

Interspecific Relationships in *Pistacia* Based on RAPD Fingerprinting

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Abstract. Phylogenetic relationships among nine species in the genus *Pistacia* were studied by randomly amplified polymorphic DNA (RAPD) analysis. The following species were included: *P. atlantica*, *P. terebinthus*, *P. eurycarpa*, *P. vera*, *P. integerrima*, *P. mexicana*, *P. palaestina*, *P. lentiscus*, and *P. khinjuk*. Genomic DNA was extracted from leaf tissue and RAPD analysis was performed using 20 primers. A total of 242 fragments were generated and 228 bands were polymorphic at the inter-specific level. Subjecting these data to phylogenetic analysis yielded a shortest cladogram that is 338 steps long, featuring two main groups. *P. vera*, *P. khinjuk*, *P. eurycarpa*, *P. atlantica*, and *P. integerrima* were included in one group, while *P. terebinthus*, *P. palaestina*, *P. mexicana*, and *P. lentiscus* formed the second group. The first group included species with single-trunked and big trees, whereas the species included in the second group mostly grow as shrubs or small trees. The cladogram showed that the closest pairs of species were *P. terebinthus* and *P. palaestina*, *P. eurycarpa* and *P. atlantica*, *P. vera* and *P. khinjuk*, and *P. mexicana* and *P. lentiscus*. We suggest that *P. palaestina* is in fact a variety of *P. terebinthus* in view of the small genetic distance between them. This study also showed that *P. eurycarpa* (syn. *P. atlantica* var. *kurdica*) is a distinct species from *P. atlantica*, rather than a variety within the same species.

Studies on the phylogeny of the genus *Pistacia* L. (Anacardiaceae) are rather few, and are mostly based on morphological characterization. The first monographic study of the genus was made by Engler (Zohary, 1952) who listed eight species and a few varieties. After Engler, several species have been added by different authors. So far, the most complete taxonomic study was done by Zohary (1952), who included 11 species in the genus *Pistacia* and divided them into four sections: 1) Section Lenticella Zoh.—*P. mexicana* HBK, *P. texana* Swingle; 2) Section Eu-Lentiscus Zoh.—*P. lentiscus* L., *P. weinmannifolia* Poisson, *P. saportae* Burnat; 3) Section Butmela Zoh.—*P. atlantica* Desf.; and 4) Section Eu-terebinthus—*P. terebinthus* L., *P. palaestina* Bois., *P. khinjuk* Stocks, *P. vera* L., and *P. chinensis* Bge. Following Zohary, several other authors described and classified *Pistacia* species based on their morphology. Yaltirik (1967a) classified *Pistacia* species in Turkey and added a new species, *P. eurycarpa* Yalt., that had been designated as *P. atlantica* var. *kurdica* by Zohary (1952). Kafkas and Perl-Treves (2001) suggested, based on morphological and molecular evidence, that *P. eurycarpa* is a close relative of *P. atlantica*, but is a different species. Zohary considered *P. palaestina* as a

separate species, whereas Yaltirik (1967a) retained it as a variety of *P. terebinthus*. Zohary (1972) described five *Pistacia* species in Israel. Grundwag and Werker (1976) described wood anatomy of *Pistacia* species in Israel, and Dong and Baas (1993) performed a similar study in China. Al-Yafi (1978) divided *P. atlantica* into four subspecies according to their leaf morphology. Kokwaro and Gillet (1980) described a new *Pistacia* species in East Africa, *P. aethiopica* Kokwaro, based on leaf morphology and tree size. The latter authors considered this species to be synonymous with *P. lentiscus* var. *emarginata* Engl., or a variety of *P. atlantica*. Lin et al. (1984) characterized leaf morphology, photosynthesis and leaf conductance of nine *Pistacia* species. El-Oqlah (1996) described *Pistacia* species in Jordan morphologically and anatomically.

Among these species, *Pistacia vera*, the pistachio, has edible nuts and commercial importance. The other species grow in the wild and their seeds are used mainly as a rootstock seed source and rarely used for fresh consumption, oil extraction, and soap production. All *Pistacia* species are dioecious and wind-pollinated. *P. vera* is believed to be the most ancestral species and the other species are probably its derivatives (Zohary, 1952). There are two centers of diversity of *Pistacia*: one comprises the Mediterranean region of Europe, Northern Africa, and the Middle East countries. The second comprises the Eastern part of Zagros mountains from Crimea to the

Caspian Sea. Pistachio cultivation extended westward from its center of origin to Italy, Spain, and other Mediterranean regions of Southern Europe, North Africa, and the Middle East, as well as eastward to China, and more recently to the United States and Australia (Hormaza et al., 1994, 1998; Maggs, 1973). Currently, Iran, United States, Turkey, and Syria are the main pistachio producers in the world, contributing over 90% of the world production (FAO 2000).

