



Fingerprint and Molecular Identification of Some Iraqi Date Palm Cultivars Using Inter-Simple Sequence Repeat (ISSR) Markers.

Ahmed Usaff Lafta Aqeel Hadi Abdulwahid¹ Ali Hussan Ali

Dep of Horticulture College of Agriculture University of Basrah

Aqeelhadi6@gmail.com¹, mob: +964-7802174940

Abstract

many of molecular markers used to determination the fingerprint and diversity in date palm. the ISSR markers are more suitable and efficiency to use for this aim, in this paper we are used a seven ISSR markers to find the fingerprinting and diversity for seven Iraqi date palm, which are not mentioned in other papers by this technique, after the DNA extraction from young leaves with reliable quality and quantity, the genomic DNA and ISSR markers were amplified by PCR Special program, then a DNA fragment amplification cultivars were electrophoresis, then calculating the number of mono- and polymorphic bands, which used to calculate a similarity and genetic distance between cultivar, then it are used to build a phylogenetic tree and cluster analysis.

Introduction

Dates palm is one of the most important fruit in Iraq, the Arabian Gulf and a large part of the Arab world, because it is tolerance of the dry and semi-dry environment, as well as it represents a big thing in Iraqis and Arabs live because it is mentioned in the Quran and the Prophet Muhammad peace be upon him, It has great nutritional and economic benefits, fruit may represent an integrated food, the other part of the tree can enter in manufacturing industry useful to humans (Abdulwahid,2011; ,2011) Jaradat).

Date palm tree is a delicious plant, Thus, the process of cross-pollination was necessary to give an economic crop, but the resulting plants from the seed which come from cross-pollination process are non-similar to the parents, therefore there are many cultivars were produced from it. The Al Baker,1972, mention there are more than 600 cultivars can be found in Iraq land.(Abdulwahid,2011).

The diversity of dates palm varieties and their spread over large areas was the reason that a large number of these varieties did not have the right to study and diagnosis, some of them are similar in appearance and fruit traits, but their cultivars take a different name. The other cultivars were similar in names but differ in genetics and appearance. Because the phenotypic trait was affected by environment, their phenotypic trait have made it a weak tool for diagnosis and identification of varieties. Therefore the researcher have discovered a new good tool to identify the varieties depend on the molecular markers which have not been affected by environment or age tissue and give a high polymorphism (Hammadi et al.,2009) ..

DNA-based molecular markers have differed in power, speed, and cost. These techniques include RAPD, AFLP, ISSR, STR and SNP others. In the current study, ISSR is one of the most powerful techniques adopted by many researchers in distinguishing date palm varieties for strength, ease and cost (Zehdi et al.2004 ; Al-Khateeb and Jubrael,2006 Munshi and Osman,2010)

In order to preserve date palm cultivars scattered in Iraq from loss without knowing its genetic and phenotypic characteristics and to prevent the confusion of labels and the formation of a relational database to resolve disputes over naming and to preserve ownership

of varieties, this research was carried out to document the genetic distance and draw the fingerprint for each cultivars, we are tack a seven uniq cultivare (Aumrani, Jamal aldeen, Stwi, Umalblaliz, Sultini, Azraq Azraq and Asrasi) of date palm which mentioned a few in researchs.

Materials and Methods

The genomic DNA was extracted from the young leaves of seven dates palm trees (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi) using a ready- plant kit to extract the DNA from the leaves. The younger white leaves which near the center of sout tip of date palm trees, were selected for low concentrations of chlorophyll and low fiber. They were transferred to the laboratory and sterilized with ethylene alcohol 70% using medical cotton and then cut into small pieces and grind to powder by liquid nitrogen, then transfer in to vial til used.

Extraction of genomic DNA using peq_LAP kit from the VWR US company, as described in Instruction Book. The genome extract was trsted by electrophoreses with 2% agarose. The purity of the genome produced using Nanodrop was evaluated at a wavelength of 260 and 280 nanometers.

PCR program with ISSR markers:

Seven ISSR primer which used (814, 844, hb09, hb10, hb12, Is02 and IS71). As the instraction book company for reaction mixture, showed that the sample size of the reaction was 25ml containing Green Mix, 12.5 ml, 2 ml of primer, 8 ml genomic DNA and then complete the volume to 25 microliters with free ions distilled water. The PCR program was descpied in table (1) .The PCR product was electrophoreses by 1.5% agarose.

Table (1) The PCR program

T	stage	Temperature	Duration time	no of circle
1	Pre – Denaturation	95	5 min	1
2	Denaturation	95	1 min	40
3	Annealing	36	1 min	
4	Extension	72	1 min	
5	Final extension	72	5 min	1

The agarose gel results were analyzed by photocapt programe package, the expressed of spand, when the band is appare give (1), while the absence band of the was expressed as (0), the Polymorphism , monomorphesim, efficiency, diagnostic power and number of unique pands were calculated and reselt was descased.

Results and discussion

Primer 814

The results showed Fig (1) that the primer 814 gave a 40 total bands, the F3 and F1 cultivars were given a maximum number of bands (7 bands), whereas the f7 cultivar was given a minimum bands, the maximum average of band by cultivars equal 5.71 band/cv, the cluster f5 and f3 which contain two cultivars was equal in genetic distance with cluster f6 and f7 which reach 1.73, this is a nearest distance between two cultivars, but the long distance between two cultivars were recorded by f2 with f3, this is very clear in PCA fig. These results were agreed with Yonis et al.(2008) on Egypt date palm when found there were a high genetic diversity between seven cultivars and found there was 87% of this cultivars were difference between them and The results are consistent in one aspect with the researchers (Hussein, 2005 ; Eissa, 2009; Hamza, 2011 and Haider, 2012).

