

Polymorphism of Protein Fractions as Biochemical Markers for Identification of Wheat Varieties

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Abstract

الملخص

The electrophoretic banding patterns of gliadin, albumin and globulin in wheat varieties grown in Sulaimanyah were determined by acid-polyacrylamide gel electrophoresis (Acid-PAGE) and Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE). The varieties of bread wheat used in the study showed 21 different patterns: six of them were identified to be corresponding to Ω -gliadin, three to the γ -gliadin, six to the β -gliadin, and other six to the α -gliadin. The gliadin patterns of wheat bread varieties greatly differed from the patterns of Italian wheat variety (Costantino), and the variation detected in wheat varieties from Sulaimanyah was limited to forty seven patterns. Only Tammuz showed the presence of gamma 47. The gliadin gamma 47 is only genetic markers for technological properties in wheat. The electrophoretic banding patterns showed variations between the bread wheat varieties. The highest value of similarity was observed between the varieties Cham 4 and Cham 6 for gliadin analysis. The gliadin pattern revealed more polymorphisms than the albumin and globulin pattern. On the other hand, in durum wheat varieties, two types of omega gliadin: omega gliadin 33 and 35 and two types of gamma gliadin were found: gamma gliadin 42 and gamma gliadin 45. A total of 84 band patterns were identified (durum wheat), of which 2 different mobility bands were in the region of omega gliadins, 1 in the region of gamma gliadins, 2 in the region of beta gliadin and 2 in the region of alpha gliadins. Genetic diversity index was highest in Creso Kurde and followed by Ovanto.

تتضمن الدراسة تحديد التركيب النموذجي لبروتينات كليادين والبومين وكلوبولين لستة اصناف من الحنطة الناعمة المزروعة في السليمانية بوساطة الفصل الكهربائي لهلام الحامض متعدد الاكرييلاميدي. وتم اكتشاف 21 نموذجاً او تراكيباً مختلفة منها 6 نماذج للكليادين اوميكا و3 نماذج للكليادين كما و6 نماذج للكليادين بيتاً 6 نماذج للكليادين الفا. ان التركيب النموذجي للكليادين الموجود في الاصناف الحنطة الناعمة المزروعة في السليمانية تختلف عن تركيب النموذجي للكليادين الموجود في صنف الإيطالي. ان بذور الصنف تموز هو الوحيد الذي يحتوى على كليادين كما و47 التي تعتبر دليل ايجابي لنوعية الحنطة. ان التركيب النموذجي لبروتينات كليادين والبومين وكلوبولين اظهرت الاختلاف الوراثي بين الاصناف المدروسة كما لوحظت اوجه التشابه بين بذور الصنفين شام 4 وشام 6 واكتشافت النتائج ان التحليل النموذجي للكليادين اظهرت اختلافاً واضحأً بين الاصناف المدروسة مقارنة بالتحليل النموذجي للبومين وكلوبولين. وفي الحنطة الخشناء، تم اكتشاف نوعين من كليادين اوميكا 33 و 35 ونوعين من كليادين كما و42 و 45. وجدت 2 نماذج مختلفة للكليادين في المنطقة اوميكا ونموذج واحد في منطقة كما و تركيبين مختلفين في المنطقة بيتاً و تركيبين مختلفين في المنطقة الفا. ان الاصناف كريزو كردى و او凡تو من الاصناف الاكثر اختلافاً عن باقي الاصناف المدروسة. كما ان تركيب النموذجي لبروتين البومين وكلوبولين برزت الاختلاف الوراثي بين الاصناف المدروسة.

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Keywords: Bread wheat, Durum wheat, Gliadins, Albumin, Globulin, Acid-PAGE, SDS-PAGE.

