Biochemical, Morphological and Molecular Evaluation of Nine Fenugreek Landraces

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8 ABSTRACT

Aim: Identification of plant genotypes is an important process to register the plant cultivars, protect breeder's right, maintain the genotype genetic purity, perform the field inspection as a supportive method to seed analysis and protect seed industry. So, the objective of this work was to distinguish among nine landraces of fenugreek (Trigonella foenum graecum L.) at the seedling, chemical, biochemical, and molecular levels.

Methodology: Germination percentage and seedling vigor characteristics were tested using ISTA rules, seed chemical composition content was measured. SDS-PAGE and RAPD-PCR methods were used for biochemical and molecular differentiation among the genotypes, respectively.

Results: The results of seedling characteristics revealed that there is no significant difference among the genotypes in the germination percentage. Genotype-8 had the highest seedling vigor index, while genotype-10 had the lowest one. Chemical composition such as moisture content, crude protein content, oil content, ash content, crude fiber contents, and carbohydrates were analyzed. SDS-PAGE revealed a total of 21 bands with molecular weight (mw) ranging from 241.7 to 6.5 kDa. Eleven out of 21 were polymorphic bands and seven unique markers were found, four of them were positive and the others were negative. RAPD-PCR revealed a total number of 103 DNA bands were detected as generated by 8 random primers, in which 64 were polymorphic bands. Twenty two unique RAPD markers were found, which all of them were positive.

Conclusion: Present investigation provided the information about seed germination, seed characters, biochemical and molecular differences of nine Egyptian fenugreek landraces. The results showed that L8 performed well with respect to seedling vigor index and fiber content, while L10 and L14 performed well with respect to protein and oil content, respectively. So, these landraces could be used in the breeding programs for developing the fenugreek.

Keywords: fenugreek, Trigonella foenum graecum L., RAPD, SDS-PAGE, Seed vigor, Chemical 10 analvsis.

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1. INTRODUCTION

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17 Fenugreek (Trigonella foenum graecum L.) is one of the old legumes used as a food and medicinal 18 plant in the Mediterranean region. Actually, it is being widely cultivated in many countries 19 (Petropoulos, 2002). The fenugreek is a high value but low volume crop with multipurpose 20 applications. It is popularly used as spice and its medicinal value is also highly appreciated for diabetes and heart ailments (Suresh Kumar et al., 2005). Although its cultivation was mostly 21 22 concentrated in Asia and the Mediterranean region, it is now widely cultivated in northern Africa and 23 central Europe (Petropoulos, 2002; Basu et al., 2014).

24 Genetic diversity in plant materials results from variations in DNA sequences and environmental 25 effects. In addition, it is used as a resource for re-vegetation of disturbed sites to allow natural 26 selection and adaptation to occur. Therefore, estimation of the genetic diversity among plants is 27 important for the improvement of any crop and for preserving natural variation for adaptation (Mondini et al., 2009). Genetic diversity can be determined using morphological, biochemical, and molecular 28 29 markers (Gonçalves et al., 2008). These markers differ from each other with respect to important 30 features such as genomic abundance, level of polymorphism detected, locus specificity, 31 reproducibility, technical requirements, cost, and the type of data that they generate.

32 Seed storage proteins are deposited in relatively large quantities in mature seeds and typically remain 33 more stable than other plant tissues until they germinate (Mirali et al., 2007). Therefore, proteins can 34 be easily extracted from seeds and analyzed with sodium dodecyl sulfate polyacrylamide gel 35 electrophoresis (SDS-PAGE) technique leading to separation of seed storage proteins into specific 36 banding patterns, which generates higher levels of genetic polymorphisms on the basis of differences 37 in protein intensity among genotypes (Sinha et al., 2012). Additionally, it is a method commonly used 38 to investigate genetic diversity and to classify plant varieties (Kakaei and Kahrizi, 2011), as genetic 39 markers for genetic variation, to detect genetic diversity in cultivated and wild plant species, and to 40 provide information on phylogenetic relationships among accessions (Kumar and Tata, 2010; Emre, 41 2011). The major advantages of this protein marker technique include assessments of codominance, 42 absence of epistatic and pleiotropic effects, ease of use, and a comparatively inexpensive yet 43 powerful method of measuring allele frequencies for specific genes (Mondini et al., 2009). 44 Electrophoretic makers appear to be due to neutral genes which are not linked to any loci that affect 45 the cultivar and value (Vishwanath et al., 2011). Shazia et al. (2011) used SDS-PAGE to analyze 46 seed proteins of 28 fenugreek genotypes. Considerable variation in seed protein composition within 47 most cultivars complicated the use of SDS-PAGE for characterizing cultivars using protein seeds. 48 Even though, there were differences in protein patterns among the genotypes.

49 Molecular markers, particularly DNA genetic markers, are valuable in that they show genetic 50 differences on a more detailed level without interference from environmental influences (Kumar et al., 51 2009), and involve techniques that provide fast results detailing genetic variation and reflecting 52 underlying genetic diversity (Mamatha, et al., 2017). Furthermore, DNA polymorphisms have become the markers of choice for investigating phylogenetic relationships among various plant varieties 53 54 (Martosa et al., 2005), genome identification (Plomion et al., 1995), molecular characterization (Singh et al., 2010) and in development of unique molecular signatures (Sudheer-Pamidimarri et al., 2009). 55 56 RAPD markers are most useful because of low cost, speed and no need of radioactivity (Mohammadi 57 and Prasanna, 2003). It is also used plant population genetic study (Rana and Bhat, 2002), phylogeny, gene tagging, gene mapping (Naghia et al., 2002) assessing genetic variations and 58 identifying hybrids (Jug et al., 2004). Previous studies evaluated genetic diversity among fenugreek 59 60 accessions using molecular markers such as rapid amplified polymorphic DNA (RAPD) and inter-61 simple sequence repeats (ISSRs) (Harish et al., 2011; Sundaram and Purwar, 2011; Sharda et al., 62 2013).

