

Morphological and biochemical diversity among wild-grown carob trees (*Ceratonia siliqua* L.)

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ABSTRACT

Tree, leaf, pod and seed morphology, as well as pod biochemistry of 36 wild-grown carob genotypes sampled from rural areas in Marmaris district located at western Turkey, were investigated. Leaf and pod dimensions, pod and seed weight, seed ratio, pod and seed colour and shape and surface traits were investigated. Soluble solid content (SSC), titratable acidity, vitamin C and protein and dietary fibre contents were also detected. Results showed significant differences for all quantitative traits, although differences are more pronounced for some pod (weight, width, length and thickness) and seed characteristics (weight, dimensions and ratio). Pod and seed colour, shape and surface qualitative traits were found to be quite variable among genotypes. The majority of genotypes had an open tree growth habit. Leaf length and width were found to be between 8.04 cm (M19) and 11.60 cm (M12) and 8.40 cm (M2) and 12.04 cm (M12) among genotypes. Pod weight ranged from 8.3 g (M35) to 29.5 g (M3) in the wild genotypes. The average pod dimensions (width, length and thickness) were between 14.27 and 23.38 mm, 12.54 and 21.67 cm and 4.80 and 8.37 mm, respectively. The SSC ranged from 49.36 to 69.36% in the pods of wild carob genotypes. The results of this study indicate a good genetic resource potential of Turkish wild carob populations for future breeding programmes.

Keywords: crop wild relatives, genetic diversity, pod dimensions, soluble solid content, vitamin C

INTRODUCTION

Plant genetic resources include cultivars, landraces, unnamed genotypes, accessions, wild species closely related to cultivated varieties, breeder's elite lines and mutants. The loss of genetic diversity caused by the practice of agriculture and the availability of genetic information has resulted in a great effort dedicated to the study and collection of plant genetic resources (Kafkas et al., 2008; Ercisli et al., 2011; Van et al., 2011; Zia-Ul-Haq et al., 2014; Eyduzan et al., 2015; Akin et al., 2016).

In the last 30 years, the importance of plant genetic resources has become more evident as a result of growing concerns about biodiversity, conservation and genetic erosion. Global food production and food

security are still a major challenge for the future of mankind. Therefore, securing plant genetic resources for future generations has become a priority not only in developing countries but also throughout the world (Sahin et al., 2002; Altindag et al., 2006; Gepts, 2006; Ercisli et al., 2008; Kafkas et al., 2018).

The characterisation of fruit genetic resources by analysing plant morphology is the simplest and easiest approach (Ercisli et al., 2012; Butiuc-Keul et al., 2019; Fazenda et al., 2019). Because morphological markers, such as growth habit, fruit and seed colour, fruit and seed weight, fruit and seed shape, harvest dates, taste, plant height, disease response, and so on, are scored visually.

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All those traits are generally scored quickly, simply and without laboratory equipment (Bhat et al., 2010).

Three phytogeographical regions, such as Euro-Siberian, Mediterranean and Irano-Turanian, overlap in Turkey. Euro-Siberian region stretches along most of North Anatolia and Europe. Historically, Turkey has been a pathway for many civilisations and hosted many of them. The movement of communities has contributed to the enrichment of genetic diversity by transferring mainly the cultivated species and the seeds of wild plants from one place to another (Ercisli, 2004; Halasz et al., 2010). The topography of Turkey exhibits significant variety where ecological factors change frequently over a short distance. Asian section is a large, roughly rectangular peninsula situated like a bridge between Europe and Asia.

Carob tree (*Ceratonia siliqua* L.) is one of the oldest trees in the world and has been grown since ancient times in most Mediterranean basin countries, and it has an important value from an economic and environmental point of view. The species is a flowering evergreen shrub and it is popular for its sweet edible pods. The carob tree can be found growing extensively in the wild of the Eastern Mediterranean region. Its fruits are legumes commonly referred to as pods that are elongated, compressed or curved (Batlle and Tous, 1997). Carob pods have active constituents, which include calcium, phosphorus, potassium, sodium and magnesium. The carob fruit is rich in proteins, carbohydrates and fibres. The combination of these compounds provides relatively high nutritional values leading to the claimed health benefits of carob (Turhan et al., 2006; Tous et al., 2009; Gezer, 2018).

The main carob producer in the world is Spain followed by Italy, Portugal, Greece, Morocco and Turkey. Turkey shares approximately 10% of the carob production in the world (FAO, 2018). Turkey has a rich natural carob population obtained from seeds, and trees

thrive together with several other species of the maquis in the Mediterranean and Aegean regions. Carob has been a neglected species and accepted forest tree in Turkey. The country does not have commercial carob orchards, although some new ones have recently been established. Wild carob trees show great diversity and concentrate along the Mediterranean and Aegean regions (Figure 1) (Pazir and Alper, 2018; Durmaz and Ozel, 2019).

