

## Bio-Texturisation of Soy Protein Isolate

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### ABSTRACT

The possibility of using microbial transglutaminase (MTGase, 2.3.2.13) on bio-texturising of soy protein isolate is the main object of this study. Covalent cross-linking of proteins by MTGase is easy and simple for application, safe and well controlled process than commercial methods used for protein texturisation. Two commercial preparations of MTGase were used for restructuring soy protein isolate (SPI). The investigation of the product (gel) formed was evaluated using textural profile analysis (TPA). The results indicated that using of MTGase for texturising of SPI enhance greatly most of TPA parameters. The most affected parameters were hardness, cohesiveness and gumminess. Bio- texturising of proteins is a promising tool for restructuring and modifying the structure net work of protein molecules. The cross-linking reactions promoted by the action of MTGase allowing for the development and widespread use of new and improved functional properties in food industry.

**Keywords:** Bio-texturisation, microbial transglutaminase, plant proteins, soy protein isolate, textural parameters.

### INTRODUCTION

Proteins are one of the main food components; hence their modifications via chemical, physical or enzymatic methods are alternative available tools for improvement and/ or development of new functional properties of protein-based foods (Gaspar & Góes-Favoni, 2015).

Plant proteins occupy an important place in human nutrition, especially in the diets of low-income earners of developing countries. Furthermore, animal proteins are scarce in many countries, and vegetable proteins have a lower price than muscle proteins. Besides, it is well known that meat contains cholesterol and a higher proportion of saturated fatty acids than polyunsaturated fatty acids, which exert suppressive effects on pathogenesis of many diseases (Asgar *et al.*, 2010).

The functional properties of proteins are the physiochemical properties that affect their behavior in food systems, contributing to its quality and sensory or acceptability i.e., texture, appearance and flavour (Gaspar & Góes-Favoni, 2015).

Enzymatic texturising of protein or bio-texturising of protein is a promising tool for modification of protein molecules. The commercial enzyme, (MTGase), demonstrated its potential application in vegetable products such as soy which is little

exploited in the market (Góes-Favoni & Bueno, 2014). MTGase still rarely been used in texturisation of plant proteins. However, there is an increasing demand for vegetarian foods in many branches of the food industry (Dube *et al.*, 2007, Piestrasiak *et al.*, 2007, Chang *et al.*, 2011). The enzyme MTGase catalyses acyl transfer reaction forming inter- and intramolecular cross-links between amino acid residues of glutamine and lysine of protein molecules. The MTGase has several characteristics which covering many applications in food technology. The enzyme does not alter the pH, colour or flavour of food and it is active in a wide range of pH ranging from 4.0 to 9.0 (Ando *et al.*, 1989, Cui *et al.*, 2007).

With respect to substrate specificity, the enzyme has a broad specificity, so most food proteins could be cross-linked by MTGase (Nonaka *et al.*, 1997). Besides, the enzyme is capable of incorporating amino acids or peptides covalently into proteins, which improve nutritive value of food proteins (Yokoyama & Kikuchi, 2004). A practical application of this enzyme is the formation of heat-stable and irreversible gels, which could be obtained in different substrates even at relatively low protein concentrations (Góes-Favoni & Bueno, 2014). Covalent conjugation of two or more proteins of different origins may also be triggered by MTGase, generating a new protein with properties

different from those of original proteins (Gerrard, 2002). Several studies have reported that MTGase does not pose health risks and has no microbial antigenic potential (Seguro *et al.*, 1996, Kuraishi *et al.*, 2001).

