

Supplementary**Morphological and genetic diversity in olive (*Olea europaea* subsp. *europaea* L.) clones and varieties****Supplementary Table S1a: Phenotypic traits of the olive leaf and fruit and their description**

Traits*	
Leaf (1 trait)	
Shape (FFE)	1- Elliptic (L/W < 4) 2- Elliptic-lanceolate (L/W 4-6) 3- Lanceolate (L/W > 6)
Fruit (4 traits)	
Shape (position A) (FOF)	1- Spherical (L/W < 1.25) 2- Ovoid (L/W 1.25-1.45) 3- Elongated (L/W > 1.45)
Position of maximum transverse diameter (position B) (DIF)	1- Towards base 2- Central 3- Towards apex
Nipple (MUF)	1- Absent 2- Tenuous 3- Obvious
Location of start of colour change (VRF)	1- From the base Uniformly 2- Across the whole epidermis 3- From the apex

*Abbreviations: FFE- Leaf shape; FOF - Fruit shape (position A); DIF - Position of maximum transverse diameter (position B) in fruit; MUF - Nipple of fruit; VRF - Location of start of fruit colour change; L/W- Fruit length/width

Supplementary Table S1b: Phenotypic traits of endocarpe and their description

Endocarpe (10 caractères)*	
Shape (position A) (FOE)	1- Spherical (L/W < 1.4) 2- Ovoid (L/W 1.4-1.8) 3- Elliptic (L/W 1.8-2.2) 4- Elongated (L/W > 2.2)
Symmetry (position A) (SAE)	1- Symmetric 2- Slightly asymmetric 3- Asymétric
Symmetry (position B) (SBE)	1- Symétric 2- Slightly asymmetric
Position of maximum transverse diameter (position B) (DIE)	1- Towards base 2- Central 3- Towards apex
Apex (position A) (APE)	1- Pointed 2- Rounded
Base (BAE)	1- Truncate 2- Pointed 3- Rounded
Surface (position B) (SUE)	1- Smooth 2- Rugose 3- Scabrous
Number of grooves (NSE)	1- Low (< 7) 2- Medium (7-10) 3- High (> 7)
Distribution of the grooves (DSE)	1- Regular 2- Grouped around the suture
Termination of the apex (position A) (TAE)	1- Without mucron 2- With mucron

*Abbreviations: FOE- Endocarp shape (position A) ; SAE - Endocarp symmetry (position A); SBE - Endocarp symmetry (position B); DIE - Position of maximum transverse diameter (position B) in endocarp; APE - Endocarp apex (position A); BAE - Endocarp base; SUE - Endocarp surface (position B) ; NSE - Number of endocarp grooves; DES - Distribution of endocarp grooves; TAE - Termination of the endocarp apex (position A); L/W - Endocarp length/width

Supplementary Table S2: Morphological characters recorded

Genotypes	FFE	FOF	DIF	MUF	VRF	FOE	SAE	SBE	DIE	APE	BAE	SUE	NSE	DES	TAE
Picholine de Languedoc	Elliptic (L/W<4)	Elongated	Central	Absent	From the apex	Elongated	Asymetric	Symetric	Towards apex	Pointed	Truncate	Rugose	Medium (7-10)	Regular	With mucro
Leccino	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Asymetric	symetric	Central	Rounded	Rounded	Rugose	Medium (7-10)	regular Grouped around the suture	With mucro
Branquita	Elliptic (L/W<4)	Spherical	Central	Absent	From the apex	Ovoid	Symetric	symetric	Towards apex	Rounded	Truncate	Smooth	Low (< 7)		With mucro
Ayvalik	Lanceolate (L/W >6)	Ovoid	Central	Absent	From the apex	Ovoid	Symetric	symetric	Central	Rounded	Rounded	Rugose	Medium (7-10)	regular	With mucro
Carolea	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Obvious	across the whole epidermis	Elliptic	Slightly asymetric	symetric	Towards apex	Rounded	Truncate	Rugose	Medium (7-10)	regular	With mucro
Sorani	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the base	Uniformly, Elliptic	Slightly asymetric	symetric	Towards apex	Pointed	Truncate	Rugose	Medium (7-10)	regular	Without mucro
Arbequine	Elliptic (L/W<4)	Spherical	Towards base	Absent	across the whole epidermis	Ovoid	Symetric	symetric	Central	Rounded	Rounded	Rugose	Medium (7-10)	regular	Without mucro
Pich Marocaine	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Slightly asymetric	Symetric	Central	Pointed	Rounded	Rugose	High (>10)	Regular Grouped around the suture	With mucro
Menara	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Slightly asymetric	Symetric	Central	Pointed	Rounded	Rugose	High (>10)		With mucro
Haouzia	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Slightly asymetric	Symetric	Central	Pointed	Rounded	Rugose	Medium (7-10)	Regular	With mucro
S7	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Tenuous	From the apex	Elliptic	Asymetric	Slightly asymetric	Central	Pointed	Pointed	Rugose	Medium (7-10)	Regular	With mucro
S8	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Tenuous	From the apex	Elliptic	Asymetric	Symetric	Central	Pointed	Rounded	Rugose	Medium (7-10)	Regular Grouped around the suture	With mucro
M14	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Symetric	Symetric	Central	Pointed	Pointed	Scabrous	Medium (7-10)	Regular Grouped around the suture	Without mucro
M16	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Elliptic	Slightly asymetric	Symetric	Central	Pointed	Pointed	Rugose	Medium (7-10)	Regular Grouped around the suture	Without mucro
M26	Elliptic lanceolate (L/W 4-6)	Ovoid	Central	Absent	From the apex	Ovoid	Slightly asymetric	Symetric	Central	Pointed	Rounded	Rugose	Medium (7-10)	Regular Grouped around the suture	With mucro
Bakhboukh beldi	Lanceolate (L/W >6)	Ovoid	Central	Tenuous	across the whole epidermis	Ovoid	Slightly asymetric	Symetric	Central	Rounded	Rounded	Rugose	Medium (7-10)	Regular Grouped around the suture	With mucro
Bouhouk Rkik	Lanceolate (L/W >6)	Ovoid	Central	Tenuous	From the apex	Elliptic	Slightly asymetric	Symetric	Towrds apex	Pointed	Pointed	Smooth	Medium (7-10)	Regular	Without mucro
Bouhouk Laghlid	Lanceolate (L/W >6)	Elongated	Towards apex	Tenuous	across the whole epidermis	Ovoid	Slightly asymetric	Symetric	Towrds apex	Pointed	Rounded	Rugose	Medium (7-10)	Regular	Without mucro
Bouhouika	Lanceolate (L/W >6)	Elongated	Central	Obvious	across the whole epidermis	Ovoid	Slightly asymetric	Symetric	Towrds apex	Pointed	Pointed	Smooth	Medium (7-10)	Regular	Without mucro

