

Analysis of Seed Proteins in Groundnut Cultivars (*Arachis hypogaea* L.)

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ABSTRACT

The seed protein contents and protein banding pattern were studied in commonly cultivated groundnut cultivars. The groundnut cultivars such as ICGV00351, TMV-7, CO-4, CO-6 and TG-374 were used for quantitative and qualitative analysis of seed proteins. The protein contents varied among the different varieties of groundnut. The maximum protein content was observed in CO-6 followed by CO-4, TMV-7, ICGV00351 and TG-374. There was a slight difference in protein content among the different cultivars. All the five cultivars of groundnut were subjected to SDS-PAGE analysis. The results revealed that the variation in total number of bands and MW-Rf values. The maximum number of MW-Rf value was noticed in TG-374 and ICGV00351, and the minimum MW-Rf value was 11 recorded in CO-6 and TMV-7.

Keywords: Seed proteins, Spectrophotometric and Electrophoretic analysis

I. INTRODUCTION

Groundnut (*Arachis hypogaea* L.) is an annual warm-season plant of the legume family that originated in South America and groundnut has been cultivated since ancient times. The seeds contain 40-50% oil, 20-30% proteins and are an excellent source of vitamins B. Large seeded varieties are used for roasting and confections and small-seeded types are used for oil. The groundnut meal is rich in protein and can be used as food or feed [1].

Many workers used seed protein electrophoresis for characterization of cultivated and wild species of groundnut [2-9]. The seed storage protein analysis helps in identification and characterization of diversity in crop varieties, cultivars and their wild varieties and also provides information on phylogenetic relationship of the accession[10-12]. Seed storage protein profiles have also been used to study evolutionary relation of several crop plants [13]. The objective of this study was the estimate the seed storage proteins and know the banding pattern of proteins in some of the commonly cultivated groundnut varieties.

II. MATERIALS AND METHODS

2.1. Seed Material

Five commonly cultivated groundnut cultivars such as ICGV00351, TMV-7, CO-4, CO-6 and TG-374 were obtained from Tamil Nadu Agricultural University, Coimbatore and used as an experimental material.

2.2. Protein Estimation

The protein estimation was carried out by using Lowry *et al.*, (1951).

2.3. SDS-PAGE Analysis:

The Sodium dodecyl sulphate polyacrylamide gel electrophoresis Laemmli, (1970) was followed in this study.

III. RESULTS

3.1. Protein content in different groundnut varieties

The protein content was estimated in seeds of five groundnut varieties. The seed storage protein content was maximum in Co-6 (193mg/g/fw) followed by 187mg/g/fw in ICGV00351. At the same time the protein trend was 176mg/g/fw in TMV-7 and 162 mg/g/fw in CO-6 and 144 mg/g/fw in TG-374. The protein analysis of groundnut seeds indicated the differences among the five varieties (Table- 1).

3.2. SDS - PAGE analysis of seed proteins:

The seed storage proteins were analyzed by SDS-PAGE for all the five varieties. The maximum number of bands observed was 21 for all the varieties. In TMV-7 and Co-6 MW-Rf was 11. But in TG-374 and ICGV00351 recorded highest MW-Rf of 17. At the same time in Co-4 the MW-Rf of 12 was noticed. The following proteins with MW-Rf values of 0.457, 0.642, 0.685, 0.714, 0.785, 0.814, 0.857 and 0.928 were commonly noticed in all the five varieties. The MW-Rf 0.528 was present in ICGV00351, TG-374 and Co-6 but absent in Co-4 and TMV-7. Like that 0.514

was expressed in ICGV00351, TMV-7 and TG-374 but not found in Co-4 and Co-6. The MW-Rf 0.557, 0.585 and 0.742 were observed in ICGV00351 and TG-374. The MW-Rf 0.342 present in all the varieties except ICGV00351. The MW-Rf of 0.57, 0.842 and 0.871 recorded in TG-

374 but absent in all the four varieties. The MW-Rf of 0.328 and 0.385 were present only in ICGV00351. The MW-Rf 0.900 was expressed in ICGV00351 and Co-6 but not noticed in remaining three varieties. The MW-Rf of 0.971 was present in all the varieties except Co-6.