In Turkey, particularly in the Aegean region, carob production is still carried out using seed-propagated genotypes. Thus, in the market, carob fruits are composed of very mixed trees. Therefore, carob industry and consumers experience certain difficulties in obtaining products of standard quality. It is necessary to conduct further breeding studies using different seed-propagated carob trees and select them to increase yield and quality. The success of a breeding programme depends mainly on sound information about the breeding material.

To our knowledge, no studies are reporting morphological and biochemical analysis of seed-propagated carobs grown in the Aegean region of Turkey. Therefore, this study aimed to characterise and evaluate the diversity of 36 seed-propagated carob genotypes and develop strategies for preserving the endangered genetic resources of this species.

MATERIALS AND METHODS

A total of 36 seed-propagated wild-grown carob genotypes were used, and mature pods were collected from rural areas of Marmaris district between 2016 and 2017 (Figure 1). A total of 30 pods from each genotype were randomly chosen to measure the different parameters. The morphological parameters determined were tree growth habit, leaf width, leaf length, number of leaflets/leave, pod colour, pod shape, pod surface, pod weight, pod width, pod length, pod thickness, seed colour, seed shape, seed surface, seed weight and seed

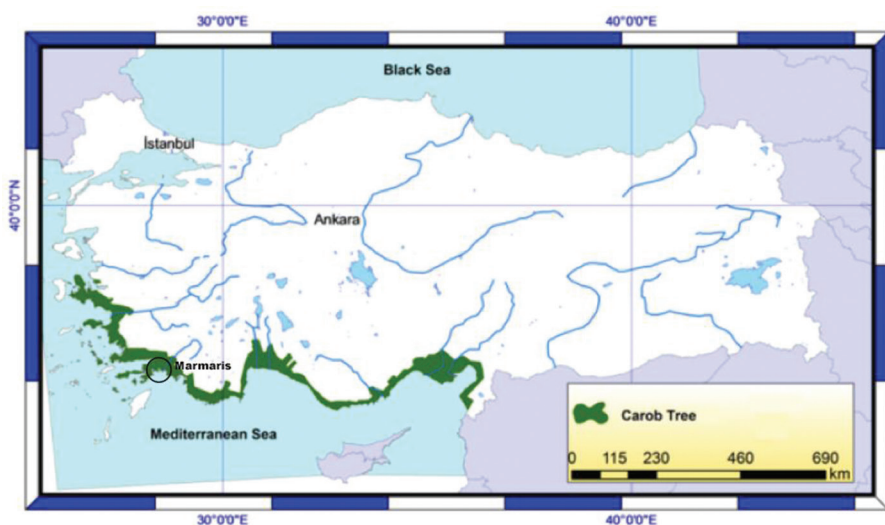


Figure 1. Natural distribution of carob trees in Turkey and our sampling location (Marmaris district) (Sahin and Tasligil, 2016).

ratio. Quantitative measurements were done using a calliper with a precision of ± 0.1 mm and an electronic balance (SCALTEC SBA33, Heiligenstadt, Germany). The dimensions of pod, seed and leaf were measured using a digital calliper. Biochemical parameters were soluble solid content (SSC), titratable acidity, vitamin C, protein and dietary fibre contents. Seed ratio was calculated using whole pod and seed weight. For chemical analysis, pulps were crushed with a blender (Philips Hr2653/90 Viva Collection Promix, The Netherlands) and then passed through a 35-mesh sieve. About 10 g of pulp was introduced with 40 ml water in a baker and homogenised in a homogeniser (IKA, Deutschland) for 5 min. SSC of samples was measured at 25 ± 0.5 °C using a refractometer (KEM, Kyoto, Japan). Titratable acidity was determined by titration with 0.1 N NaOH and calculated as % citric acid (anhydrous) (AOAC, 2007). Vitamin C was determined by RQFlex (Merc Co, Darmstadt, Germany). Total nitrogen of carob powder was determined according to the AOAC official method and was converted to protein content using the conversion factor 6.25 (AOAC, 2007) with Macro Kjeldahl digestion and distillation apparatus. For dietary fibre analysis, 4 g of carob powder were digested with 200 ml of 5% HCL for 30 min. The mixture was filtered and washed with hot water. Then, the residue was digested with 200 ml of 5% NaOH under reflux for 30 min. The mixture was filtered and washed with distilled water until pH neutrality. The material was washed with 20 ml ethyl alcohol and 20 ml ethyl ether. Finally, the residue was dried at 100 °C for 2 h, and the residual mass was considered as fibres (De Padua et al., 2004). Protein and fibre contents were expressed as %.