Primer 844

Fig (2) shown the 844 primer gave 51 total bands, the average band per cultivar was 7.28 (band / class). The F2 cultivar was given a maximum bands reach 9 bands whereas the f1 cultivar given a minimum band (6 bands), all of these bands was a polymorphic band and non recorded the monomorphic band at this cultivar when tested with primer 844. The cluster of f1 and f2 have a nearest distance between two cultivars reach 1.73, then the cluster of f3 and f4 cultivar with distance reach 2.00, while the long distance between two cultivars were noted at f7 with f1 and f3 which recorded 3.60. from principle component analysis fig. noted the long distance between (f5 , f6 and f7) cultivars and (f1, f2, f3 and f4) because the Principal component analysis (PCA) is a statistical analysis used an orthogonal transformation to reducing a number of observations of possibly correlated variables in to linearly value variables can represented in two dimensional fig. These results were agreed with Rania et al., (2008) when using this primer on seven Dates palm cultivars in Egypt, the maximum band is 7 band, but using this primer In Qatar on several date palm cultivars by Ahmed and al Qaradawi (2008) who conclude their ISSR primer 844 is a good tool to cultivars diagnostics.

ISSR 814

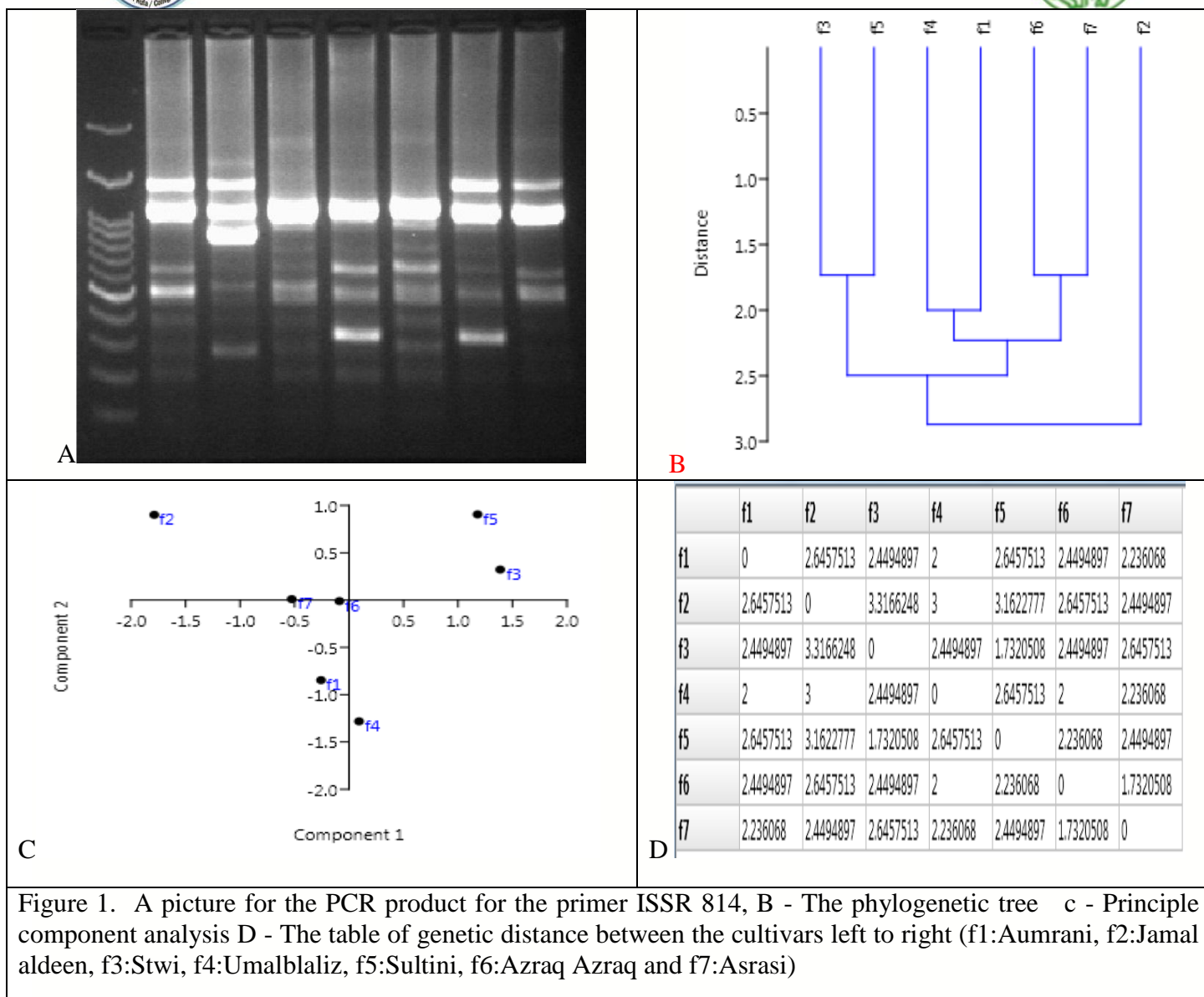


Figure 1. A picture for the PCR product for the primer ISSR 814, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