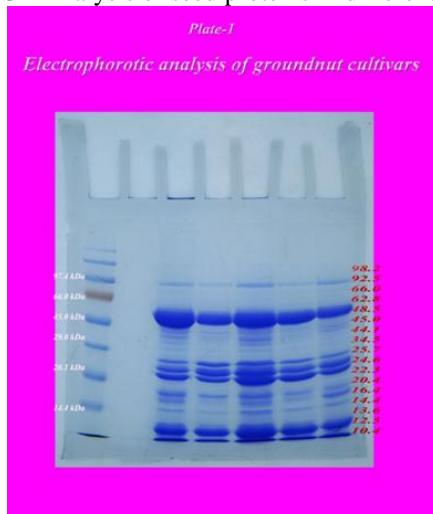
Table -1 Protein content in seeds of groundnut cultivars

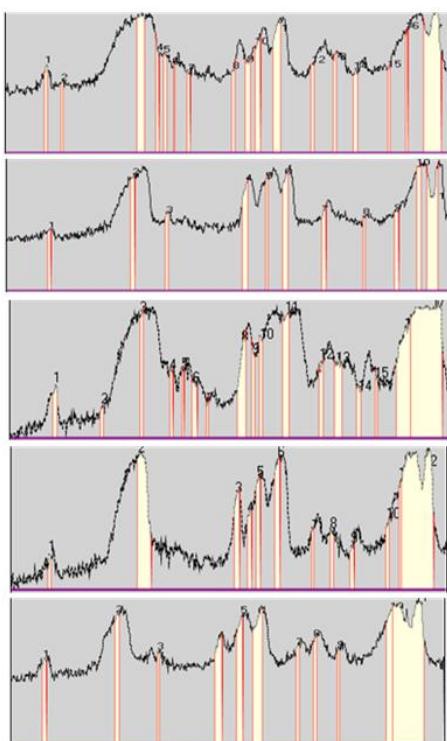
Serial No.	Name of the sample/variety	Protein content mg/g/fw
1	ICGV00351	187
2	TMV-7	176
3	CO-6	193
4	C0-4	162
5	TG374	144

Table :2 Rf Value on different protein bands of groundnut cultivars based on SDS-PAGE

S.NO	Rf-Value	ICGV00351	TMV-7	TG-374	CO-4	CO-6
1	0.328	+	-	-	-	-
2	0.342		+	+	+	+
3	0.385	+	-	-	-	-
4	0.457	+	+	+	+	+
5	0.514	+	+	+	-	-
6	0.528	+	-	+	-	+
7	0.557	+	-	+	-	-
8	0.585	+	-	+	-	-
9	0.642	+	+	+	+	+
10	0.685	+	+	+	+	+
11	0.714	+	+	+	+	+
12	0.742	+	-	+	-	-
13	0.757		-	+	-	-
14	0.785	+	+	+	+	+
15	0.814	+	+	+	+	+
16	0.842	-	-	+	-	-
17	0.857	+	+	+	+	+
18	0.871	-	-	+	-	-
19	0.900	+	-	-	-	+
20	0.928	+	+	+	+	+
21	0.971	+	+	+	+	-

Figure 1- SDS-PAGE Analysis of seed proteins in different groundnut cultivars





Densitogram for seed proteins of different groundnut cultivars

IV. DISCUSSION

The quantitative estimation of seed protein contents showed variations among the five varieties of groundnut. Like that of our present study [14] studied the chemical composition of five varieties of groundnut. They suggested that protein content is genetically controlled. It is also influenced by nitrogen fertilizer application and agronomies practices. Similar studies were reported by [15-17] in bambara groundnut (*Vigna subterranean*)[18]. analyzed the changes in protein contents associated with storage of summer groundnut seeds.

In this research work the seed storage proteins were analyzed by SDS PAGE. The SDS-PAGE technique is mostly thought as a reliable means for the reason that total seed proteins are mainly free of environmental variations [19]. Genetic diversity can be easily evaluated through biological markers [20,21]. Protein types and their diversity varied among a variety of crop species, which may assist us for the early detection of species at seed level and to acquire the information on clarity genetic assets [22]

Seed storage have been used as genetic markers obtained by electrophoresis to resolve the taxonomic and evolutionary problems of several crop plants [23]. Sodium Dodecyl Sulphate Polyacrylamide Gel Electrophoresis (SDS-PAGE) is used due to its validity and simplicity for describing genetic structure of crop germplasm, but its implication has been limited mainly to cereals

due to less polymorphism in most of the legumes [24]. Like that of our research work,[25] reported electrophoretic studies on seed protein profile of blackgram, *Vignamungo* (L.) hepper.

