Effect of degree of hydrolysis on the functional properties of some oilseed proteins

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1. INTRODUCTION

Proteolytic or enzymatic modification of food proteins is an ancient technology by which mankind has been able to improve palatability and storage stability of the available protein resources. The enzymatic hydrolysis or modification of proteins results in two types of hydrolysates. First, extensively hydrolyzed products with increased solubility, but these have few if any functional properties. They also possess a bitter taste. Recently, it became clear that the extent of proteolysis must be limited if some functional properties are to be maintained or enhanced. Second, partially hydrolyzed products. This group encompasses a wide range of products with varying degree of hydrolysis. The degree of hydrolysis (DH) is a measure of the extent of hydrolytic degradation of proteins and is the most widely used indicator among different protein hydrolysates (1,2). Partially hydrolyzed proteins are prepared by stopping the hydrolysis process at a desired time. By this method a variety of products ranging in size from the original protein to small peptides or free amino acids can be obtained, based on the specificity of enzyme used and the relative rates of peptide bonds hydrolyzed. Another method of producing a controlled size-peptides is through the use of highly specific proteases so that only a few bonds are hydrolyzed and the further reaction stops (2).

Apart from their nutritional properties, the functional properties of protein or protein products must be taken into account as stated by Finch (3). Pour-El (4) had broadly defined functionality as any property of a food or food ingredient except its nutritional ones that affected its utilization. A listing of the key physico-chemical and related functional properties of protein hydrolysates include; solubility, viscosity, emulsification, foaming, thermostability, and flavor attributes are common to intact protein, but they differ from those of intact proteins by virtue of the DH (3,5). The functional properties of protein hydrolysates are also influenced by the specificity of enzyme used, the physical and chemical nature of intact protein, and hydrolysis conditions (5-8). The optimum conditions for the hydrolysis of soybean,
sesame, and rice bran meal proteins using papain or bromelain were elucidated in previous paper (9).

The aim of the present study was to investigate the effect of the DH on the functional properties of partially hydrolysed proteins. To achieve this goal the same enzymes and substrates investigated in previous paper (9) were used. The rate of enzymatic degradation was first determined followed by carrying partial hydrolysis by taking aliquots during the hydrolysis process, then determining their DH. The functional properties of some chosen hydrolysates were then determined and the relation between the DH and functional properties were discussed.

2. MATERIALS AND METHODS

2.1. Materials

Soybean and sesame seed meal was prepared by dehulling, grinding and defatting with n-hexane using a soxhlet apparatus. This extraction was carried out until the residual oil in the meal did not exceed 1%. Rice bran was treated similarly with the omission of the dehulling step. The prepared meals were spread to dry at room temperature, and ground to pass an 80-mesh screen.

Enzymes

Papain: was from Difco Laboratories (MICHIGAN USA) and Bromelain: from Sigma (Missouri USA). All reagents were of analytical grade.

2.2. Methods

2.2.1. Functional properties

2.2.1.1. Nitrogen Solubility Index as developed by Smith and Circle (10)

In a 400ml beaker were mixed 5g sample and 200ml water and the mixture was continuously stirred at 120 rpm for 120 minutes at 30°C. The mixture was transferred quantitatively in a 250ml volumetric flask, 2 drops of antifoaming agent added and diluted to mark with distilled water and mixed thoroughly. It was allowed to stand for a few minutes, then 40 ml were decanted in a 50ml centrifuge tube and centrifuged for 10 minutes at 1500rpm and supernatant filtered through a funnel containing a plug of glass wool. The clear filtrate was collected in a 100ml beaker and aliquots were taken for nitrogen determination

\[
\% \text{ Water soluble protein} = \frac{(S-B) \times N \times 0.014 \times 100}{\text{Wt. of sample}}
\]

Where:

- \( S \) = ml of alkali back titration of sample
- \( B \) = ml of alkali back titration of blank
- \( N \) = normality of alkali

\[
\text{Nitrogen Solubility Index (NSI)} = \frac{\% \text{water soluble nitrogen} \times 100}{\% \text{Total nitrogen}}
\]