Statistical analysis

No differences were found between years thus, the data of both years were pooled. All data were analysed using SPSS software and procedures. Tables of analysis of variance were constructed using the least significant difference (LSD) method at $p < 0.05$. The principal coordinate analysis (PCoA) was performed to show the relationships and differentiation of the carob genotypes in a three-dimensional array of eigenvectors using the DCENTER and EIGEN modules of NTSYS-pc 2.10e software.

RESULTS AND DISCUSSION

Morphological analysis of carob leaf, pod and seed quantitative traits revealed significant variations among genotypes. Qualitative traits (colour, shape and surface aspect of pods and seeds) also showed variation among genotypes (Tables 1–3).

The genotypes exhibited three tree growth habits, such as open, weeping and erect, and among them open growth habit was dominant followed by weeping (Table 1). Batlle and Tous (1997) reported that open growth habit was dominant among carob cultivars from

Spain, Italy, Portugal, Greece, Tunisia and the USA followed by weeping growth habit.

The genotypes differed from each other statistically ($p < 0.05$) in terms of leaf width and length. However, the number of leaflets/leave was non-significant (Table 1). Leaf width and length were ranged from 8.40 cm (M2) to 12.04 cm (M12) and from 8.04 cm (M19) to 11.60 cm (M12), respectively (Table 1). The number of leaflets/leave was insignificant among carob genotypes and was found between 5.9 and 7.1. In the literature, there was a limited report about leaf dimensions of carob because most of the morphological studies are concentrated on pod and seed characteristics. Ahmed et al. (2019) found leaf width and leaf length of carobs in Morocco between 9.53 and 10.77 cm and 10.19 and 11.45 cm, respectively. They also reported the number of leaflets per leave between 6.5 and 7.4 indicating similarities with our study.

Pod characteristics (colour, shape, surface, weight, width, length and thickness) are provided in Table 2.

The majority of genotypes had dark-brown pod colour (47.22%) followed by reddish brown (27.78%) and clear brown (25.00%). Dark-brown, clear-brown and reddish-brown pod colours were previously reported in carob cultivars and genotypes (Batlle and Tous, 1997; Ait Chitt et al., 2007). Curved pod shape was dominant (61.11%) among genotypes followed by equally spiral and straight shape (19.44%). Batlle and Tous (1997) reported a curved, spiral and straight pod shape among carob cultivars and genotypes, which is in accordance with our study. In terms of pod surface, 19 genotypes had a wrinkled surface (52.78%) followed by 15 genotypes with a smooth surface (41.67%) and 2 genotypes with a rough surface (5.56%) (Table 2). Batlle and Tous (1997) reported that the pod surface varied from smooth to wrinkled in carobs.

Pod weight and pod dimensions (width, length, length and thickness) are shown in Table 2, and we found statistically significant differences among genotypes ($p < 0.05$) for pod weight, pod width, pod length and pod thickness.

Pod weight of 36 carob genotypes was quite variable and ranged from 8.3 g (M35) to 29.5 g (M3) among genotypes. The average pod dimensions (width, length and thickness) were found between 14.27 and 23.38 mm, 12.54 and 21.67 cm and 4.80–8.37 mm for the wild genotypes (Table 2). La Malfa et al. (2012) investigated morphological and technological characteristics of eight carob cultivars grown in Sicily, Italy, and reported average pod weight, pod width, pod length and pod thickness as 13.7–33.4 g, 19.3–26.8 mm, 14.9–22.9 cm and 6.8–14.0 mm, respectively. In Algeria, a study examined wild-grown carobs in 10 regions and reported average pod weight, pod width, pod length and pod thickness as 7.04–30.57 g, 18.10–31.80 mm, 10.30–18.75 cm and 4.50–8.20 mm, respectively (Boublenza et al., 2019). Barracosa et al. (2007) stated pod weight, pod width, pod length and pod thickness of 15 carob cultivars from Portugal as 13.20–26.39 g;