63 The present study aimed to: i) characterize nine fenugreek landraces at the seedling, chemical, 64 biochemical, and molecular levels, ii) examine the genetic variation and polymorphisms among the 65 landraces understudy using SDS-PAGE and RAPD techniques, and iii) estimate the genetic 66 relationships among these landraces.

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2. MATERIAL AND METHODS 68

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70 2.1 Plant material

71 Seeds of nine Fenugreek (Trigonella foenum graecum L.) landraces were provided from the Legume 72 Crops Research Department, Field Crops Research Institute, Agricultural Research Center, Giza,

73 Egypt. These landraces were collected from Beni Suef (L3 and L7), Menia (L5), Asuit (L8), Sohag (9),

74 Giza (L10, L13, and L14), and Fayoum (L11).

76 2.2 Seedling vigor characteristics

77 To estimate the germination percentage and seedling characteristics of the fenugreek, 50 randomly 78 seeds of each genotype were tested as recommended by ISTA (1999). All seeds were surface 79 sterilized by immersion in 0.5% sodium hypochlorite (NaOCI) solution for 5 min to prevent fungal 80 infections and then rinsed three times with sterile water to remove any residual from NaOCI. The 81 sterilized seeds were then scattered on the upper surface of two sheets of sterile Whatman No. 1 filter 82 paper that had been pre-moistened with 10 mL of sterile, distilled water and placed in separate sterile 83 Petri plates (150 mm in diameter x 15 mm deep). The plates containing the seeds were placed in a 84 controlled environment chamber at 20 ± 2 °C for germination. Seed germination was observed daily 85 with water added to each Petri plate as necessary to maintain moisture levels. Seedling development 86 was measured at 15 days after transfer into the Petri plates by monitoring seed germination (ISTA, 87 1999), by measuring seedling stem and root lengths, and determining seedling fresh and dry weights 88 of ten randomly selected seedlings. Seedling vigor index was calculated following the procedure 89 (seedling length in cm x germination percentage) outlined by ISTA (1999). Seedling dry weights were 90 determined after drying the plant seedlings to a constant weight in a hot air oven at 85°C (12 h) 91 (Krishnasamy and Seshu, 1990).

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93 **2.3 Seed chemical composition analysis**

94 The seed chemical composition content (Moisture, protein, oil, fibers, ash and carbohydrate) of the 95 fenugreek genotypes under investigation was measured according to the proceeding outlined by 96 A.O.A.C.(1990).

97 **2.4 SDS- protein electrophoresis**

Sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) technique was used to
 characterize the different genotypes by their protein fingerprint. Protein profiling was carried out
 according to Laemmli (1970) as modified by Studier (1973).

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102 2.5 DNA extraction

103 DNA was extracted from 100 mg of young leaves for each genotype using mi-Plant Genomic 104 DNA Isolation Kit (metabion). The concentration and purity were determined by spectrophotometer.

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106 2.6 RAPD analysis

107 RAPD analysis was carried out according to Williams et al., (1990) using 10-mer oligonucleotide
 108 primers. Eight primers were selected as potentially useful. The codes and sequences of the used
 109 primers are shown in Table (1).

110 PCR reactions were optimized and mixtures (25 µl total volume) were composed of dNTPs (200 µM), 111 Mg Cl2 (1.5 mM), 1x buffer, primer (0.2 µM), DNA (50 ng), and Taq DNA polymerase (2 units). 112 Amplification was carried out in a Thermo Cycler (PTC 200) programmed for 94 °C for 3 min (one 113 cycle); followed by 94 °C for 30 sec, 36 °C for 1 min and 72 °C for 2 min (36 cycle); 72 °C for 10 min 114 (one cycle), then 4 °C (infinitive). Amplification products (15 µl) were mixed with 3 µl loading buffer 115 and separated on 1.3% agarose gel and stained with 0.5 µg/ml ethidium bromide, and visualized 116 under ultraviolet light and photographed. DNA fragment sizes were determined by comparisons with 117 the 100 bp DNA Ladder plus. 118

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121	Table 1. Sequences of the 10-mer RAPD primers (5'-3').
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No.	Code name	5'-3' Sequences	
1	OPC-1	TTCGAGCCAG	
2	OPC-10	TGTCTGGGTG	
3	OPF-4	GAATGCGGAG	
4	OPF-10	GGGCCACTCA	
5	OPA-17	GACCGCTTGT	~
6	OPG-05	CTGACGTCAC	
7	OPAM-01	TCACGTACGG	
8	OPP-05	CCCCGGTAAC	

123 2.7 Data analysis

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The results of SDS-PAGE and RAPD analysis were entered in a computer file as binary matrices where 0 stands for the absence of a band and 1 stands for the presence of a band in each individual sample. Similarity coefficients were calculated according to Dice matrix (Nei and Li 1979). Construction of the dendrogram tree was performed using the unweighted pair group method based on arithmetic mean (UPGMA) as implemented in the SPSS program version 10.