2.2.1.2. Protein Dispersibility Index was determined according to Smith and Circle (10)

20g sample were weighed. A 300 ml volumetric flask was filled with distilled water at 25°C. About 50 ml water were poured into a blender cup. The sample was transferred to the blender cup and stirred with a spatula to form a paste. The remaining water was added in increments with stirring to form a smooth slurry. Blend at 8,500 rpm for 10 minutes. The slurry was poured into a 600ml beaker and after the slurry has separated decant or pipette a portion into 50 ml centrifuge tube and centrifuge for 10 minutes at 2,700 rpm. Pipette 15ml of supernatant into a kjedahl flask and determine protein (15ml=1g sample).

\[
\% \text{Water dispersible protein} = \frac{(S-B) \times N \times 0.014 \times 100 \times 6.25}{\text{Wt. of sample}}
\]

Where:

- \( S \) = ml of alkali back titration of sample
- \( B \) = ml of alkali back titration of blank
- \( N \) = normality of alkali

\[
\text{Protein Dispersibility Index (PDI)} = \frac{\% \text{water dispersible protein}}{\% \text{total protein}}
\]

2.2.1.3. Water absorption capacity was determined as described by Sosulski (11)

5g sample was weighed into a 50 ml centrifuge tube and 40 ml of distilled water were added from a burette the mixture was stirred with a stirring rod until homogenous. Centrifugation was carried out at 2000 rpm for 5 minutes. Decant the clear liquor back into the burette, which must contain water to at least the lowest graduation point. Determine the volume of the decanted liquor.

\[
\text{Ml of water absorbed} = 40 - \text{ml of decanted liquor}
\]

\[
\% \text{Water absorption} = \frac{\text{ml of water absorbed} \times 100}{\text{Wt. of sample}}
\]

2.2.1.4. Wettability was determined as described by Lucas Meyer (12)

Wettability means the ability of a powder to be wetted. It is expressed as the times in seconds a
certain quantity of powder needs to penetrate into a calm water surface. A 400 ml wide mouth beaker was filled with 150 ml water and covered with a glass plate. A glass hopper was adjusted above the glass plate with the lower aperture (30mm) on the plate. 1g of the powder sample was poured into the hopper then the glass plate was drawn aside and powder was allowed to drop onto surface of the water. Time was stopped in seconds when powder was wetted.

2.2.1.5. Flowability were carried out according to the method of Lucas Meyer (12)

It signifies the ability of a powder to flow, it was measured in seconds as the time a certain powder required to leave a funnel 50° wall inclination and 30mm diameter lower opening.

2.2.1.6. Bulk Density was also determined according to Lucas Meyer (12)

In order to choose appropriate packaging units, it is advisable to determine the bulk density of any instantized product. 100g powder was poured into a 250ml measuring cylinder quickly and the cm-quantity was read immediately. Bulk density was stated in g/cm. The powder volume is usually indicated as cm/100g powder and can be converted into powder bulk density by the following formula:

\[
\text{Bulk density in g/cm} = \frac{100}{\text{cm/100g powder}}
\]

2.2.1.7. Oil Holding Capacity measured as described by Childs and Forte (13)

0.1g of sample was added to a previously weighed 15 ml glass centrifuge tube and then 5 ml cottonseed oil was added. The tube and contents were agitated with a Vortex mixer for 1 min and then centrifuged for 15 minutes at 1000 rpm. The supernatant oil was decanted and the weight if oil bound per gram sample was calculated.

2.2.1.8. Gelation was carried out according to Circle et al. (14)

A series of solutions at 2, 3, 4, 5, 6, 7, 8, 10, 12 and 14% concentration for the protein sample was added to glass stoppered tubes. The tubes were heated to 75 - 80°C for 30 minutes in a water bath, then were placed in test-tube racks in ice and left in a cold room at 4°C for 60 minutes. At the end of the time interval the racks were removed from the ice and each tube was taken to test the stability of gel. The tubes containing stable gels were inverted and left in the rack for 30 minutes. The results were reported in terms of the lowest concentration of protein that remained stable after 30 minutes at room temperature.

