Simple Sequence Repeat Identification and Endocarp Characterization of Olive Tree Accessions in a Tunisian Germplasm Collection

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Abstract. Most traditional olive-producing countries possess a diversified genetic patrimony in *Olea europaea* L. Since the emergence of modern olive growing system, the identification, classification, and conservation of autochthonous olive cultivars is a priority for these countries. In this work, a total of 84 accessions belonging to the "Boughrara"-Sfax olive germplasm collection located in Tunisia have been screened using a powerful set of eight simple sequence repeat markers (SSRs). The study revealed a high genetic variability among the collection and detected a total of 64 alleles. For better management of the mentioned germplasm bank, an improved classification of the entries, including new denominations, has been proposed. In addition, several cases of mislabeling, synonymy, and homonymy have been clarified. Genetic relationships among cultivars have been analyzed showing four major clusters. Finally, a correspondence factor analysis demonstrated that cultivars tend to cluster depending on their main use as oil or table olives. No clear clustering tendencies were observed when the geographical origin of cultivars was used as the criteria for the analysis. All results obtained by SSR screening and classification were in accordance with classification based on morphological traits of fruit endocarps.

Olive (Olea europaea L.) is one of the oldest cultivated species in the Mediterranean basin. Tunisia has played an important role in the establishment of the crop in this area as a result of its dynamic history and geographical location (Trigui et al., 2006). Olive still plays a significant role in both the economy and society of Tunisia today. It occupies onethird of the cultivated land (1,667,000 ha), represents the predominant activity of 29% of farmers, and affects one million Tunisian people's revenues (Jardak, 2006). Tunisia is considered the second producer and exporter of olive oil in the world, only preceded by the European Union. Tunisian production and exporting averages reach 185,000 tons and 139,600 tons, respectively (averages 2004 to 2009; International Olive Council, 2010). Like most of traditional olive-producing countries

(Rallo et al., 2005), Tunisia includes diversified genetic resources, which are mainly formed by small distributed autochthonous cultivars. The characterization of this diversity is essential for the better management of the national germplasm and for genetic breeding programs.

The classical approach in the identification of olive cultivars has been based on agronomical and/or morphological studies. It consists of measures such as production average, precocity, vigor, etc. and the direct observation of several distinctive traits on branches, fruits, endocarps, leaves, and inflorescences (Barranco and Rallo, 1984; Barranco et al., 2005). Among the aforementioned morphological characters, endocarps are considered one of the most discriminative traits. These traits are highly inheritable, have low sensitivity to environmental conditions, and are easily observed on collected and stored material (Rallo et al., 2005). A number of analyses has been conducted on Tunisian germplasm material from different regions of the country using morphological and/or agronomical characters (Mehri et al., 1997; Taamalli et al., 2006; Trigui et al., 2002). Several collections of olive cultivars have been created in Tunisia. Among them, the "Boughrara"-Sfax olive germplasm collection is considered the most important in terms of cultivar numbers. Although the morphological and agronomical studies are very useful for both exploration surveys (Barranco et al., 2000, 2005; Barranco and Rallo, 1984; Cimato et al., 1993) and germplasm bank characterization (Caballero et al., 2006), results can be influenced by several non-genetic factors. As a consequence, molecular methods are very suitable to reach a better understanding of the material's genetic diversity.

Molecular studies have started with the use of isoenzyme markers (Ouazzani et al., 1993; Trujillo et al., 1995) and at a later stage have been carried out using DNA markers as restriction fragment length polymorphisms (Besnard et al., 2001a), random amplified polymorphic DNAs (Belaj et al., 2001; Fabbri et al., 1995), amplified fragment length polymorphisms (Angiolillo et al., 1999) and microsatellite markers [simple sequence repeats (SSRs)] (Baldoni et al., 2009; La Mantia et al., 2005).

The microsatellite technique is one of the most reliable methods used in olive cultivar characterization. It has revealed a high informative level because the markers are polymorphic, multiallelic, and codominant. Moreover, it consists of a relatively simple methodology that permits an easy interpretation of results (Rafalski et al., 1996). SSR markers have been successfully used in germplasm bank classification and contributed to better management of several olive collections around the world (Khadari et al., 2003; Muzzalupo et al., 2006). To provide a better worldwide applicable tool for olive DNA typing, a list of 11 SSR markers has been recently selected among 37 microsatellites available for olive cultivar characterization. The mentioned list has been based on the reproducibility, quality of scoring, information content, independent segregation,

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and discriminative capacity of the markers (Baldoni et al., 2009).

Some studies using microsatellite markers have been achieved on Tunisian cultivars. They have been carried out in olive trees belonging to different collections (Taamalli et al., 2008), on trees of commercial interest original from different geographical regions of Tunisia (Rekik et al., 2008), or to study the differences between cultivated varieties of olive and Oleaster trees mainly from the northern region of the country (Hannachi et al., 2008).