ISSR844

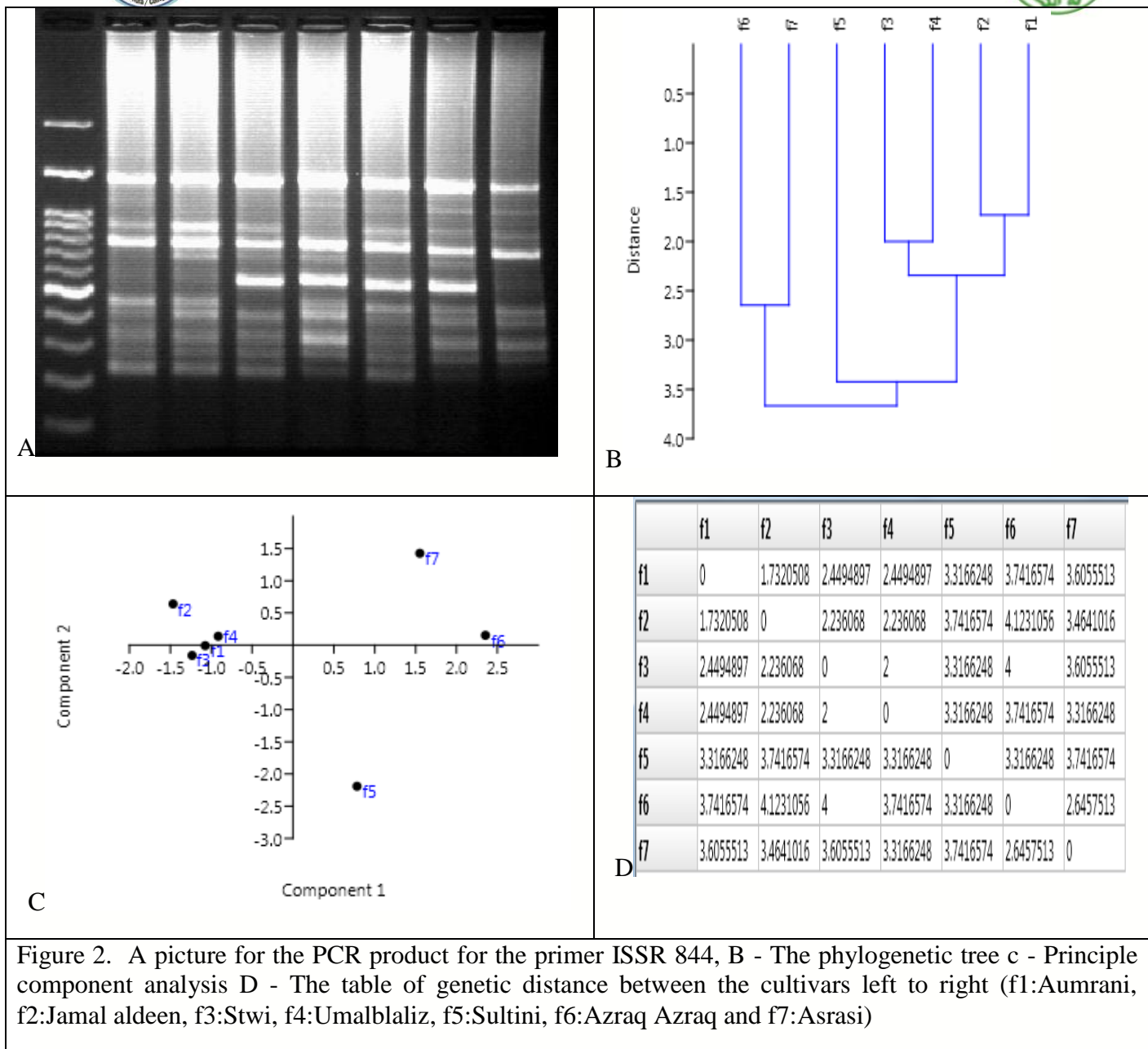


Figure 2. A picture for the PCR product for the primer ISSR 844, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umablalaz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

primers hb9, hb10 and hb12

Figures (3, 4 and 5) show the primers hb9, hb10 and hb12. The total bands recorded were 52, 33, and 61 with average 7.42, 4.71 and 8.71 respectively. The nearest distance was recorded by f3 and f5 in primer hb10 which means there are no differences between these cultivars and the primer hb10 did not recognize the difference between these cultivars, because they have 5 similar bands between 425-1100pb. Then the second genetic distance recorded by primer hb9 between f1 and f6, which reached 1.00 then f6 and f7 in hb12 which reached 1.41. The long distance was recorded between f3 and f7 when used the hb9 reached 3.60, then the second long distance (3.31) was recorded by more on cultivar under three primers which were used. The results are consistent in one aspect with the researchers (Karim et al, 2010 ; Moghaieb, 2010 and Srivastav, 2014).