Textural profile analysis (TPA) covered a wide range of food textural properties, which are usually divided into two main groups; primary and secondary. The primary parameters are those which could be measured directly from the TPA curve including; hardness (firmness), fracturability (brittleness), cohesiveness (stickiness), adhesiveness and springiness (elasticity). The secondary parameters, gumminess and chewiness are those that could be adequately calculated from two or more of the primary parameters. Generally, TPA could be used correctly for solid and semisolid foods. All of the TPA parameters must not present for all tested samples. Gumminess is only applicable to semisolids and is mutually exclusive with chewiness. Also, the parameters of viscosity and brittleness are found to be mutually exclusive. Only solids can have the characteristic of brittleness, whereas viscosity is limited, in the classification of liquid and semisolid products. The TPA test is simple and the textual parameters deduced intuitively understandable, it has become very popular. The result can be very misleading if the test is not conducted with the proper operational settings. Comparisons between TPA results are only likely to be valid if identical test protocols including the test geometry, speed and percentage of compression, are all kept constant (Bourne, 1978, Rosenthal, 2010, Trinh & Glasgow, 2012).

The present study focuses on the bio-texturising of soy protein isolate using MTGase and its opportunities to modify and improve many of the main functional properties of proteins. The effect of two commercially MTGase preparation on cross-linking of soy protein isolate was studied. The main textural properties of the gel formed were evaluated using texture profile test.

## MATERIALS AND METHODS

### Materials

Soy protein isolate, SPI, (SSPI-90D1W, Linyi Shansong Biological Products Co., Ltd. China) was used in the present study. The SPI was stored in a closed container at 4°C until used. Two commercial preparations of microbial transglutaminase obtained

from Ajinomoto Foods Europe, S.A.S. France and Mühlenchemie GmbH & Co.KG, Germany. They were designated as E<sub>1</sub> and E<sub>2</sub>, respectively. All the chemicals used were analytical grade.

### Methods

#### MTGase activity

The enzyme activity was determined using the colorimetric hydroxamate procedure with N-Carboxy-L-glycine. A calibration curve was prepared using  $\gamma$ -glutamic acid  $\gamma$ -monohydroxamate (Macedo *et al.*, 2007).

#### Preparation of gel samples

Soy protein isolate (30g) were mixed with 45ml of distilled water containing 0.6g MTGase, which representing 2% w/w of the amount of SPI used (Min & Green, 2008). After mixing, the resulted paste was placed into Petri dishes and incubated at 37°C for 30, 60, 90 and 120 min. These treatments were assigned as T2, T3, T4 and T5 for each enzyme preparation. For enzyme free samples (control, T1), the same process was followed, but no enzyme was added (Taylor & Walsh, 2001, Chang *et al.*, 2011).

#### Texture Profile Analysis (TPA)

TPA of the gel formed was performed on the texture analyzer (TexturePro CTV1.2 Build 9, Brookfield Engineering Labs. Inc. England). At zero time and at the end of each incubation period, a piece (3x 3 cm) of the gel formed (height 1.5 cm) was removed from the Petri dishes and placed on the platform (TA-RT-KI) with 10 Kg load cell, trigger load 0.07 N and test speed 1.5 mm/s using cylindrical plunger (TA 10). The samples were compressed to 50 % of its original height at crosshead in two bites (Yuan & Chang, 2007). The tested TPA parameters were; hardness, cohesiveness, springiness and gumminess. The measurements performed on three replicates.

#### Statistical procedures

F-test and analysis of variance of treatments difference was performed according to Steel & Torrie (1980). Statistical analysis was done by, ANOVA, F-test, and least significant difference (LSD) procedures available within the SAS software package, version 9.13 2007. The following model was used to analyze data obtained in factorial experiment.

$$Y_{ij} = \mu + \alpha_i + B_j + \alpha B_{ij} + e_{ij}$$

Where;  $Y_{ij}$  = effect of enzymes with different incubation time;  $\mu$  = overall mean;  $\alpha_i$  = the effect of enzyme,  $B_j$  = the effect of time,  $\alpha B_{ij}$  = interaction effect and  $e_{ij}$  = random error assumed to be independently and randomly distributed.