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131 **3. RESULTS AND DISCUSSION**

133 **3.1 Germination and seedling characteristics**

134 Variations in seed germination, shoot and radicle length, fresh and dry weights, and seedling vigor 135 among the nine investigated fenugreek landraces are presented in Table 2. Seed germination ranged 136 from a low of 96% in genotype10 to a high of 100% in the L3, L7, L8, L11 and L14. Results indicated 137 that the root length of genotype 8 was the highest value (8.8 cm), while L14 gave the lowest value 138 (6.1 cm). Shoot length values of genotypes under study indicated that the highest value was recorded for L11 (5.8 cm), while the lowest value was found for L10 (4.5 cm). The highest fresh 139 140 weight value was recorded for L13 (173.2 mg), while the lowest fresh weight value was found for L10 141 (104.2 mg). The dry weight value of the landraces understudy ranged from 10.1 to 13.1 mg for the L3 142 and L7, respectively. Regarding to seedling vigor index, L8 had the highest value (1440), while L10 143 had the lowest value (1047). The variations in germination characteristics and chemical composition 144 could be attributed to the genotype of fenugreek and/or the differences in the environmental 145 conditions, the time of harvesting and the storage conditions. Previous studies (Naidu et al 2011, Farahbakhsh 2012 and Ritu 2016) on different characteristics for fenugreek characterization have 146 147 also reported similar results on the same characters

148 **3.2 Seed chemical composition**

Results in Table (3) showed the seed chemical composition content of nine fenugreek landraces. The highest moisture content was recorded for L 9 (12.51%), while the lowest moisture content was found for L 3 (11.25%). The results indicated that L 10 had the highest protein content (26.23%), while L 7 gave the lowest value (22.6%). The oil results showed that the highest oil content was found for L 14 (6.53%), while the lowest oil content was recorded for L 10 (3.46%). Regarding to the ash content, the results showed that L 11 gave the highest values (7.88%), while L 10 had the lowest value (5.65%). Also, the highest fiber content value was recorded for L 8 (7.46%), while the lowest fiber content

value was found for L 7 (4.48 %). Results indicated that the highest value of carbohydrate content
was recorded for L 7 (50.52 %), while the lowest value was found for L 11 (42.48 %).

Constyne	Germination	Radicle	Shoot	Seedling	Seedling	Seedling vigor	
Genotype	(%)	length (cm)	length (cm)	fresh weight (mg)	dry weight (mg)	index	
L3	100	6.2	5.0	137.2	10.1	1120	
L5	98	6.8	5.3	129.2	10.6	1185	
L7	100	6.7	4.8	144.5	13.1	1150	
L8	100	8.8	5.6	126.5	11.7	1440	
L9	97	6.6	5.2	112.1	11.6	1145	
L10	96	6.4	4.5	104.2	10.2	1047	
L11	100	6.9	5.8	136.9	11.3	1270	
L13	96	7.1	5.5	173.2	12.6	1210	
L14	100	6.1	5.6	141.2	12.2	1170	

158 **Table 2. Germination and seedling characteristics of fenugreek landraces.**

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160 Table 3. Chemical composition analysis of Fenugreek seeds.

Genotype	Moisture	Protein	Oil	Ash	Fiber	Carbohydrate
L3	11.25	23.86	5.73	6.95	5.53	46.74
L5	12.10	24.04	3.68	6.90	5.52	47.76
L7	12.13	22.60	3.63	6.67	4.48	50.52
L8	12.28	24.19	3.51	7.11	7.46	45.45
L9	12.51	24.71	4.04	7.26	5.95	45.53
L10	12.06	26.23	3.46	5,65	5.88	46.72
L11	12.20	25.26	4.86	7.88	7.32	42.48
L13	12.16	23.81	5.91	6.87	4.72	46.53
L14	11.75	22.74	6.53	6.66	4.61	47.71

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Many investigators (Singh et al., 2010; Sumayya et al., 2012; Jignesh et al., 2015) have also reported similar results for the same traits of different fenugreek genotypes. As mentioned previously, carbohydrates, proteins, and lipids are the main component of the seeds, and they are mostly responsible for the functional properties that have made them new ingredients in the development of 166 new products. Total crude protein content is also affected by several factors including genetic factors,

soil type, climatic conditions, region, and fertilizers (Deshpande and Damodaran 1990).

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169 3.3 SDS-PAGE analysis

Protein banding patterns of the studied fenugreek landraces as revealed by SDS-PAGE for the total seed protein are shown in Tables (4 and 5). The data showed that the total numbers of bands in all of the studied genotypes were 21 bands. The total number of bands among genotypes ranged from 12 for L 8 to 18 for L 13. L5, L7, L9, L11 and L14 gave similar number of bands (16 bands). Meanwhile, L 3 and L10 showed similar number of bands (15 bands).