1. Introduction

The gluten proteins of wheat provide unique attributes to flour. The gluten proteins are classified based on their electrophoretic mobilities into glutenins and gliadins. Gliadins are monomeric proteins of molecular weight ranges between 30, 000 – 75, 000 Daltons. Gliadins are known to have extensive genotypic polymorphism (Branlard *et al.*, 1993). Based on the electrophoretic

mobilities, the gliadins are classified into four different group's alpha (α), beta (β), gamma (γ), and omega (Ω) or (ω) gliadins. The genes coding for these proteins are present on the short arm of chromosome 1 and chromosome 6 of wheat. They are tightly linked genes present on the three homologous loci of chromosome 1 as Gli-A1, Gli-B1 and Gli-D1. In chromosome 6, they are present as Gli-A2, Gli-B2 and Gli-D2 (Wrigley and Shepard, 1973; Brown and Flavell, 1981). The Gli-1 genes code for the ω and γ -gliadins and the Gli-2 genes code for α and β -gliadins. The Gli-1 locus is tightly linked to the LMW glutenin loci Glu-3 in chromosome1. The gliadin

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loci are known to be inherited in simple Mendelian inheritance and multiple-allelism has been reported for both the Gli-1 and Gli-2 loci (Metakovsky *et al.*, 1984; Metakovsky, 1991).

Gliadin components have high content of glutamine amino acid. For instance ω -gliadins have more than 50% glutamine (Lasztity, 1984). The proline content is also high in gliadins exceeding what found in HMW glutenins. The high proline content plays an important role in the secondary structure of gliadins (Makarenko *et al.*, 2002).

However, gliadins are poor in other basic amino acids such as lysine, arginine and histidine (Kasarda *et al.*, 1974). The determination of the N-terminal sequence of the gliadins has supported the theory that a large number of gliadins is due to a genetic mutation of a common precursor during natural evolution of bread wheat. Most of α , β -gliadins and some of the γ gliadins have similar N-terminal sequence. The α , β , γ -gliadins have six to eight cysteine residues, as a result, three to four intramolecular disulphide bonds occur (Kasarda *et al.*, 1984). Although the HMW and LMW glutenins form the disulphide cross-linked gluten matrix a small proportion (5-10%) of α and γ -gliadins occupy the matrix cross-link function. The ω gliadins also may take part in the polymer formation (Kukataite *et al.*, 2004). The gliadins, like the LMW glutenin, can also acts as the chain terminators. The gliadins are generally contributed to the viscosity and extensibility of the dough. The glutenin: gliadin ratio is thought to be an important determinant of quality. The higher glutenin to gliadin ratio is shown to contribute to greater dough strength (MacRitchie, 1985). Another study suggests that an increase in the relative gliadin content is associated with increase extensibility and loss of dough strength (Edwards *et al.*, 2001). Glutenin enrichment can circumvent this and increase dough strength. Although gliadins have been associated to certain parameters of bread making quality, they are considered unimportant to dough strength (Gianibelli *et al.*, 2001). Changes in the glutenin to gliadin ratios modify the bread making quality. In the previous studies deletions in the Gli-1 loci were found to exhibit greater dough strength (MacRitchie & Lafiandra 2001). This study aimed to examine the variations of the gliadin, albumin and globulin patterns in wheat varieties.

2. Materials and Methods

2.1. Plant Samples

One hundred gram of grains of each wheat varieties (bread wheat: Costantino, Tammuz, Aras, Rabia, Cham 4, Cham 6 and durum wheat: Creso kurde, Creso Italy, Ovanto, Bakrajo 1, Acsad 65, Cemmitto) were collected from the department of Agriculture of Sulaimanyah. All varieties used in this study are genetically pure.

2.2. Gliadin Extraction and Electrophoresis

Gliadin patterns were determined by A-PAGE. Twenty gram of wheat grains was hammer-milled to a fine powder. Forty mg of flour was mixed with 200 μ l of 60% (v/v) aqueous ethanol. The mixture was incubated in a closed microtube for 2h at 60°C and centrifuged for 10 min, at

14000 rpm. The supernatant was transferred to a microtube and diluted with 1 ml of acetone. The mixture was incubated for 10 min at RT. After centrifugation for 10 min at 14000 rpm, the pellet was resuspended in buffer A (30% Glycerol, 6 M Urea, 25 mM acetic acid, 0.01% w/v pyronin) (Sewa *et al.*, 2005).

