



Characterization of genetic diversity of cactus species (*Opuntia spp.*) in Morocco by morphological traits and molecular markers

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Abstract

The genetic diversity within and among 124 accessions of *Opuntia spp.* collected from different regions of Morocco was assessed using morphological descriptors and molecular markers. Based on 10 morphological traits, the accessions were separated into 3 main clusters; each cluster was containing accessions from different regions and species. Polymerase chain reaction (PCR) was then performed on 22 accessions from different regions and species, with 10 inter-simple sequence repeat (ISSR) primers and one random amplified polymorphic DNA (RAPD) primer. ISSR primers produced 66 bands overall, 64 (96.9 %) of which were polymorphic while 6 bands were generated by the RAPD marker, all polymorphic. The polymorphic information content (PIC) values ranged from 0.62 to 0.97, with an average of 0.82. The dendrogram of genetic differences generated using the unweighted pair-group method using arithmetic averages (UPGMA) method showed 7 different clusters at a similarity of 0.76, which was confirmed by the principal component analysis (PCA). The main conclusion of our work is the high genetic similarity between *Opuntia ficus indica* and *Opuntia megacantha* species in Morocco. Our results will be useful for plant breeding and genetic resource conservation programs.





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Introduction

The cactus pear (*Opuntia spp.*) is a plant genus native to arid and semi-arid regions of Mexico¹. It belongs to the family Cactaceae, which is consisting of about 1500 species³⁰ and it is widely distributed throughout temperate, subtropical and cold regions (Feugang *et al.* 2006). In Morocco, the cactus pear is one of the most important crop plants in the semi-arid zones. It covers an area of about 120000 ha⁵ and it is represented mainly by the species *Opuntia ficus indica*. The accessions of this species are classified into three distinct populations: (i) the Christians' nopal with prickly cladodes used as a field fence; (ii) the Muslims' nopal with inermis cladodes used as a green fodder for cattle and (iii) the Moses' nopal with large inermis cladodes that produce big pears²⁰. The cactus pear population is present throughout the country, from Eastern to Southern Morocco¹⁸, with predominance in Guelmim-Sidi Ifni and Haouz-El Kelâa des Sraghnas regions⁵. Increasing the cactus pear area of cultivation is one of the strategies of the Moroccan Ministry of Agriculture. The cactus pear is a tree with high economic and ecologic value and various uses. It is cultivated for human consumption since it is characterized by important nutritional components⁴². The cactus pear has also many beneficial effects on health, including anti-cancer effect, anti-viral effect and anti-diabetic-effect among others²³. On the other hand, the cactus pear may be utilized as forage due to its nutritional characteristics; along this line, Andrade-Montemayor *et al.* (2011)⁴ reported that cactus cladodes could be used for goats feeding in semi-arid regions. The cactus pear was also successfully used for feeding Santa Inês lambs¹³ and dromedary camel¹⁵. Furthermore, it was reported that some cactus constituents could be extracted and used as food additives or for pharmaceutical and cosmetic purposes³². The process of agricultural intensification and modern agricultural practices has led to genetic diversity degradation and gene pools loss²⁻²⁶. Determining the level of genetic diversity among and within plant populations is primordial for genetic resources conservation and breeding programs²⁴⁻²⁸. The use of morphological descriptors is of great help for genetic diversity assessment¹⁴. In the recent years, molecular markers have been proven powerful to determine the genetic diversity among and within species¹² and could be used as complementary methods to morphological descriptors¹⁴. Along

this line, inter-simple sequence repeat (ISSR) and random amplified polymorphic DNA (RAPD) markers have been widely used to assess the genetic diversity in many species. Both the techniques are characterized by their simplicity, rapidity, ability to show high polymorphism with no prior knowledge of the genome studied³⁶⁻⁴¹. In the recent years, several studies have been carried out on the genetic diversity of the cactus pear. For example, Valadez-Moctezuma *et al.* (2014)³⁹ studied the genetic diversity within and among Mexican *Opuntia* species using RAPD and ISSR markers. In Italy, Labra *et al.* (2003)²⁷ evaluated the genetic diversity in *Opuntia* species with cpSSR and AFLP markers. Zoghalmi *et al.* (2007)⁴⁴ have assessed the genetic diversity of *Opuntia ficus indica* L. in Tunisia using RAPD markers. More recently, Bendhifi *et al.* (2013)⁶ published another work on the genetic diversity of the same species in Tunisia using morphological descriptors and RAPD markers. Despite the high importance of the cactus pear in Morocco, its genetic diversity has been scarcely studied. Therefore, the aim of this investigation was to assess the genetic diversity within and among *Opuntia spp.* from different regions of Morocco, using morphological descriptors as well as ISSR and RAPD markers.

Materials and Methods

Plant Material

In the following investigation, we used 124 *Opuntia spp.* accessions belonging to different species: *O. ficus indica*; *O. megacantha*; *O. robusta*; *O. aequatorialis*; *O. dillenii*; *O. leucotricha* and *O. inermis* (identified for the first time in Morocco; Table 1). The accessions are planted in the experimental site called Ain Nzagh of National Institute of Agronomic Research-Regional Research Centre of Settat, (INRA-CRRA Settat, Morocco). Originally, the accessions were collected from different geographical sites of Morocco as shown in Table 1 and Figure 1 then planted in the experimental site in June 2011. All the accessions have undergone the same culture edaphoclimatic conditions.

Morphological Analysis

In this study, 10 quantitative morphological traits including plant height, plant diameter, average number of cladodes per plant, cladode height, cladode width, index of cladode shape (calculated as the ration of one-year-old cladodes length divided

by their width), cladode thickness, mean distance between areoles, average number of spines per areole and mean length of the longest spine per areole. Data were recorded from 4 randomly chosen plants (2 years old) for each accession. Morphological data were subjected to analysis of variance (ANOVA) and means were separated with the Student-Newman-Keuls test at a 0.05 probability level. A proximity matrix was generated using Squared Euclidean distance then clustering of accessions was performed using Ward's method. All statistical analyses were performed with SPSS v 16.0 software (IBM, Chicago, IL, USA).

Molecular Analysis

DNA Extraction

In this study, we used 22 ecotypes from different geographic origins in Morocco, and which represent seven species of *Opuntia* identified from the morphological traits; for example, for *O. ficus indica*, we used 8 accessions, each one was collected from a different region. Total genomic DNA was extracted according to the protocol of Doyle and Doyle (1990)¹⁷ with some slight modifications, using 1 g of lyophilized and ground cladode tissue. The DNA concentration was then diluted to 20 ng/ μ L with TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH 8) using a biophotometer (Eppendorf).

Issr Analysis

Fourteen ISSR primers (Table 2) purchased either from Operon Technologies Inc. (Alameda, CA, USA) or Qiagen (Valencia, CA, USA) were tested in this study; and 2.5 μ L of each primer was added to a PCR mixture containing 3 μ L DNA template, 12.3 μ L sterile distilled water, 3.5 μ L (25 mM) MgCl₂, 1 μ L (10 mM) dNTPs, 0.2 μ L (5u/ μ L) Taq polymerase and 2.5 μ L Taq buffer (All purchased from Promega, Madison, WI, USA). DNA amplifications were performed using a DNA thermal cycler (Techne) with the following parameters: initial denaturation at 94°C for 5 min, 45 cycles of 15 s of denaturation at 94 °C, 30 s of annealing at the specific temperature of each primer (Table 2), and 60 s of extension at 72°C; then a final extension at 72 °C for 7 min.

Rapd Analysis

For RAPD analysis, 34 RAPD primers (Table 2) purchased from The University of British Columbia (UBC; Vancouver, Canada) were tested. The PCR

mixture was containing the same above-mentioned components for ISSR primers, but with substituting ISSR primers with 0.4 μ M of each RAPD primer. Amplification conditions were as follows: initial denaturation at 94°C for 5 min followed by 30 cycles of denaturation at 94 °C for 60 s, annealing at 36 °C for 60 s, and extension at 72 °C for 90 s; with a final extension at 72 °C for 15 min.

Issr and Rapd Gel Electrophoresis

After amplification, 5 μ L bromophenol blue (Sigma, St. Louis, MO, USA) was added to each amplification product. Afterwards, 8 μ L of each sample was electrophoresed at 90 V, 5 W for 1 h on 1.8 % agarose gel (Promega, Madison, WI, USA) in 100 mL TBE buffer (90 mL sterile distilled water, 10 mL TBE buffer 10X; Promega, Madison, WI, USA) supplemented with 6.6 μ L ethidium bromide (Sigma, St. Louis, MO, USA). Molecular weight of the amplified bands was estimated by comparison with lambda DNA/*EcoR* I and Hind III markers (Promega, Madison, WI, USA) as standard. The samples were then visualized and photographed with a gel documentation system (Fisher Scientific).