175 Table 4. Molecular weight of SDS-PAGE seed storage protein of fenugreek landraces.

No. bands	M.W	L3	L5	L7	L8	L9	L10	L11	L13	L14
1	241.7	-	-	-	-	+	-	-		
2	225.4	-	-	-	-	-	+	+	+	_
3	203.6	+	+	+	+	+	+	+	+	+
4	185.4	+	-	+	+	+	+	+	+	+
5	154.2	-	+	-	-	- <	2		-	-
6	107.5	+	+	+	+	+	+	+	+	+
7	92.9	+	+	+	+	+	+	+	+	+
8	86.1	-	-	+	-	-	-	-	-	-
9	79.7	+	+	+	-	+	+	+	+	+
10	66.7	+	+	+	+	+	+	+	+	+
11	59.9	+	+	+	+	+	+	+	+	+
12	49.7	÷	+	+	+	+	+	+	+	+
13	36.2	+	+	-	-	+	-	+	+	+
14	28.1	+	+	+	-	+	+	-	+	+
15	24.9	+	+	+	+	+	+	+	+	+
16	21.8	-	+	+	+	-	-	+	+	+
17	16.6	+	+	+	+	+	+	+	+	+
18	13.7	+	+	+	+	+	+	+	+	+
19	11.7	+	+	+	+	+	+	+	+	+
20	9.5	+	+	+	-	+	+	+	+	+
21	6.5	-	-	-	-	-	-	-	+	-

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(+) = band present and (-) = Band absent

177 The molecular weight (MW) of bands ranged from 241.7 kDa for L9 to 6.5 kDa for L13. Also, there are 178 twelve common bands that were found in all landraces. Some landraces contained specific bands 179 which could be used to identify and characterize them among others. For example, each of L9, L5, 180 L7, and L13 had a unique band, which has molecular weight of 241.7, 154.2, 86.1 and 6.5 kDa, 181 respectively. However, band with MW of about 225.4 kDa is present only in L10, L11, and L13. These 182 obtained results could be considered as positive unique marker (PUM). Meanwhile, bands with MW of 183 about 79.7 and 9.5 kDa were found in all landraces except L8. Similarly, bands with MW of about 28.1 184 kDa are found in all landraces except L8 and L11. Also, band with MW of about 36.2 kDa is present in all landraces except L7, L8 and L10. This could be considered as negative unique marker (NUM). The 185 186 data obtained in the present study showed distinct protein polymorphisms in each fenugreek genotype, which may result from base changes in DNA altering protein sites. Therefore, these 187 188 polymorphisms may serve as genetic markers because they can be highly polymorphic and their 189 variability is generally highly heritable. Previous studies (Ahmed et al., 2010; Cheema et al., 2010; 190 Jignesh et al., 2015) found different patterns among fenugreek genotypes using SDS-PAGE.

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Genotype	High MW	Low MW	Total bands	Positive marker	Negative marker
	(kDa)	(kDa)	number		
L3	203.6	9.5	15	$\mathbf{\nabla}$	
L5	203.6	9.5	16	1(154.2)	
L7	203.6	9.5	16	1 (86.1)	
L8	203.6	11.7	12		2 (79.7 and 9.5)
L9	241.7	9.5	16	1 (241.7)	
L10	225.4	9.5	15		
L11	225.4	9.5	16		
L13	225.4	6.5	18	1 (6.5)	
L14	203.6	9.5	16		

Table 5. Total number of bands and the MW of the highest and the lowest bands for the SDS seed proteins in fenugreek landraces.

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196 3.4 RAPD analysis

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198 The eight RAPD primers used in this study displayed marked amplification with distinct bands. The 199 RAPD markers generated by these primers revealed characteristic profiles for each genotype in terms 200 of number and position of RAPD bands (Tables 6 and 7, and Fig. 1). A total number of 103 DNA 201 bands were detected as generated by the 8 random primers for the nine landraces used in the 202 present study, in which 64 (62.12%) were polymorphic bands. However, 39 bands were common 203 (monomorphic) for all landraces. Primer OPF-4 gave the lowest number of bands (5 bands) in which 204 all of them were monomorphic bands, while primer OPAM-01 gave the largest number of bands (18 205 bands) in which 16 out of them were polymorphic with percentage 88.89%. The results revealed 22 206 unique positive markers for all the landraces. Primers OPC-01, OPC-10 and OPF-04 did not show any 207 kind of markers. No negative markers were scored with any primer. These genotype-specific markers 208 can be used in subsequent experiments to detect molecular markers for polymorphic genes with 209 economic importance among these and other genotypes. Hahn et al., (1995) reported that even 210 though RAPD markers are useful for grouping inbred lines with different genetic backgrounds, RFLPs 211 are better for determining the genetic relatedness between lines. Beaumont et al., (1996) reported 212 that the RAPD technique was found to be a powerful method to provide improved probes coverage on 213 a previously created RFLP map and to locate markers linked to chromosomal regions of interest. 214 RAPD markers have been useful in evaluation of genetic diversity and markers assisted selection

215 offers a great opportunity and effectiveness in selecting valuable plant genotypes (Young and Cho

216 2002; Harris, 1999).

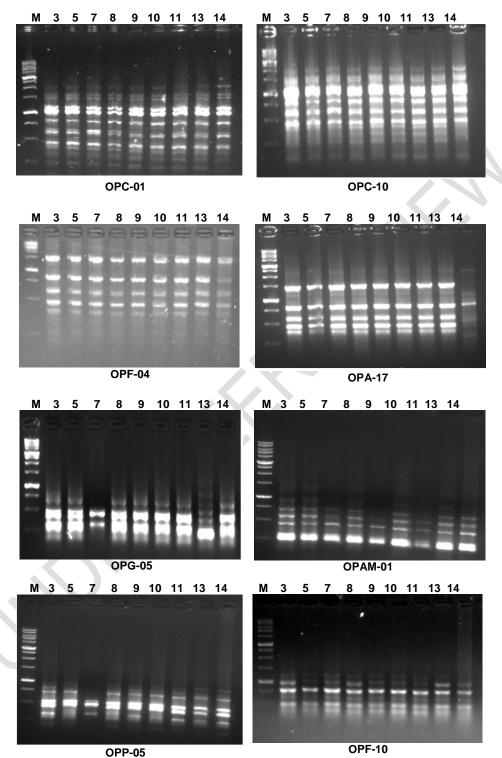


Figure (1): Agarose gel (1.2%) in TAE buffer stained with ethidium bromide showing RAPD-PCR polymorphism of DNA for nine fenugreek landraces (3, 5, 7, 8, 9, 10, 11, 13, and 14, respectively) using eight random primers. M refers to 100 bp DNA Ladder plus.