## RESULTS AND DISCUSSION

### Overall view

The measured activity of the two enzymes used,  $E_1$  and  $E_2$ , were  $7.5 \pm 0.2$  U/g of each enzyme preparation. So, an equal amount of enzymes were used. An identical protocol including; test geometry, compression speed, percentage of compression and other operating conditions are all kept constant for all tested samples so the results are likely to be valid (Rosenthal, 2010). The samples used in the present study were treated as semisolid materials; therefore gumminess rather than chewiness are presented in the results (Anonymous, 2012, Trinh & Glasgow, 2012). A graphic of the two-bite, force distance texture profile analysis curves of the two MTGase used on bio-texturizing of soy protein isolate is presented in Fig. (1). For reason of space, the data at zero time and after 90 min of incubation are only presented. The main features of these TPA curves are summarized in the following points: (1) the curves show a steep initial slope which indicates that the samples tested have high hardness, (2) the smooth curves,

sharp peaks and absence of shoulders suggests no fracturability and/ or no point of rupture or gross mechanical failure of the gel formed, (3) for adhesiveness, a low adhesiveness value was observed only in the control sample, while no noticeable values of adhesiveness were observed for all samples containing enzymes. This indicated that cross-linking of proteins by MTGase leads to overcome the low cohesiveness of the enzyme free samples (Piestrasik *et al.*, 2007, Asgar *et al.*, 2010, Rosenthal, 2010, Anonymous, 2012, Trinh & Glasgow, 2012, Nishinari *et al.*, 2013, Banjare *et al.*, 2015).

The statistical analysis of the data obtained is presented in Table (1). These data revealed that there were significant differences within all the TPA parameters tested. The main observations are: (1) the effect of  $E_1$  was generally higher than  $E_2$ , Table (1-A), (2) for time of incubation, Table (1-B) showed an increasing of all TPA tested parameters by increasing the time of incubation for both enzymes, the highest increase was noticed through the first 30 min of incubation for both enzymes, (3) the interaction between the enzyme and the time of incubation, Table (1-C), indicated that addition of MTGase significantly increased all the tested TPA parameters of the gel formed.

### Interpretation of the TPA parameters

**Hardness** refers to the force required to cause a given deformation. It represents the highest point

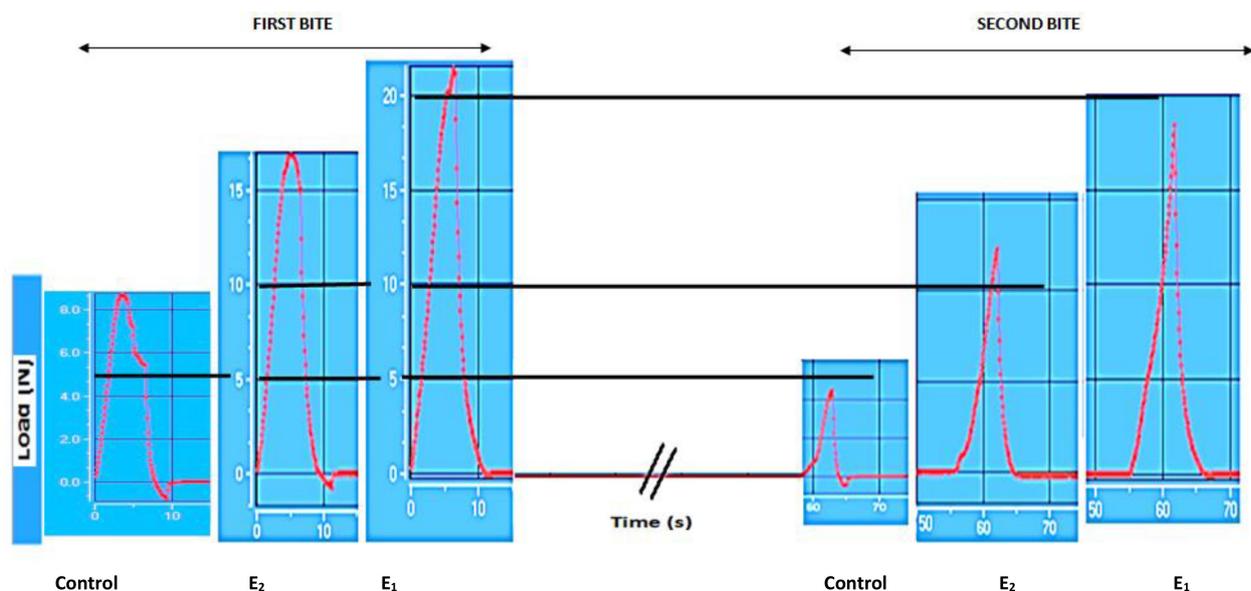


Fig. 1: A graphic for the two – bite force time texture profile analysis curves of bio – texturised SPI samples.