Table 1. Tree and leaf characteristics of carob genotypes

Genotypes	Tree growth habit	Leaf width (cm)	Leaf length (cm)	Number of leaflets/leave
M1	Open	11.10 ab	9.60 ab	6.0 ^{NS}
M2	Weeping	8.40 e	8.80 ab	6.2
M3	Open	11.50 ab	10.78 ab	5.9
M4	Open	11.30 ab	11.10 ab	6.3
M5	Erect	9.85 cd	9.40 ab	7.0
M6	Open	9.04 de	8.28 ab	7.1
M7	Open	11.70 ab	11.50 ab	6.4
M8	Open	8.90 de	8.82 ab	6.3
M9	Weeping	10.42 bc	10.30 ab	5.9
M10	Open	9.20 d	9.10 ab	6.2
M11	Open	9.28 bc	9.07 ab	6.4
M12	Open	12.04 a	11.60 a	6.5
M13	Erect	9.33 cd	9.00 ab	7.1
M14	Weeping	11.10 b	10.74 ab	7.0
M15	Open	10.80 bc	10.50 ab	6.8
M16	Open	11.74 ab	10.95 ab	6.8
M17	Open	11.38 ab	11.05 ab	5.9
M18	Weeping	11.90 ab	11.40 ab	6.2
M19	Open	9.00 de	8.04 b	6.1
M20	Open	10.25 bc	9.64 ab	5.9
M21	Open	9.60 cd	9.30 ab	7.0
M22	Weeping	9.40 cd	9.20 ab	7.0
M23	Open	9.18 d	8.80 ab	6.5
M24	Open	10.02 cd	9.15 ab	6.8
M25	Open	9.56 cd	9.28 ab	6.8
M26	Erect	9.11 de	9.02 ab	6.9
M27	Weeping	10.17 c	9.78 ab	6.3
M28	Open	9.59 cd	9.19 ab	6.0
M29	Open	10.86 bc	10.11 ab	5.9
M30	Open	9.45 cd	8.29 ab	6.7
M31	Open	11.41 ab	10.15 ab	6.9
M32	Weeping	9.50 cd	8.90 ab	5.9
M33	Erect	9.33 cd	9.02 ab	6.4
M34	Open	10.35 bc	10.07 ab	6.1
M35	Open	9.20 d	9.02 ab	6.3
M36	Open	10.60 bc	10.10 ab	6.0

Means with different letters in the same column differ significantly ($p < 0.05$).

NS, Non-significant.

16.16–23.38 mm; 12.95–20.35 cm and 7.19–10.86 mm, respectively. Albanell et al. (1991) reported the highest pod weight (14.88 g), pod length (15.83 cm) and pod width (21.10 mm) among 182 common carob trees from various areas in Spain. Haddarah et al. (2013) reported pod weight (8.93–36.85 g), pod length (11.42–24.25 cm), pod width (17.30–27.40 mm) and pod thickness (4.8–9.2 mm) among wild carob genotypes in Lebanon. Russo and Polignano (1996) found mean pod length and width as 17.1 cm and 22.7 mm, respectively. Gharnit et al. (2006) reported average pod length, width and thickness as 13.5 cm, 19.5 mm and 6.9 mm in Morocco. Our results are in good agreement with the aforementioned studies. The main selection criteria in carobs have traditionally focussed on large pod size, high pulp and sugar content. Carob pod size is also important to withstand strong

winds during spring to prevent premature fruit drop (Batlle and Tous, 1997).

Seed colour, seed shape, seed surface, total seed weight and seed ratio are shown in Table 3. Most of the genotypes had clear-brown pod colour (44.4%) followed by dark brown (33.3%), reddish brown (11.1%) and blackish brown (11.1%). Rounded pod shape was dominant (55.5%) among genotypes followed by elliptical (30.5%) and oval seed shape (13.8%). In terms of seed surface, 55.5% of the genotypes had smooth seed surface, 36.1% of the genotypes had wrinkled surface, 5.55% of the genotypes had very rough and 2.77% of the genotypes had rough seed surface (Table 3). Albanell et al. (1991) and Batlle and Tous (1997) reported that the majority of carob seeds had a smooth seed surface.

