezultatima količina i vrijeme dodatka MTG značajno su utjecali na porast volume- na, topljenje, viskoznost, osjet hladnoće, čvrstoću, glatkoću, punoću okusa u ustima, boju, izgled, okus, miris te na mikrostrukturu sladoleda ($p<0,01$). Također se pokazalo da se MTG može koristiti skupa sa stabilizatorima, no u tom slučaju se treba dodavati nakon toplinske obrade sladoledne smjese. Zaključno, uzorak B se prema rezultatima senzorske analize pokazao najsličnijim kontrolnom uzorku.

Ključne riječi: sladoled smanjenog udjela masti, transglutaminaza, stabilnost, kvaliteta, mikrostruktura

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