Statistical Analysis of Molecular Data

For each gel, we recorded the presence (1) or absence (0) of bands in individual lanes; consequently, a binary data matrix was created. For each marker, we calculated the polymorphic

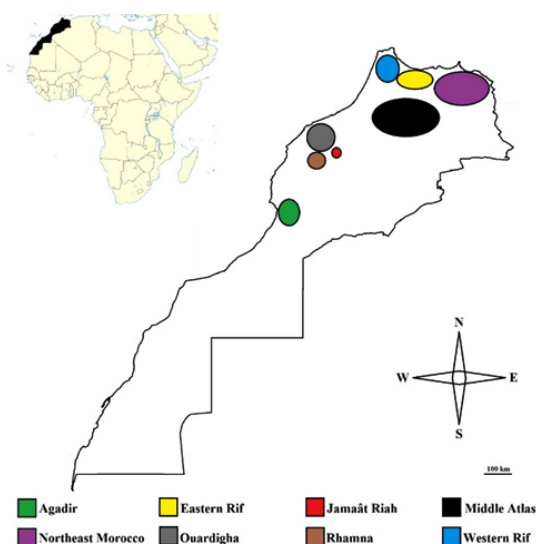


Fig. 1: Map of Morocco showing the geographic origin of the 124 accessions of *Opuntia* spp. used in this study.

information content (PIC) using the formula developed by Botstein *et al.* (1980)⁷:

$$PIC = 1 - \sum_{i=1}^n p_i^2 - 2 \sum_{i=1}^{n-1} \sum_{j=i+1}^n p_i^2 p_j^2$$

where p_i and p_j are the frequencies of the i th and j th alleles and n is the total number of such alleles.

The Dice's coefficients were calculated and a dendrogram was generated based on the similarity matrix and the SAHN module using unweighted pair-group method of arithmetic analysis (UPGMA). Afterwards, a principal component analysis (PCA) was performed. All these analyses were conducted using NTSYSpc v 2.1 software³⁴.

Results and Discussion

Morphological Traits to Assess Genetic Variability

Genetic Variability Within *Opuntia ficus indica*

Our investigation showed that *Opuntia ficus indica* is present in all the studied areas. However, our results demonstrated clear differences among the 78 accessions with regard to morphological descriptors. For example, the plant height was ranging from 21.33 cm in Ofi-531 (from Eastern Rif) to 48.25 cm in Ofi-762 (from Agadir) and the plant diameter was ranging from 16.43 cm in Ofi-275 (from Ouardigha) to 71.38 in Ofi-163 (from Rhamna). In some cases,

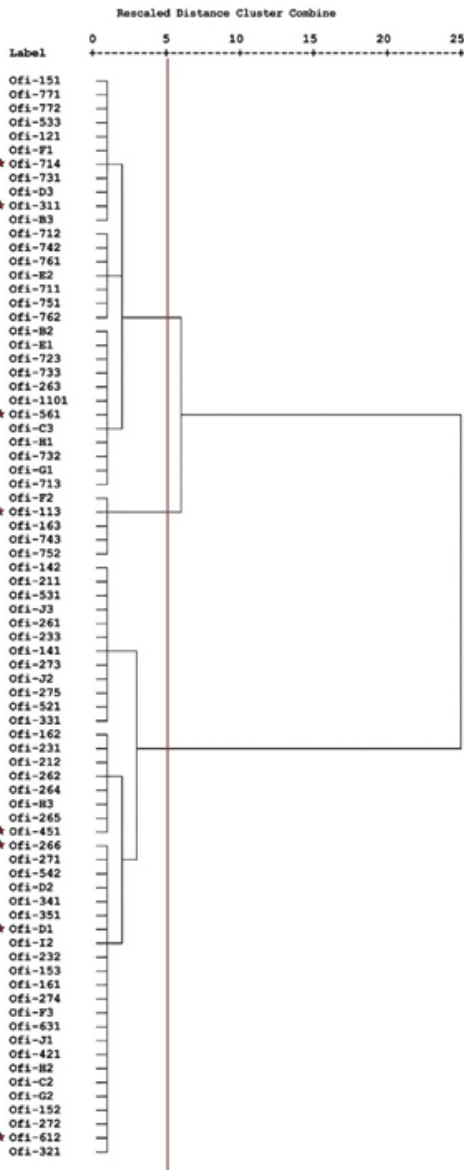


Fig. 2: Dendrogram grouping 78 accessions of *Opuntia ficus indica* based on 10 morphological traits and Ward's method. The red stars indicate accessions used for molecular analysis.

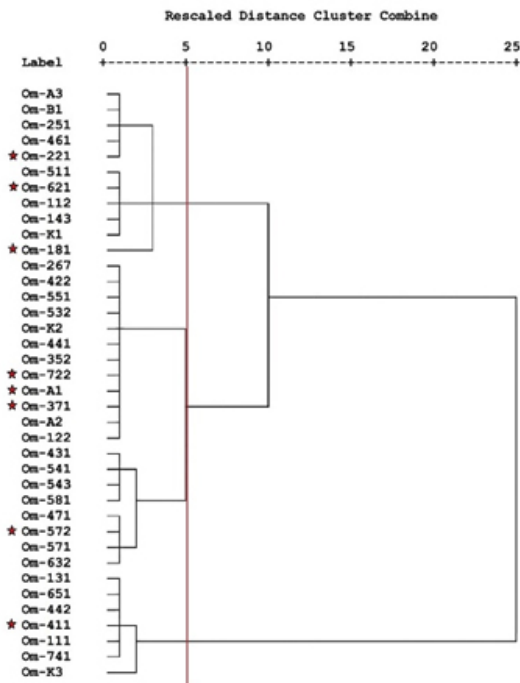


Fig. 3: Dendrogram grouping 38 accessions of *Opuntia megacantha* based on 10 morphological traits and Ward's method. The red stars indicate accessions used for molecular analysis.

Table 1: Means (+SD) of morphological traits in different accessions of *Opuntia* spp. in Morocco
 *Mean values with the same letters are not significantly different (p> 0.05)
 (-): Absence of spine