Table 2. Pod characteristics of carob genotypes

Genotypes	Pod colour	Pod shape	Pod surface	Pod weight (g)	Pod width (mm)	Pod length (cm)	Pod thickness (mm)
M1	Dark brown	Curved	Wrinkled	28.2 ab	22.33 ab	20.41 ab	8.07 ab
M2	Dark brown	Spiral	Wrinkled	12.2 cd	14.27 d	12.54 c	4.82 bc
M3	Reddish brown	Curved	Smooth	29.5 a	21.67 ab	18.94 ab	8.11 ab
M4	Clear brown	Curved	Wrinkled	25.6 ab	22.02 ab	17.68 b	7.97 ab
M5	Reddish brown	Straight	Smooth	15.4 cd	17.44 c	16.38 bc	5.18 bc
M6	Dark brown	Curved	Wrinkled	16.1 cd	17.66 bc	15.91 bc	5.63 bc
M7	Clear brown	Curved	Smooth	25.5 ab	19.33 bc	18.27 ab	7.08 ab
M8	Clear brown	Curved	Smooth	17.7 bc	16.94 cd	14.03 bc	4.88 bc
M9	Dark brown	Spiral	Rough	23.4 b	21.14 ab	13.98 bc	5.94 bc
M10	Dark brown	Curved	Wrinkled	16.6 cd	17.76 bc	16.47 bc	5.56 bc
M11	Clear brown	Straight	Wrinkled	15.2 cd	19.56 bc	16.87 nc	6.36 bc
M12	Dark brown	Curved	Wrinkled	26.3 ab	23.08 a	19.83 ab	8.37 a
M13	Reddish brown	Curved	Smooth	16.2 cd	20.50 ab	19.33 ab	6.02 bc
M14	Dark brown	Straight	Wrinkled	25.6 ab	18.15 bc	19.98 ab	7.11 ab
M15	Reddish brown	Straight	Smooth	22.1 bc	19.10 bc	18.55 ab	6.20 bc
M16	Clear brown	Curved	Smooth	25.1 ab	22.48 ab	20.02 ab	8.02 ab
M17	Dark brown	Curved	Rough	28.4 ab	20.83 ab	21.67 a	7.92 ab
M18	Dark brown	Spiral	Wrinkled	25.3 ab	20.21 b	20.90 ab	7.62 ab
M19	Reddish brown	Curved	Wrinkled	15.6 cd	16.56 cd	17.10 bc	5.35 bc
M20	Reddish brown	Spiral	Smooth	21.3 bc	19.61 bc	19.03 ab	6.40 bc
M21	Dark brown	Curved	Wrinkled	19.7 bc	19.23 bc	17.90 ab	5.95 bc
M22	Clear brown	Spiral	Smooth	17.7 bc	18.78 bc	19.04 ab	6.03 bc
M23	Dark brown	Curved	Wrinkled	10.9 d	18.52 bc	14.21 bc	4.77 bc
M24	Dark brown	Curved	Smooth	22.3 bc	20.07 bc	19.43 ab	5.81 bc
M25	Dark brown	Straight	Smooth	17.4 c	18.20 bc	16.84 bc	5.70 bc
M26	Reddish brown	Straight	Wrinkled	19.4 bc	20.22 b	14.98 bc	6.35 ab
M27	Clear brown	Curved	Wrinkled	21.6 bc	18.41 bc	18.95 ab	6.82 ab
M28	Dark brown	Straight	Wrinkled	12.6 cd	18.10 bc	12.88 c	5.91 bc
M29	Reddish brown	Curved	Smooth	11.4 bc	19.07 bc	18.23 ab	7.30 ab
M30	Clear brown	Curved	Smooth	9.2 de	15.44 cd	16.67 bc	6.58 b
M31	Dark brown	Curved	Wrinkled	25.5 ab	20.18 b	19.19 ab	6.24 bc
M32	Reddish brown	Curved	Smooth	15.0 cd	17.60 bc	17.94 ab	5.59 bc
M33	Reddish brown	Spiral	Smooth	10.3 de	15.46 cd	14.73 bc	4.65 c
M34	Dark brown	Spiral	Wrinkled	21.9 bc	20.70 ab	20.19 ab	6.47 b
M35	Clear brown	Curved	Wrinkled	8.3 e	16.11 cd	14.98 bc	4.91 bc
M36	Dark brown	Curved	Wrinkled	22.6 bc	18.64 bc	19.23 ab	7.71 ab

Means with different letters in the same column differ significantly ($p < 0.05$).

There were statistically significant differences ($p < 0.05$) among genotypes in terms of total seed weight and seed ratio (%) (Table 3). The highest total seed weight was observed in M1 genotype as 3.41 g, followed by M3 as 3.20 g and M4 as 3.07 g, while the lowest total seed weight was obtained from M30 genotype as 1.10 g (Table 3). The seed ratio was found between 6.29% (M26) and 13.98% (M35) (Table 3). La Malfa et al. (2012) reported total seed weight and seed ratio between 1.57 and 2.34 g and 4.9 and 16.9%, respectively. Barracosa et al. (2007) stated the total seed weight among 15 carob cultivars in Portugal as 1.83–2.99 g. Boublenza et al. (2019) investigated carobs from Algeria and determined the total seed weight and seed ratio as 0.98–2.30 g and 7.35–14.58%, respectively. Albanell et al. (1991) reported seed ratio (12.11%) among 182 common carob trees from various areas in Spain.