Molecular studies addressing the genus *Pistacia* are few. The pollen isozyme patterns of nine different enzymes were studied by Louskas and Pontikis (1979) in *P. vera*, *P. terebinthus* and *P. lentiscus*, to assay their inter-specific relationships. They found a closer phylogenetic relationship between *P. vera* and *P. terebinthus* than between *P. vera* and *P. lentiscus*. In our previous study (Kafkas and Perl-Treves, 2001), we mainly focused on the intra-specific relationship in the germplasm of three species (*P. atlantica*, *P. terebinthus*, and *P. eurycarpa*). Parfitt and Badenes (1997) were the first to provide a classification of 10 *Pistacia* species at the molecular level, and characterized these species based on chloroplast DNA profiles and subdivided the genus into two sections, 'Terebinthus' (*P. vera*, *P. khinjuk*, *P. atlantica*, *P. integerrima*, *P. chinensis*, *P. terebinthus*) and 'Lentiscus' (*P. mexicana*, *P. texana*, *P. weinmannifolia*, *P. lentiscus*). The latter authors could not discriminate *P. vera* from *P. khinjuk*, and *P. texana* from *P. mexicana*. In the present study, we aimed to clarify the taxonomic relationships among nine species in the genus *Pistacia* by RAPD analysis, which targets the nuclear rather than organellar genome. Two of the species, *P. palaestina* and *P. eurycarpa*, were not included in the study of Parfitt and Badenes (1997).

Materials and Methods

Plant material. Plant material for this study was collected from three different countries: Turkey, Israel, and the United States (Table 1).

Genomic DNA extraction and RAPD reactions. Leaf samples were collected from a single representative accession per species, frozen in liquid nitrogen and stored at -70°C until use. Genomic DNA was extracted from leaf tissue by the CTAB method of Doyle and Doyle (1987) with minor modifications (Kafkas and Perl-Treves, 2001).

Randomly amplified polymorphic DNA (RAPD) analysis was performed according to Williams et al. (1990) with minor modifications. Amplification reactions were carried out in a 25 μL volume containing 10 mM Tris-HCl, pH 9.0, 50 mM KCl, 1.5 mM MgCl_2 , 0.1% Triton X-100, 0.2 μM primer, 100 μM each of dATP, dGTP, dCTP and dTTP, 1 unit of Taq DNA polymerase and 10 ng of genomic DNA. Each reaction mixture was overlaid with mineral oil. DNA reactions were performed in a PTC-100 thermal cycler (MJ-Research, Watertown, Mass.). The program included 1 cycle of 2 min at 94°C , followed by 35 cycles of 45 s at 94°C , 1 min at 36°C , and

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Table 1. List of the plant materials used in this study

Species	No.	Country	Accession no.	Province	County	Village
<i>P. eurycarpa</i>	1	Turkey	56-K-07	Siirt	Aydinlar	Ikizbaglar
<i>P. atlantica</i>	1	Turkey	01-A-04	Adana	Center	Kuyumcular
<i>P. terebinthus</i>	3	Turkey	09-T-02	Adana	Center	Balcali
<i>P. integerrima</i>	1	Turkey	PI-006	Adana	Univ. of Çukurova	
<i>P. vera</i>	1	Turkey	Cv Siirt	Gaziantep	Pistachio Research Institute	
<i>P. palaestina</i>	3	Israel	---	Jerusalem hills		
<i>P. lentiscus</i>	1	Israel	---	Jerusalem hills		
<i>P. khinjuk</i>	1	USA	No:7	Dr. Parfitt, Univ. of California, Davis		
<i>P. mexicana</i>	1	USA	DPIS0027	National Clonal Germplasm Repository, USDA-ARS, Davis, Calif.		

2 min at 72 °C, for denaturing, annealing and primer extension, respectively. The last cycle was followed by a final incubation for 5 min at 72 °C and the PCR products were stored at 4 °C prior to analysis. Amplification products were analyzed by gel electrophoresis in 1.8% agarose in 1× TBE buffer, stained with ethidium bromide and photographed under UV light. From a preliminary screen of 200 RAPD primers, the 20 most polymorphic primers (Univ. of British Columbia primers #147, 165, 189, 302, 304, 308, 319, 322, 327, 338, 345, 346, 348, 353, 354, 356, 376, 381, 383, 396) were selected and used for fingerprinting and characterization of the nine *Pistacia* species.