ISSRhb09

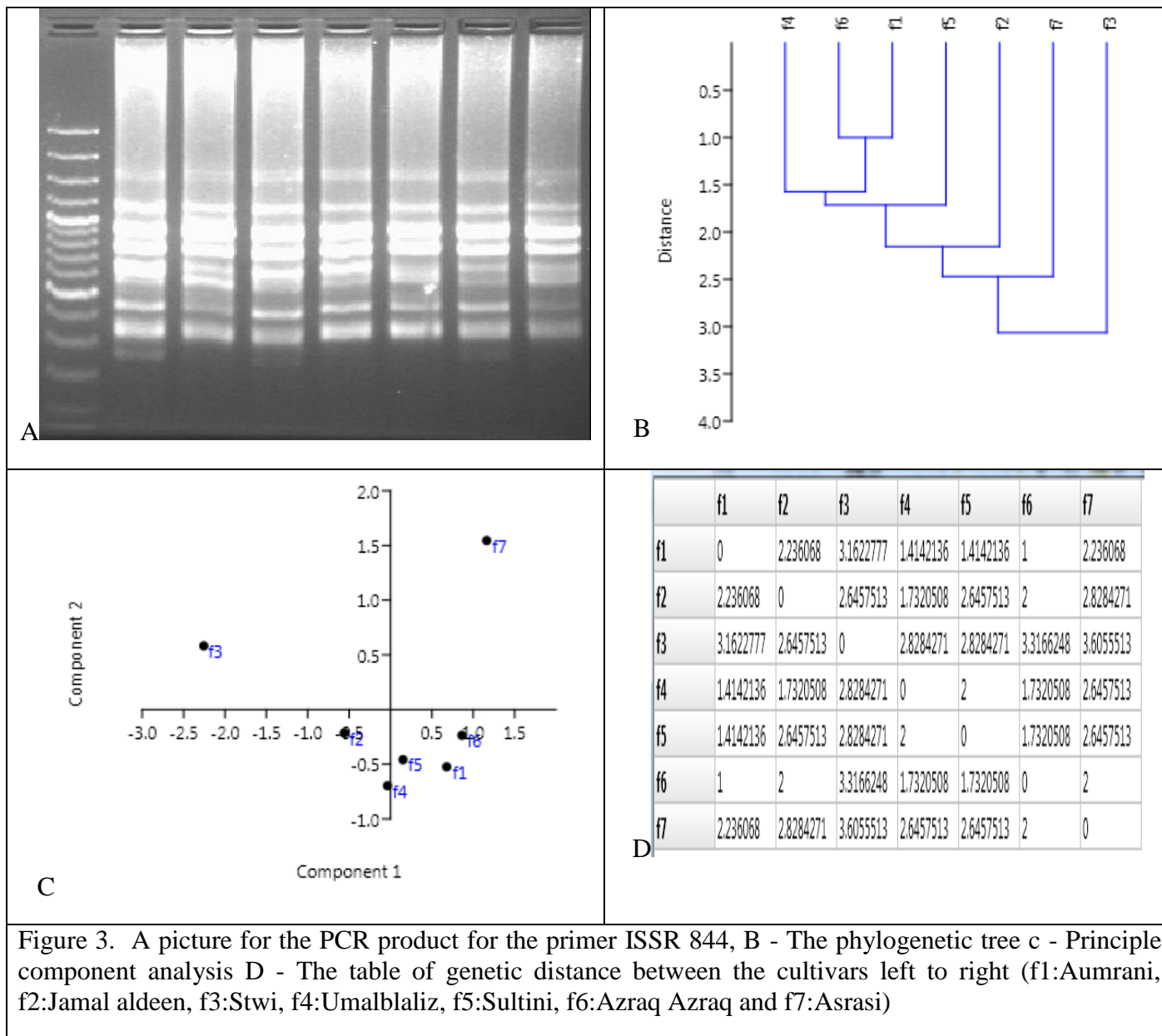


Figure 3. A picture for the PCR product for the primer ISSR 844, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

ISSR hb10

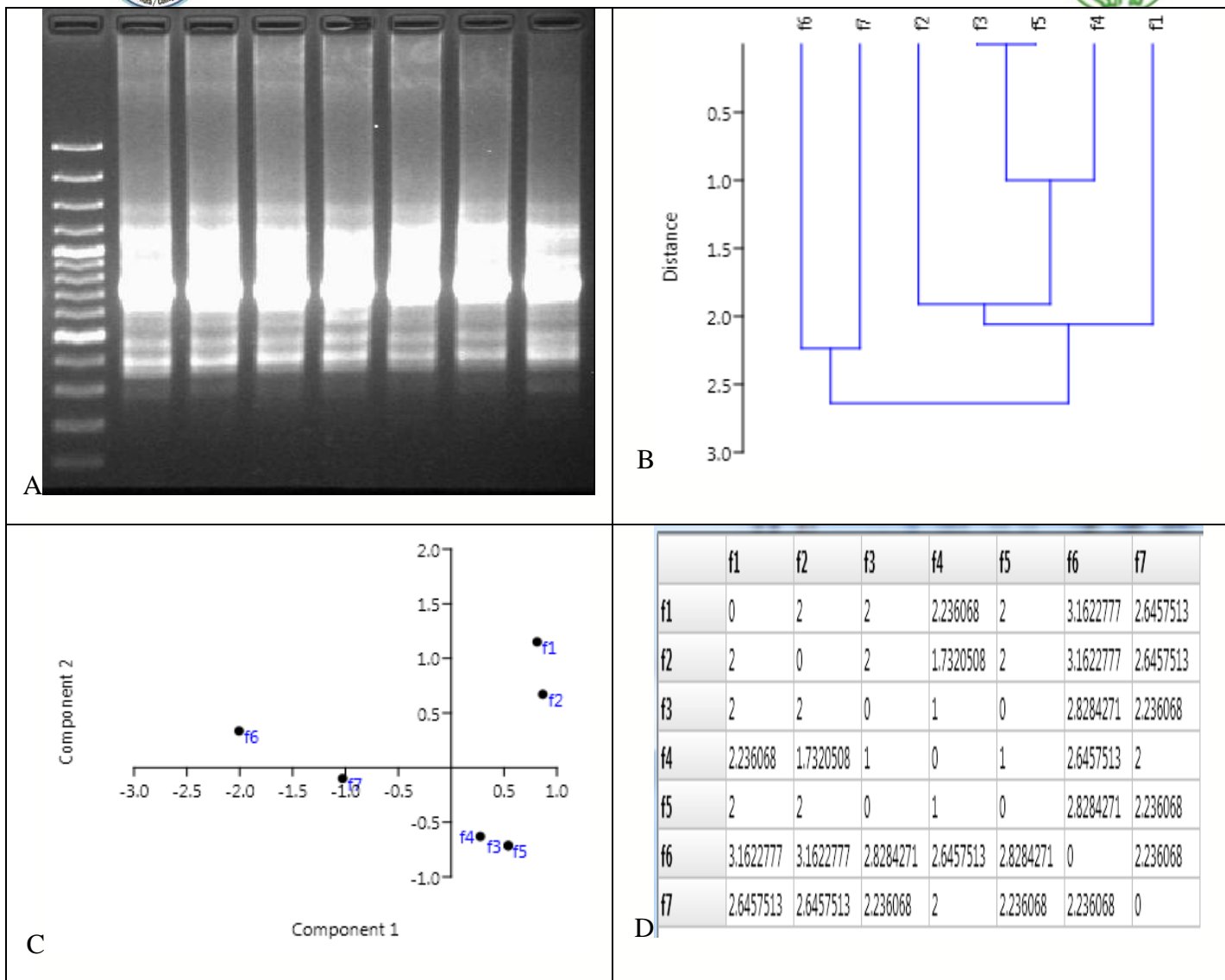


Figure 4. A picture for the PCR product for the primer ISSR hb10, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