Haddarah et al. (2013) reported total seed weight (1.36–3.10 g) and seed ratio (8.41–20.10%) among wild carob genotypes in Lebanon. Gharnit et al. (2006) reported average total seed weight and seed ratio between 1.88 and 28 g and 16.6 and 22.0% in Morocco. These studies indicate similarities with our results.

The biochemical characteristics of carob genotypes fruits are shown in Table 4. SSC, titratable acidity, protein and dietary fibre contents differed significantly among genotypes ($p < 0.05$). However, no differences were found among genotypes in terms of vitamin C (Table 4).

The highest SSC content was seen in genotype M12 as 69.36% followed by M3 (68.41%) and M7 (68.23%), while the lowest SSC content was obtained from M23 genotype as 49.36% (Table 4). Titratable acidity and vitamin C were less variable than SSC and ranged from

Table 3. Seed characteristics of carob genotypes

Genotypes	Seed colour	Seed shape	Seed surface	Total seed weight (g)	Seed ratio (%)
M1	Dark brown	Rounded	Wrinkled	3.41 a	12.09 ab
M2	Dark brown	Elliptical	Smooth	1.24f g	10.16 ab
M3	Clear brown	Rounded	Smooth	3.20 ab	10.85 ab
M4	Clear brown	Rounded	Wrinkled	3.07 ab	11.99 ab
M5	Reddish brown	Oval	Smooth	1.85 de	12.01 ab
M6	Dark brown	Rounded	Wrinkled	1.72 e	10.68 ab
M7	Clear brown	Rounded	Smooth	3.01 b	11.80 ab
M8	Clear brown	Rounded	Smooth	1.60 ef	9.04 b
M9	Dark brown	Elliptical	Rough	2.30 cd	9.83 ab
M10	Dark brown	Rounded	Wrinkled	1.57 ef	9.46 ab
M11	Clear brown	Elliptical	Smooth	1.62 ef	10.66 ab
M12	Dark brown	Rounded	Very rough	2.90 bc	11.03 ab
M13	Clear brown	Oval	Smooth	1.48 ef	9.14 ab
M14	Dark brown	Elliptical	Wrinkled	2.87 bc	11.21 ab
M15	Reddish brown	Elliptical	Smooth	2.20 cd	9.95 ab
M16	Clear brown	Rounded	Smooth	2.69 bc	10.72 ab
M17	Dark brown	Rounded	Very rough	2.55 c	8.98 bc
M18	Dark brown	Elliptical	Wrinkled	2.45 cd	9.68 ab
M19	Reddish brown	Oval	Wrinkled	1.35 f	8.65 bc
M20	Clear brown	Elliptical	Smooth	2.36 cd	11.08 ab
M21	Blackish brown	Rounded	Wrinkled	1.40 ef	7.11 bc
M22	Clear brown	Oval	Smooth	1.58 ef	8.93 bc
M23	Dark brown	Rounded	Wrinkled	1.19 fg	10.91 ab
M24	Blackish brown	Rounded	Smooth	1.98 de	8.88 bc
M25	Dark brown	Elliptical	Smooth	1.48 ef	8.51 bc
M26	Reddish brown	Elliptical	Wrinkled	1.22 fg	6.29 c
M27	Clear brown	Rounded	Wrinkled	2.15 d	9.95 ab
M28	Dark brown	Oval	Wrinkled	1.16 fg	9.21 ab
M29	Clear brown	Rounded	Smooth	2.26 cd	10.56 ab
M30	Clear brown	Rounded	Smooth	1.10 g	11.96 ab
M31	Blackish brown	Rounded	Wrinkled	2.61 bc	10.24 ab
M32	Clear brown	Rounded	Smooth	1.28 fg	8.53 bc
M33	Clear brown	Elliptical	Smooth	1.20 fg	11.6 5 ab
M34	Clear brown	Elliptical	Smooth	2.60 bc	11.87 ab
M35	Clear brown	Rounded	Smooth	1.16 fg	13.98 a
M36	Blackish brown	Rounded	Smooth	2.05 de	9.07 b

Means with different letters in the same column differ significantly ($p < 0.05$).

0.45% (M13) to 1.12% (M9) and from $5.9 \text{ mg} \cdot 100 \text{ g}^{-1}$ (M19 and M22) to $10.2 \text{ mg} \cdot 100 \text{ g}^{-1}$ (M12), respectively (Table 4). SSC values of carob genotypes have been reported to be between 32 and 70% (Marakis et al., 1988).