In our study, the MW-Rf values were significantly varied among the five varieties of groundnut. The diversity in seed storage protein have been reported by [26] for wheat varieties. According to [27] in mature seeds, the type and amount of proteins are more constant than other plant tissues. So , the SDS-PAGE pattern of seed storage proteins of groundnut showed polymorphism on the basis of difference in protein intensity among the varieties. [28] reported that the presence are absence of protein bands has also been applied for detection of polymorphism of *Brassica* cultivars.

The present investigation revealed that differences in seed protein profiles. The different groundnut cultivars showed some variations. Similar to our finding the result of differentiation of yellow sarson and brown seeds types of *Brassica* clearly separated the yellow seeds and brown seeded varieties by SDS PAGE [29].

Like that of our present study, several researchers such as [30,31] reported significant variation among different species of groundnut for seed storage proteins, therefore it is suggested to use different species of groundnut to increase the genetic diversity of the germplasm. [32] reported low genetic diversity in *Arachis hypogaea*.

V. CONCLUSION

The present study clearly indicated the seed storage proteins in terms of quantitative and qualitative analysis of groundnut cultivars. The MW-Rf values were significantly varied among the cultivars. The presence or absence of MW-Rf values are useful for identification of particular variety of groundnut.

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REFERENCES

- [1]. Pardee, Peanut. Encarta Encyclopedia, Microsoft Corporation, 2002,USA.
- [2]. R.L.Ory and J.P.Cherry, Protein from peanut cultivars (*Arachis hypogaea*) grown in different areas. V. Biochemical observations on electrophoretic patterns of proteins and enzymes. Journal of American Peanut Research and Education Association, 4(1);1972,32-40.
- [3]. R.Dawson and A.D.McIntosh, Varietal and environmental differences in the proteins of the groundnut (*Arachis hypogaea*). Journal of the Science of Food and Agriculture, 24(5);1973,597-609.
- [4]. J.P.Cherry, Comparative studies of seed proteins and enzymes of species and collections of Arahis by gel electrophoresis. Peanut Science, 2(2);1975,57-65.
- [5]. C.F.Savoy, D.Stoker and S.K.Pancholy, Relatedness of "Jenkins", "Hawthorne" and "Storers Jumbo" peanuts. Proceedings of American Peanut Research and Education Association, Inc., 10(1):1978,76.
- [6]. E.Klozova, J.Svachulova, J. Smartt, E.Hadac, V.Turkova and V.Hadacova, The comparison of seed protein patterns within the genus *Arachis* by polyacrylamide gel electrophoresis. Biologia Plantarum, 25(4);1983,266-273.
- [7]. Y.YChiou, C.S.Hsu, F.Z.Tyan and T.T.Tsai, Physical and chemical study of proteins in various groundnut cultivars grown in Taiwan. Journal of the Chinese Agricultural Chemical Society, 26(2);1988,189-196.
- [8]. T.G.Krishna and R.Mitra, The probable genome donors to *Arachis hypogaea* L. Based on arachin seed storage protein. Euphytica, 37(1)1998,47-52.
- [9]. A.K.Singh, S.Gurtu and R.Jambunathan, Phylogenetic relationships in the genus *Arachis* based on seed protein profiles. Euphytica, 74(3);1993,219-225.
- [10]. M.Nisar, A. Ghafoor, M.R. Khan, H.Ahmad, A.S. Ahmad, Qureshi and H.Ali, Genetic diversity and geographic relationship among local and exotic chickpea germplasm. Pak. J. Bot., 39;2007,1575-1581.
- [11]. S.D.Tanksley and R.A.Jones, Application of alcohol dehydrogenase allozymes in testing the genetic purity of F1 hybrid of tomato. Hort. Sci., 16;1981, 179-1871.
- [12]. V.O.C.Thanh and Y.Hirata , Seed storage protein diversity of three rice species in the Mekong Datta. Biosphere conservation 4;2000, 59-67.
- [13]. M.Ravi, S.Geethanjali, F.Sameefarheen and M.Maheswara, Molecular marker based on genetic diversity analysis in rice (*Oriza sativa* L.)using RAPD and SSR markers. Euphytica. 133;2003,243-252.
- [14]. Mst. Farhana Nazneen Chowdhury, Md.Delwar Hosen and Md. Sanjedur Rahman, Comparative study on chemical composition of five varieties of groundnut(*Arachis hypogaea*L.). W.J. Agri. Sci. 11(5)2015,247-254.
- [15]. N.H.Poulter, Properties of same proteins fractions from Bambara groundnut. J.Sci.Food. Agric.(32;1981, 44-50.
- [16]. I.A.A.M.Onimawo, and A.Usman, Proximate composition and functional properties of four cultivars of bambara groundnut (*Voandzeia subterraneam*). Plant Foods Human natr.53;1998,153-158.
- [17]. J.E.M.Ferrao, A.M.B.C.Ferrao and A.M.G.Antunes, Bambara groundnut (*Vinga subterraneam*) aspects of its nutritional value. Gracia Deorta Serie de Estudios Agronomicos.(14);1987, b35-39.
- [18]. V.B.Yadav, R.W.Bharur, and D.R.Nagawade, Biochemical changes associated with storage of summer groundnut (*Arachis hypogaea* L.) seeds. Journal of Crop Science.5(1);2014,112-115.
- [19]. A.Javaid, A.Ghafoor, and R.Anawar, Seed storage protein electrophoresis in groundnut for evaluating genetic diversity. Pak. J. Bot., 36;2004,25-29.
- [20]. M.A.Rabbani, A.A.Qureshi, M.Afzal, R.Anwar and S.Komatsu, Characterization of mustard (*Brassica juncea* (L.) Cezern. & Cross) germplasm by SDS-PAGE of