The present work aims to identify and classify 84 olive accessions obtained by a national exploration survey and introduced in the "Boughrara"-Sfax olive germplasm collection (South of Tunisia). The work has been carried out using eight microsatellite markers, seven of them mentioned as some of the most reliable existing microsatellite markers for olive cultivar characterization (Baldoni et al., 2009). Furthermore, this work emphasizes the possibility of integration of both molecular and morphological approaches for germplasm classification because endocarp traits have been used to corroborate all conclusions obtained by the SSR screening.

Table 1. Morphological	characters of	of the endocarps of	on 84 an	alyzed accession	s.
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					Position				Number of	Distribution	
					of the				vascular	of vascular	
			Symmetry	Symmetry	maximum		Form	Roughness	bundles	bundles	Presence
			of position	of position	transversal	Form of	of the	of the	over the	over the	of
	Wt ^z	Form ^y	A ^x	B^w	diam ^v	the apex ^u	baset	surface ^s	surface ^r	surface ^q	mucro ^p
Chemlali Sfax	L	EP	SA	S	С	R	R	S	М	G	Р
Chemlali Sfax COI	L	EP	SA	S	С	R	R	S	М	G	Р
Sig 3PG	L	EP	SA	S	С	R	R	S	М	G	Р
Baldi	L	EP	SA	S	С	R	R	S	М	G	Р
Chargia BL8											
BL24	L	EP	SA	S	С	R	R	S	М	G	Р
Kotti K18	L	EP	SA	S	Ċ	R	R	S	М	G	Р
CHLOuled	Ē	EP	SA	ŝ	Č	R	R	ŝ	M	Ğ	P
Youssef Sig 13	_			-	-			-			-
Hchichina 10	L	EP	SA	S	С	R	R	S	М	G	Р
Sehli Sned 7	L	EP	SA	S	С	R	R	S	М	G	Р
Kotti K12	L	EP	SA	S	С	R	R	S	М	G	Р
K17	L	EP	SA	S	С	R	R	S	М	G	Р
Hchichina 6	L	EP	SA	S	С	R	R	S	М	G	Р
BL26	Ē	EP	SA	ŝ	Č	R	R	ŝ	M	Ğ	P
Chemlali Biawa	Ľ	EP	SA	Š	Č	R	R	Š	M	G	p
Zalmati Zarzis	T	EP	SA SA	S	C	R	R	S	M	G	P
Sig 15	I	EI	SA	S	C	D	D	S	M	G	D
Sig 15 Sahli Cafaa	L T	EL	SA	5	C	R D	R D	5	IVI M	C	r D
Dawla1	L	EP	SA	5	C	ĸ	К _	3	IVI	G	P
BL23	L	EP	SA	S	С	R	R	S	Μ	G	Р
BL20	L	EP	SA	S	С	R	R	S	Μ	G	Р
Horr Sfax	L	EP	SA	SA	С	R	R	S	М	G	Р
BL16 G	L	EP	SA	S	С	R	R	S	М	G	Р
Hchichina 7	L	EP	SA	S	С	R	R	S	Μ	G	Р
Kotti K9G	L	EP	SA	S	С	R	R	S	Μ	G	Р
Kotti k21	L	EP	SA	S	С	R	R	S	М	G	Р
Kotti K30G	L	EP	SA	S	С	R	R	S	Μ	G	Р
Sig 113 G	L	EP	SA	S	С	R	R	S	Μ	G	Р
Sehli Jbeniana BL15 Bent	L	EP	SA	S	С	R	R	S	М	G	Р
LOUZII DI 27	т	EI	C A	C	C	р	р	C	м	C	D
DL3/	L	EL	SA	5	C	K D	K D	5	IVI N	G	P
Youssef	L	EP	SA	5	C	ĸ	K	5	IVI	G	P
Sahli Gafsa autre que Gtar	L	EP	SA	S	С	R	R	S	М	G	Р
Horr Blettech BL11 PG	L	EL	SA	S	С	R	R	S	М	G	Р
Khchinet Sig	L	EP	SA	SA	А	R	R	S	L	G	Р
SIG19	L	EP	SA	SA	А	R	R	S	L	G	Р
Bou Bazzoula BL19PG	L	EP	SA	SA	А	R	R	S	L	G	Р
Kotti K23 (avec mam)	L	EP	SA	SA	А	R	R	S	L	G	Р
BL2	L	EP	А	S	В	R	R	S	М	R	А
BL31	Ľ	EP	A	Š	B	R	R	Š	M	R	A
Chemlali	I	FP	Δ	SA	A	R	P	S	I	G	P
Zarzis	T	ED		0.1		D	n n	0	L	0	I D
Hchichina H9	L	EP	А	5A	А	К	Р	8	L	G	Ч
Chemchali Gtar 3	М	EP	SA	S	С	R	R	R	М	R	Р
Jemri Bouchouka	М	EP	SA	S	С	R	R	R	М	R	Р
Kotti K30 PG	Μ	EP	SA	S	С	R	R	R	Μ	R	Р

(Continued on next page)