Protein and dietary fibre contents were found between 3.73% (M6) and 6.95% (M12) and 4.9% (M11 and M17) and 7.7% (M3), respectively (Table 4). Haddarah et al. (2013) reported protein content between 3.61 and 4.82% and dietary fibre content between 4.80 and 7.74% among wild carob genotypes in Lebanon. Ozcan et al. (2007) found that carob pods had 4.71% protein content and 9.69% crude fibre content.

The study showed variation among wild carob populations for leaf, pod and seed morphology. In the perspective of identifying the best genotypes for breeding programmes and/or accession collection, additional descriptors should be examined such as regular bearing

(Keles and Bilir, 2015); gender and harvest period or fruit ripening (Tous et al., 2009), pedicel length as a criterion of pod abscission (Tous et al., 2009); trunk cross-section and canopy volume (Tous et al., 2009), fruit sweetness, flowering and fruiting phenology and precocity (Haddarah et al., 2013).

PCoA was applied to the data using NTSYS 2.10e software, and the contribution rates of the first three principal coordinates were 47.1, 22.4 and 9.2%, respectively, accounting for 78.7% of the variance. The first principal coordinate (PCoA1), which explains 47.1% of the overall variance, is identified with the pod weight, pulp weight and seed weight, while the second principal coordinate (PCoA2) is related to the leaf and leaflet dimensions, pod length and pulp and seed ratio (Table 5). Carob genotypes samples were partitioned into four distinct groups. PCoA Groups 1, 2, 3 and 4 included 10, 8, 10 and 8 samples, respectively

Table 4. Biochemical characteristics of carob pods

Genotypes	SSC (%)	Acidity (%)	Vitamin C (mg · 100 g ⁻¹)	Protein (%)	Dietary fibres (%)
M1	67.21 ab	0.76 bc	10.1 ^{NS}	3.88 bc	7.0 b
M2	53.12 d	0.88 ab	9.3	6.50 ab	6.0 cd
M3	68.41 ab	0.60 bc	8.3	4.77 bc	7.7 a
M4	65.45 ab	0.80 bc	9.4	4.59 bc	5.8 cd
M5	59.56 c	0.68 bc	7.6	5.25 bc	5.5 de
M6	57.23 cd	0.55 bc	9.8	3.73 c	5.0 de
M7	68.23 ab	0.92 ab	7.1	5.40 ab	6.8 bc
M8	54.14 cd	0.85 b	7.5	5.04 bc	6.3 c
M9	64.11 b	1.12 a	7.6	4.94 bc	6.1 cd
M10	58.28 cd	1.02 ab	8.0	3.96 bc	5.0 de
M11	57.45 cd	1.05 ab	6.6	4.11 bc	4.9 e
M12	69.36 a	0.55 bc	10.2	6.95 a	5.3 de
M13	59.98 bc	0.45 c	9.4	4.20 bc	6.6 bc
M14	67.56 ab	0.78 bc	8.7	5.35 b	5.8 cd
M15	63.44 bc	0.83 b	7.6	4.30 bc	5.3 de
M16	65.45 ab	0.90 ab	7.5	4.22 bc	5.6 d
M17	68.22 ab	0.66 bc	8.0	6.02 ab	4.9 e
M18	66.44 ab	0.59 bc	8.4	6.18 ab	5.0 de
M19	51.56 de	0.62 bc	5.9	5.44 ab	5.3 de
M20	63.30 bc	0.75 bc	6.2	5.50 ab	7.1 ab
M21	58.12 cd	0.84 b	5.8	4.33 bc	5.4 de
M22	57.64 cd	0.90 ab	5.9	5.10 bc	5.7 cd
M23	49.36 e	1.00 ab	7.1	4.60 bc	7.0 b
M24	62.55 bc	1.05 ab	7.7	4.11 bc	5.9 cd
M25	56.35 cd	0.96 ab	8.5	4.70 bc	5.6 d
M26	55.83 cd	0.92 ab	9.2	4.15 bc	6.4 bc
M27	63.13 bc	0.71 bc	7.4	6.44 ab	7.0 b
M28	52.35 de	0.77 bc	8.1	5.33 b	6.2 cd
M29	61.21 bc	0.84 b	7.7	5.30 b	5.0 de
M30	50.45 de	0.55 bc	6.8	5.04 bc	5.5 de
M31	67.40 ab	0.80 bc	7.3	4.27 bc	7.4 ab
M32	51.80 de	0.93 ab	6.4	4.41 bc	6.4 bc
M33	49.74 de	0.76 bc	7.7	4.50 bc	6.0 cd
M34	60.80 bc	0.64 bc	8.2	4.90 bc	5.8 cd
M35	49.56 de	0.68 bc	8.0	3.95 bc	5.4 de
M36	62.26 bc	0.91 ab	7.5	4.44 bc	5.0 de

Means with different letters in the same column differ significantly ($p < 0.05$).