*Abbreviations: See the footnotes of supplementary Tables S1a and S1b

Supplementary Table S3: Size of the amplified fragment, number alleles and observed heterozygosity at 20 microsatellite loci in the 19 olive genotypes

Locus	Primer sequences (5'→3')	Annealing temperature (°C)	Size range (bp)	Number of alleles	Observed heterozygosity (<i>H_o</i>)
DCA3 ^a	F: CCCAAGCGGAGGTGTATATTGTTAC R: TGCTTTTGTCTGTTTGAGATGTTG	55	230-251	6	0.79
DCA5 ^a	F: AACAAATCCCATACGAACTGCC R: CGTGTGCTGTGAAGAAAATCG	55	-	1	-
UDO36 ^c	F: AACACTGTGCCACCTCAACA R: GAACCCAACCCCATCTTAC	55	141-162	7	1
DCA15 ^a	F: GATCTTGTCTGTATATCCACAC R: TATACCTTTTCCATCTTGACGC	55	-	1	-
GAPU71B ^b	F: GATCAAAGGAAGAAGGGGATAAA R: ACAACAAATCCGTACGCTTG	55	120-153	5	1
UDO12 ^c	F: TCACCATTCTTAACTTCACACCA R: TCAAGCAATTCCACGCTATG	55	-	1	-
GAPU71A ^b	F: GATCATTTAAAATATTAGAGAGAGAGA R: TCCATCCATGCTGAACTT	55	192-209	3	0.42
UDO14 ^c	F: TTCCCCTTATTCATGTGAACC R: ACTGCAGTTTGGGAATCAAA	55	90-120	5	1
UDO17 ^c	F: GCCCACAACCTCTTGAACC R: GCGATTTTTCCCTGTATTTAGGT	55	152-168	5	0.47
DCA17 ^a	F: GATCAAATTCTACCAAAAATATA R: TAATTTTTGGCAGTAGTATTGG	55	103-210	9	0.84
UDO27 ^c	F: TCCGTGCAAACCATGAAATA R: TTGATGACTAGCACACATGTAGGA	55	-	1	-
UDO34 ^c	F: CTCTCGGGCATGTATCATT R: TTGCATATTTGTATGATTCATT	50	-	1	-
GAPU59 ^b	F: CCCTGCTTTGGTCTTGCTAA R: CAAAGGTGCACTTTCTCTCG	55	205-237	3	0.89
UDO44 ^c	F: AATCCGACAAGTTGTGTGTG R: CACAGCACCCAACCAGATT	55	-	1	-
UDO6 ^c	F: TCAGTTTGTGTCCTTTAGTGGA R: TTGTAATATGCCATGTAACCTCGAT	55	146-189	5	0.74
DCA 1 ^a	F: CCTCTGAAAATCTACACTCACATCC R: TGAACAGAAAGAAGTGAACAATGC	55	-	1	-
DCA 7 ^a	F: GGACATAAAACATAGAGTGCTGGGG R: AGGGTAGTCCAACCTGCTAATAGACG	55	-	1	-
DCA 4 ^a	F: CTTAACTTTGTGCTTCTCCATATCC R: AGTGACAAAAGCAAAAGACTAAAGC	55	-	1	-
DCA 11 ^a	F: GATCAAACACTGCACGAGAGAG R: TTGTCTCAGTGAACCCTTAAACC	55	-	1	-
DCA 13 ^a	F: GATCAGATTAATGAAGATTTGGG R: AACTGAACCTGTGTATCTTGCATCC	55	-	1	-

^aSefc et al. (2000); ^bCarriero et al. (2002) and ^cCipriani et al. (2002)