To polymerize the gel, one drop (about 200 μ l) of fresh 30% H₂O₂ was added to 75 ml of cold gel solution (8.3% acrylamide, 0.415% N,N'-methylenebis-acrylamide, 0.1% ascorbic acid, 0.00067% Fe₂(SO₄)₃ in 5.1 mM Allactic buffer, pH 3.1). Each sample (30 μ l) was loaded onto a gel sized 180 (length) X 140 (width) X 2 (thickness) mm. Electrophoresis from the anode (the upper buffer) to the cathode (the lower buffer) was performed at 500 V for about 4 h until the tracking dye, pyronin, passed to the gel bottom. The gel was immersed into the staining solution containing 0.05% Coomassie Brilliant Blue R-250 (w/v) and 8% trichloroacetic acid for 5–16 h, and destained with distilled water (Metakovsky, 1991).

2.3. Albumin Extraction and Electrophoresis

Twenty gram of wheat grains was hammer-milled to a fine powder. Forty mg of flour was mixed with 400 μ l of 0.5 M NaCl. The mixture was incubated in a closed microtube for 2h at 4°C, and centrifuged at 14000 rpm for 10 min; the supernatant was precipitated in 1 ml acetone. The pellet was suspended in solution [28.5% sample buffer (7% SDS, Tris-HCl 0.01 M, pH 6.8, 30% glycerol, 0.001% comassie bleu) and 5% 2-mercaptoethanol]. The samples were incubated at 80°C for 20 min before loading. Samples of 30 μ l were loaded each the slots of SDS-PAGE. The gel consisted of a 15% separating gel, pH 8.4, beneath a 3% stacking gel, pH 6.8. Electrophoresis was carried out at room 27°C using a locally-made vertical electrophoresis apparatus. The running was performed at 15 mA/gel for 18 hours. Subsequently, the gels were stained by immersion in a solution consisting of 12.5% (w/v) trichloroacetic acid, 0.01% (w/v) Coomasie Brilliant Blue R250 and destained with distilled water (Tanaka *et al.*, 2003).

3. Data Analysis

Electrophoregrams for each variety of wheat were scored and the presence (1) or absence (0) of each bands noted. Presence and absence of bands were entered in a binary data matrix. Analysis was carried out using a statistical package SPSS-PC, version 15, by using the Dice similarity.

4. Results and Discussion

4.1. Variation in Gliadin pattern

4.1.1. Bread wheat

Electrophoretic separation of gliadin components exists in the bread wheat material (Figure 1). The Acid-PAGE analysis showed alpha, beta, gamma and omega namely three classes of gliadins. According to the nomenclature of Bushuk and Zillman (1978), the reference Costantino (reference variety) possessed omega gliadin Ω -35, 36, 38 (Table 1). Cham 4, Cham 6 and Rabia possessed only omega gliadin Ω -36. Aras possessed omega gliadin Ω -35