ISSR hb12

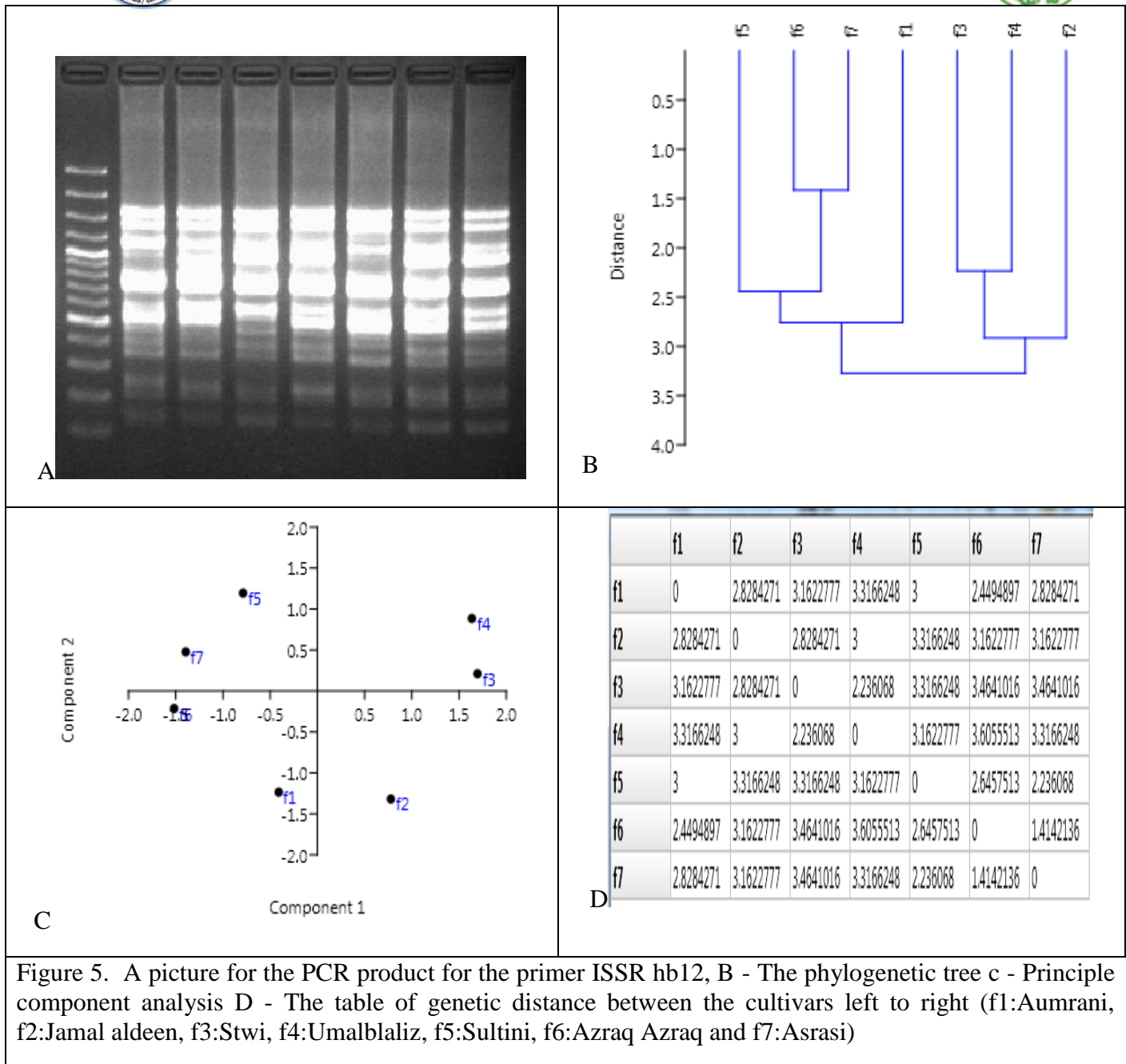


Figure 5. A picture for the PCR product for the primer ISSR hb12, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

Primer IS02

The primer IS02 and IS71 in figer (6 and 7) were given total pand 43 and 77 respectively, with averag 6.13 and 11.00 bands/cv. the nearest cultivars in primer IS02 was f4 and f5 which recorded a genetic distance 1.00 , wheareas the nearest distance in primer IS71 between f5 and f7 cultivars wthich recorded 2.00, This is illustrated by the PCA fig. It shows that two Close points are genetically close, and farther points are genetically longer distant. The results are consistent in one aspect with the researchers (Mohamed et al, 2014 ; Ibrahim, 2014 and Elsheikh, 2014).