Species	Geographic origin	Accession	Plant height (cm)	Plant diameter (cm)	Average number of cladodes per plant	Cladode height (cm)	Cladode width (cm)	Index of cladode shape (cm)	Cladode thickness (cm)	Mean distance between areoles (cm)	Average number of spines per areole	Mean length of the longest spine per areole (cm)
<i>O. ficus indica</i>	Jamaât Riah	Off-B2	36.80 +3.16 ^{abcd}	45.08 +1.92 ^{abcd}	3.00 +2.16 ^{def}	26.63 +2.36 ^{bcdefghijkl}	13.63 +2.41 ^{bc}	0.51 + 0.09 ^{bcdef}	0.83 + 0.22 ^{abc}	2.70 + 0.36 ^{bcdefgh}	-	-
	Jamaât Riah	Off-B3	37.10 +2.68 ^{abcd}	47.13 +9.91 ^{abcd}	2.75 +0.96 ^{def}	21.98 +5.54 ^{defghijkl}	10.93 +2.44 ^{bcd}	0.50 +0.06 ^{bcdef}	0.68 +0.21 ^{abc}	2.58 +0.15 ^{cd}	-	-
	Jamaât Riah	Off-C2	33.38 +8.84 ^{abcd}	37.93 +16.12 ^{abcd}	3.00 +0.82 ^{def}	21.80 +0.98 ^{defghijkl}	13.10 +0.33 ^{bc}	0.59 +0.01 ^{bcdef}	0.98 +0.05 ^{abc}	2.90 +0.08 ^{bcdefgh}	-	-
	Jamaât Riah	Off-C3	36.25 +9.81 ^{abcd}	37.00 +6.77 ^{abcd}	1.75 +0.50 ^{def}	24.88 +1.84 ^{bcdefghijkl}	13.25 +1.56 ^{bc}	0.56 +0.02 ^{bcdef}	0.73 +0.13 ^{abc}	2.93 +0.50 ^{bcdefgh}	-	-
	Jamaât Riah	Off-D1	29.50 +4.55 ^{abcd}	28.38 +5.76 ^{abcd}	2.50 +0.58 ^{def}	20.08 +1.05 ^{defghijkl}	11.35 +0.29 ^{bcd}	0.57 +0.02 ^{bcdef}	1.10 +0.00 ^{abc}	2.45 +0.10 ^{def}	-	-
	Jamaât Riah	Off-D2	30.13 +14.77 ^{abcd}	32.38 +14.51 ^{abcd}	2.00 +0.82 ^{def}	28.90 +3.53 ^{bcdefghijkl}	13.19 +3.05 ^{bc}	0.47 +0.05 ^{bcdef}	1.20 +0.00 ^{abc}	2.68 +0.49 ^{bcdefgh}	-	-
	Jamaât Riah	Off-D3	36.38 +3.42 ^{abcd}	51.38 +7.61 ^{abcd}	3.75 +0.50 ^{def}	26.63 +3.38 ^{bcdefghijkl}	13.85 +1.64 ^{bc}	0.52 +0.03 ^{bcdef}	0.90 +0.47 ^{abc}	2.68 +0.31 ^{bcdefgh}	-	-
	Jamaât Riah	Off-E1	37.10 +10.54 ^{abcd}	43.33 +4.19 ^{abcd}	2.25 +1.50 ^{def}	25.13 +2.78 ^{bcdefghijkl}	13.30 +1.72 ^{bc}	0.62 +0.16 ^{bcde}	0.80 +0.29 ^{abc}	3.05 +0.33 ^{bcdefgh}	-	-
	Jamaât Riah	Off-E2	42.75 +4.86 ^{abc}	52.25 +25.16 ^{abcd}	3.75 +1.89 ^{def}	27.43 +1.92 ^{bcdefghijkl}	13.80 +0.68 ^{bc}	0.50 +0.05 ^{bcdef}	0.75 +0.41 ^{abc}	2.90 +0.29 ^{bcdefgh}	-	-
	Jamaât Riah	Off-F1	27.45 +12.12 ^{abcd}	51.95 +32.28 ^{abcd}	3.50 +1.73 ^{def}	28.25 +3.77 ^{bcdefghijkl}	14.28 +1.81 ^{bc}	0.49 +0.04 ^{bcdef}	0.75 +0.41 ^{abc}	3.75 +0.29 ^{abc}	-	-
	Jamaât Riah	Off-F2	37.25 +7.58 ^{abcd}	66.00 +12.93 ^{abcd}	4.00 +1.15 ^{def}	28.15 +4.60 ^{bcdefghijkl}	14.25 +1.94 ^{bc}	0.51 +0.04 ^{bcdef}	0.78 +0.39 ^{abc}	3.18 +0.82 ^{bcdefgh}	-	-
	Jamaât Riah	Off-F3	31.88 +1.89 ^{abcd}	34.00 +5.58 ^{abcd}	2.25 +0.50 ^{def}	22.10 +2.68 ^{bcdefghijkl}	12.50 +0.71 ^{bcd}	0.55 +0.05 ^{bcdef}	0.60 +0.18 ^{abc}	2.78 +0.26 ^{bcdefgh}	-	-
	Jamaât Riah	Off-G1	41.25 +14.31 ^{abcd}	37.28 +25.85 ^{abcd}	3.50 +2.08 ^{def}	27.55 +0.90 ^{bcdefghijkl}	10.13 +2.25 ^{cd}	0.41 +0.08 ^{bcdef}	1.05 +0.12 ^{abc}	3.05 +0.29 ^{bcdefgh}	-	-
	Jamaât Riah	Off-G2	32.63 +1.70 ^{abcd}	30.13 +4.01 ^{abcd}	1.50 +0.58 ^{def}	23.75 +3.77 ^{bcdefghijkl}	13.15 +2.21 ^{bc}	0.55 +0.02 ^{bcdef}	0.63 +0.13 ^{abc}	2.80 +0.24 ^{bcdefgh}	-	-
	Jamaât Riah	Off-H1	39.38 +6.63 ^{abcd}	39.68 +1.89 ^{abcd}	2.50 +0.58 ^{def}	26.45 +4.79 ^{bcdefghijkl}	13.73 +1.39 ^{bc}	0.52 +0.08 ^{bcdef}	0.98 +0.32 ^{abc}	3.10 +0.20 ^{bcdefgh}	-	-
	Jamaât Riah	Off-H2	30.00 +7.67 ^{abcd}	35.75 +5.74 ^{abcd}	1.50 +0.58 ^{def}	24.00 +2.04 ^{bcdefghijkl}	12.85 +0.53 ^{bc}	0.53 +0.02 ^{bcdef}	1.10 +0.16 ^{abc}	2.65 +0.04 ^{bcdefgh}	-	-
	Jamaât Riah	Off-H3	26.30 +6.53 ^{abcd}	37.25 +2.33 ^{abcd}	2.00 +0.00 ^{def}	21.60 +1.96 ^{bcdefghijkl}	10.60 +1.55 ^{bcd}	0.48 +0.03 ^{bcdef}	1.20 +0.08 ^{abc}	2.60 +0.24 ^{bcdefgh}	-	-
	Jamaât Riah	Off-I2	29.88 +1.03 ^{abcd}	29.00 +11.05 ^{abcd}	1.75 +0.96 ^{def}	18.17 +6.18 ^{bcdefghijkl}	11.60 +1.10 ^{bcd}	0.72 +0.25 ^b	0.70 +0.24 ^{abc}	2.80 +0.43 ^{bcdefgh}	-	-
	Jamaât Riah	Off-J1	30.93 +2.69 ^{abcd}	33.00 +8.77 ^{abcd}	2.50 +1.00 ^{def}	21.33 +1.43 ^{bcdefghijkl}	10.43 +1.25 ^{cd}	0.49 +0.04 ^{bcdef}	0.70 +0.22 ^{abc}	2.73 +0.19 ^{bcdefgh}	-	-
	Jamaât Riah	Off-J2	24.90 +10.25 ^{abcd}	27.23 +11.98 ^{abcd}	3.00 +0.82 ^{def}	19.50 +0.41 ^{bcdefghijkl}	10.50 +1.22 ^{cd}	0.53 +0.05 ^{bcdef}	0.90 +0.08 ^{abc}	2.50 +0.41 ^{abc}	-	-
	Jamaât Riah	Off-J3	22.83 +4.28 ^{abcd}	30.50 +3.67 ^{abcd}	2.75 +0.50 ^{def}	18.83 +1.09 ^{bcdefghijkl}	9.88 +2.29 ^{cd}	0.53 +0.14 ^{bcdef}	0.98 +0.22 ^{abc}	2.55 +0.13 ^{cd}	-	-
Rhanna	Off-113	35.00 +3.34 ^{abcd}	62.88 +15.93 ^{abcd}	4.25 +1.50 ^{def}	25.95 +3.09 ^{bcdefghijkl}	13.33 +1.48 ^{bc}	0.51 +0.03 ^{bcdef}	0.75 +0.41 ^{abc}	2.75 +0.31 ^{bcdefgh}	-	-	
Rhanna	Off-121	29.83 +2.17 ^{abcd}	46.75 +10.20 ^{abcd}	6.25 +1.71 ^{ab}	25.28 +3.05 ^{bcdefghijkl}	12.58 +1.26 ^{bcd}	0.50 +0.02 ^{bcdef}	0.88 +0.43 ^{abc}	2.90 +0.22 ^{bcdefgh}	-	-	
Rhanna	Off-141	25.50 +0.00 ^{bcd}	26.50 +3.27 ^{ghi}	1.00 +0.00 ^{def}	16.30 +0.24 ^{bcdefghijkl}	9.50 +0.41 ^{cd}	0.57 +0.05 ^{bcdef}	0.90 +0.08 ^{abc}	2.60 +0.16 ^{bcdefgh}	-	-	
Rhanna	Off-142	24.75 +6.50 ^{bcd}	28.03 +4.01 ^{ghi}	1.00 +0.00 ^{def}	22.60 +1.96 ^{bcdefghijkl}	12.85 +2.16 ^{bc}	0.56 +0.05 ^{bcdef}	1.15 +0.04 ^{abc}	2.65 +0.53 ^{bcdefgh}	-	-	
Rhanna	Off-151	33.93 +3.99 ^{abcd}	45.45 +3.32 ^{abcd}	2.25 +1.26 ^{def}	26.70 +2.52 ^{bcdefghijkl}	12.00 +1.36 ^{bcd}	0.45 +0.08 ^{bcdef}	0.78 +0.22 ^{abc}	3.00 +0.14 ^{bcdefgh}	-	-	
Rhanna	Off-152	31.48 +5.12 ^{abcd}	29.93 +1.34 ^{abcd}	2.50 +0.58 ^{def}	24.10 +3.39 ^{bcdefghijkl}	11.30 +2.43 ^{bcd}	0.45 +0.04 ^{bcdef}	0.80 +0.38 ^{abc}	2.83 +0.46 ^{bcdefgh}	-	-	
Rhanna	Off-153	27.10 +8.23 ^{abcd}	32.78 +20.66 ^{abcd}	2.00 +0.82 ^{def}	22.50 +0.82 ^{bcdefghijkl}	11.29 +2.47 ^{bcd}	0.47 +0.05 ^{bcdef}	1.00 +0.24 ^{abc}	3.23 +0.66 ^{bcdefgh}	-	-	
Rhanna	Off-161	26.28 +5.33 ^{abcd}	31.33 +4.49 ^{abcd}	1.25 +0.50 ^{def}	20.43 +3.81 ^{bcdefghijkl}	11.60 +2.05 ^{bcd}	0.57 +0.06 ^{bcdef}	0.78 +0.36 ^{abc}	2.78 +0.32 ^{bcdefgh}	-	-	
Rhanna	Off-162	27.25 +2.30 ^{abcd}	43.05 +11.40 ^{abcd}	2.50 +0.58 ^{def}	22.37 +1.69 ^{bcdefghijkl}	12.77 +0.38 ^{bcd}	0.57 +0.03 ^{bcdef}	0.80 +0.28 ^{abc}	2.80 +0.14 ^{bcdefgh}	-	-	
Rhanna	Off-163	42.13 +5.39 ^{abcd}	71.38 +5.02 ^{abcd}	4.75 +1.89 ^{def}	29.13 +4.21 ^{bcdefghijkl}	14.53 +1.75 ^{bc}	0.50 +0.05 ^{bcdef}	0.88 +0.17 ^{abc}	2.85 +0.87 ^{bcdefgh}	-	-	
Rhanna	Off-1101	38.00 +4.32 ^{abcd}	36.50 +7.08 ^{abcd}	2.25 +0.50 ^{def}	27.70 +2.95 ^{bcdefghijkl}	13.48 +1.19 ^{bc}	0.49 +0.04 ^{bcdef}	0.95 +0.33 ^{abc}	3.00 +0.56 ^{bcdefgh}	-	-	
Ouardigha	Off-211	25.38 +9.62 ^{bcd}	24.98 +9.72 ^{ghi}	1.00 +0.00 ^{def}	22.25 +1.02 ^{bcdefghijkl}	12.00 +0.00 ^{bcd}	0.54 +0.02 ^{bcdef}	1.15 +0.12 ^{abc}	2.40 +0.08 ^{def}	-	-	