2.2.1.9. Emulsifying Capacity was determined according to Swift et al. (15)

100mg of protein product were mixed with 30 ml of 1N-NaCl and heated at 80°C for 5 minutes, then blended in an Omni Mixer at 1000 rpm for 2 minutes after the addition of 20 ml of corn oil. Cutting and mixing at 13000 rpm for 30 seconds. Further addition of oil was continued at a rate of 0.8 ml/second until the emulsion broke and the phase transition occurred. Emulsifying capacity was reported as ml oil added to 100 ml protein product.

2.2.1.10. Thermostability was measured by the method of Kramer and Kwee (16)

Suspensions of protein samples and enzyme modified protein samples (2%, pH 6.8) were agitated on a magnetic stirrer for 15 minutes and 2 aliquots (10 ml each) were placed in screw capped test tubes and heated in a boiling water bath for 20 minutes. After rapid cooling to 22°C and centrifugation at 2000 rpm for 20 minutes, the protein content of the supernatant was assayed. Thermal aggregation of enzyme modified hydrolysate was represented in the difference in protein content of supernatant caused by heating.

2.2.1.11. Foam Stability was determined according to AOAC official methods of analysis (17)

1% protein dispersion was prepared and 25 ml of this dispersion were taken and placed in a quick fit measuring cylinder and closed with stopper the solution was shaken until it turned into a foam, the time was recorded. Another 25 ml of the protein dispersion were shaken for a period of time less than which turned it all into foam. The time of shaking and volume after shaking were recorded.

\[
\text{Foam Stability} = \frac{t}{2.303 \times \log \left( \frac{b}{c} \right)}
\]

Where:
\[
t = \text{time in seconds}
\]
\[
b = \text{volume of solution settling from foam during time t}
\]
\[
c = \text{volume of solution remaining in foam during time t}
\]

2.2.1.12. Nitrogen Solubility was determined according to Lyman et al. (18)

1g sample was weighed in 300 ml Erlenmeyer flask to which 100 ml of 0.02 N-NaOH solution were
added and the flask was shaken on a mechanical shaker in an incubator at 30°C for 1 hour. The material was then centrifuged at 5000 rpm for 10 minutes. 1 ml aliquots were taken from supernatant for nitrogen determination. Nitrogen solubility was expressed as percent total nitrogen which was soluble in 0.02 N-NaOH

\[
\% \text{ Nitrogen Solubility} = \frac{\% \text{ nitrogen soluble in 0.02 N NaOH} \times 100}{\text{total nitrogen}}
\]

2.3. Experimental

2.3.1. Enzymatic hydrolysis of protein

In a typical experiment a glass beaker containing a distilled water dispersion of the meal (calculated to contain 5% meal protein) was placed in a thermostatic water bath. The dispersion was continuously stirred with an electric stirrer, the pH and temperature were adjusted as previously determined by Taha et al. (9). The enzyme was added and stirring continued for the required time after which the reaction was stopped with 0.05 M trichloro acetic acid. The pH and temperature for each enzyme on each substrate as determined previously (9) were: when using papain on soybean meal (E/S 0.06, pH 7.2, temp. 50°C) on sesame meal (E/S 0.29, pH 7.0, temp. 50°C) on rice bran meal (E/S 0.10, pH 7.0, temp. 50°C) and when using bromelain on soybean meal (E/S 0.07, pH 6.0, temp. 45°C) on sesame meal (E/S 0.058, pH 6.0, temp. 45°C) on rice bran meal (E/S 0.21, pH 6.0, temp 45°C).

2.3.2. The Rate of Enzyme Degradation

This was determined by taking 10 ml aliquots of hydrolysates at 5, 10, 15, 20, 25, 30, 60, 120 minutes, and carrying formol titration on the aliquots. The amount of 0.1N NaOH consumed by the formol titration (19) was plotted against the time of the reaction. Ml NaOH represent the amount of – COOH group present in the 10 ml hydrolysate. The acidity as shown by the titration is an approximate measure of the amino nitrogen present.

2.3.3. Partial hydrolysis of meal protein

The experiment was carried as in experiments 1 and 2, then removing 25 ml aliquots at 5 minutes intervals and determining their degree of hydrolysis using the Formol titration method .The hydrolysates with different DH were spray dried and the powder kept in a dessicator for analysis of the functional properties.