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221 Although RAPD analysis is guick and well adapted for the efficient non-radioactive DNA fingerprinting 222 of genotypes (Thorman et al., 1994), problems with reproducibility of amplification and with scoring of 223 error data have been reported for RAPDs (Demeke et al., 1997 and Karp et al., 1997). Powell et al., 224 (1996) and Pejic et al., (1998) found the lowest correlations among RAPDs and other marker systems 225 (SSRs, AFLPs, and ISSRs). Pejic et al., (1998) reported that the other DNA markers provide 226 consistent information for germplasm identification and pedigree validation.

227 In conclusion, when we use another PCR-based marker technique such as ISSR, SSR, and AFLP, we 228 might obtain higher information content and consequently higher distinguishably among the used 229 genotypes.

Primer Name	MW (bp)	L3	L5	L7	L8	L9	L10	L11	L13	L1
OPC-01	1399.7	+	+	+	+	+	+	+	+	+
		•	•	•		•	•	\sim		
	1168.3	+	+	+	+	+	+	+	+	+
	1069.0	+	+	+	+	+	+	+	+	+
	848.1	+	+	+	+	+	+	+	+	+
	756.6	+	+	+	+	+	+	+	+	+
	594.6	+	+	+	t	+	+	+	+	+
	467.2	+	+	+	+	+	+	+	+	+
	333.9	+	+	+	+	+	+	+	+	+
	294.1	+	+	+	+	+	+	+	+	+
	237.8	Ŧ	+	+	+	+	+	+	+	+
	209.2	-	-	-	-	+	+	+	+	-
	188.5	+	+	+	+	-	-	-	-	•
	167.7	-	-	-	-	+	+	+	+	•
OPC-10	1449.1	+	+	+	+	+	+	+	+	+
	1297.0	+	+	+	+	+	+	+	+	+
N.	1221.2	+	+	+	+	+	+	+	+	+
	909.6	+	+	+	+	+	+	+	+	+
	737.9	+	+	+	+	+	+	+	+	+
	569.1	+	+	+	+	+	+	+	+	+
	466.1	+	+	+	+	+	+	+	+	+
	412.0	+	+	+	+	+	+	+	+	+

	370.0	-	-	-	-	-	+	-	+	-
	354.3	+	+	+	+	+	-	+	+	-
	304.2	+	+	+	+	+	+	-	-	-
	202.6	+	+	+	+	+	+	+	+	+
OPF-04	1676.7	+	+	+	+	+	+	+	+	+
	985.0	+	+	+	+	+	+	+	+	+
	653.2	+	+	+	+	+	+	+	+	+
	469.7	+	+	+	+	+	+	+	+	+
	367.4	+	+	+	+	+	+	+	+	+
OPA-17	1278.4	+	+	+	+	+	+	+	+	+
	959.5	-	-	-	-	-		-	+	-
	931.1	-	-	-		0	+	+	-	-
	915.7	-	-	-	+	-	-	-	-	-
	900.6	-	-	+		+	-	-	-	-
	882.7	•	+		-	-	-	-	-	-
	836.8	5	X	-	-	-	-	-	-	-
	703.5	+	+	+	+	+	+	+	+	+
	509.0	+	+	+	+	+	+	+	+	+
	377.0	+	+	+	+	+	+	+	+	+
	318.0	+	+	+	+	+	+	+	+	+
	275.7	-	+	+	-	-	-	+	+	-
	265.5	+	-	-	+	+	+	-	-	-
V.	242.7	-	-	-	-	-	-	-	-	+
OPG-05	1481.1	+	-	-	-	-	-	-	-	-
	1464.5	-	-	-	+	-	-	-	-	-
	1448.1	-	-	+	-	-	-	-	-	-
	1405.2	-	-	-	-	-	-	+	-	-
	1375.1	-	-	-	-	+	+	-	-	-

OPAM-01 7 * </th <th></th> <th>1184.3</th> <th></th> <th></th> <th>_</th> <th></th> <th>_</th> <th>-</th> <th>-</th> <th>_</th> <th>-</th>		1184.3			_		_	-	-	_	-
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$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	OPAM-01	724.5	+	-	-	-			-	-	-
613.5 .<		687.6	-	+	-	7	-	-	-	-	-
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$		635.6	-	-	-		-	+	-	-	-
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$		613.5	-	-	-	-	-	-	-	+	-
478.8 - - - + + + -<		528.7	+	+		-	-	-	-	-	-
428.3 - - - - - + + + 410.2 + + + + - - - - - 391.4 - - + + - - - - - 391.4 - - + + - - - - - 360.7 + - - - + + - - - 345.2 - + + - - - + + + + 331.6 - - - + + - - - 311.0 - - - - + + + + + + 300.1 - - - - +		497.9	V	-	-	+	-	-	-	-	-
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		478.8	-	-	-	-	-	+	-	-	-
$\begin{array}{cccccccccccccccccccccccccccccccccccc$		428.3	-	-	-	-	-	-	-	+	+
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$		410.2	+	+	+	-	-	-	-	-	-
$\begin{array}{c ccccccccccccccccccccccccccccccccccc$		391.4	-	-	-	+	-	-	-	-	-
331.6 - - + -<		360.7	+	-	-	-	-	+	+	-	-
311.0 - - + - - - 300.1 - - - + + + - - 289.7 + + + + + + + + +	V	345.2	-	+	+	-	-	-	-	+	+
300.1 - - - + + - - 289.7 + + + + + + + +		331.6	-	-	-	+	-	-	-	-	-
289.7 + + + + + + + + +		311.0	-	-	-	-	+	-	-	-	-
		300.1	-	-	-	-	-	+	+	-	-
279.3 + +		289.7	+	+	+	+	+	+	+	+	+
		279.3	-	-	-	-	-	-	-	+	+