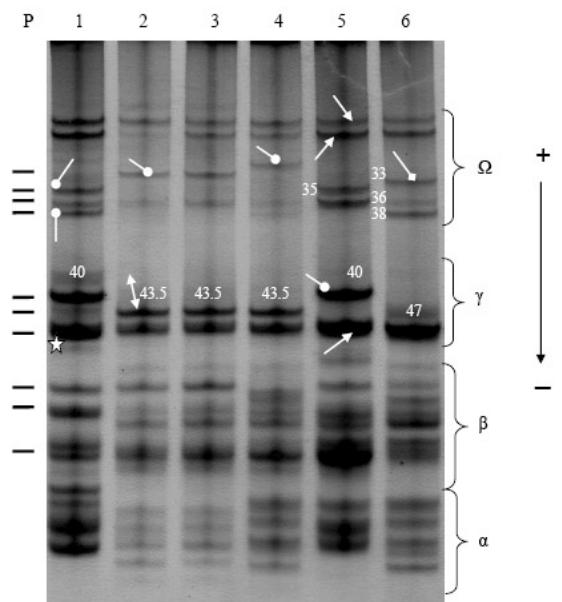


Figure 1. Electrophorogram separation of gliadin components in bread wheat varieties. 1: Costantino; 2: Cham 6; 3: Cham 4; 4: Rabia; 5: Aras; 6: Tammuz. The subunits of gliadins are designed according to Metakovskiy, 1991, Bushuk and Zillman, 1978. Arrows correspond to the protein encoded by Gli-D1, oval arrow refers to the protein encoded by Gli-B1, diamond arrow refers to the protein encoded by Gli-A3, star refers to the protein encoded by Gli-A1, double arrows refer to the protein encoded by Gli-B1. P: polymorphic bands.

Table 1. Gliadin subunits composition of six varieties of bread wheat.

Varieties	Omega gliadin (Ω)	Gamma gliadin (γ)
Costantino	35, 36, 38	40
Cham 6	36	43.5
Cham 4	36	43.5
Rabia	36, 38	43.5
Aras	35, 36	40
Tammuz	33, 36, 38	47

and 36. On the other hand, Tammuz showed three omega gliadin Ω -33, 36, and 38. According to the nomenclature of Pogna *et al.*, (1995) and Metakovskiy (1991), Costantino carries alleles Gli-A1a, Gli-A3a, Gli-B1m, Gli-B3a and Gli-D1k (Table 2). Cham 4 and Cham 6 contain the same alleles in all loci (Table 2). The alleles Gli-B3a are represented in all varieties. The allele composition at the Gli-1 loci of the genotypes analyzed here is reported in Table 1. According to Pagne *et al.*, (1987) the omega gliadins and the D-subunits of glutenin, coded by the Gli-B3 locus, are expressed in low amount and allelic variation is rather limited. Bread wheat Aras and Costantino possess the gamma gliadin component designated 40, whereas Cham 4, Cham 6, and Rabia possess the gamma gliadin 43.5. In addition, Tammuz possesses the gamma gliadin 47 (Table 1). This gamma locus is controlled by the Gli-B1 gene. The studies (Damidaux *et al.*, 1987; Payne, 1984; Joppa *et al.*, 1983) demonstrated the usefulness of gliadin gamma 45 and gamma 42, encoded at Gli-B1 locus as biochemical marker of good and poor pasta quality,

respectively. Alpha gliadin patterns varied among the varieties.

Table 2. Allele's composition at the Gli-1 and 3 loci of bread wheat varieties. The alleles are designed According to Pogna *et al.*, 1995 and Metakovskiy *et al.*, 1984.

Varieties	Gli-A1	Gli-A3	Gli-B1	Gli-B3	Gli-D1
Costantino	a	a	m	a	k
Cham 6	m	b	b	a	b
Cham 4	m	b	b	a	b
Rabia	b	b	c	a	b
Aras	b	b	d	a	k
Tammuz	m	a	e	a	k

The viscoelastic properties of gluten in both durum wheat and common wheat are influenced each of gliadin and glutenin patterns. Significant associations have been also detected between wheat quality and specific gliadin components (Damidaux *et al.*, 1987; Pogna *et al.*, 1982; Sozinov and Popereya, 1980). Pagne *et al.*, (1987) and Pogna *et al.*, (1990) suggested that allelic variation at the Glu-3 loci coding for LMW subunits of glutenin is probably responsible for differences in gluten quality, previously thought to be associated with the closely linked Gli-1 loci coding for gliadins. However, alleles at Gli-1 maintain their interest as genetic markers of quality. Dachevitch T. *et al.*, (1993) showed the genetic background of gliadins coded by the group 1 chromosomes in bread wheat Costantino. The authors identified some alleles coded to gliadins subunits in Costantino.

Result of Acid-PAGE also revealed the variations among the varieties of wheat according to the beta gliadin patterns.