Band scoring and parsimony analysis. Only the most clear and strong bands were used for phylogenetic analysis. Reproducibility of the patterns was tested by running the reactions in duplicates or in triplicates. Parsimony analysis was performed using the PAUP 3.1 program (Swofford, 1993) with different Heuristic search-options, and a 50% majority rule consensus tree was constructed from 100 bootstrap replicates of the same data. In addition, pair-wise genetic distances between all pair-wise combinations of the species were calculated by the same program. Such values represent the proportion of different bands between all the possible pairs of species.

Results

The RAPD technique was used to characterize the nine *Pistacia* species and clarify the relationships between them. A total of 242 fragments were generated by 20 arbitrary-sequence primers, and 228 bands out of these were polymorphic at the inter-specific level. RAPD fingerprinting patterns of *Pistacia* genotypes using primer BC348 are shown as an example in Fig. 1. Data were analyzed using the PAUP program, to obtain cladograms that depict the likely relationship among the species. The two shortest trees were 338 steps long, and one of them is shown in Fig. 2. A 50% consensus-cladogram resulting from bootstrap analysis of the same data with 100 replicates is given in parenthesis. Pair-wise genetic distances between all the species were calculated by the same program (Table 2).

According to these cladograms, the nine *Pistacia* species are well separated from each other, and form two main groups or clusters. The two groups are separated by a branch of 14 steps, appearing in 99% of the bootstrap replicates. One group includes *P. vera*, *P. khinjuk*, *P. eurycarpa*, and *P. atlantica*, while *P.*

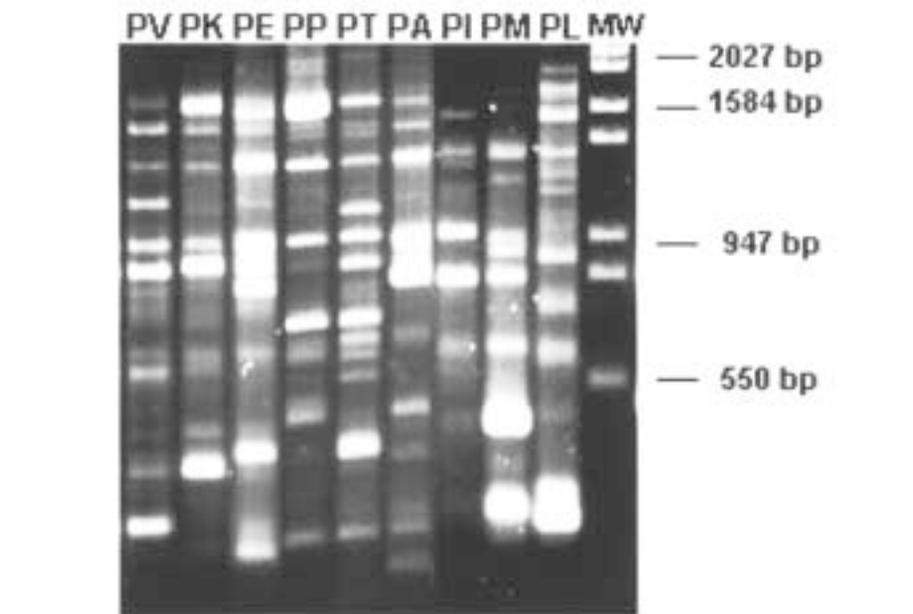


Fig. 1. RAPD banding patterns of nine *Pistacia* species using primer BC348. MW = molecular weight standards; the size of fragments is given in base pairs. PV = *P. vera*, PK = *P. khinjuk*, PE = *P. eurycarpa*, PA = *P. atlantica*, PI = *P. integerrima*, PP = *P. palaestina*, PT = *P. terebinthus*, PM = *P. mexicana*, PL = *P. lentiscus*.

terebinthus, *P. palaestina*, *P. mexicana*, and *P. lentiscus* are included in the second group. *P. integerrima*, however, does not cluster within any of the two groups and is well-separated from both of them. The cladograms also show that *P. terebinthus* and *P. palaestina* are a close pair of species, as are *P. eurycarpa* and *P. atlantica*, *P. vera* and *P. khinjuk*, and *P. mexicana* and *P. lentiscus*. A branch of 20 steps appearing in 87% of the bootstrap replicates separates *P. vera* and *P. khinjuk* from the rest of the tree suggesting that, among the wild species, *P. khinjuk* is the closest relative of cultivated pistachio. A branch of 20 steps appearing in 78% of the bootstrap replicates separates *P. eurycarpa* and *P. atlantica* from *P. vera* and *P. khinjuk*, and all four species are separated by a branch of 13 steps, appearing in 100% replicates, from the rest of the tree.