Ouardigha	Ofi-212	26.63 +3.68	abcd	36.00 +6.16	bcdelgkhi	1.50 +0.58	ef	24.40 +2.57	bcdelgkhi	12.17 +1.03	bcd	0.50 +0.05	bcdel	1.13 +0.17	abc	2.93 +0.45	bcdelg
Ouardigha	Ofi-231	29.60 +3.93	abcd	42.13 +12.68	abceghij	2.50 +0.58	def	21.10 +4.66	cdelgkhi	11.05 +3.14	bcd	0.52 +0.04	bcdel	0.58 +0.21	abc	2.53 +0.62	defg
Ouardigha	Ofi-232	28.25 +3.97	abcd	31.00 +4.32	cdelgkhi	2.00 +0.82	def	18.00 +1.63	ghijk	10.30 +0.24	cd	0.57 +0.06	bcdel	0.70 +0.16	abc	2.80 +0.16	bcdelg
Ouardigha	Ofi-233	24.13 +9.08	bcd	29.00 +5.55	efghi	1.75 +0.96	ef	18.57 +1.41	ghijkl	9.10 +0.65	cd	0.49 +0.02	bcdel	0.73 +0.12	abc	2.57 +0.49	cdelg
Ouardigha	Ofi-261	22.30 +3.19	cd	29.45 +6.44	deghij	1.00 +0.00	f	19.60 +3.59	ghijkl	10.65 +1.67	bcd	0.54 +0.02	bcdel	0.85 +0.12	abc	2.70 +0.16	bcdelg
Ouardigha	Ofi-262	26.70 +3.85	abcd	35.63 +4.17	bcdelgkhi	1.50 +0.58	ef	22.81 +3.35	cdelgkhi	11.43 +0.52	bcd	0.49 +0.03	bcdel	0.87 +0.34	abc	2.73 +0.21	bcdelg
Ouardigha	Ofi-263	39.13 +4.21	abcd	44.60 +4.33	abceghij	1.75 +0.50	ef	30.45 +1.56	abce	13.50 +0.71	bc	0.50 +0.12	bcdel	0.93 +0.46	abc	3.15 +0.24	bcdelg
Ouardigha	Ofi-264	28.33 +4.44	abcd	36.88 +9.19	abceghij	2.50 +1.00	def	23.90 +1.34	cdelgkhi	11.27 +0.56	bcd	0.47 +0.03	bcdel	0.47 +0.10	c	2.40 +0.26	efg
Ouardigha	Ofi-265	24.25 +0.61	bcd	35.25 +5.92	bcdelgkhi	3.00 +0.82	def	23.20 +0.16	cdelgkhi	10.20 +0.16	cd	0.43 +0.08	cdel	1.10 +0.08	abc	2.90 +0.73	bcdelg
Ouardigha	Ofi-266	29.88 +5.02	abcd	27.75 +1.32	ghijk	1.50 +0.58	ef	27.23 +6.65	bcdelgkhi	13.77 +1.45	bc	0.52 +0.06	bcdel	0.77 +0.09	abc	3.33 +0.24	bcdelg
Ouardigha	Ofi-271	27.17 +7.71	abcd	28.83 +8.66	efghi	1.50 +0.58	ef	26.20 +0.82	bcdelgkhi	12.45 +0.37	bcd	0.47 +0.00	bcdel	1.05 +0.04	abc	3.00 +0.00	bcdelg
Ouardigha	Ofi-272	35.88 +11.35	abcd	30.75 +8.07	cdelgkhi	2.00 +0.82	def	22.48 +3.26	cdelgkhi	12.18 +0.96	bcd	0.54 +0.06	bcdel	1.03 +0.10	abc	2.68 +0.28	bcdelg
Ouardigha	Ofi-273	24.33 +5.09	bcd	27.38 +12.15	ghijk	1.50 +0.58	ef	17.53 +2.50	ghijkl	9.87 +0.26	cd	0.57 +0.10	bcdel	0.63 +0.05	abc	2.47 +0.39	cdelg
Ouardigha	Ofi-274	28.38 +6.80	abcd	35.00 +9.70	bcdelgkhi	2.00 +0.82	def	19.73 +6.88	ghijkl	10.25 +1.19	cd	0.56 +0.18	bcdel	0.73 +0.10	abc	2.43 +0.40	efg
Ouardigha	Ofi-275	21.55 +8.86	cd	16.43 +3.51	l	1.50 +0.58	ef	18.55 +3.23	ghijkl	12.30 +0.90	bcd	0.67 +0.07	bcd	1.15 +0.04	abc	2.70 +0.24	bcdelg
Middle atlas	Ofi-311	35.68 +7.92	abcd	50.88 +3.45	abceghij	2.25 +0.50	def	24.15 +6.68	bcdelgkhi	11.65 +1.08	bcd	0.50 +0.08	bcdel	0.65 +0.10	abc	2.53 +0.10	cdelg
Middle atlas	Ofi-321	38.50 +19.07	abcd	29.13 +12.80	efghi	3.75 +0.96	bcdel	13.47 +2.17	cdelgkhi	12.87 +1.62	bc	0.54 +0.03	bcdel	0.70 +0.14	abc	3.20 +0.14	bcdelg
Middle atlas	Ofi-331	31.88 +8.23	abcd	22.25 +1.66	ghijk	1.00 +0.00	l	33.70 +0.00	l	10.70 +0.00	bcd	0.65 +0.11	bcd	0.80 +0.00	abc	2.55 +0.04	cdelg
Middle atlas	Ofi-341	30.63 +6.94	abcd	31.38 +7.06	cdelgkhi	1.50 +0.58	ef	19.27 +6.73	ghijkl	13.00 +0.50	bc	0.51 +0.05	bcdel	0.83 +0.17	abc	3.13 +0.26	bcdelg
Middle atlas	Ofi-351	31.00 +8.13	abcd	30.13 +10.01	cdelgkhi	1.75 +0.50	ef	21.07 +2.08	ghijkl	11.93 +0.99	bcd	0.56 +0.01	bcdel	0.77 +0.29	abc	3.03 +0.48	bcdelg
Northeast Morocco	Ofi-421	31.00 +7.43	abcd	32.68 +9.46	cdelgkhi	1.50 +0.58	ef	23.48 +1.07	cdelgkhi	11.53 +1.02	bcd	0.52 +0.06	bcdel	0.73 +0.21	abc	2.45 +0.26	cdelg
Northeast Morocco	Ofi-451	24.50 +1.78	bcd	39.63 +12.96	abceghij	2.50 +0.58	def	23.60 +1.31	cdelgkhi	11.30 +0.41	bcd	0.47 +0.01	bcdel	0.70 +0.00	abc	2.40 +0.08	efg
Eastern Rif	Ofi-521	21.88 +6.51	cd	21.50 +10.01	hi	1.00 +0.00	l	17.00 +1.63	ghijkl	12.00 +1.63	bcd	0.70 +0.16	bc	1.10 +0.08	abc	3.50 +0.41	abce
Eastern Rif	Ofi-531	21.33 +6.55	cd	23.33 +7.32	ghijk	1.00 +0.00	l	22.50 +0.41	cdelgkhi	12.00 +1.63	bcd	0.53 +0.02	bcdel	1.00 +0.16	abc	2.60 +0.33	cdelg
Eastern Rif	Ofi-533	32.40 +14.02	abcd	42.30 +23.93	abceghij	3.75 +2.36	bcdel	26.93 +1.37	bcdelgkhi	12.90 +0.51	bc	0.48 +0.04	bcdel	0.73 +0.21	abc	4.17 +1.31	a
Eastern Rif	Ofi-542	26.38 +15.21	abcd	29.75 +21.63	cdelgkhi	2.00 +0.82	def	29.05 +1.02	bcdel	14.85 +0.78	bc	0.51 +0.01	bcdel	0.85 +0.04	abc	3.70 +0.08	abc
Eastern Rif	Ofi-561	37.25 +5.84	abcd	34.93 +14.14	bcdelgkhi	1.75 +0.96	ef	26.20 +6.49	bcdelgkhi	12.75 +3.52	bc	0.48 +0.05	bcdel	0.78 +0.30	abc	2.70 +0.48	bcdelg
Western Rif	Ofi-612	36.25 +8.87	abcd	33.63 +9.32	cdelgkhi	2.75 +0.50	def	23.10 +2.21	cdelgkhi	11.57 +1.05	bcd	0.50 +0.03	bcdel	1.03 +0.09	abc	2.37 +0.45	efg
Western Rif	Ofi-631	30.20 +9.41	abcd	33.88 +13.14	cdelgkhi	3.50 +0.58	def	21.00 +0.41	cdelgkhi	12.25 +0.20	bcd	0.58 +0.00	bcdel	0.75 +0.04	abc	2.80 +0.00	bcdelg
Agadir	Ofi-711	41.70 +5.77	abcd	50.80 +9.72	abceghij	2.75 +1.26	def	28.38 +2.84	bcdelgkhi	12.90 +2.37	bc	0.49 +0.15	bcdel	0.55 +0.19	bc	3.03 +0.90	bcdelg
Agadir	Ofi-712	40.17 +6.54	abcd	57.50 +20.00	abceghij	4.25 +2.63	bcdel	26.03 +2.53	bcdelgkhi	15.47 +2.17	bc	0.56 +0.04	bcdel	0.87 +0.17	abc	2.70 +0.22	bcdelg
Agadir	Ofi-713	45.43 +4.64	ab	38.75 +8.21	abceghij	2.25 +1.26	def	34.00 +3.58	ab	20.23 +10.73	a	0.58 +0.25	bcdel	0.83 +0.26	abc	3.00 +0.08	bcdelg
Agadir	Ofi-714	35.78 +6.63	abcd	47.43 +8.25	abceghij	2.50 +0.58	def	25.15 +9.11	bcdelgkhi	15.78 +3.26	bc	0.67 +0.21	bcd	0.86 +0.21	abc	2.85 +0.24	bcdelg
Agadir	Ofi-731	39.03 +11.53	abcd	43.88 +15.29	abceghij	2.25 +0.58	def	28.45 +2.19	bcdelgkhi	15.08 +0.87	bc	0.53 +0.04	bcdel	1.00 +0.16	abc	3.05 +0.10	bcdelg
Agadir	Ofi-731	36.85 +4.98	abcd	48.38 +7.82	abceghij	2.00 +0.82	def	27.43 +1.54	bcdelgkhi	14.13 +1.26	bc	0.51 +0.03	bcdel	0.88 +0.28	abc	2.95 +0.54	bcdelg
Agadir	Ofi-732	42.68 +8.01	abcd	40.58 +8.39	abceghij	2.50 +0.58	def	29.00 +3.24	bcdel	14.80 +0.99	bc	0.51 +0.03	bcdel	1.10 +0.22	abc	3.13 +0.25	bcdelg
Agadir	Ofi-733	38.25 +10.44	abcd	42.65 +16.01	abceghij	2.25 +0.50	def	27.65 +8.00	bcdelgkhi	13.48 +2.67	bc	0.48 +0.05	bcdel	1.13 +0.56	abc	3.00 +0.28	bcdelg
Agadir	Ofi-742	42.93 +9.94	abcd	56.25 +23.09	abceghij	3.75 +1.71	bcdel	27.00 +2.27	bcdelgkhi	13.33 +1.68	bc	0.49 +0.07	bcdel	0.75 +0.24	abc	2.98 +0.24	bcdelg
Agadir	Ofi-743	39.75 +10.43	abcd	69.88 +26.42	ab	4.25 +3.30	bcdel	25.98 +4.29	bcdelgkhi	12.95 +1.32	bc	0.50 +0.06	bcdel	0.63 +0.29	abc	3.03 +0.40	bcdelg
Agadir	Ofi-751	45.75 +2.72	ab	50.88 +19.84	abceghij	4.00 +1.63	bcdel	32.68 +10.79	ab	13.13 +1.11	bc	0.44 +0.18	cdel	1.00 +0.26	abc	3.05 +0.33	bcdelg