	Table 1.	(Continued) Morphological	characters	of the endocar	ps on 8	4 analyzed	accessions
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					Position				Number	Distribution	
			Symmetry	Symmetry	of the maximum	Form	Form	Roughness	of vascular bundles	of vascular bundles	Presence
	Wt	Form	of position A	of position B	transversal diam	of the apex	of the base	of the surface	over the surface	over the surface	of mucro
Injassi	L	EP	SA	SA	А	R	Р	S	L	G	Р
Hchichina											
Chemlali Mahares	L	EP	SA	SA	А	R	Р	S	L	G	Р
Chetoui 2	L	EL	А	S	В	Р	R	R	М	G	Р
Chetoui 2 à mam	L	EL	А	S	В	Р	R	R	М	G	Р
Marsaline	Н	EP	S	S	С	R	R	R	М	G	А
Meski Bjawa	Н	EP	S	S	C	R	R	R	М	G	A
Fakhari Tataouine	Н	0	S	S	C	R	R	W	M	G	A
Meski Zarzis	H	0	S	S	С	R	R	W	M	G	A
Chemlali	L	EL	А	8	C	R	R	R	М	G	Р
Chemlali	T	EI	Δ	S	С	P	P	R	М	G	P
Tataouine10	L	LL	71	5	C	K	ĸ	K	101	0	1
Chemlali Sig	L	EL	А	S	С	R	R	S	М	G	р
Kotti K14	Ľ	EL	A	Š	č	R	R	Š	M	G	P
Sig 14 PG	L	EL	SA	ŝ	Č	R	R	ŝ	М	Ğ	P
Besbessi	VH	EP	SA	ŝ	Č	R	Т	Ŵ	М	R	P
Lattout Sned 3	L	0	S	ŝ	B	Р	R	W	L	G	A
CHL Sig 12PG	L	EL	SA	S	С	R	R	S	М	G	Р
Chemlali Ontha Tataouine	L	EP	SA	S	С	R	R	S	Н	R	Р
Chemlali Bouchouka	L	EL	SA	S	С	R	Р	S	М	G	Р
Horr Louzir BL32 PG	L	EL	А	S	С	R	R	S	М	G	Р
Chahla	L	0	S	SA	А	R	Р	R	М	G	Р
Semni Jbeniana Sig 0	L	EP	А	S	С	R	Р	S	М	R	Р
Jeddaria Chaal	L	EP	А	SA	А	R	R	S	М	G	Р
Mallahi ElMouammar Sig 1G	Н	EL	А	S	С	Р	Т	W	L	G	Р
Tounsi Gafsa	Т	FP	Δ	S	С	R	R	W	М	G	р
autre Zarrazi	Ľ	EP	S	S	C	R	R	S	M	G	P
Ech- Chahla /g	Ē	0	SA	SA	Ă	R	Р	R	M	G	P
Neb Tataouine	Ē	EP	A	SA	A	R	P	S	M	Ğ	P
Kotti K9 PG	L	0	SA	S	С	R	R	S	М	G	Р
Lqam El Kotti Kotti K3	L	EP	S	S	С	R	R	S	L	G	А
Kotti K56 G	L	EP	S	S	С	R	R	R	М	G	А
Chemlali Barrani	L	EL	S	S	С	R	Р	R	М	G	А
K56PG	L	EL	SA	S	С	Р	Р	S	L	G	Р
Chemlali Balhi	Μ	EP	SA	SA	С	R	R	S	L	G	Р
Sig 112PG	L	EL	А	S	С	R	R	S	Н	R	Р
Zeitoun Boubazzoula COI	М	EP	SA	S	С	R	R	R	М	G	Р
Kotti K32	L	EL	А	S	С	Р	Р	S	L	G	Р
BL32 G	Ē	0	SA	ŝ	č	R	R	Ř	M	Ğ	Ă
Zarrazi Kgh	M	õ	SA	ŝ	č	R	R	W	L	Ğ	P
Zeitoun	L	EP	S	SA	Ā	R	R	S	L	Ğ	P
Boubazzoula COI										-	
Dhokar Nafti	L	0	SA	S	С	R	R	S	L	G	Р
Zarbout Louzir Kotti K11	L L	EL EP	A S	S S	C C	P R	P R	S S	H M	G G	P P

^zWeight: low = L (less than 0.3 g); medium = M (0.3 to 0.45 g); high = H (0.45 to 0.7 g); very high = VH (greater than 0.7 g). ^yForm: spherical = S (length/width less than 1.4); ovoid = O (length/width 1.4 to 1.8); elliptic = EP (length/width 1.8 to 2.2); elongated = EL (length/width greater than 0.7 g). than 2.2).

^xSymmetry of position A: symmetric = S; slightly asymmetric = SA; asymmetric = A. ^wSymmetry of position B: symmetric = S; slightly asymmetric = SA.

^vPosition of the maximum transversal diameter: toward the base = B; centered = C; toward the apex = A.

^uForm of the apex: pointed = P; rounded = R.

^tForm of the base: pointed = P; truncated = T; rounded = R.

^sRoughness of the surface: smooth = S; rough = R; wrinkled = W.

Number of vascular bundles over the surface: low = L (less than 7); medium = M (7 to 10); high = H (greater than 10).

^aDistribution of vascular bundles over the surface: regular = R; grouped together in the suture = G.

^pPresence of mucro: present = P; absent = A.