	202.6	+	+	+	+	+	+	+	+	+
OPP-05	477.5	+	+	-	-	-	-	-	-	-
	437.3	+	-	-	+	+	-	-	-	-
	412.3	-	-	-	-	-	+	-	+	-
	397.4	+	+	-	-	+	-	+	-	+
	370.6	-	-	+	+	-	-	-	-	
	359.1	-	-	-	-	+	+	+		-
	330.3	-	-	-	-	-	-	+	+	+
	307.2	+	+	+	+	+	+ ,	+	+	+
	281.0	+	+	+	+	+	+	+	+	+
	244.2	-	+	-	-	-	-	-	-	-
	225.3	+	+	+	+	0		-	-	-
	205.1	+	+	-	~	+	+	-	-	-
	190.1	-	-	-	+	-	-	+	-	-
	180.3	-			-	-	-	-	+	+
OPF-10	573.6		$\overline{}$	+	-	-	-	-	-	-
	562.8	+	-	-	-	-	-	-	-	-
	547.4	-	-	-	-	+	-	-	+	-
	533.2	-	-	-	+	-	+	+	-	-
	474.0	+	+	+	+	+	+	+	+	+
	389.3	+	+	+	+	+	+	+	+	+
	325.3	+	-	+	-	+	-	-	-	-
\mathbf{V}	315.1	-	-	-	+	-	-	+	-	-
	304.4	-	-	-	-	-	+	-	+	-
	280.7	+	+	+	+	+	+	+	+	+
	234.2	+	+	+	+	+	+	+	+	+

Table 7. Total number of bands, monomorphic bands, polymorphic bands, positive markers, negative markers and polymorphism % of nine fenugreek landraces using eight RAPD primers.

Primer	Range	Total	Monomorphic	Polymorphic	Positive	Negative	Polymorphism
Code	size	number	bands	bands	marker	marker	%
	of band	of bands					
	(bp)						
OPC-01	1399.7-	13	10	3	0	0	23.08%
	167.7						
OPC-10	1449.1-	12	9	3	0	0	25.00%
	202.6						
OPF-04	1676.7-	5	5	0	0	0	0
	367.4						
OPA-17	1278.4-	14	5	9	5	0	64.29%
	242.7					\sim	
OPG-05	1481.1-	16	2	14	6	0	87.5%
	312.4					J.	
OPAM-	724.5-	18	2	16	8	0	88.89%
01	202.6						
OPP-05	477.5-	14	2	12	1	0	85.71%
	180.3						
OPF-10	573.6-	11	4	7	2	0	63.64%
	234.2						
Total		103	39	64	22	0	62.12%
Average		12.9	4.9	8	2.8		

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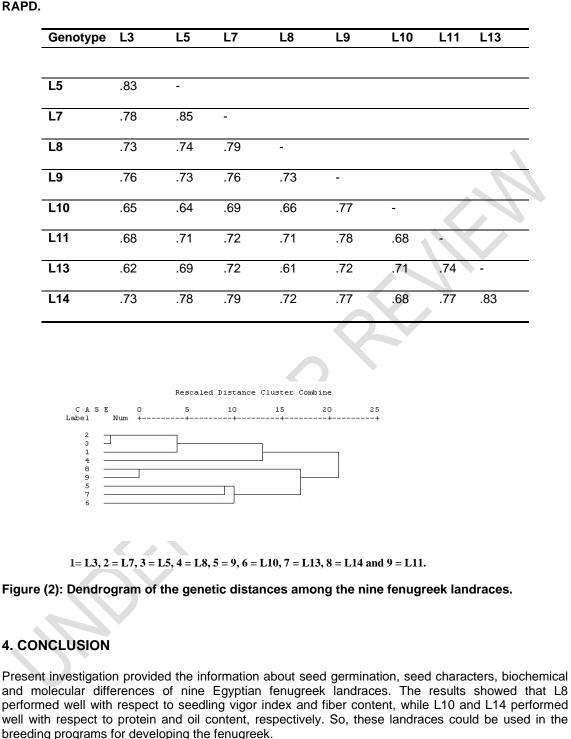
The similarity indices and the dendrogram tree among genotypes utilizing the two methods SDS-PAGE and RAPD are shown in Table (8) and Fig. (2), respectively. The highest percentage of similarity (85%) was scored between L5 and L7, while the lowest percentage of similarity (61%) was scored between L8 and L13. The dendrogram tree divided the nine fenugreek genotypes into two clusters. The first cluster included L3, L5, L7, and L8, while the rest of genotypes were grouped in the second cluster.