Agadir	Oli-752	46.18 +3.94 _{ab}	64.63 +6.42 _{abcde}	5.25 +0.50 _{bc}	31.15 +2.56 _{abcd}	15.63 +0.75 _{bc}	0.50 +0.06 _{bcdef}	0.73 +0.24 _{abc}	2.88 +0.48 _{bcdefg}	-
Agadir	Oli-761	40.00 +10.58 _{abcd}	55.38 +19.74 _{abcde}	3.50 +1.29 _{bcdef}	22.45 +6.94 _{bcdefgh}	13.43 +3.06 _{bc}	0.63 +0.22 _{bcde}	0.73 +0.40 _{abc}	2.68 +0.24 _{bcdefg}	-
Agadir	Oli-762	48.25 + 5.12 _a	45.00 +13.20 _{abcde}	3.00 +1.41 _{def}	23.65 +3.49 _{bcdefgh}	13.25 +1.46 _{bc}	0.56 +0.08 _{bcdef}	0.88 +0.34 _{abc}	2.68 +0.30 _{bcdefg}	-
Agadir	Oli-771	33.38 +3.25 _{abcd}	46.00 +5.07 _{abcde}	3.00 +1.41 _{def}	25.83 +1.72 _{bcdefgh}	12.05 +1.46 _{bc}	0.46 +0.05 _{bcdef}	0.90 +0.16 _{abc}	2.75 +0.65 _{bcdefg}	-
Agadir	Oli-772	33.93 +3.00 _{abcd}	47.07 +12.40 _{abcde}	2.00 +0.82 _{def}	25.73 +1.46 _{bcdefgh}	12.80 +1.12 _{bc}	0.49 +0.05 _{bcdef}	0.73 +0.25 _{abc}	2.63 +0.39 _{bcdefg}	-
O. mega cantha	Jamaat Riah Om-A1	30.43 +3.54 _{abcd}	42.05 +15.92 _{abcde}	2.75 +0.96 _{def}	25.38 +3.50 _{bcdefgh}	12.38 +0.48 _{bc}	0.49 +0.07 _{bcdef}	0.80 +0.18 _{abc}	2.68 +0.39 _{bcdefg}	2.70 +0.22 _{ghijklmno}
	Jamaat Riah Om-A2	32.15 + 2.83 _{abcd}	43.05 +13.93 _{abcde}	2.00 +0.90 _{def}	25.15 +1.65 _{bcdefgh}	14.67 +1.31 _{bc}	0.58 +0.03 _{bcdef}	0.90 +0.16 _{abc}	2.90 +0.16 _{bcdefg}	3.00 +0.50 _{ghijklmno}
	Jamaat Riah Om-A3	29.03 +5.85 _{abcd}	36.60 +6.13 _{abcde}	2.75 +0.96 _{def}	25.40 +1.55 _{bcdefgh}	12.80 +0.98 _{bc}	0.50 +0.01 _{bcdef}	1.05 +0.04 _{abc}	3.10 +0.33 _{bcdefg}	2.15 +0.29 _{no}
	Jamaat Riah Om-B1	29.03 +5.85 _{abcd}	36.60 +6.13 _{abcde}	1.75 +0.50 _{def}	25.25 +1.56 _{bcdefgh}	12.35 +1.11 _{bc}	0.49 +0.02 _{bcdef}	0.98 +0.10 _{abc}	2.95 +0.37 _{bcdefg}	2.38 +0.39 _{ijklmno}
	Jamaat Riah Om-K1	25.00 +10.39 _{abcd}	21.65 +12.88 _{abcde}	1.00 +0.00 _{def}	23.53 +0.38 _{bcdefgh}	13.00 +1.63 _{bc}	0.55 +0.04 _{bcdef}	1.30 +0.08 _{abc}	3.00 +0.41 _{bcdefg}	2.10 +0.08 _{op}
	Jamaat Riah Om-K2	33.93 +9.72 _{abcd}	41.88 +10.36 _{abcde}	3.50 +1.00 _{def}	26.30 +3.50 _{bcdefgh}	12.83 +2.42 _{bc}	0.48 +0.03 _{bcdef}	1.40 +1.33 _a	2.90 +0.36 _{bcdefg}	3.07 +0.51 _{bcdefghijkl}
	Jamaat Riah Om-K3	32.63 +7.87 _{abcd}	65.38 +14.10 _{abcde}	4.25 +0.98 _{def}	26.28 +1.21 _{bcdefgh}	14.05 +1.17 _{bc}	0.53 +0.03 _{bcdef}	0.68 +0.33 _{abc}	3.05 +0.17 _{bcdefg}	2.60 +0.66 _{ijklmno}
Rhamna	Om-111	29.50 + 3.27 _{abcd}	50.05 +3.23 _{abcde}	2.50 +0.58 _{def}	24.85 +1.51 _{bcdefgh}	11.00 +0.00 _{bc}	0.44 +0.02 _{bcdef}	0.80 +0.00 _{abc}	2.85 +0.12 _{bcdefg}	2.70 +0.00 _{ghijklmno}
Rhamna	Om-112	26.38 +2.06 _{abcd}	30.13 +7.88 _{abcde}	1.50 +0.58 _{def}	19.83 +4.19 _{bcdefgh}	9.40 +1.04 _{cd}	0.48 +0.06 _{bcdef}	0.53 +0.05 _c	2.67 +0.47 _{bcdefg}	2.27 +0.12 _{mnop}
Rhamna	Om-122	28.90 +5.32 _{abcd}	42.58 +8.63 _{abcde}	3.00 +1.41 _{def}	23.17 +2.05 _{bcdefgh}	14.33 +3.52 _{bc}	0.51 +0.01 _{bcdef}	0.80 +0.33 _{abc}	2.70 +0.50 _{bcdefg}	2.77 +0.46 _{ghijklmno}
Rhamna	Om-131	37.75 +3.88 _{abcd}	55.38 +20.27 _{abcde}	3.75 +2.36 _{def}	23.75 +4.17 _{bcdefgh}	13.30 +2.03 _{bc}	0.56 +0.03 _{bcdef}	1.00 +0.47 _{abc}	3.03 +0.67 _{bcdefg}	3.80 +0.58 _{ab}
Rhamna	Om-143	24.87 +0.62 _{abcd}	34.07 +3.35 _{abcde}	2.50 +1.73 _{def}	22.70 +3.84 _{bcdefgh}	9.75 +0.20 _{cd}	0.45 +0.09 _{bcdef}	0.70 +0.16 _{abc}	2.55 +0.04 _{bcdefg}	2.30 +0.08 _{lmnop}
Rhamna	Om-181	27.28 +4.30 _{abcd}	32.88 +5.42 _{abcde}	1.00 +0.00 _{def}	37.00 +1.63 _{bcdefgh}	12.50 +0.41 _{bc}	0.33 +0.02 _{bcdef}	0.90 +0.08 _{abc}	3.00 +0.08 _{bcdefg}	1.70 +0.16 _p
Quardigha	Om-221	25.13 +8.13 _{abcd}	39.25 +7.71 _{abcde}	2.00 +0.82 _{def}	20.63 +1.25 _{bcdefgh}	10.88 +0.75 _{bc}	0.52 +0.03 _{bcdef}	0.75 +0.26 _{abc}	2.38 +0.15 _{bcdefg}	2.98 +0.66 _{bcdefghijkl}
Quardigha	Om-251	29.03 + 4.45 _{abcd}	37.75 +11.26 _{abcde}	2.50 +0.58 _{def}	21.70 +0.79 _{bcdefgh}	12.50 +0.82 _{bc}	0.57 +0.02 _{bcdef}	1.00 +0.08 _{abc}	3.00 +0.00 _{bcdefg}	2.30 +0.00 _{lmnop}
Middle atlas	Om-267	31.63 +2.17 _{abcd}	43.38 +8.36 _{abcde}	2.75 +2.36 _{def}	20.93 +2.81 _{bcdefgh}	11.15 +1.35 _{bc}	0.53 +0.04 _{bcdef}	0.70 +0.18 _{abc}	2.60 +0.34 _{bcdefg}	2.75 +0.80 _{ghijklmno}
Middle atlas	Om-352	31.88 +6.76 _{abcd}	42.15 +13.78 _{abcde}	2.75 +0.50 _{def}	22.83 +2.25 _{bcdefgh}	12.70 +0.92 _{bc}	0.55 +0.02 _{bcdef}	0.77 +0.26 _{abc}	2.90 +0.43 _{bcdefg}	2.83 +0.40 _{bcdefghijkl}
Middle atlas	Om-371	32.50 +6.54 _{abcd}	42.38 +12.26 _{abcde}	2.25 +1.26 _{def}	25.50 +3.97 _{bcdefgh}	12.75 +1.85 _{bc}	0.57 +0.18 _{bcdef}	0.83 +0.15 _{abc}	2.75 +0.33 _{bcdefg}	3.08 +0.49 _{bcdefghijkl}
Northeast	Om-411	39.73 +8.94 _{abcd}	58.50 +7.93 _{abcde}	3.25 +0.50 _{def}	25.30 +2.89 _{bcdefgh}	13.95 +2.42 _{bc}	0.55 +0.05 _{bcdef}	0.73 +0.24 _{abc}	3.15 +0.83 _{bcdefg}	3.13 +0.57 _{bcdefghijkl}
Northeast	Om-422	30.63 +2.02 _{abcd}	44.63 +7.38 _{abcde}	2.50 +1.73 _{def}	22.88 +1.11 _{bcdefgh}	11.85 +2.28 _{bc}	0.52 +0.07 _{bcdef}	0.80 +0.32 _{abc}	2.73 +0.25 _{bcdefg}	3.35 +0.13 _{bcdefghijkl}
Morocco	Northeast	Om-431	38.85 +7.35 _{abcd}	40.95 +19.31 _{abcde}	3.00 +0.82 _{def}	13.23 +0.76 _{bc}	0.55 +0.03 _{bcdef}	0.80 +0.22 _{abc}	2.75 +0.21 _{bcdefg}	3.57 +0.39 _{abcd}
Morocco	Northeast	Om-441	36.63 + 7.23 _{abcd}	43.35 +10.61 _{abcde}	1.75 +0.96 _{def}	12.97 +0.54 _{bc}	0.47 +0.06 _{bcdef}	1.20 +0.85 _{abc}	2.63 +0.26 _{bcdefg}	2.97 +0.12 _{bcdefghijkl}
Morocco	Northeast	Om-442	35.25 +3.57 _{abcd}	54.50 +7.33 _{abcde}	2.00 +0.00 _{def}	13.60 +0.90 _{bc}	0.51 +0.01 _{bcdef}	0.75 +0.21 _{abc}	2.85 +0.39 _{bcdefg}	3.28 +1.03 _{bcdefghijkl}
Morocco	Northeast	Om-461	29.80 +12.09 _{abcd}	34.15 +15.09 _{abcde}	2.50 +0.58 _{def}	12.40 +1.73 _{bc}	0.54 +0.02 _{bcdef}	0.87 +0.17 _{abc}	2.43 +0.31 _{bcdefg}	2.53 +0.34 _{bcdefghijkl}
Morocco	Northeast	Om-471	35.00 +5.72 _{abcd}	33.00 +8.16 _{abcde}	1.00 +0.00 _{def}	13.75 +1.02 _{bc}	0.57 +0.01 _{bcdef}	0.85 +0.04 _{abc}	3.05 +0.04 _{bcdefg}	2.80 +0.08 _{bcdefghijkl}
Eastern Rif	Om-511	28.13 +8.93 _{abcd}	30.20 +10.41 _{abcde}	3.50 +0.58 _{def}	24.25 +3.06 _{bcdefgh}	12.90 + 1.96 _{bc}	0.52 +0.01 _{bcdef}	1.10 +0.00 _{abc}	2.50 +0.00 _{bcdefg}	2.85 +0.53 _{bcdefghijkl}
Eastern Rif	Om-532	36.50 +6.16 _{abcd}	45.40 +20.76 _{abcde}	3.00 +1.15 _{def}	20.98 +3.81 _{bcdefgh}	12.40 +2.09 _{bc}	0.59 +0.05 _{bcdef}	0.80 +0.20 _{abc}	2.75 +0.21 _{bcdefg}	2.43 +0.22 _{bcdefghijkl}
Eastern Rif	Om-541	37.38 +6.14 _{abcd}	40.75 +15.02 _{abcde}	2.00 +0.82 _{def}	22.75 +3.84 _{bcdefgh}	12.38 +1.60 _{bc}	0.55 +0.06 _{bcdef}	0.80 +0.29 _{abc}	2.73 +0.46 _{bcdefg}	2.43 +0.49 _{bcdefghijkl}
Eastern Rif	Om-543	39.13 + 6.14 _{abcd}	38.58 +9.14 _{abcde}	2.25 +0.50 _{def}	20.90 +2.31 _{bcdefgh}	11.03 +1.86 _{bc}	0.52 +0.06 _{bcdef}	0.85 +0.13 _{abc}	2.13 +0.25 _{bcdefg}	2.88 +0.25 _{bcdefghijkl}
Eastern Rif	Om-551	31.75 +5.39 _{abcd}	44.50 +7.82 _{abcde}	2.50 +1.73 _{def}	20.50 +4.02 _{bcdefgh}	13.85 +7.67 _{bc}	0.70 +0.19 _{bc}	0.90 +0.16 _{abc}	2.30 +0.38 _{bcdefg}	2.35 +0.64 _{bcdefghijkl}