ISSR IS02

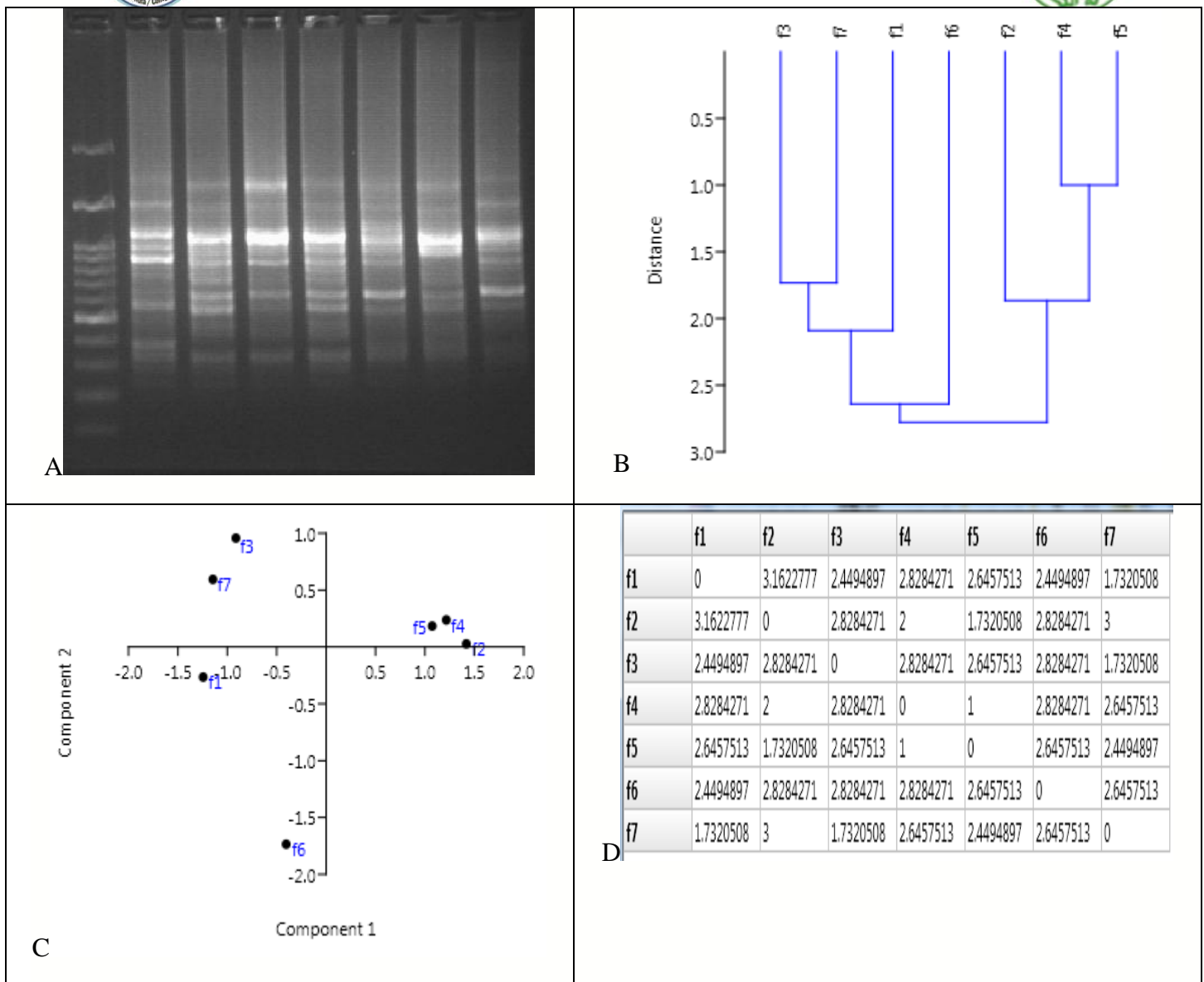


Figure 6. A picture for the PCR product for the primer ISSR IS02, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