4.1.2. Durum wheat

The Acid-PAGE analysis showed four classes of gliadins namely alpha, beta, gamma and omega (Figure 2). The varieties contain two types of omega: 33 and 35. Only Creso Kurde contained omega 33 and 35 and the rest of varieties possessed omega 35 (Table 3). On the other hand, the acid-PAGE showed one type of gamma gliadin: 45 which is considered as a good marker for pasta making quality (Pogna, 1990). On the other hand, the polymorphism of omega and gamma gliadin is revealed by one and two dimensional gel (Nieto-Taladriz, 1993). Similar studies (Damidaux *et al.*, 1987; Payne, 1984; Joppa *et al.*, 1983) demonstrated the usefulness of gliadin gamma 45 and gamma 42, encoded at Gli-B1 locus as biochemical marker of good and poor pasta quality, respectively. The viscoelastic properties of gluten in both durum and common wheat are influenced each of gliadin and glutenin patterns. Significant associations were also detected between wheat quality and specific gliadin components (Damidaux *et al.*, 1987; Pogna *et al.*, 1982; Sozinov and Popereya, 1980). Pagne *et al.*, (1987) and Pogna *et al.*, (1990) suggested that allelic variation at the Glu-3 loci coding for LMW subunits of glutenin is probably responsible for differences in gluten quality, previously thought to be associated with the closely linked Gli-1 loci coding for gliadins.

In this study, Acid-PAGE of grain storage proteins gliadin was performed in order to analyze molecular weight of gliadin subunits and investigate genetic diversity

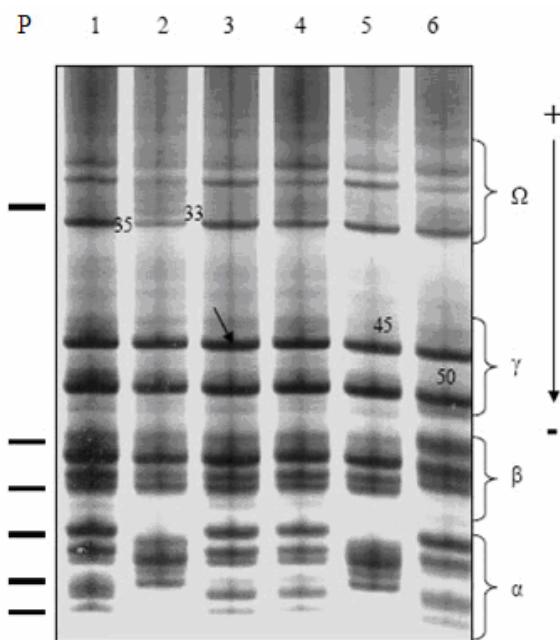


Figure 2. Electrophorogram showing banding pattern of durum wheat protein (Gliadin). 1: Creso Italy; 2: Creso Kurde; 3: Cemmitto; 4: Ascad 65; 5: Ovanto; 6: Bakrajo 1. The subunits of gliadins are designed according to Pogna *et al.* 1990. Arrow corresponds to the protein encoded by locus Gli-B1-2. P: Polymorphic bands.

Table 3. Omega and gamma gliadin subunits composition and locus of gliadin of durum wheat varieties.

Varieties	Omega gliadin	Gamma gliadin	Locus
Creso Italy	35	γ-45 and 50	Gli-B1-2
Creso Kurde	33, 35	γ-45 and 50	Gli-B1-2
Cemmitto	35	γ-45 and 50	Gli-B1-2
Ascad 65	35	γ-45 and 50	Gli-B1-2
Ovanto	35	γ-45 and 50	Gli-B1-2
Bakrajo 1	35	γ-45 and 50	Gli-B1-2

Among different wheat varieties. The electrophorograms showing proteins banding pattern of different wheat varieties are given in figure 2. Eighty four bands were obtained among which 6 bands showed variation, but the other bands were common in all varieties. The results revealed that Ovanto showed only one present band (Figure 2). At low molecular weight of gliadin (alpha gliadin), there are three bands polymorphic, reflecting more diversity.