(The data shown is that at zero time ( control) and after 90 min of incubation for each enzyme  $E_1$  and  $E_2$ ).

**Table 1. Means of the texture profile parameters as influenced by enzymes, time of incubation and their interaction in soy protein isolate gel treatments.**

Factor	Hardness (N)	Cohesiveness	Springiness (mm)	Gumminess (N)
<b>A. Enzyme</b>				
E1	16.46 <sup>a</sup>	0.4427 <sup>a</sup>	8.82 <sup>a</sup>	7.68 <sup>a</sup>
E2	14.49 <sup>b</sup>	0.3780 <sup>b</sup>	8.31 <sup>a</sup>	5.66 <sup>b</sup>
<b>B. Time of incubation (min)</b>				
T1(0)	8.69 <sup>e</sup>	0.208 <sup>e</sup>	5.400 <sup>b</sup>	1.84 <sup>e</sup>
T2(30)	13.55 <sup>d</sup>	0.350 <sup>d</sup>	9.290 <sup>a</sup>	5.08 <sup>d</sup>
T3(60)	16.10 <sup>c</sup>	0.435 <sup>c</sup>	9.015 <sup>a</sup>	6.62 <sup>c</sup>
T4(90)	19.14 <sup>b</sup>	0.475 <sup>b</sup>	9.478 <sup>a</sup>	10.10 <sup>a</sup>
T5(120)	20.42 <sup>a</sup>	0.515 <sup>a</sup>	9.645 <sup>a</sup>	9.78 <sup>b</sup>
<b>C. Enzyme * Time of incubation</b>				
E1 * T1	8.69 <sup>i</sup>	0.206 <sup>h</sup>	7.07 <sup>e</sup>	1.84 <sup>i</sup>
E1 * T2	14.66 <sup>g</sup>	0.360 <sup>g</sup>	9.14 <sup>c</sup>	5.25 <sup>h</sup>
E1 * T3	16.45 <sup>e</sup>	0.460 <sup>c</sup>	8.56 <sup>d</sup>	7.58 <sup>d</sup>
E1 * T4	21.40 <sup>a</sup>	0.600 <sup>a</sup>	9.79 <sup>a</sup>	12.89 <sup>a</sup>
E1 * T5	21.11 <sup>b</sup>	0.510 <sup>b</sup>	9.54 <sup>ab</sup>	10.85 <sup>b</sup>
E2 * T1	8.69 <sup>i</sup>	0.210 <sup>h</sup>	7.07 <sup>e</sup>	1.84 <sup>i</sup>
E2 * T2	12.44 <sup>h</sup>	0.400 <sup>f</sup>	9.44 <sup>abc</sup>	4.91 <sup>g</sup>
E2 * T3	15.75 <sup>f</sup>	0.410 <sup>ef</sup>	9.47 <sup>abc</sup>	5.66 <sup>f</sup>
E2 * T4	16.87 <sup>d</sup>	0.430 <sup>de</sup>	9.16 <sup>bc</sup>	7.18 <sup>e</sup>
E2 * T5	19.72 <sup>c</sup>	0.440 <sup>cd</sup>	9.75 <sup>a</sup>	8.71 <sup>c</sup>
LSD0.05	.034	0.0244	.311	.017

Means followed by the same letter(s) are not significant, but different  
Letters are significant at 0.05 level of probability according to LSD method.