Primer IS71

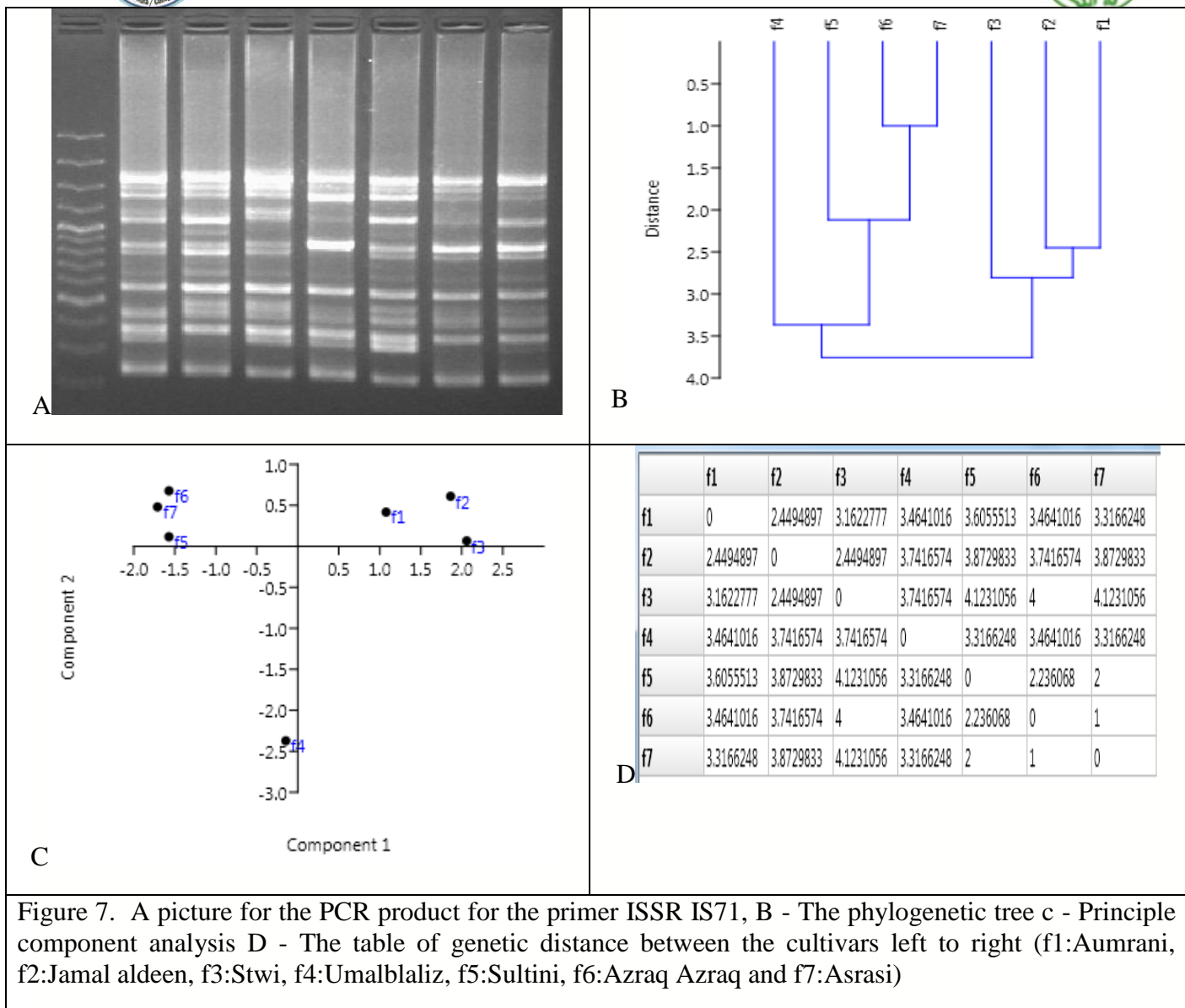


Figure 7. A picture for the PCR product for the primer ISSR IS71, B - The phylogenetic tree c - Principle component analysis D - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umablalaz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

Figure (8) and table (2) shown the cultivares were divided into two clusters, the first cluster was contained f6 and f7 cultivars, whereas the second cluster was contained three sub clusters, the f1 and f2 were a nearest cultivars in one sub cluster, a genetic distance reach 6.55 then the f4 and f5 in another cluster reach 6.70, but the f3 cultivar take the individual cluster. the table (1) shows that the total number of bands produced from the PCR product for the seven primers of the ISSR technique is (357) bands, the hb12 recorded a highest number of band reach 61, whereas the primer 814 recorded a lower number of band reach 40 band, the primer 844 was recorded a highest polymorphic band percent (100%) and good discrimination power. the better unique bands were recorded by primer 844 reach 8 bands with 17.02 unique percent. The results of the paper carry out in Syria on 23 date palm cultivars using the ISSR markers, showed that the phylogenetic tree divided into two clusters, first group (Khastawi, Deglet Noor, and Maktoom) at genetic distance reach 0.30, and the second group included the other cultivars, the researcher addition the ISSR markers is a good tool to find diversity and fingerprint between date palm cultivars (Haider et al.2012).

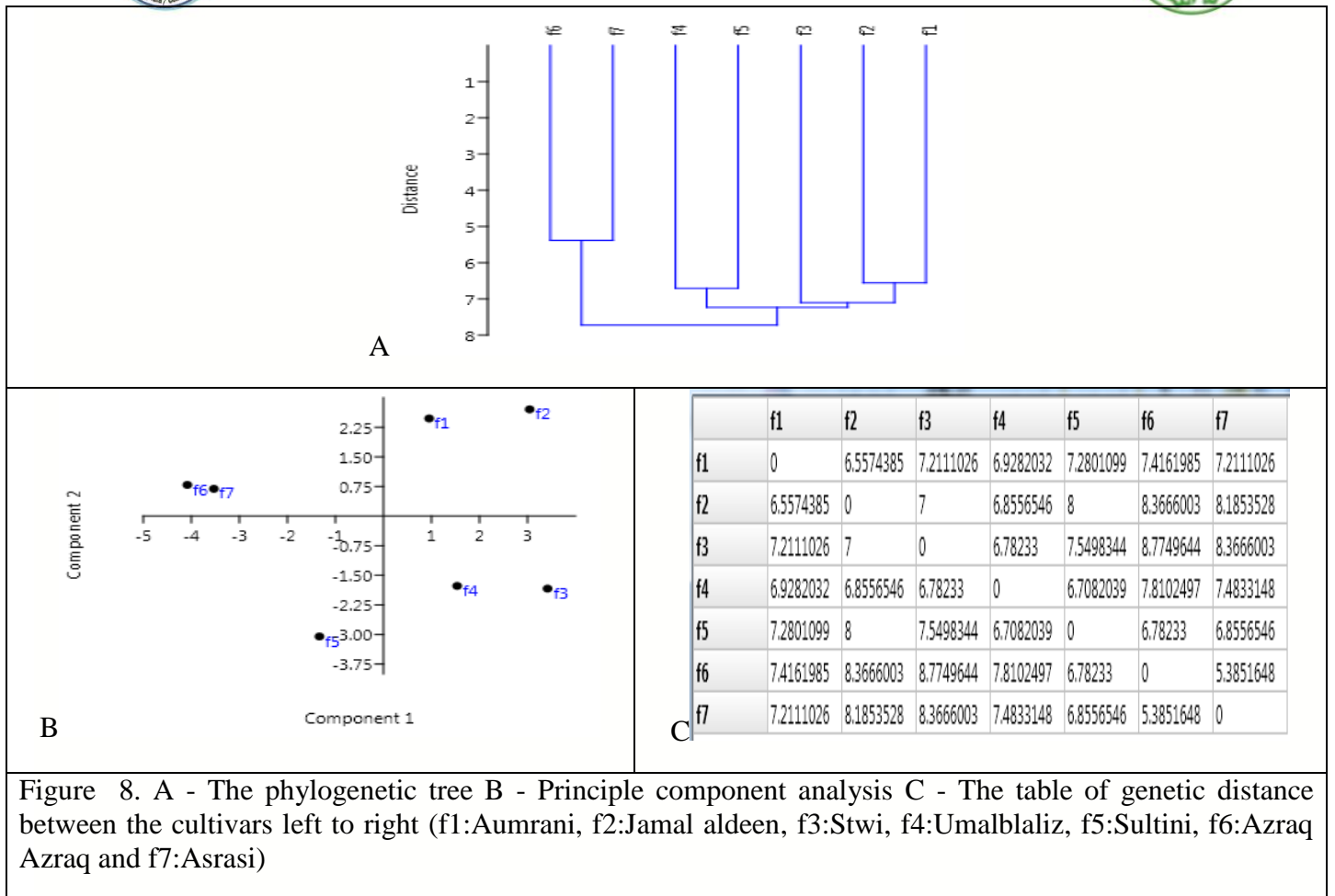


Figure 8. A - The phylogenetic tree B - Principle component analysis C - The table of genetic distance between the cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