Material and Methods

Plant material and DNA extraction. Samples from 84 accessions were obtained from the national olive collection: "Conservatoire National de l'Olivier de 'Boughrara'-Sfax" (south of Tunisia: long. 34°59'05" N, lat. 10°32'56"; elevation: 125 m). Total genomic DNA was isolated from fresh, young leaves following a CTAB protocol originally developed by Murry and Tompson (1980) and further modified by De la Rosa et al. (2002).

Microsatellite assay. A set of eight microsatellite markers (ssrOeUA-DCA3, ssrOeUA-DCA9, ssrOeUA-DCA15, ssrOeUA-DCA16, ssrOeUA-DCA18, GAPU71B, GAPU101, and UDO99-043) were selected (Carriero et al., 2002; Cipriani et al., 2002; Sefc et al., 2000). The majority of the selected markers has been described to be very efficient for olive cultivar identification and population genetic studies (Baldoni et al., 2009). Polymerase chain reactions (PCRs) were carried out in a T-Gradient Thermoblock (Biometra, Goettingen, Germany) thermal cycler in a total volume of 20 µL containing 5 ng of genomic DNA, 1.5 mm MgCl₂, 60 µM dNTPs, 0.028 U/µL Taq polymerase (Biotools, Madrid, Spain), 0.2 µM forward primer (either FAM, HEX, or NED fluorescently labeled), and 0.2 µM of the corresponding unlabelled reverse primer. PCR steps were: initial denaturation at 95 °C for 5 min, 35 cycles with three steps: of 95 °C for 20 s (denaturation), 50 to 52 °C depending on the primers combination for 30 s (annealing), and 72 °C for 30 s (extension). A final extension step of 72 °C for 8 min was carried out. PCR products were separated in an automatic capillary sequencer, ABI PRISM 3100-AVANT GENETIC ANALYSER (Applied Biosystems/ Hitachi, Tokyo, Japan), using a definite size marker: GeneScan-500 ROX Size Standard (Applied Biosystems, Warrington, U.K.).

Data analysis. Amplified fragments were analyzed and scored using the GENEMAPPER 3.0 and GENOTYPER 3.7 software (Applied Biosystems). Both the expected and the observed heterozygosity (H_E/H_O) indices were calculated by using the CERVUS 3.0 software (Marshall et al., 1998) according to the equilibrium law of Hardy-Weinberg (Nei, 1987) as well as the polymorphic information content (PIC) (Botstein et al., 1980) and the frequency

of null alleles (Pemberton et al., 1995). The probability of identity (Waits et al., 2001) was computed by using the GIMLET 1.3.3 program (Valiere, 2002). The genetic relationships among the identified cultivars and the cophenetic correlations were analyzed by NTSYS-PC 2.02 software (Rohlf, 1998) using the DICE coefficient and the unweighted pair group method with arithmetic mean (UPGMA) (Sneath and Sokal, 1973). Factor analysis was carried out through the DARWIN 5.0.155 program (Perrier et al., 2003) and also by adopting the DICE coefficient for the calculation of the genetic distances.

Morphological description of endocarps. For endocarp description, 11 characters were selected from the pomological pattern widely used for olive cultivar characterization (Barranco et al., 2005). They are described in Table 1.

Results and Discussion

Microsatellites diversity in the "Boughrara"-Sfax germplasm collection. All accessions were screened for the eight selected microsatellite markers. Amplification products were obtained for the majority of them (98.8%). In the remaining samples, amplification failure was observed in three accessions at GAPU 101 locus, two accessions at UDO99-43 locus, and only one accession at both ssrOeUA-DCA15 and ssrOeUA-DCA16 locus. Microsatellites showed a high level of polymorphism throughout the analyzed accessions and revealed the existence of 64 alleles. The number of alleles per locus ranged from five (ssrOeUA-DCA15 and GAPU71B loci) to 12 (UDO99-43 locus), with an average of eight alleles per locus (Table 2). Such a high average reveals the elevated level of variability within the "Boughrara"-Sfax germplasm collection when compared with similar studies. The lower averages obtained in other studies can be explained by either the use of less polymorphic microsatellite markers (Belaj et al., 2004; Montemurro et al., 2005; Rekik et al., 2008) or by the use of less diversified plant material (Carriero et al., 2002; Noormohammadi et al., 2007). Higher averages have been reported by other workers (Baldoni et al., 2009; Taamalli et al., 2008) most likely as a consequence of using more

diversified genotypes. The analysis of the distinctive endocarp characters also revealed high variability in the current study. The diversity in Tunisian olive germplasm has been already reported by using morphological and agronomical traits (Mehri et al., 1997; Trigui et al., 2002) and molecular analysis as well (Grati-Kamoun et al., 2006; Rekik et al., 2008; Taamalli et al., 2006, 2008).