2.3.4. Degree of Hydrolysis (DH)

From experiment 2.3.3.

\[
\% \text{ DH} = \frac{(B_2 - B_1) \times 14}{SW \times TN} \times 100
\]

\(B_1\) mL 0.1 N-NaOH consumed by sample at certain time
\(B_2\) mL 0.1N-NaOH consumed by blank
14= atomic weight of nitrogen
SW=amino nitrogen assayed by formol titration
TN=total nitrogen assayed by micro kjeldahl method.

3. RESULTS AND DISCUSSION

3.1. Rate of Enzymatic Degradation

Figures 1 and 2 are diagrammatic representations of the rate of enzymatic degradation of soybean, sesame, and rice bran meal proteins with papain and bromelain enzymes, respectively, under the optimum conditions of E/S, pH, and temperature previously determined (9), and at different time intervals.

Figure 1 shows that when soybean meal protein was hydrolyzed with papain the rate of degradation kept increasing up to 20 minutes after which it became constant. While on hydrolysis of sesame and rice bran meal protein with papain the rate of degradation kept increasing until one hour after which it remained unchanged. It is clear from Figure 2 that when using the enzyme bromelain for the three oilseed meal protein substrates; the rate of enzymatic degradation kept increasing until 30 minutes after which a plateau was reached.

![Figure 1](image-url)

Figure 1
Rate of enzymatic degradation of the three substrates hydrolyzed with papain.
3.2. Degree of Hydrolysis

Degree of hydrolysis is defined as the percentage of peptide bonds cleaved during the reaction.

Tables III and IV give the DH for the partially hydrolyzed soybean, sesame, and rice bran meal proteins when proteolytic enzymes papain and bromelain were used, respectively. The values of DH in tables I and II may be useful in preparing hydrolysates with certain functional properties, since the DH is the principle determinant of protein hydrolysate properties (1,2). From Tables I and II, twelve hydrolysates were chosen for further study. These hydrolysates were those hydrolysates of soybean, sesame, and rice bran with the least DH using both papain and bromelain, that corresponded to 5 minutes of hydrolysis. This choice was based on the fact that at the beginning of the hydrolysis process, hydrolysates with low DH will exhibit low bitterness (20). The other chosen hydrolysates were those resulting after 30 minutes hydrolysis, where the DH in most cases did not show much increase after 30 minutes, except in the case of rice bran hydrolysed with bromelain where there was a clear increase in DH between 30 and 120 minutes hydrolysis. For simplification, the hydrolysates of the three substrates chosen after 30 minutes of hydrolysis were:

### Table I
Degree of hydrolysis (%DH) during partial hydrolysis of the three substrates when using papain enzyme

<table>
<thead>
<tr>
<th>Substrate</th>
<th>Time (min.)</th>
<th>%DH</th>
</tr>
</thead>
<tbody>
<tr>
<td>Soybean</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sesame</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Rice bran</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

### Table II
Degree of hydrolysis (%DH) during partial hydrolysis of the three substrates when using bromelain enzyme

<table>
<thead>
<tr>
<th>Substrates</th>
<th>Time (minutes)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Soybean</td>
<td></td>
</tr>
<tr>
<td>Sesame</td>
<td></td>
</tr>
<tr>
<td>Rice bran</td>
<td></td>
</tr>
</tbody>
</table>

3.3. Functional Properties and their relation to the degree of hydrolysis

Certain functional properties of the protein hydrolysates play a more dominant role than others and dictate the choice of hydrolysates for a specific end use. The range of desirable and attractive functional properties that should be looked for is almost as broad as the range of foods themselves.

Table III gives the determined values for the functional properties of the chosen hydrolysates under investigation.
Wettability: The data in table III reveals that enzymatic hydrolysis at both high and low DH generally results in improvement of the wettability property when using both enzymes. In general low DH gave better results of wettability than high DH when using the same enzyme and substrate. Papain lowered the wettability time of soybean and rice bran hydrolysates compared to bromelain. Hydrolysing soybean meal with papain for 5 minutes resulted in 75% improvement in wettability over the non hydrolyzed soybean meal. Hydrolysing sesame meal with bromelain for 30 minutes improved the wettability of hydrolysate 97% over the non hydrolyzed sesame meal. Best improvement in wettability for rice bran was achieved with papain at low DH. Thus enzymatic hydrolysis is recommended to improve wettability of soybean and sesame meal. Papain at lower DH gave better results of wettability than high DH when using both enzymes. In general low DH generally results in improvement of the wettability property of the hydrolysate. On the other hand hydrolysing it with bromelain 30 minutes resulted in improvement of wettability. Both low and high DH are recommended to improve flowing properties of soybean and sesame protein hydrolysates, yet high DH resulted in more improvement in flowability.