NS, Non-significant.

(Figure 2). The first group characterised by high pod and seed weight, larger and longer leaves, high seed size and high SSC content. The second and third groups had medium-sized leaves, pods and seeds and medium sugar content. The last group had lower pod and seed weight, low leaf dimensions and low sugar content. PCoA revealed useful information on the characterisation and comparison of carob germplasm collections in terms of morphological and biochemical data. At the same time, PCoA indicated some common features of wild carob genotypes, thanks to which it is possible to divide the analysed wild carobs into bigger fruited ones, more sweet ones, or those with a higher content of pulp and seed ratio. The substantial dispersion of Turkish wild carob genotypes in the PCoA plot suggests a high level of diversity, which can make them attractive for future

breeding programmes and long-term conservation strategies.

CONCLUSION

Our findings showed that morphological characteristics and biochemical composition were considerably influenced by the genotype factor. Even though such variation did not seem to be geographically structured because all genotypes are from similar ecological and soil conditions, and the analysed traits could be considered under an agronomic and an industrial perspective (seed yield and pulp weight). Such descriptors, together with additional ones, would contribute to the characterisation of carob genetic resources and guide the choice of populations for *in situ* conservation or be a source of

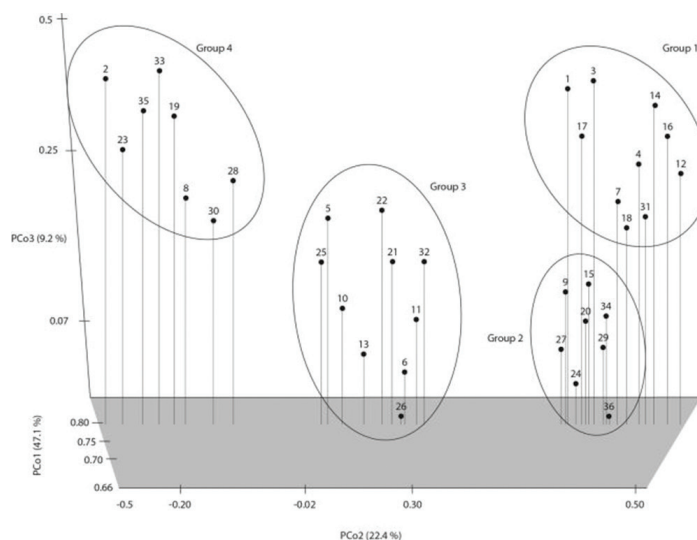


Figure 2. PCoA plot for the 36 analyzed wild carob genotypes (the genotype number codes on the plot refer to Tables 1–4).

Table 5. Factor loadings for each variable on the components of PCoA analysis

Variable	PC1	PC2	PC3
Leaf length	5.54	13.11	1.20
Leaf width	1.12	18.74	1.03
Number of leaflets	4.13	10.11	0.17
Length of leaflets	0.03	16.13	1.67
Pod weight	22.71	0.10	0.55
Pod width	9.22	0.50	9.15
Pod length	2.45	11.23	4.66
Pod thickness	7.11	0.08	7.57
Pulp weight	22.13	0.80	6.21
Seed weight	10.06	0.57	14.13
Seed number	7.06	0.09	25.11
Pulp weight	3.41	2.23	8.62
Seed ratio	1.45	12.09	15.37
Pulp ratio	2.23	10.53	4.22
pH	0.34	1.03	0.07
SSC	0.49	0.90	0.05
Acidity	0.13	0.56	0.13
Vitamin C	0.04	0.08	0.00
Protein	0.23	0.09	0.05
Dietary fibre	0.12	0.03	0.04
Eigenvalue	5.43	2.98	1.67
% of variance	47.1	22.4	9.2
Cumulative variance	47.1	69.5	74.7

material for restoration, breeding programmes and germplasm collections.

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AUTHOR CONTRIBUTIONS

N.K. and S.E. designed the study and performed the field work. M.A. and S.P.E. was involved in the statistical analysis. A.K., G.I. and H.I.S performed the laboratory study. All authors contributed to data analysis, discussion of results and the writing and editing of the manuscript.

CONFLICT OF INTEREST

The authors declare that there are no potential conflicts of interest regarding the research, authorship and publication of this manuscript.

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