Table (2) characteristic of seven ISSR primers on seven date palm cultivars left to right (f1:Aumrani, f2:Jamal aldeen, f3:Stwi, f4:Umalblaliz, f5:Sultini, f6:Azraq Azraq and f7:Asrasi)

No	Peimer Name	Total band	Monomorph bands	Monomorph band percent	Polymorphic bands	Polymorphic bands percent	Unique band	Unique Band percent	Efficient of primer	Discrimination power
1	814	40	2	0.04	45	95.74	6	12.76	11.20	14.24
2	844	51	0	0	47	100	8	17.02	14.28	14.87
3	Hb09	52	2	0.04	45	95.74	4	8.51	14.56	14.24
4	Hb10	33	1	0.02	46	97.87	5	10.63	9.24	14.55
5	Hb12	61	3	0.06	44	93.61	6	12.76	17.08	13.92
6	Is02	43	2	0.04	45	95.74	6	12.76	12.04	14.24
7	IS71	77	3	0.06	44	93.61	6	12.76	21.56	13.92
	total	357	13		316				100	

Refrances:

1. Al-Khateeb T. A. and J. M. Jubrael, (2006). The use of RAPD markers for sex and male cultivars identification in (*Phoenix Dactylifera* L.) in Iraq. *Acta Hort.*, vol. 736, pp. 162–170.
2. Eissa E, Abd El-Razek A, El-Sharabasy S, Rizk R.(2009). Morphological and molecular genetic characterization of soft date palm (*Phoenix dactylifera* L.) cultivars in Egypt. *Egyptian Journal Genetics and Cytology*. 38: 269-284.
3. Elsheikh, M. H.; Abd El- Motty, E. Z.; Elsabagh, A. S. and Yassin, R.(2014). Fruit properties and molecular characterization using ISSR markers of six Libyan date palm cultivars. *Research Journal of Agriculture and Biological Sciences*, 10(1): 47-52.
4. Haider, N. ; Nabulsi, I. and Mir Ali, N. (2012). Phylogenetic relationships among date palm (*Phoenix dactylifera* L.) cultivars in Syria using RAPD and ISSR markers. *Journal of Plant Biology Research*, 1(2): 12-24.
5. Hammadi, H.; Mokhtar, R.; Mokhtar, E.and Ali, F.(2009). New approach for the Morphological Identifcation of date palm (*Phoenix dactylifera* L.) cultivars from Tunisia. *Pak. J. Bot.*, 41(6) : 2671-2681.
6. Hamza, H. ; Elbakkay, M.; Ben Abederrahim, M. A. and Ferchichi Ali, A. (2011). Molecular and morphological analyses of date palm (*Phoenix dactylifera* L.) subpopulations in southern Tunisia. *Spanish Journal of Agricultural Research* , 9(2): 484-493.
7. Hussein, E. H. A., S. S. Adawy, S. E. Ismail, and H. A. El-Itriby (2005). "Molecular characterization of some egyptian date palm germplasm using RAPD and ISSR markers," *Arab J. Biotech.*, vol. 8, pp. 83-98, 2005.
8. Ibrahim, I. A. ; Hashem, M. H. ; Hemeida, A. A.; Hassan, M. M. and Maksoud, A..A.(2014). Characterization of genetic diversity of Date palm (*Phoenix dactylifera* L.) cultivars collected from New Valley governorate (El-Kharga and Dakhleh) based on morphological variability and molecular markers. *Life Science Journal* ,11(11).



9. Karim, K., B. Chokri, H. Amel, H. Wafa, H. Richid, and D. Nouredine (2010). "Genetic diversity of Tunisian date palm germplasm using ISSR markers," *Int. J. Bot.*, vol. 6, pp. 182-186.
10. Moghaieb REA, Abdel-Hadi AA, Ahmed MRA, Hassan AGM. (2010). Genetic diversity and sex determination in date palms (*Phoenix dactylifera* L.) based on DNA markers. *Arab Journal of Biotechnology*. 132: 143- 156.
11. Mohamed, H.; Elsheikh, A.S. ; Abd allah, E. and Ahmed Said Elsabagh (2014). Morphological characterization and genetic analysis by using RAPD and ISSR markers of Some Olive Cultivars grown in egypt. *World Applied Sciences Journal* , 30 (4): 420-427.
12. Munshi and G. Osman,(2010). "Investigation on molecular phylogeny of some date palm (*Phoenix Dactylifra* L.) cultivars by protein, RAPD and ISSR markers in Saudi Arabia," *Aust. J. Crop Sci.*, vol. 4, pp. 23-28.
13. Srivashtav, V. S.; Solanki, V. H.; Patel, H. K.and Kansara, R. V.(2014). Molecular characterization of Date Palm (*Phoenix dactylifera*) using combined marker analysis grownin kutch region of India. *Journal of Cell and Tissue Research* , 14(1) 4083-4088.
14. Zehdi, S. H. Sakka, A. Rhouma, S. A. Ould Mohamed, M. Marrakchi, and M. Trifi, (2004). "Analysis of Tunisian date palm germplasm using simple sequence repeat primers," *Afr. J. Biotechnol.*, vol. 3, pp. 215-219.