Expected heterozygosity values ranged between 0.54 (ssrOeUA-DCA15) and 0.82 (UDO99-43) with an average of 0.68, whereas H_o varied from 0.31 (ssrOeUA-DCA15) to 0.96 (in both ssrOeUA-DCA18 and UDO99-43) with an average of 0.77 (Table 2). Except for the ssrOeUA-DCA15 and ssrOeUA-DCA9 loci, H_E was lower than H_O . In the case of the ssrOeUA-DCA15 and ssrOeUA-DCA9 loci, H_E higher value of H_O is most likely the result of the positive probability of null alleles ($F_0 =$ 0.3 and $F_0 = 0.08$, respectively), whereas F_0 was negative in all the remaining cases. Null alleles usually exist when primers cannot anneal with DNA during PCR reaction as a result of point mutations (Pemberton et al., 1995) and thus it affects the estimated heterozygosis level in the correspondent locus. Positive null allele's frequencies have been also reported for the ssrOeUA-DCA15 locus by Baldoni et al. (2009) and Noormohammadi et al. (2007). Therefore, we recommend the mentioned marker to be discarded for further uses. The high H_0 average (0.77) confirms the important diversity among the "Boughrara"-Sfax germplasm collection. Similar averages (0.74 and 0.62) have been reported by Rekik et al. (2008) and Taamalli et al. (2008) in Tunisian cultivars.

Polymorphic information content (PIC) calculated from H_E and allele frequencies showed an average value of 0.63. This index ranged from 0.48 (ssrOeUA-DCA15) to 0.79 (UDO99-43) (Table 2). Similar PIC values have been reported in other studies (Bandelj et al., 2004; Noormohammadi et al., 2007). Except for the ssrOeUA-DCA15, all PIC values were higher than 0.5, which demonstrates the powerfulness of the markers. Two of them presented values over 0.7; hence, they can be considered particularly useful for genetic mapping (Bandelj et al., 2004). The results recently obtained by Baldoni et al. (2009), who recommend a consensus list of SSRs for worldwide olive

Table 2. Characteristics of the alleles observed after	er amplification with the	e eight simple sequence repeat	combinations and different genetic indexes
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	Size range	No. of alleles	No. of unique alleles	No. of allele patterns	H_E^z	H _O ^y	F_0^x	PIC ^w	PI^{v}
SsrOeUA-DCA18	166-180	7	0	13	0.72	0.96	-0.16	0.67	0.12
SsrOeUA-DCA3	228-249	6	0	11	0.65	0.72	-0.05	0.58	0.18
SsrOeUA-DCA15	241-264	5	1	9	0.54	0.31	+0.3	0.48	0.26
SsrOeUA-DCA16	120-172	11	2	19	0.74	0.86	-0.09	0.69	0.10
GAPU71B	117-141	5	0	11	0.66	0.91	-0.17	0.60	0.17
SsrOeUA-DCA9	161-208	11	1	16	0.60	0.52	+0.08	0.56	0.19
GAPU101	183-217	7	0	15	0.75	0.91	-0.1	0.70	0.10
UDO99-43	166-214	12	2	21	0.82	0.96	-0.08	0.79	0.05
Mean ^u		64	6	115	0.68	0.77	_	0.63	0.15

 ${}^{z}H_{E}$ = expected heterozygosity.

 ${}^{y}H_{O}$ = observed heterozygosity.

 ${}^{x}F_{0}$ = probability of null alleles.

^wPIC = polymorphic information content.

^vPI = probability of identity.

"Number of alleles, unique alleles, and allele patterns are total.

DNA typing, are in complete accordance with the current work. Seven of the eight SSRs used here for the analysis of "Boughrara"-Sfax germplasm collection together with the GAPU103A (not used in this work) are currently considered the most reliable markers for olive cultivar discrimination and for population genetic studies (Baldoni et al., 2009).

Discrimination and identification of Tunisian cultivars in the germplasm collection. The probability of identity (PI) can be considered as an index of the microsatellite discriminative power. In the present study, it ranged from 0.05 (UDO99-43) to 0.26 (ssrOeUA-DCA15) with an average of 0.15 (Table 2). In a similar study, Noormohammadi et al. (2007) obtained a higher average (0.19) for this index because they used a less discriminative set of microsatellites. In the current work, the cumulative PI value was $1.23 \ 10^{-7}$, which confirms the high discriminative level of the used microsatellite set. UDO99-43, GAPU101 ssrOeUA-DCA16, and ssrOeUA-DCA18 were the most informative markers (cumulative PI: 7.6 10^{-5}). They were able by themselves to discriminate 82.5% of the whole genotypes identified by the eight microsatellites altogether. UDO99-43 alone allows the discrimination of 21 genotypes, which means 52.5% of the entire identified genotypes. The mentioned marker showed a very low probability of identity (5. 10^{-2}). It can be considered the most powerful marker used in this work. The same observation has been reported by other workers (Baldoni et al., 2009; Belaj et al., 2004; Noormohammadi et al., 2007) and thus UDO99-43 is a highly recommended marker for olive cultivar discrimination.

Numerous duplications (44) were observed within the "Boughrara"-Sfax germplasm collection, whereas only 40 genotypes were identified among the 84 analyzed accessions (Table 3). The highest number of duplications (30) was detected regarding the

Table 3. Accessions studied, geographical origin, and proposed denominations.