Bulk Density: Is an important factor since it helps in choosing the appropriate packaging units, it is stated as g/cm³. Soybean and sesame meals treated with both papain and bromelain show an increase in bulk density over their corresponding untreated meals, which means larger packing units. On the other hand rice bran hydrolysates show a decrease in the bulk density over nonhydrolysed meal, meaning smaller packing units. No relation could be drawn between DH and bulk density.

Nitrogen Solubility Index (NSI): Is a very important measure of the functionality of the proteins in different food systems, especially in fortifying nutritious beverages. There is a great increase in the NSI of all hydrolysates over the non hydrolysed meals from which they were prepared (TABLE III). This result is confirmed by the nitrogen solubility carried according to Lyman (18) technique in table III. A direct relation between DH and nitrogen solubility could be drawn, increasing the DH results in an increase in nitrogen solubility.

Increased solubility of diverse proteins through enzymatic hydrolysis has been extensively reported by (4, 21-26). These studies concluded that even partial or limited hydrolysis produced substantially increased solubility of the resulting hydrolysates, particularly at the isoelectric point of the parent protein. The enhanced solubility of the hydrolysates is due to their smaller molecular size (24,27), and the newly exposed ionizable amino and carboxyl groups that increase the hydrolysate's hydrophobicity (28).

Results of wettability, dispersibility, flowability, and nitrogen solubility, predicted the good performance of the soybean, sesame, and rice bran hydrolysates over the nonhydrolysed meals, which means larger packing units. On the other hand rice bran hydrolysates show a decrease in the bulk density over nonhydrolysed meal, meaning smaller packing units. No relation could be drawn between DH and bulk density.
of the chosen hydrolysates in instant food products and beverages.

**Water Absorption Capacity (WAC):** Is the ability of a product to absorb water or swell. This property is important in the manufacture of bakery products, pastas, doughnuts and others. The hydrolysates of soybean with both enzymes show a slight improvement in their WAC over soybean meal. Also hydrolysates of sesamme using papain show increase in WAC, but sesame hydrolysed with bromelain and all rice bran hydrolysates show a decrease in WAC over the original meals prepared therefrom. This suggests that only soybean hydrolysates and sesame hydrolysed with papain can be used for bakery and pasta fortification. No relation could be drawn between DH and WAC. Generally enzymatic hydrolysis is not recommended to improve WAC.

**Oil Holding Capacity (OHC):** Is the ability of a protein to bind with oil. Is an important criterion in the meat industry (sausages, hamburgers etc.) and also in the doughnut industry. Table III shows that hydrolyzing soybean meal with papain and bromelain for 5 minutes increases the OHC from 1.9% for nonhydrolysed soy meal to 3.6% and 3.7% for soybean hydrolysed with papain and bromelain, respectively, while hydrolysing soybean with both enzymes for 30 minutes decreases the OHC. All sesame hydrolysates showed decrease in their OHC over their nonhydrolysed meal. Rice bran hydrolysed with papain for 5 minutes showed increase in the OHC over rice bran meal. No relation could be drawn between DH and OHC. The results indicate that only soybean hydrolysed with both enzymes for 5 minutes and rice bran hydrolysed with papain for 5 minutes can perform well in meat systems.

Choi and Rhee (29) studied enzymic hydrolysis of conventionally produced soy protein isolate, and of 7S and 11S rich fractions with pronase and alcalase. They found that changes in oil absorption capacity after modification varied with protein fraction and enzyme. Ponnamplan et al. (30) working with oat flour found that proteolysis reduced emulsifying stability, fat binding capacity, and heat coagulability when compared to soy flour.