4.2. Genetic Diversity and Cluster Analysis

4.2.1. Bread wheat

The electrophorogram proteins banding pattern, for seeds of all varieties of wheat detected by Acid-PAGE (Figure 1), showed segregation for gliadin.

Thirty four bands were obtained, and 11 bands were showed variations, but the other bands were common in all varieties. Cluster analysis of wheat grain storage proteins was carried out, depending on the results of Acid-PAGE

and using the UPGMA analysis to find the diversity among the given varieties of wheat as shown in the dendrogram (Figure 3). The diagram indicated five main groups: group 1= Cham 6 and Cham 4, group 2= Rabia, group 3= Tammuz, group 4= Costantino, group 5= Aras. At Dice dissimilarity of distance 1, Cham 6 and Cham 4 showed more similarity than others varieties.

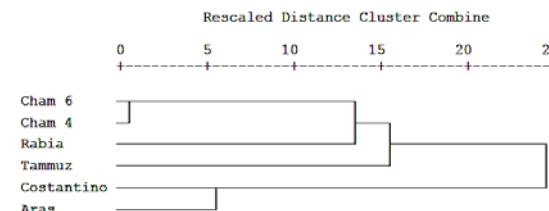


Figure 3. Dendrogram of bread wheat varieties showing the dissimilarity among the varieties based on A-PAGE- Gliadin.

At distance 13, there are 3 groups: group 1= Cham 6, Cham 4, Rabia, group 2= Tammuz, group 3= Costantino and Aras. At distance 17, Cham 6, Cham 4, Rabia and Tammuz are in the group 1 and Costantino and Aras are in the group 2. At dissimilarity distance 24 the diversity divides into two groups: group 1= Cham 6, Cham 4, Rabia and Tammuz and group 2= Costantino and Aras. Costantino and Aras showed more dissimilarity distance with the rest of the varieties.

4.2.2. Durum wheat

Dendrogram cluster analysis of wheat grain storage proteins (Gliadins) resulted from Acid-PAGE (Figure 2), using the UPGMA analysis, showed the diversity among the given varieties on the bases of dissimilarity distance of Dice (Figure 4). The diagram revealed four main groups: group 1= Creso Kurde and Ovanto, group 2= Cemmitto and Ascad 65, group 3= Bakrajo 1, group 4= Creso Italy. At Dice dissimilarity of distance 1, the varieties Creso Kurde and Ovanto and the varieties Cemmitto and Ascad 65 showed more similarity than others varieties.

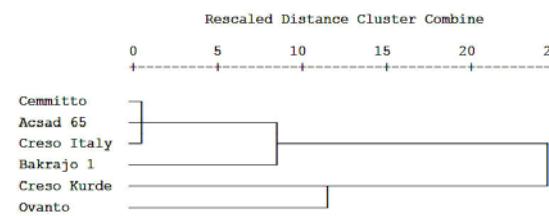


Figure 4. Dendrogram of durum wheat varieties showing the dissimilarity among the varieties based on A-PAGE- Gliadin.

At distance 5, there are 3 groups: group 1= Creso Kurde and Ovanto, group 2= Cemmitto, Ascad 65 and Bakrajo 1, group 3= Creso Italy. At distance 15, the varieties Creso Kurde and Ovanto are in the group 1 and the varieties Cemmitto, Ascad 65, Bakrajo 1 and Creso Italy are in the group 2. At dissimilarity distance 22, the diversity divides into two groups: group 1= Creso Kurde and Ovanto, group 2= Cemmitto, Ascad 65, Bakrajo 1, Creso Italy. The varieties Creso Kurde and Ovanto showed more dissimilarity distance with the rest of the varieties.