	001	0 / 1	1				
Accession name	Origin	Diffusion level	Proposed name	Accession name	Origin	Diffusion level	Proposed name
Chemlali Sfax ^z	Sfax	Primary	Chemlali Sfax	Injassi Hchichina	Sfax	Secondary	Injassi Hchichina
Chemlali Sfax COI	Sfax	Primary	Chemlali Sfax	Chemlali Mahares ^r	Sfax	Secondary	Injassi Hchichina
Sig 3PG	Sfax	Primary	Chemlali Sfax	Chétoui 2	Tunis	Primary	Chetoui
Baldi Charqia BL8 ^v	Sfax	Primary	Chemlali Sfax	Chétoui 2 à mam	Tunis	Primary	Chetoui
BL24	Sfax	Primary	Chemlali Sfax	Marsaline	Tunis	Primary	Marsaline
Kotti K18	Sfax	Primary	Chemlali Sfax	Meski Bjawa ^q	Tunis	Primary	Marsaline
CHL Ouled Youssef Sig 13	Sfax	Primary	Chemlali Sfax	Fakhari Tataouine	Tataouine	Primary	Fakhari Tataouine
Hchichina 10	Sfax	Primary	Chemlali Sfax	Meski Zarzis ^p	Mednine	Primary	Fakhari Tataouine
Sehli Sned 7 ^v	Gafsa	Primary	Chemlali Sfax	Chemlali Chouamekh ^z	Mednine	Primary	Chemlali Chouamekh
Kotti K12	Sfax	Primary	Chemlali Sfax	Chemlali Tataouine10°	Tataouine	Primary	Chemlali Chouamekh
K17	Sfax	Primary	Chemlali Sfax	Chemlali Sig ^z	Sfax	Secondary	Chemlali Sig
Hchichina 6	Sfax	Primary	Chemlali Sfax	Kotti K14	Sfax	Secondary	Chemlali Sig
BL26	Sfax	Primary	Chemlali Sfax	Sig 14 PG	Sfax	Secondary	Chemlali Sig
Chemlali Bjawa ^v	Tunis	Primary	Chemlali Sfax	Besbessi	Tunis	Primary	Besbessi
Zalmati Zarzis ^v	Mednine	Primary	Chemlali Sfax	Lattout Sned 3	Gafsa	Primary	Lattout Sned
Sig 15	Sfax	Primary	Chemlali Sfax	CHL Sig 12PG ^z	Sfax	Tertiary	Horr Sig
Sehli Gafsa Dawla1 ^v	Gafsa	Primary	Chemlali Sfax	Chemlali Ontha	Tataouine	Secondary	Chemlali Ontha
				Tataouine ^z			Tataouine
BL23	Sfax	Primary	Chemlali Sfax	Chemlali Bouchouka ^z	Sfax	Tertiary	Chemlali Bouchouka
BL20	Sfax	Primary	Chemlali Sfax	Horr Louzir BL32 PG	Sfax	Tertiary	Horr Louzir
Horr Sfax ^v	Sfax	Primary	Chemlali Sfax	Chahla ^y	Sfax	Tertiary	Chahleya
BL16 G	Sfax	Primary	Chemlali Sfax	Semni Jbeniana Sig 0	Sfax	Tertiary	Semni Jbeniana
Hchichina 7	Sfax	Primary	Chemlali Sfax	Jeddaria Chaal	Sfax	Tertiary	Jeddaria Chaal
Kotti K9G	Sfax	Primary	Chemlali Sfax	Mallahi El Mouammar	Sfax	Tertiary	Mallahi El Mouammar
Kotti k21	Sfav	Drimory	Chemlali Sfay	Tounsi Gafea	Gafea	Secondary	Tounci Gafea
Kotti K20G	Sfax	Primary	Chemlali Sfax	Autre Zarrazi ^w	Madnina	Secondary	Zarrazi Ejibal
Sig 113 G	Sfax	Primary	Chemlali Sfax	Fch-Chabla/gy	Sfax	Tertiary	Ech-Chahla
Sehli IbenianaBI 15	Sfax	Primary	Chemlali Sfax	Neb Tataouine	Tataouine	Secondary	Neh Tataouine
Bent Louzir ^v	Siux	1 minur y	Chemian Slax	iveo rataoune	Tataballie	Secondary	iveo rataoune
BL37	Sfax	Primary	Chemlali Sfax	Kotti K9 PG	Sfax	Tertiary	Horr El Kotti
Chemlali Ouled Youssef ^v	Sfax	Primary	Chemlali Sfax	Lqam El Kotti Kotti K3	Sfax	Tertiary	Lqam El Kotti
Sahli Gafsa autre que Gtar ^v	Gafsa	Primary	Chemlali Sfax	Kotti K56 G	Sfax	Tertiary	Horr El Mouammar
Horr Blettech BL11 PG	Sfax	Primary	Chemlali Sfax	Chemlali Barrani ^z	Mednine	Secondary	Chemlali Barrani
Khechinet Sig	Sfax	Secondary	Khechinet Sig	K56PG	Sfax	Tertiary	Lqam El Mouammar
SIG19	Sfax	Secondary	Khechinet Sig	Chemlali Balhi ^z	Mednine	Secondary	Chemlali Balhi
Bou Bazzoula BL19PG ^u	Sfax	Secondary	Khechinet Sig	Sig 112PG	Sfax	Tertiary	Zeitoun Khdim El Bey
Kotti K23 (avec mam)	Sfax	Secondary	Khechinet Sig	Zeitoun Boubazzoula COI ^x	Sfax	Tertiary	Zeitoun Boubazzoula
BL2	Sfax	Tertiary	Horr Charqia	Kotti K32	Sfax	Tertiary	Zeitoun El Manacher
BL31	Sfax	Tertiary	Horr Charqia	BL32 G	Sfax	Tertiary	Chemlali Lach'hab
Chemlali Zarzis ^z	Mednine	Primary	Chemlali Zarzis	Zarrazi Kgh ^w	Mednine	Primary	Zarrazi Kgh
Mfartah Hchichina H9 ^t	Sfax	Primary	Chemlali Zarzis	Zeitoun Boubazzoula COI ^x	Sfax	Secondary	Mbazzel Kbir
Jemri Bouchouka	Tataouine	Primary	Jemri Bouchouka	Dhokar Nafti	Mednine	Secondary	Dhokar Nafti
Chemchali Gtar 3 ^s	Gafsa	Primary	Jemri Bouchouka	Zarbout Louzir	Sfax	Secondary	Zarbout Louzir
Kotti K30 PG	Sfax	Primary	Jemri Bouchouka	Kotti K11	Sfax	Tertiary	Zeyeti El Kotti