of the peak in the first compression cycle (Bourne, 1978). Hardness is the most representative parameters of TPA. It is the direct effect of action of MT-Gase on cross-linking of proteins (Gutt *et al.*, 2014). Fig. (1) showed that the second peak is smaller than the first indicating some weakening of the internal structure of the gel formed. This effect is consistent with TPA as technique (Rosenthal, 2010). The data present in Table (1-C) and Fig. (2-A) showed that the two MTGases have a noticeable effect on the hardness of the gel formed with priority of E<sub>1</sub> than E<sub>2</sub>. The hardness was increased by increasing the incubation time to 90 min for E<sub>1</sub> and extended to 120 min for E<sub>2</sub>. The highest value of hardness (21.40 N) was that of E<sub>1</sub> after 90 min incubation which is about 2.5 times higher than that of the control. These results indicated that the cross-linking of proteins by MTGase strength the network structure of protein molecules and hence increase the gel hardness. These results are in agreement with that reported by Kuraishi *et al.* (2001).

**Cohesiveness** is a direct indication to how the test material adheres to itself, so its structural integrity opposes successfully compressive or tensile stress, it is affected by the chemical structure of the tested material (Anonymous, 2012). As shown in Table (1-C) and Fig. (2-B), the cohesiveness rose up from 0.206 to 0.600 for E<sub>1</sub>, while it increased from 0.210 to 0.430 for E<sub>2</sub> for both the control samples and after 90 min incubation, respectively. Gaspar & Góes-Favoni (2015) stated that the use of MTGase improved the textural properties including hardness and cohesiveness resulting in strong gels with a compact and ordered structural conformation.

**Springiness** is an indication of how much a deformed sample returns to its original size and shape. The less a product is destroyed, the more springiness it will be observed (Trinh & Glasgow, 2012). For the two MTGase tested, springiness was moderately increased through the first 30 min of