The presence of some patterns may correlate with a higher adaptive value of germplasm to the particular environment (Metakovsky *et al.*, 1991; Sewa *et al.*, 2005). The association between genotypes of wheat and their

environment has also been reported by Nevo *et al.*, (1988, 1995). However, no significant relationship between genotypes of wheat and their environments was reported (Dreisigacker *et al.*, 2004). Different combinations of gliadin patterns were prevalent in different regions, suggesting the adaptive properties of individual alleles or the chromosome segments, in which these alleles reside (Nevo *et al.*, 1995; Metakovsky and Branlard, 1998). Different gliadins might have some advantage over other gliadins in adaptation to the conditions, prevailing in these zones or these are closely linked with genes having adaptive values to the specific environment, though that needs to be confirmed by genetic analysis (Sewa *et al.* 2005).

4.3. Variation at The Albumin and Globulin Subunit

4.3.1. Bread wheat

Albumins and globulins were characterized by rich protein pattern. The numbers of band varied from 13 to 15, and they were defined by molecular weight 106–2 kDa (Figure 5). The protein pattern of albumins and globulins was divided into two relatively wide areas 66–23 kDa and 16–2 kDa. The electrophoretic pattern of tested varieties showed high similarity. The differentiating areas were located at the four zones with molecular weights of 92–87 kDa, 68–58 kDa, 49–39 kDa and 10–4 kDa for the 1st, 2nd, 3rd and the 4th zone, respectively.

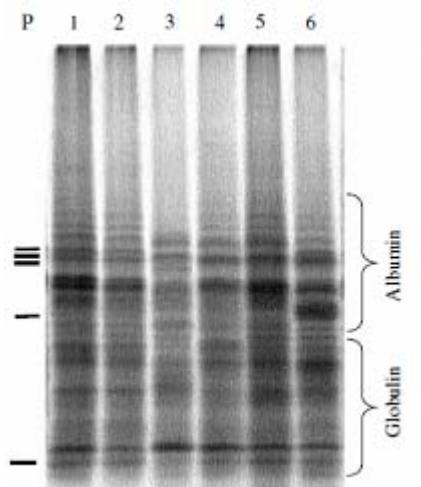


Figure 5. Electrophorogram showing banding patterns of bread wheat protein soluble (Albumin and globulin). 1: Costantino; 2: Cham 6; 3: Cham 4; 4: Rabia; 5: Aras; 6: Tammuz. P: polymorphic bands.

The similarity matrix and the resulting dendrogram (Figure 6) characterized similarity coefficients of the tested varieties. Although the results showed high similarity, the tested varieties were firmly identified. Tammuz possesses the five bands polymorphic, and Cham 4 possesses only one subunit. The highest similarity was found between Rabia and Aras. On the other hand, the lowest value of the similarity coefficient was found between Cham 4 and the rest of varieties. The diversity, in high molecular weight protein subunits, is the result of gene silencing in some varieties encoding these proteins (Lawrence and shephred, 1980).

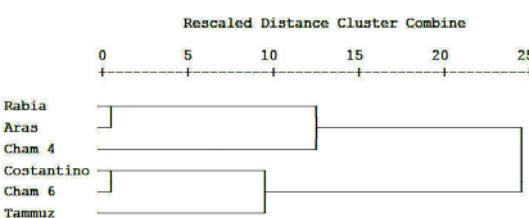


Figure 6. Dendrogram of bread wheat varieties showing the dissimilarity among the varieties based on SDS-PAGE- Albumin and globulin.

4.3.2. Durum wheat

Albumins and globulins were characterized in durum wheat by SDS-PAGE (Figure 7). The differentiating areas were located to zone: 40–60kDa. The SDS-PAGE gel did not show any difference among the varieties at the level globulin. The number of bands present is similar for all varieties.