3.5 The genetic distance among genotypes

- 253 254
- ___
- 255

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Table 8. Similarity matrix among the genotypes based on combined analysis of SDS-PAGE and



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Ahmed MF, Iqbal M, Masood MS, Rabbani MA, Munir M. Assessment of genetic diversity among
Pakistani wheat (*Triticum aestivum* L.) advanced breeding lines using RAPD and SDSPAGE.
Electronic J. Biotech, 2010;13 (3):1-10.

- 283 AOAC. Official Methods of analysis, of the Assoc. of Official Analytical Chem. U.S.A; 1990.
- Basu A, Basu SK, Kumar A, Sharma M, Chalghoumi R, Hedi A, Francisco SS, Morufat OB, Elsayed
 EH, Cetzal-Ix W. Fenugreek (*Trigonellafoenum-Ggraecum* L.), A potential new crop For Latin
 America. American Journal of Social Issues and Humanities. 2014; 4: 145-162.
- Beaumont VH, Mantet J, Rocheford TR, Widholm JM. Comparison of RAPD and RFLP markers for mapping F₂ generations in maize (*Zea mays* L.). Theor. Appl. Genet., 1996; 93: 606-612.
- Cheema NM, Malik MA, Qadir G, Rabbani MA. Characterization of castor bean genotypes under
 various environments using SDS-PAGE of total seed storage proteins. Pakistan J. Bot. 2010; 42
 (3):1797-1805.
- 292 Demeke T, Sasikumar B, Hucl P, Chibbar RN. Random amplified polymorphic DNA (RAPD) in cereal 293 improvement. Maydica, 1997; 42: 133-142.
- Deshpande SS, Damodaran S. Food legumes: chemistry and technology. Adv. Cereal Sri. Techno.
 1990; 10: 147-241.
- Emre I. Determination of genetic diversity in the *Vicia* L. (Section *Vicia*) by using SDS-PAGE. Pak. J. Bot. 2011; 43: 1429-1432.
- Farahbakhsh H. Germination and seedling growth in un-primed and primed seeds of Fennel as affected by reduced water potential induced by NaCl.Int. Res. J. Appl. Basic. Sci. 2012; 3(4):737-744.
- Hahn V, Blankenhorn K, Schwall M, Melchinger AE. Relationships among early European maize
 inbreeds: III. Genetic diversity revealed with RFLP and pedigree data. Maydica, 1995; 40: 299-310.
- Harish AKG, Ram K, Singh B, Phulwaria M. Molecular and biochemical characterization of
 differentaccessions of fenugreek (*Trigonella foenum-graecum* L.). Libyan Agr. Res. Cent. J. Int. 2011;
 2: 150-154.
- Harris SA. RAPDs in Systematics- A useful methodology? In: Hollingworth P.M., Bateman R.M.,
 Gornall R.J. (Eds.); Molecular systematic and plant evolution pp. 211-228, Taylor and Fransis,London,
 U.K; 1999.
- ISTA. International rules for seed testing. Seed Science & Technol. Proc. Int. Seed Test. Assoc. 1999;
 31(1):1-152.
- Jignesh Patel J, Dhruve J, Talati JG. Biomolecular Characterization of Different Fenugreek Genotypes
 (*Trigonellafoenum-graecum*L.).Int. J. Curr. Microbiol. App. Sci, 2015; 4(6): 201-210.
- Jug T, Dovc P, Pohar J, Snoj A. RAPD analysis as a tool for discriminating (marble trout X brown trout) from hybrid in the zones of hybridization. J. Anim. Breeding Genet. 2004; 121: 156-162.
- Kakaei M, Kahrizi D. Study of seed proteins pattern of *Brassica napus*varieties via sodium dodecyl
 sulfatepolyacrylamide gel electrophoresis. Int. Res. J. Biotechnol. 2011; 2: 26-28.
- Karp A, Edwards K, Bruford M, Vosman B, Morgante M, Seberg O, Kremer A, Boursot P, Arctander P,
 Tautz D, and Hewitt G. Newer molecular technologies for biodiversity evaluation: opportunities and
 challenges. Nature Biotechnol. 1997; 15: 625-628.
- Krishnasmy V, and Seshu DV. Phosphine fumigatio influence on rice seed germination and vigor.
 Crop Sci. 1990; 30:28- 85.
- Kumar OA, Tata SS. SDS-PAGE seed storage protein profiles in chili peppers (*Capsicum* L.). Not.
 Sci. Biol. 2010; 2: 86-90.