- total seed protein. Pak. J. Bot, 33;2001,173-179.
- [21]. M.Akhtar, Phylogenetic relationship among species based on agronomic and biochemical analysis. M. Phil. Thesis, Department of Biological Sciences, Quaid-I-Azam University, Islamabad, Pakistan, 2001,99p.
 - [22]. M.A.Rahman and Y.Hirata, Genetic diversity in Brassica species using SDS-PAGE analysis. Biol.Sci. 4;2002,239-242.
 - [23]. G.Ladizinsky and T.Hymowitz, Seed protein electrophoresis in taxonomic and evolutionary studies. Elet.J.Bio.5(1);1979,1-4.
 - [24]. A.Ghafoor, Z.Ahmad, A.S.Qureshi and M.Bashir, Genetic relationship in Vigna mungo (L.) Hepper and V. Radiata (L.) Wilczek, R., based on morphological traits and SDS-PAGE. Euphytica, 123;2002,367-378.
 - [25]. C.Kole, Indian J.Genet.,62;2002,345-346.
 - [26]. M.F.Khan, E.Schumann and W.E.Weber, Characterization of Pakistan wheat varieties for general cultivation in the mountainous regions of Azad Kashmir. Asian journal of Plant Science, (16); 2000,699-702.
 - [27]. A.Zeb, A. Zahir, T.Ahmad and A.Abdumanon, Physiological characteristics of wheat varieties growing in the same and different ecological regions of Pakistan. Pakistan journal of Biological science, 9(9);2006, 1823-1828.
 - [28]. M.Sadia, A.M.Salman, M.A.Rabbini and S.R.Pearce, Electrophoretic characterization and the relationship between some Brassica species. Electronic journal of Biology, 5(1);2009,1-4.
 - [29]. S.Das and K.K.Mukherjee, Comparative study on seed proteins of Ipomoea. Seed Science and Technology,23(2);1995,501-509.
 - [30]. P.G.Lahman, B.P. Forster, P.McNicol, J.P.Moss and W.Powell, Seed storage protein variation in Arachis species. Genome, 37(3);1994,487-496.
 - [31]. M.R. Berotzo and J.F.M.Valls, Seed storage protein electrophoresis in Arachis pintoi and A.repens (Leguminosae) for evaluating genetic diversity. Genetic Resources and Crop evolution. 48(2); 2001,121-130.
 - [32]. A.K.Singh, S. Gurtu and R.Jambunathan, Phylogenetic relationships in the genus Arachis based on seed protein profiles. Euphytica, 74(3);1993, 219-225.