Accession	35.50 +10.45	40.33 +5.37	42.55 +9.17	45.88 +1.11	40.98 +8.04	50.00 +8.99	2.50 +1.73	23.80 +2.03	13.23 +1.34	0.95 +0.10	2.33 +0.15	4.25 +0.50	2.40 +0.12
Eastern Rif	Om-571	40.33 +5.37	42.55 +9.17	45.88 +1.11	40.98 +8.04	50.00 +8.99	2.50 +1.73	23.80 +2.03	13.23 +1.34	0.95 +0.10	2.33 +0.15	4.25 +0.50	2.40 +0.12
Eastern Rif	Om-572	37.50 +11.02	35.00 +4.49	28.50 +3.67	19.85 +2.16	37.63 +3.35	1.00 +0.00	25.35 +4.20	13.85 +3.55	1.05 +0.20	2.30 +0.41	3.50 +0.58	2.30 +0.08
Eastern Rif	Om-581	35.63 +9.07	40.38 +7.80	50.50 +6.81	51.95 +13.07	50.00 +8.99	4.00 +1.83	21.13 +1.18	11.40 +0.86	0.85 +0.06	2.05 +0.31	4.25 +0.50	2.85 +0.10
Western Rif	Om-632	37.45 +4.59	30.68 +7.81	28.13 +7.20	54.00 +6.87	41.50 +8.50	3.00 +0.82	17.30 +4.87	11.50 +1.35	0.70 +0.20	2.48 +0.13	4.00 +0.00	2.30 +0.14
Western Rif	Om-621	29.00 +7.82	28.13 +7.20	28.13 +7.20	54.00 +6.87	41.50 +8.50	2.50 +1.29	21.75 +0.20	11.00 +0.00	0.90 +0.08	2.35 +0.29	3.50 +0.58	2.35 +0.12
Western Rif	Om-651	38.13 +7.40	28.13 +7.20	28.13 +7.20	54.00 +6.87	41.50 +8.50	2.75 +0.50	22.93 +3.20	13.40 +2.09	0.80 +0.22	2.63 +0.26	4.00 +0.00	2.95 +0.50
Agadir	Om-722	31.98 +4.04	41.50 +8.50	57.43 +6.95	50.00 +8.99	42.20 +6.37	3.50 +0.58	24.23 +1.93	12.56 +1.70	0.80 +0.24	2.63 +0.26	4.00 +0.00	3.03 +0.05
Agadir	Om-741	31.63 +3.12	42.20 +6.37	57.43 +6.95	50.00 +8.99	42.20 +6.37	3.75 +1.26	23.75 +1.44	11.95 +1.14	0.85 +0.47	2.08 +0.61	3.75 +0.50	2.63 +0.62
Northeast	Or-412	29.28 +6.52	42.20 +6.37	57.43 +6.95	50.00 +8.99	42.20 +6.37	2.75 +0.96	17.90 +3.59	17.33 +3.45	1.38 +0.05	3.60 +0.45	4.50 +0.58	4.03 +0.05
Morocco													
O. aequ	Rhanna	39.88 +6.94	34.63 +4.70	28.50 +3.67	19.85 +2.16	37.63 +3.35	3.25 +0.50	30.63 +2.95	14.03 +9.99	1.08 +0.39	2.43 +0.33	5.00 +0.82	3.48 +0.32
O. dillenii	Ouardigha	19.85 +2.16	28.50 +3.67	19.85 +2.16	19.85 +2.16	37.63 +3.35	7.75 +3.50	13.75 +1.84	6.25 +0.86	0.45 +0.00	2.90 +0.16	-	-
O. leuco	Middle atlas	37.63 +3.35	53.33 +9.51	53.33 +9.51	53.33 +9.51	53.33 +9.51	4.00 +0.82	24.75 +3.10	14.38 +0.25	1.13 +0.31	2.70 +0.26	5.00 +0.82	3.85 +0.54
O. in-	tricha	42.55 +9.17	42.93 +18.69	42.93 +18.69	42.93 +18.69	42.93 +18.69	2.25 +1.26	24.40 +5.63	13.18 +2.19	0.93 +0.32	2.90 +0.12	4.50 +1.29	3.53 +1.01
O. in-	ermis	45.88 +1.11	50.50 +6.81	50.50 +6.81	50.50 +6.81	50.50 +6.81	4.00 +1.63	27.30 +3.50	13.63 +1.49	0.50 +0.02	2.73 +0.38	4.25 +0.50	3.65 +0.84
		40.98 +8.04	51.95 +13.07	51.95 +13.07	51.95 +13.07	51.95 +13.07	3.25 +0.50	23.95 +5.17	13.45 +2.09	0.85 +0.24	2.62 +0.26	4.75 +0.96	3.10 +1.05
		35.63 +8.94	50.00 +8.99	50.00 +8.99	50.00 +8.99	50.00 +8.99	5.75 +0.96	20.73 +3.26	9.60 +1.82	0.46 +0.02	2.33 +0.25	-	-