- Kumar P, Gupta VK, Misra AK, Modi DR. Potential of molecular markers in plant biotechnology. Plant
 Omics J. 2009; 2: 141-162.
- Laemmli MK. Cleavage of structure protein during assembly of the head bacteriophage T4. Nature, 1970; 227: 680-685.
- Martosa V, Royob C, Rharrabtia Y, Garcia del Morala LF. Using AFLPs to determine phylogenetic
 relationships and genetic erosion indurum wheat cultivars released in Italy and Spain throughout the
 20thcentury. Field Crops Res. 2005; 91: 107-116.
- Mirali N, El-Khouri S, Rizq F. Genetic diversity and relationships in some *Vicia*species as determined
 by SDSPAGE of seed proteins. Biol. Plantarum, 2007; 51: 660-666.
- Mohammadi SA, Prasanna BM. Analysis of Genetic diversity incrop plants Salient Statistical tools
 and considerations. Crop Sci. 2003; 43: 1235-1248.
- Mondini L, Noorani A, Pagnotta MA. Assessing plant genetic diversity by molecular tools. Diversity,
 2009; 1: 19-35.
- Naghia PT, Malik JPS, Pandey MP, Singh NK. Application of RAPD markers for genetic distance
 Analysis of hybrid rice parental lines. Indian J. Genet. 2002; 62(1): 1-4.
- Naidu, MM, Shyamala BN, Naik PJ, Sulochanamma G, Srinivas P. Chemical composition and antioxidant activity of the husk and endosperm of fenugreek seeds. LWT Food Sci. Technol. 2011; 44: 451 456.
- Nei M, Li WH. Mathematical model of studying genetic variation in terms of restriction endonucleases.
 Proc. Natl. Acad. Sci. USA, 1979; 76: 5269-5273.
- Pejic I, Ajmone-Marsan P, Morgante M, Kozumplick V, Castiglioni P, Taramino G, Motto M.
 Comparative analysis of genetic similarity among maize inbred lines detected by RFLPs, RAPDs,
 SSRs, and AFLPs. Theor. Appl. Genet. 1998; 97: 1248-1255.
- Petropoulos GA. Fenugreek- the Genus *Trigonella*. Taylor and Francis, London and NewYork, 2002;
 Pages: 200.
- Plomion C, O'Malley DM, and Durel CE. Genomic analysis in maritime pine (*Pinus pinaster*).
 Comparison of two RAPD maps using selfed and open-pollinated seeds of the same individual.
 Theor. Appl.Genet. 1995; 90 (7-8): 1028-1034.
- Powell W, Morgante M, Andre C, Hanafey M, Vogel J, Tingey S, Rafalsky A. The comparison of RFLP, RAPD, AFLP, and SSR markers for germplasm analysis. Mol. Breed. 1996; 2: 225-238.
- Rana MK, Bhat KV. Genetic diversity analysis in Indian diploid cotton (*Gossipiumspp.*) using RAPD
 markers. Indian J. Genet. 2002; 62(1): 11-14.
- Ritu G. Effect of salt stress on seed germination and seedling growth of *trigonellafoenum-graecum*.
 Int. J. Mendel, 2016; 33(1-2): 3-4.
- Sharda C, Meena RS, Singh R, Vishal MK, Choudhary V, Panwar A. Assessment of genetic diversity
 among Indian fenugreek (*Trigonellafoenum-graecum* L.) Varieties using morphological and RAPD
 markers. Legume Res. 2013; 36 (4): 289-298.
- 360 Shazia E, Anwar R, Masood S. Evaluation of kasurimethi*trigonellafoenumgraecum*l.var.to establish gi 361 right of Pakistan. Pakistan J. Agric. Res. 2011; 24: 1-4.
- Singh P, Singh U, Shukla M, Singh RL. Variation of some phytochemicals in methi and saunfplants at
 different stages of development. J. Her-bal Medicine Toxicol. 2010; 4(2): 93-99.

- Singh P, Singh S, Mishra SP, and Bhatia SK. Molecular characterization of genetic diversity in *Jatropha curcas* L. Genes Genomics, 2010; 4: 1-8.
- 366 Sinha KN, Singh M, Kumar C. Electrophoretic study of seed storage protein in five species of 367 Bauhinia. J. Pharm. Biol. Sci. 2012; 4: 8-11.

Sudheer-Pamidimarri DV, Singh S, Mastan SG, Patel J, Reddy MP. Molecular characterization and
 identification of markers for toxic and non-toxic varieties of *Jatropha curcas* L. using RAPD, AFLP and
 SSR markers. Mol. Biol. Rep. 2009; 36: 1357-1364.

- 371 Sundaram S, Purwar S. Assessment of genetic diversity among fenugreek (*Trigonella foenum-*372 *graecum* L.), using RAPD molecular markers. J. Med. Plants Res. 2011; 5: 1543-1548.
- Studier FW. Analysis of bacteriophage T1 early RNAs and proteins of slab gels. J. Mol. Biol. 1973; 79:
 237 -248.
- Sumayya AR, Sivagami S, Nabeelah A. Screening and biochemical quantification of phyto-chemicals in fenugreek (*Trigonella foenum-grae-cum*). Res. J. Pharm. Biol. Chem. Sci. 2012; 3(1): 165-169.
- 377 Suresh Kumar G, Shetty AK, Sambaiah K, Salimath PV. Antidiabetic property of fenugreek seed 378 mucilage and spent turmeric in streptozotocin- induced diabetic rats. Nutr. Res. 2005; 25:1021-1028.

Thorman CE, Ferreira ME, Camargo LEA, Tivange JG, and Osborn TC. Comparison of RFLP and
RAPD markers to estimate genetic relationship within and among cruciferous species. Theor. Appl.
Genet. 1994; 88: 973-980.

- 382 Vishwanath K, Prasanna KPR, Pallvi HM, Rajendra PS. Identification of tomato (*Lycopersicon* 383 esculentum) varieties through total soluble seed proteins. Res. J. Agric. Sci. 2011; 2: 8-12.
- Williams JK, Kubelik AR, Livak KJ, Rafalski JA, Tingey SV. DNA polymorphisms amplified by arbitrary
 primers are useful as genetic markers. Nucleic Acids Res. 1990;18: 6531-6535.
- 386 Young K, Cho L. Quantitative trait loci Associated with Foliar *Trigonelline* accumulation in *Glycinemax.*, J. Biomed. Biotechnol. 2002; 2(3): 151-157.