Discussion

The most comprehensive studies concerning the taxonomic relationships between *Pistacia* species was performed by Zohary (1952) and, at the molecular level, by Parfitt and Badenes (1997). The former author subdivided the genus into four sections, whereas the latter ones subdivided it into two sections,

'*Terebinthus*' and '*Lentiscus*'. According to the rules of botanical nomenclature presented in the International Code of Botanical Nomenclature (ICBN), any subgeneric taxon including the type species of a genus must be designated by the same name as the genus (the autonym). Thus, section '*Terebinthus*' would, under the ICBN, become section *Pistacia*. The primary characters for taxonomic identification of *Pistacia* species were leaf rachis wing, leaflet size and shape, number of leaflet pairs, absence or presence of terminal leaflet, leaflet apex shape, nut size, and shape. Leaf rachis wing was one of the most discriminative characteristics used by Zohary (1952), who classified *P. atlantica* in a distinct Section, *Butmela*, on the basis of such character.

In our study, we obtained two groups of species: one group comprised *P. vera*, *P. khinjuk*, *P. eurycarpa*, *P. atlantica*, and *P. integerrima*; and the second group included *P. terebinthus*, *P. palaestina*, *P. mexicana*, and *P. lentiscus*. Considering their morphology, the species in the first group form big trees, whereas the second group tends to form small trees or shrubs. Parfitt and Badenes (1997) found that *P. vera* was more closely related to *P. terebinthus* than to *P. integerrima*. *P. terebinthus* trees in California grow as single,

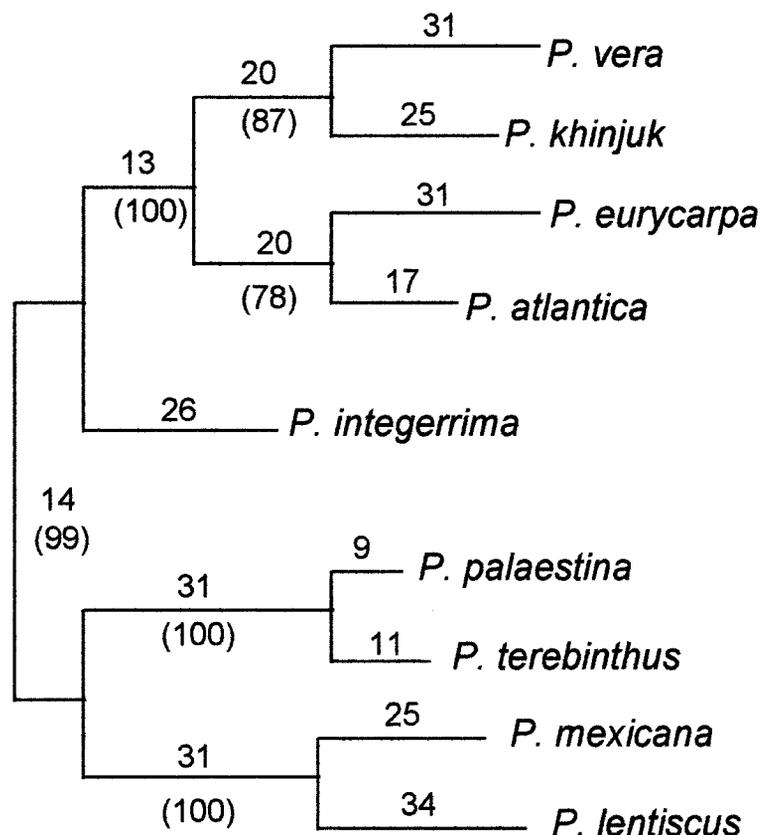


Fig. 2. Parsimony analysis of molecular fingerprinting data from nine *Pistacia* species. The database included 242 RAPD bands. A heuristic search was conducted by the PAUP software using TBR optimization, resulting in two trees of 338 steps. One of the trees is depicted. Numbers indicate the length (no. of steps) of each branch. Percentage values of a 50% majority rule consensus tree obtained from 100 Bootstrap replicates of the same data are shown in parentheses.