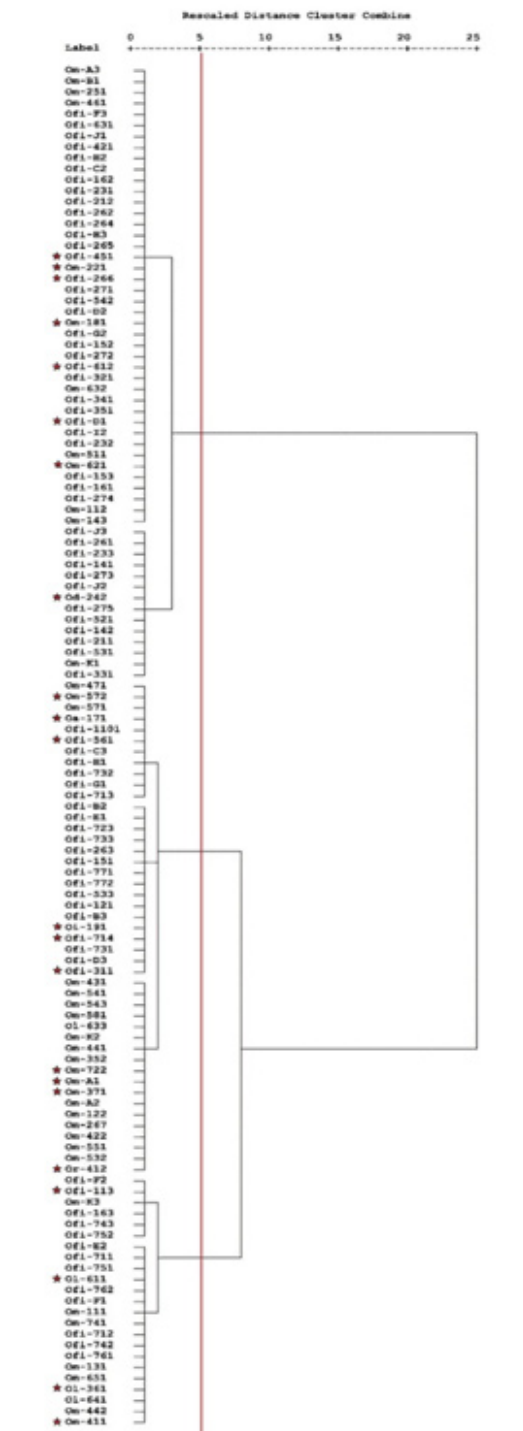


Fig. 4: Dendrogram grouping 124 accessions of *Opuntia spp.* based on 10 morphological traits and Ward's method. The red stars indicate accessions used for molecular analysis.

accessions from the same geographical site exhibited significantly different measures of morphological traits; for example, in accessions from Rhamna, cladode length was ranging from 16.30 cm in Ofi-141 to 29.13 cm in Ofi-163 (Table 1). The *O. ficus indica* accessions showed similarity coefficients ranging from 0.00 to 1.00 (proximity matrix not shown). The lowest similarity coefficient was noticed between accessions Ofi-212 and Ofi-262 from Ouardigha and between accessions Ofi-771 and Ofi-772 from Agadir while the highest similarity coefficient was observed between accessions Ofi-163 (from Rhamna) and Ofi-275 (from Ouardigha). In some cases, accessions from different regions show very low similarity coefficient (e.g. Ofi-C3 from Jamaât Riah and Ofi-561 from Eastern Rif displayed a similarity coefficient of 0.001). The dendrogram presented in Figure 2 suggested 3 major clusters. The first cluster contains 30 accessions (38.46 % of the total accessions); the second cluster contains 5 accessions (6.41 % of the total accessions) and the third cluster is comprised of 43 accessions (55.13% of the total accessions). Surprisingly, the three clusters contain accessions collected from different geographical sites.

Genetic Variability Within *Opuntia Megacantha*

The plant height of *O. megacantha* accessions ranged from 24.67 cm to 40.33 cm, plant diameter from 21.65 cm to 65.38 cm, the average number of cladodes per plant from 1.00 to 4.25, cladode height from 17.30 cm to 37.00 cm, cladode width from 9.40 cm to 14.67 cm, the index of cladode shape from 0.33 to 0.70, cladode thickness from 0.53 cm to 1.40 cm, the mean distance between areoles from 2.05 cm to 3.15 cm, the average number of spines per areole from 3.00 to 5.00 and the mean length of the longest spine per areole from 1.70 cm to 3.80 cm. Here again, some accessions from the same region displayed significantly different measures of morphological traits (Table 1). The range of similarity coefficients was from 0.00 to 1.00 (proximity matrix not shown). The lowest similarity was between Om-B1 and Om-A3, both from Jamaât Riah, while the highest similarity was between Om-K1 and Om-K3, from Jamaât Riah too. The dendrogram (Figure 3) shows 3 clusters, the first cluster consisted of 11 accessions (28.95 % of the total accessions); the second cluster contains 20 accessions (52.63 % of the total accessions) whereas the third cluster

comprised 7 accessions (18.42 % of the total accessions). Interestingly, accessions collected from different regions could be found within the same cluster.

Genetic Variability Among *Opuntia* Spp.

Taking into consideration all the studied accessions of *Opuntia* spp., the plant height ranged from 19.85 cm in Od-242 to 48.25 cm in Ofi-762, the average number of cladodes per plant from 1.00 to 7.75 in Od-242, with cladodes of different heights (from 13.70 in Ofi-331 to 37.00 in Om-181) and widths (from 6.25 cm in Od-242 to 20.23 in Ofi-713). Cladode thickness varied from 0.47 cm in Ofi-264 to 1.40 cm in Om-K2 (Table 1). Among the studied species, *O. ficus indica*, *O. dillenii* and *O. inermis* are spineless. The similarity coefficients were ranging from 0.00 to 1.00. The dendrogram shows three different clusters (Figure 4), and accessions from different species or geographic origins could be gathered into the same cluster. This shows that the clusters obtained do not fit with the species or geographical sites evaluated.

Molecular Markers to Assess Genetic Diversity

After screening 14 ISSR and 34 RAPD primers, 10 ISSR and 1 RAPD primers were selected for producing unambiguous, reproducible and intense bands. The ISSR primers generated a total of 66 bands, 64 (96.9 %) of which were polymorphic. The number of bands observed per locus varied from 4 to 9, with an average of 6.6. The RAPD primer (UBC-571) gave 6 bands, all polymorphic (Table 2). Based on the PIC values, the markers were highly informative. Indeed, the PIC values varied from 0.62 to 0.97, with an average of 0.82. The highest PIC value was observed in primer ISSR-8 while the lowest value was observed in primer ISSR-29 (Table 2). The dendrogram displayed 7 clusters at a similarity of 0.76 (Figure 5). *O. inermis* was separated from all the other species. The two accessions of *O. leucotricha* (OI-361 and OI-611) were placed into two different groups while accessions Or-412 (*O. robusta*) and Oa-171 (*O. aequatorialis*) were gathered into the same cluster. Interestingly, 14 accessions belonging to *O. ficus indica* and *O. megacantha* were gathered into the same group. This was confirmed by PCA (Figure 6), which results were similar to those obtained by the dendrogram.