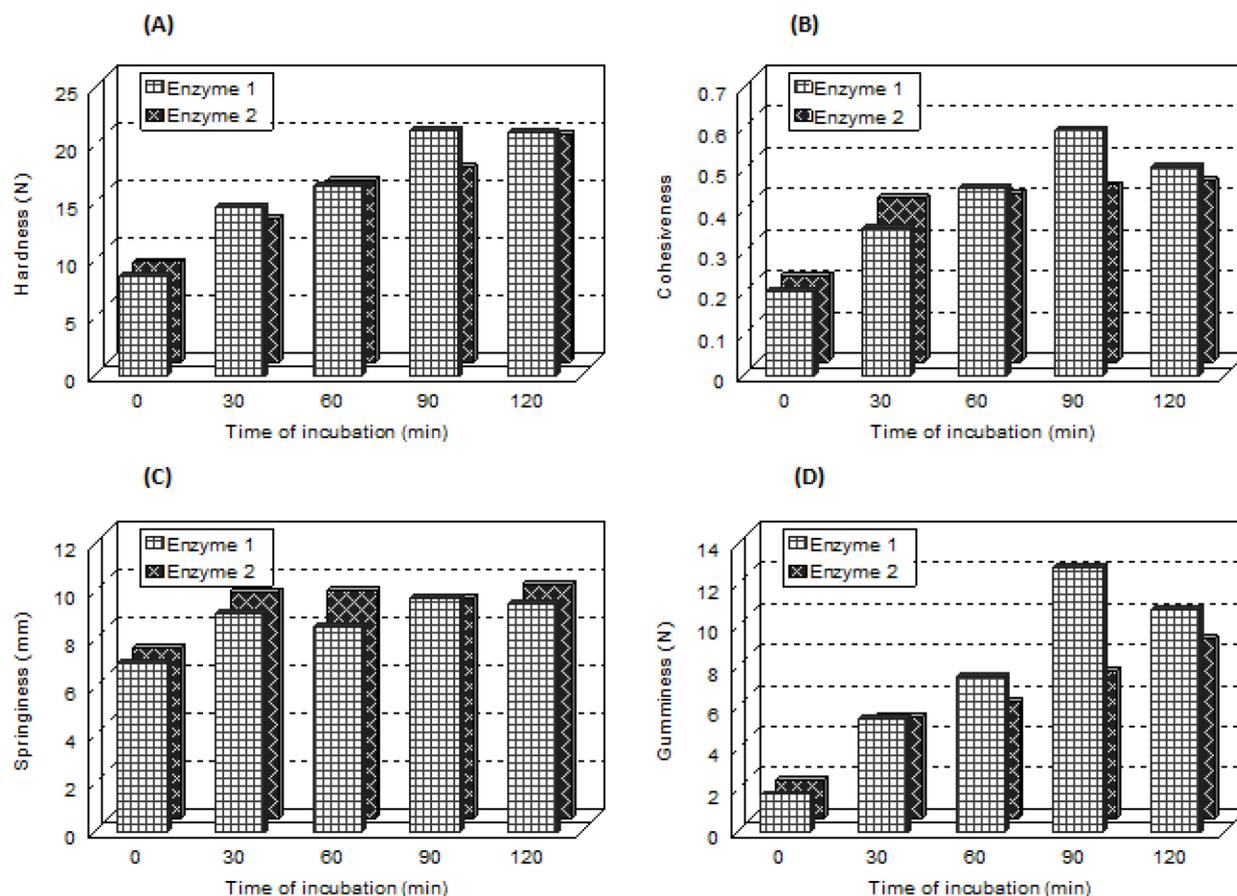


Fig. 2: Effect of incubation time on textural parameters of bio-texturised SPI treatments

incubation but less pronounced through the rest of incubation period (Table 1-C and Fig. 2-C). The springiness values of the control, after 30 and 120 min of incubation were 7.07, 9.14 and 9.54mm for E<sub>1</sub> and 7.07, 9.44 and 9.75mm for E<sub>2</sub>, respectively.

**Gumminess** is one of the secondary parameters of TPA. It mainly refers to the semisolid material. Generally the results, Table (1-C) and Figure (2-D), indicated that the gel produced by E<sub>1</sub> was gummier than that of E<sub>2</sub>. The maximum gumminess (12.89 N) was obtained after 90 min of incubation for E<sub>1</sub>, and then decreased to 10.85 N after 120 min of incubation.

## CONCLUSION

Bio-texturising of proteins is a promising tool for restructuring and modifying the structure network of protein molecules. It is very simple process compared with the commercial methods used for production of structured protein. The process doesn't alter the pH, colour or flavour of food. It will stop spontaneously when no more substrates, glutamine or lysine, available. The cross-linking reac-

tions promoted by the action of MTGase allowing for the development and wide spread use of new and improved functional properties by the food industry.

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## التشكيل البنائي الحيوي لمعزول بروتين الصويا

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الهدف الرئيسى من هذا البحث هو دراسة إمكانية استخدام إنزيم الترانسجلوتاميناز الميكروبي فى التشكيل البنائى الحيوى لمعزول بروتين الصويا . يقوم هذا الإنزيم بتكوين روابط عرضية للبروتينات بطريقة سهلة وسريعة وأمنة ويسهل التحكم فيها مقارنة بالطرق التجارية المستخدمة للتشكيل البنائى للبروتين .

تم خلال هذه الدراسة استخدام اثنين من المستحضرات التجارية لإنزيم الترانسجلوتاميناز الميكروبي لإعادة تشكيل معزول بروتين الصويا ثم دراسة خواص الجل المتكون باستخدام جهاز تقدير القوام وقد أظهرت النتائج أن استخدام هذا الإنزيم يؤدي إلى تحسين معظم خواص قوام الجل المتكون وكان التأثير الأقوى على خواص كل من خواص الصلابة، الالتصاق والمضغ .

تعتبر طريقة التشكيل الحيوى للبروتينات وسيلة واعدة لتشكيل أو تعديل البناء الشبكي لجزيئات البروتينات حيث تؤدي تفاعلات الربط العرضى الناتجة عن فعل إنزيم الترانسجلوتاميناز الميكروبي إلى تطوير وزيادة استخدامه فى تحديث و تحسين الخواص الوظيفية فى مجال التصنيع الغذائى .