Table 2. Pair-wise distances among *Pistacia* species² calculated by PAUP program.

	PV	PK	PE	PA	PI	PP	PT	PM	PL
PV	---	0.227	0.264	0.285	0.364	0.455	0.450	0.504	0.550
PK		---	0.289	0.314	0.314	0.409	0.413	0.488	0.508
PE			---	0.186	0.322	0.459	0.467	0.517	0.537
PA				---	0.244	0.393	0.409	0.463	0.479
PI					---	0.277	0.293	0.351	0.376
PP						---	0.080	0.339	0.347
PT							---	0.339	0.360
PM								---	0.240
PL									---

²PV = *P. vera*, PK = *P. khinjuk*, PE = *P. eurycarpa*, PA = *P. atlantica*, PI = *P. integerrima*, PP = *P. palaestina*, PT = *P. terebinthus*, PM = *P. mexicana*, PL = *P. lentiscus*

very large trees (D.E. Parfitt, 1997, personal communication). Whitehouse (1957) reported that, in California, *P. terebinthus* grows as a single-trunked tree and has a narrow leaf rachis wing. However, according to Zohary (1952) and to our observations under different ecological conditions, *P. terebinthus* usually grows as a shrub, rarely as a single tree, and never exhibits a leaf rachis wing. Ayfer and Serr (1961) also reported that *P. terebinthus* trees in California are similar to *P. atlantica*, based on tree and leaf characteristics, and may be a variety of *P. atlantica*. Therefore, we may conclude that the trees identified as *P. terebinthus* in California may have been misidentified, resulting in a close association of *P. terebinthus* with *P. vera*, *P. khinjuk*, and *P. atlantica* by Parfitt and Badenes (1997). We

suggest that these trees are not true *P. terebinthus*, but a further study comparing Californian *P. terebinthus* samples with our Mediterranean sample is required to test this hypothesis.

In our study, *P. terebinthus* and *P. palaestina* formed a close pair, and they have sometimes been regarded as a single species. Engler who first classified *Pistacia* species, indeed considered *P. palaestina* as a variety of *P. terebinthus* (reviewed in Zohary, 1952). Zohary, however, considered *P. palaestina* as a distinct species, due to two main distinctive characteristics. *P. palaestina* has mostly paripinnate leaves and acuminate leaflets, while *P. terebinthus* has imparipinnate leaves and obtuse or acute leaflets. Yaltirik (1967a) described two subspecies within *P. tere-*

binthus. The first, *P. terebinthus* subspecies *terebinthus*, had imparipinnate leaves with the terminal leaflet of the median leaves often as large as the lateral ones, and obtuse or ovate-oblong lateral leaflets. The second is *P. terebinthus* subsp. *palaestina* with either paripinnate and/or imparipinnate leaves, the terminal leaflet of the median leaves always smaller than the laterals or reduced to a bristle, and acuminate or oblong-lanceolate lateral leaflets. Our results support Engler's and Yaltirik's classifications. To confirm this hypothesis, we analyzed two additional *P. terebinthus* samples (from Southern and Western Turkey, respectively), and two additional *P. palaestina* samples (from Israel) using the same primers, and similar RAPD patterns were obtained between *P. palaestina* and *P. terebinthus* samples (data not shown). In our previous study (Kafkas and Perl-Treves, 2001), we also obtained similar pair-wise distances between accessions within a *Pistacia* species. All these suggest that the above leaf characters probably may not be sufficient to assign *P. palaestina* as a different species. Therefore, we find it more appropriate to consider *P. palaestina* as a variety of *P. terebinthus*.

Zohary (1952) classified *P. eurycarpa* as a variety of *P. atlantica* (var. *kurdica*) because of the presence of leaf rachis wing that narrower than in the type of *P. atlantica*. Yaltirik (1967b), on the other hand, treated this plant as a different species, because the leaves are light green on both sides (instead of being dark green above and pale below as in *P. atlantica*), and the nuts are depressed and bigger than in typical *P. atlantica*. Furthermore, the leaflets are usually thicker and never as numerous or narrow as in *P. atlantica*, and the leaf rachis wing is narrower or even absent. Kafkas and Perl-Treves (2001) retained *P. eurycarpa* as a different species by cluster analysis using both morphological and molecular data. This study also suggests that *P. eurycarpa* is closely related to *P. atlantica* but represents a distinct species, since the genetic distance between them is similar to the distance between other pairs of species.

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