Discussion

In the present study, the genetic diversity of 124 accessions of *Opuntia spp.* was evaluated based on 10 morphological descriptors. The use of morphological descriptors to determine genetic diversity within plant populations is simple, rapid and inexpensive⁹. Thus, such descriptors were used in many plant species such as date palm¹⁹, hexaploid oat¹⁶ and indigenous rice⁴³. To the best of our knowledge, this is the first work to assess genetic variability within and among *Opuntia* populations in Morocco using morphological descriptors. Studies on the genetic variability in *Opuntia* species using morphological descriptors are scarce. Peña-Valdivia *et al.* (2008)³¹ assessed genetic diversity within 55 accessions of *Opuntia spp.* in Mexico based on 65 morphological descriptors; and indicated that only few morphological traits showed significantly different measures within the studied populations. In a more recent study, Helsen *et al.* (2009)²⁵ highlighted a clear differentiation between *Opuntia* species using descriptors based on fruit, flower, seed, spine and trunk characteristics. In our case, the 124 accessions showed significant differences in the measured parameters and were distributed into 3 main distinct clusters based on Ward's method, containing 56, 45 and 23 accessions, respectively. Interestingly, accessions classified in the same

species could be located in different clusters and vice versa, even though some species contain spines while others do not. This may be explained by several factors including the environmental effect and evolutionary change¹¹⁻⁴⁰. Our results are in good agreement with Valadez-Moctezuma *et al.* (2014)³⁹ who supported the hypothesis about the existence of a smaller number of *Opuntia* species compared to those currently described, but with high intraspecific genetic variation; and with Labra *et al.* (2003)²⁷ who suggested that the presence of spines should not be considered as a parameter in *Opuntia* taxonomy. Our results showed the influence of geographical origins on the morphological traits within the same species. Indeed, in some cases, accessions of the same species but from different regions were gathered into different clusters. This might be due to the environment effect³⁷⁻³⁸. Our results confirm previous findings on *Opuntia spp.* in Sardinia and Corsica islands, which show that environmental factors such as elevation, soil drainage, temperature and rainfall affect plant morphology²¹. As far as *Opuntia ficus indica* is concerned, the use of morphological or agro-morphological traits to assess genetic diversity within its accessions is very scarce. Felker *et al.* (2005)²² compared genotypes of *O. ficus indica* from different origins based on yield and fruit characters, and significant differences were observed. In another

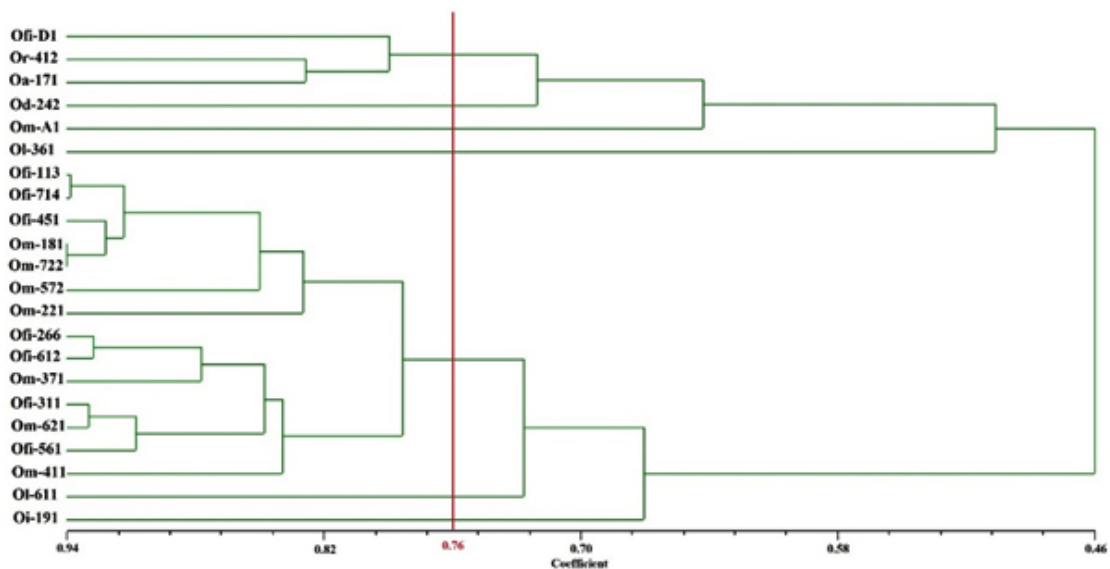


Fig. 5: Dendrogram grouping 22 accessions of *Opuntia spp.* based on 10 ISSR and 1 RAPD markers.

investigation, Bendhifi *et al.* (2013)⁶ compared 28 Tunisian accessions of *O. ficus indica* using 20 descriptors based on plant, cladode and fruit characteristics. Here again, significant differences were observed between accessions. In our case, the 10 morphological traits showed significant differences between the 78 accessions. Remarkably, some accessions from the same geographical origin displayed significantly different measures of some traits. The use of morphological markers to assess genetic diversity within populations is simple and inexpensive. However, such markers can be affected by the environment³. Therefore, in the recent years, molecular markers have been widely used to assess genetic diversity within plant populations. Indeed, it was reported that molecular markers cover a larger proportion of the genome than the morphological ones and are less influenced by the environment since genetic variability is evaluated based on genotype rather than phenotype²⁶⁻³⁸. In addition, molecular markers have been proven powerful to detect variability when it could not be possible with morphological markers; i.e. a given morphological marker can affect other traits and the usefulness of morphological markers is restricted by their limited number³. Along this line, molecular markers have been applied alone or in association with morphological descriptors to assess genetic variability in many plant species; e.g. in Phoenix *dactylifera* L.³⁵, *Olea europaea* L.⁸, *Citrus sinensis*

L.²⁹. Concerning the cactus pear, *Labra et al.* (2003)²⁷ evaluated the genetic diversity among 11 *Opuntia* species from Italy using chloroplastic simple sequence repeat (cpSSR) and amplified fragment length polymorphism (AFLP) markers, and a high genetic similarity between *O. ficus indica* and *O. megacantha* was revealed. Indeed, these authors suggested that *O. ficus indica* should be considered as a domesticated form of *O. megacantha*. Caruso *et al.* (2010)¹⁰ assessed the genetic diversity among 62 accessions belonging to 16 *Opuntia* species from different origins, using simple sequence repeat (SSR) primers, and found that the *O. ficus indica* accessions did not cluster separately from some *Opuntia* species, including *O. megacantha*. In a more recent study, Valadez-Moctezuma *et al.* (2014)³⁹ assessed the genetic variability among 12 Mexican *Opuntia* species (52 cultivars) with ISSR and RAPD markers, and reported that *O. ficus indica* and *O. megacantha* might have a common ancestry. Our results are consistent with all these authors, since a high genetic similarity was observed between *O. ficus indica* and *O. megacantha*. On the other hand, Valadez-Moctezuma *et al.* (2014)³⁹ screened 120 RAPD primers, of which five were able to generate bands (4.16 %). In our case, only one RAPD primer over 34 (2.94%) generated bands. Our findings indicated that ISSR markers are the most efficient and informative in *Opuntia* species. Recently, ISSR markers have been used to assess the genetic and

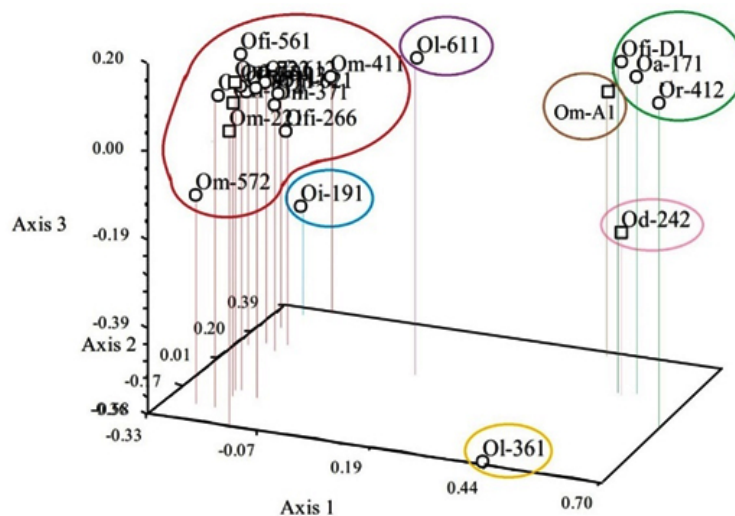


Fig. 6: 3D graphical distribution based on principal components analysis (PCA) showing the distribution of the 22 *Opuntia* spp. accessions.

phylogenetic relationships of 15 *Opuntia* species from Argentina, Bolivia, Brazil, Paraguay, and Uruguay³³. Our findings are in contrast with those of Valadez-Moctezuma *et al.* (2014)³⁹, who indicated that the assessment of genetic diversity with RAPD markers was relatively more effective than with ISSR markers. In our study, the total number of polymorphic bands (64) and the PIC values (0.62-0.97) of ISSR markers were different from those found by Valadez-Moctezuma *et al.* (2014)³⁹, in which the number of polymorphic bands was 70 and the PIC varied from 0.21 to 0.27. This difference in polymorphic bands number and PIC values could be explained by the different ISSR primers employed (different sequences), the difference among accessions and the high distance between the geographical sites (Mexico and Morocco). Regarding the species *Opuntia ficus indica*, Zoghalmi *et al.* (2007)⁴⁴ used 8 RAPD primers (over 22 screened) to assess the genetic variability within 36 Tunisian accessions, and a considerable genetic diversity was detected. In Morocco, the genetic diversity within *O. ficus indica* accessions was evaluated using RAPD markers¹⁸, and the effect of the geographical origin was revealed. In our case, findings showed high genetic similarity among the studied accessions of

O. ficus indica, even though they were collected from different regions of Morocco.

Conclusion

This paper is the first report to assess the genetic diversity within and among *Opuntia* species in Morocco using morphological descriptors and molecular markers. A low level of genetic diversity was observed among the accessions analyzed and a high similarity between *O. ficus indica* and *O. megacantha* accessions was highlighted. Our findings show also that ISSR markers are very efficient and informative in *Opuntia* species. Our results should be taken into consideration for plant breeding and genetic resource conservation programs. Further investigations are currently underway in order to determine the biochemical characteristics of cladode, flower, fruit and juice of these species, and to specify their molecular families to highlight the most appropriate use of each species.

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