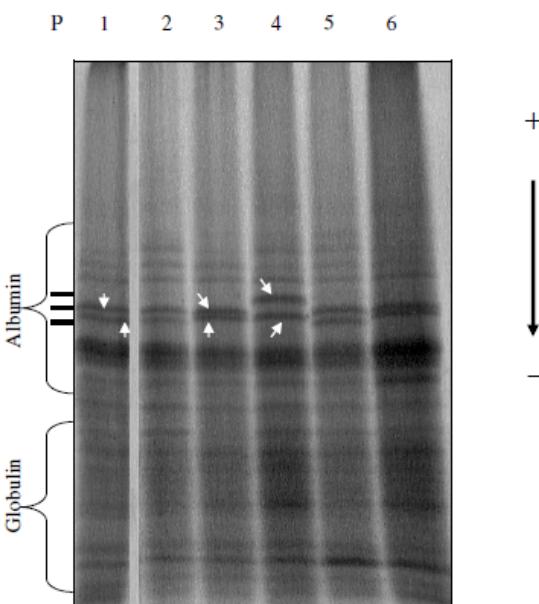


Figure 7. Electrophorogram showing banding pattern of durum wheat protein soluble (Albumin and globulin). 1: Bakrajo 1; 2: Acsad 65; 3: Creso Kurde; 4: Creso Italy; 5: Cemmitto; 6: Ovanto, P: Polymorphic bands.

The similarity matrix and the resulting dendrogram (Figure 8) characterised similarity coefficients of the tested varieties. The highest similarity was found between Creso Kurde and Ovanto and also for Acsad 65, Cemmitto and Bakrajo 1 (Figure 8). On the other hand, the lowest value of the similarity coefficient was found between the group which contains Creso Kurde, Ovanto and Creso Italy and the group containing Acsad 65, Cemmitto and Bakrajo 1.

The seed proteins are very suitable and useful genetic markers, because they are not influenced by external conditions, and so they enable to identify tested plant genotypes (Černý and Šašek, 1996). The finding predicates higher level of the intra-varieties polymorphism in these cultivars. Patterns of albumins and globulins among varieties were very similar and their characters corresponded with the detection by American authors, who used water dissolved proteins for identification of two common wheats in non-reduction conditions (Kim and Bushuk, 1995). Apart from the gluten proteins, water-

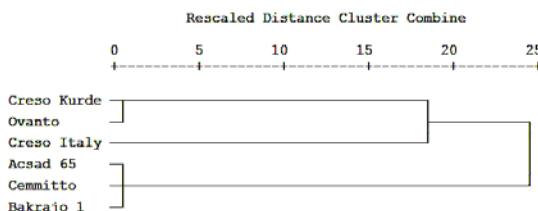


Figure 8. Dendrogram of durum wheat varieties showing the dissimilarity among the varieties based on SDS-PAGE- Albumin and globulin.

soluble albumins and salt-soluble globulins constitute from 10 to 22% of total flour protein (Singh and MacRitchie, 2001a). Albumins such as α -amylase/trypsin inhibitors (Buonocore *et al.*, 1985; Shewry *et al.*, 1984) and puroindolines (Garcia-Olmedo *et al.*, 2002) may have dual roles as nutrient reserves for the germinating embryo and as inhibitors of insects and fungal pathogens prior to germination. Puroindolines influence grain hardness (Morris, 2002). Generally, albumins and globulins are not thought to play a critical role in flour quality, although, minor importance on bread-making quality has been reported (Schofield and Booth, 1983). Both protein fractions are important from nutritional point, because of rather high amounts of essential amino acids. This experience is confirmed with the comparison of polymorphism among water soluble protein (albumins and globulins) and seed storage proteins (gliadins) of Dvořáček *et al.* (2001a).

In conclusion, the results showed that the gamma gliadin 47 which is linked to the strong quality of wheat for baking, is less frequent in the varieties tested. The two marker systems were analysed for a complex similarity evaluation of tested varieties. On the other hand, the gliadin pattern revealed more high level of polymorphisms than the albumin and globulin pattern. Identification of gliadins alleles in wheat varieties in Kurdistan may be useful for selection aims in breeding programs to determine the relationship between gluten viscoelastic properties and allelic variations at Gli-1, Glu-3 and Glu-3 in wheat varieties.

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