**Gelation:** Is important in comminuted meat products as a protein's emulsifying capacity. It is reported as the lowest concentration of protein that remained as a stable gel after 30 minutes at room temperature. Table III indicates that all hydrolysates gelled at 2% protein concentration, while the three non hydrolyzed meals gelled at 3% protein concentration. No relation could be drawn between DH and gelation, but generally enzymatic hydrolysis is not recommended to improve gelation.

Adler-Nissen and Olsen (21) and Mahmoud (31) observed no gel formation for soy protein hydrolysates and explained the inability of hydrolysates to form heat induced gels by the reduced hydrophobicity of protein hydrolysates. Mahmoud (31) also suggested that presumably the increased charge repulsion between the peptides would lead to their loss of gelation ability.

**Emulsifying Capacity (EC):** Emulsifying and film forming ability of plant proteins is essential for those proteins to perform well in meat systems. Also a protein's ability to form emulsions is critical to their application in mayonnaise, salad dressing, milks, frozen deserts. Results in Table III show an improvement in the emulsifying capacities of all hydrolysates when hydrolysed for 5 minutes with both papain and bromelain. It can be concluded that the lower the DH the better is the emulsifying property of the hydrolysates.

It is generally accepted that the emulsifying property of hydrolyzed proteins was improved by carefully controlling the extent of hydrolysis (21,31). Philips and Beuchat (28) explained the observed improvement in the emulsifying property upon very limited hydrolysis that it could be presumably attributed to exposure of the hydrophobic protein interior which enhances adsorption at the interface, forming a cohesive interfacial film, with the hydrophobic residues interacting with oil and hydrophilic residues with water. It was also reported that as the extent of hydrolysis increased the emulsifying capacity decreased (2). Arias and Felacio (22) on hydrolysing sesame meal with bacterial protease found decrease in the emulsifying capacity as hydrolysis proceeded. Bernardi et al. (25) working with soy protein concentrate found emulsifying capacity to be unchanged when hydrolysing with fungal or bacterial protease, but emulsion stability decreased. Cha and Yoon (32) studied enzymatic modification of soy protein isolate using papain and actinidin. They observed that emulsifying capacity and emulsifying stability showed a marked decrease in the pH range 3-8.

**Thermostability:** Is the ability of a protein hydrolysate to remain soluble without aggregation leading to precipitation under elevated processing and sterilization temperature. Results in Table III show that hydrolysates of soybean show a decrease in their thermostability compared to the soybean meal. Sesame hydrolyzed with papain for 30 minutes show an increase in thermostability over sesame meal, rice bran meal show a 20% thermostability, which is increased to 25% on hydrolysis with both papain and bromelain for 5 minutes, and to 44% on hydrolysis with bromelain for 30 minutes. No relation could be drawn between DH and thermostability.

Mahmoud (31) studied the thermostability of extensively hydrolyzed soy protein and casein in hypoallergenic infant formulae and found that the hydrolysates remained soluble during UHT treatment (150°C) and subsequent retort sterilization (132°C).
Foam Stability: Is the capacity to form stiff, stable foam and is an important requirement of proteins to be incorporated into a gel cakes, whipped toppings, deserts, and soufflé-like products. Table III show that all treatments resulted in a decrease in the foam stability of the hydrolysates when compared to the meals prepared therefrom. No relation was found between DH and foam stability. Thus these hydrolysates cannot be incorporated into any of the above food items.

Hermanson et al. (33), Kang (34) both reported enzymatic hydrolysis to decrease foam stability. While Bernardi et al. (25) and Cha and Yoon (32) reported that enzymatic modification of soybean protein products improved U.P. foaming property.

Bitterness: the staff of our department tasted the hydrolysates, they did not detect any bitterness.

It can be concluded from the above results of the previous work that no definite correlation could be drawn between the degree of hydrolysis and most of the functional properties investigated.

A direct correlation could be made in the following cases:

- Nitrogen solubility of hydrolysates is directly proportional to the degree of hydrolysis, the higher the degree of hydrolysis the more the protein solubilized.
- Dispersibility of hydrolysate is also directly proportional to DH, although not necessarily improved over the nonhydrolysed meals.
- Wettability, flowability, and emulsifying capacity are improved at low degree of hydrolysis then decrease with increasing degree of hydrolysis.

REFERENCES


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