^zCases of homonymy in "Chemlali" denomination.

^yCases of homonymy in "Chahla" denomination.

^xCases of homonymy in "Zeitoun Boubazzoula" denomination.

"Cases of homonymy in "Zarrazi" denomination.

vCases of synonymy in Chemlali Sfax cultivar.

^uCase of synonymy in Khechinet Sig cultivar.

^tCase of synonymy in Chemlali Zarzis cultivar.

^sCase of synonymy in Jemri Bouchouka cultivar.

^rCases of synonymy of Injassi Hchichina cultivar.

^qCase of synonymy of Marsaline cultivar.

^pCase of synonymy of Fakhari Tataouine cultivar.

°Case of synonymy of Chemlali Chouamekh cultivar.

'Chemlali Sfax' genotype explained by the wide distribution of this cultivar across the country.

The identification achieved in the present work is extremely important for better management of the "Boughrara"-Sfax germplasm collection. Furthermore, it can be useful for providing a well-identified material to olive nurseries.

Synonyms and homonyms. Several cases of synonyms (different denomination for the same genotype) and homonyms (same denomination for different genotype) were detected in the collection. The highest number of synonyms (nine) was also observed in the cultivar Chemlali Sfax, which is the most widely cultivated in the southern part of the country. Moreover, our results showed that 'Chemlali Sfax' synonyms exist in the majority of the olive-producing regions of Tunisia (Sfax, Gafsa, Mednine, and Tunis) with different denominations (Table 3). Among them, "Sehli Sned," "Sehli Gafsa," and "Sehli Jbeniana Bent Louzir" accessions are associated with the same adjective "Sehli" (which means "olive from the coast"). The remaining cases of synonymy are represented in Table 3. Similar cases of synonymy have been reported in Tunisian cultivars belonging to different collections and olive groves (Rekik et al., 2008; Taamalli et al., 2008). Endocarp characters displayed low variability among all duplicated profiles (Fig. 1). This scarce variability is most likely the result of the influences of the field and tree conditions (nutrition, water and phytosanitary status, etc.). However, more clear differences were revealed in all the endocarp characters analyzed between these individuals identified as independent genotypes.

On the other hand, a typical case of homonymy among Tunisian olive groves is that of "Chemlali," which has been reported by morphological, agronomical, and molecular studies (Grati-Kamoun et al., 2006; Mehri et al., 1997; Rekik et al., 2008; Taamalli et al., 2008; Trigui et al., 2002). This homonymy was also observed within the "Boughrara"-Sfax germplasm collection. In fact, "Chemlali" is a common and general denomination used by growers, although it has been reported that a high level of polymorphism exists within this denomination. The present work



Fig. 1. Dedrogram of the identified cultivars based on DICE coefficient and unweighted pair group method with arithmetic mean (UPGMA) cluster analysis of the simple sequence repeat (SSR) data obtained. A picture of a representative endocarp is included for each accession.

demonstrates that polymorphism among this denomination is extremely high. In some cases, only one of 16 alleles was shared between homonymous cultivars such as 'Chemlali Sfax' and 'Chemlali Balhi'. Several others cases of homonymy were detected in the collection (Table 3). Homonymous confusions are basically the result of the traditional denomination system. Cultivar names are mostly based on generic denominations like "Injassi" (pear form) or "Dhokar" (pollinator), etc. Similar cases have been reported in other olive-producing countries after using molecular markers (Belaj et al., 2001; Khadari et al., 2003; Mekuria et al., 1999; Noormohammadi et al., 2007; Trujillo et al., 2005). Homonyms were analyzed using endocarp morphology as the criteria. As an example, high variability in endocarp characteristics was detected particularly within Chemlali's denominations, in which the form and symmetry of the endocarp showed conspicuous differences as well as the form of the base, the roughness of the surface, and the number and distribution of vascular bundle over the surface.

Genetic relationships. Genetic relationships between identified genotypes were studied by means of the DICE coefficient. All duplicated samples were discarded from the analysis. The highest value of cophenetic coefficient (0.75) was obtained using the UPGMA clustering method. Therefore, the mentioned method was adopted as the most appropriate for further clustering (Fig. 1). DICE coefficient ranged from 0.0 (between cultivars like Mallahi El Mouammar and Chemlali Ontha Tataouine) to 0.96 (between 'Horr El Kotti' and 'Zitoun Boubazzoula'). The dendrogram demonstrated the presence of four major groups, five cultivars widely separated, and several very similar genotypes. Group 1 contains eight clearly discriminated genotypes. The second cluster was formed by seven genotypes, including the cultivar Chetoui, the second most widely distributed cultivar in Tunisia. 'Chetoui' was clearly separated within the cluster (0.61 DICE coefficient with 'Zeitoun Khdim El Bey', the closest cultivar). Two genotypes of this group ('Horr Sig' and 'Zeitoun Khdim El Bey') were very similar to 'Chemlali Sig' with DICE coefficients higher than 0.85. The third cluster was the most important in terms of genotype number (15 genotypes). It included the widely distributed cultivar, Chemlali Sfax, which is situated close to three genotypes with DICE similarity coefficients higher than 0.92 ('Zeyeti El Kotti', 'Lqam El Kotti', and 'Horr El Mouammar'). The fourth cluster included five genotypes. Two of them ('Lquam El Mouammar' and 'Zeitoun El Manchar') presented a DICE coefficient close to 0.93. Other workers have also reported such a high level of similarity between olive cultivars from several regions of the world (Belaj et al., 2004; Montemurro et al., 2005; Noormohammadi et al., 2007) and also among Tunisian cultivars (Rekik et al., 2008). These and other studies support the idea of these genotypes being considered as independent cultivars, although similarity



Fig. 2. Factor analysis of the identified genotypes according to their main uses.

among them is relatively high. In fact, microsatellites, as low proportions of the genome, were unable to detect intracultivar polymorphism as a result of mutations occurring in very long-living trees asexually propagated (Baldoni et al., 2009). This proposal is supported by previous agronomical observations within the collection.

On the other hand, the five cultivars Zarbout Louzir, Chemlali Chouamek, Marsaline, Mallahi El Mouammar, and Chemlali Balhi were clearly separated and distant from any other observed clusters. This can be attributed to their different origins. For example, Marsaline is a French cultivar, whereas the denomination 'Mallahi El Mouammar', which can be translated as "settler's table olive," is most likely a foreign cultivar. The morphological characters of the endocarps were in accordance with the dendrogram of the genetic relationship shown here (Fig. 1). For example, only very small differences were detected between Horr Sig and Chemlali Sig cultivars, which showed a high coefficient of DICE similarity (0.96). On the contrary, wide morphological differences were observed between the endocarps of 'Mallahi El Mouammar' and 'Chahleya' (with DICE coefficient 0.27).

Finally, factor analysis has been conducted to characterize clustering tendencies among the identified cultivars (Fig. 2). The

analysis revealed only 16.38% and 12.14% of clustering according to two principal components. Such low percentages only allow considering the clustering as tendencies and not as clear separations. The test ended with the clustering of the four olive table cultivars altogether. This is probably a result of Tunisian growers' selections during the olive cultivation history of the country. The selection has been made according to producer requirements, genotypes intended for table olive consumption, and others for olive oil and mixture uses. However, no clustering tendencies were reported according to the cultivar's geographical origin, probably as the result of the land continuity between oliveproducing regions, which promoted the genetic exchange (data not shown). Cultivar clustering in accordance to the geographical area of cultivation has been reported by different authors (Besnard et al., 2001b; Claros et al., 2000; Fabbri et al., 1995). However, it is not generally observed, even using powerful genetic markers like SSRs, except in those regions where the genetic exchange has been very limited.

In conclusion, the total number of alleles, the number of allele per locus, and the H_O demonstrate the high genetic diversity in the "Boughrara"-Sfax germplasm collection located in Tunisia. A high number of mislabeled accessions and several synonymous and homonymous cases were reported. These

results put emphasis on the necessity of an improved classification within this collection, containing a very important part of the Tunisian olive genetic resources.

With the exception of ssOeUA-DCA15, all the microsatellites used confirmed to be reliable for olive cultivar classification in a germplasm collection. Therefore, the set used here can be recommended for DNA fingerprinting in other Tunisian collections and thus contribute to the establishment of a universal molecular database of olive genetic resources based on standard SSR markers as proposed by Baldoni et al. (2009).

The results of the SSR classification were in good agreement with the differences observed after analyzing morphological characters of the endocarps. We therefore suggest that the incorporation of both approaches is extremely advisable for accurate characterization of olive germplasm.

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