

BIOCHEMICAL AND BIOACTIVE CONTENT IN FRUITS OF WALNUT (*JUGLANS REGIA* L.) GENOTYPES FROM TURKEY

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ABSTRACT

Walnuts (*Juglans regia* L.) are a good source of dietary minerals and contain significant amounts of potassium, phosphorus, magnesium and iron. In Turkey, walnut orchards are found in the Çoruh Valley and are generally composed of local varieties. Since chemical and physical properties of walnuts are very important in determining their quality, in this study, nut and kernel characteristics, fatty acid composition and phenolic composition were determined for 26 walnut genotypes collected from the valley. Total fat content was found to be between 53.75% and 71.43%, while the crude protein amount ranged from 10.21% to 20.71%. Total carbohydrates were calculated between 14.31-27.52% and ash content was found to be between 1.64% and 3.32%. The oleic acid content and linoleic acid content ranged from 18.34 to 25.58% and 37.09% to 87.51%, respectively. The linolenic acid content was determined between 5.52% and 11.03%. Our study showed that, differences in phenolic component, crude protein and fatty acid content of walnuts were caused by different growing conditions. Genotypes have a rich and stable structure in terms of fruit quality parameters, minerals, organic acids, phenolic composition, antioxidant and fatty acids amounts and amino acids. In other words, these characteristics did not show much variation in genotypes grown in the Valley. However, there was a great variation in hormone levels among genotypes. In terms of the parameters examined, GN2, GN10, GN15, GN24, GN25 and GN26 genotypes were determined as rich and stable genotypes.

KEYWORDS:

Amino acid, hormone, nutrient element, antioxidant and fatty acids, biplot analysis, walnut

INTRODUCTION

Walnut (*Juglans regia* L.) is the most economically important species and the oldest cultivated fruit in the world. Walnut is native to a wide region in USA, European and Asian countries [1]. Total world production of walnuts is 3 million 663 thousand tons. The production of China, the world's largest walnut producer, reached 1.6 million tons in 2018. China realized 43.3 percent of the total world walnut production in 2018. Iran follows China with 16.7 percent share. The third biggest producer is the United States of America with 620 thousand tons which corresponds to 11.2 percent of the global walnut production. Turkey ranks fourth with 5.9 percent share and 215 thousand tons of production [2].

Turkey is a germplasm center of walnut since Anatolia was on the historical migration routes. Many different genotypes were brought to Anatolia throughout the history via traveling traders along the Silk Road and walnut trees are exceptionally abundant within almost all regions in Anatolia. The local walnut populations are not scattered, rather grown in indoor gardens in the Çoruh Valley. Continuous seed propagation for thousands of years in Turkey has given rise to a great number of seedling walnut trees, which stands as a valuable walnut gene resource [3]. The seeds from very ancient times showed different characteristics and in time adapted to the ecological conditions of the region, which formed a wide genetic variation. In this genetic variation, it is extremely important to ensure that the types that are highly qualified are protected by reproduction. Owing to its geological, geomorphological and climatic characteristics, Çoruh Valley has a rich biological diversity. The Valley is located within the Caucasus Ecoregion, which is among the 200 ecological regions of the world designated by the World Wildlife Fund (WWF) [4,5]. Çoruh Basin (Figure 1) has a surface area of 19 748 km² and located in the northeast of Turkey (39°40' to 42°35' longitude and 39°52' to 41°32' latitudes).

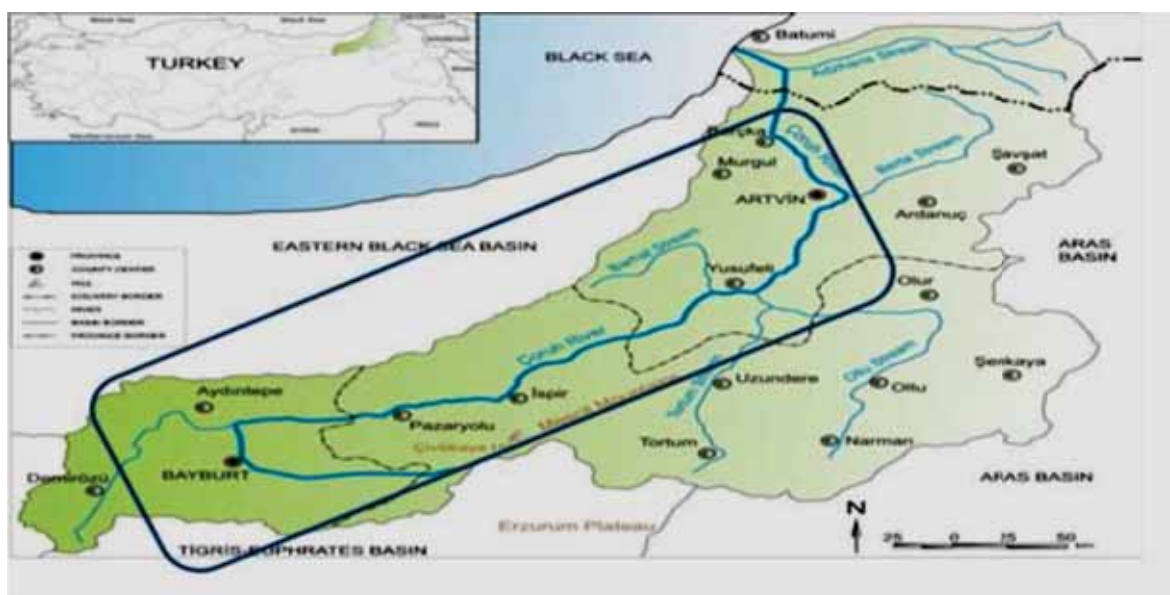


FIGURE 1
Çoruh Valley

Walnut trees have significant economic value due to the quality of the wood and nutritious nuts. The leaves and the nuts also have medicinal importance for their significant pharmacological value [6]. Walnuts are rich in vitamin E [7], ω -3 fatty acids [8] and dietary fibers [6], which provide many health benefits when consumed in sufficient amounts [9, 10, 11].

Walnut oil (up to 740 g kg⁻¹ in some commercial varieties) is largely composed of unsaturated fatty acids, particularly α -linolenic acid (ALA; 18:3n-3) and linoleic acid (LA; 18:2n-6), and can be consumed without refining [12]. Tocopherols are the phenolic compounds found in walnuts that possess vitamin E (antioxidant) activity [13]. Walnuts are also rich in phytosterols which are important dietary components that contribute to reducing serum cholesterol levels [12].

Chemical and mineral contents of walnuts can vary by variety, genotype, ecology, agricultural practices, climate, and soil conditions. Some genotypes may have a better effect on human metabolism due to their nutrient compositions. The aim of this study was to determine walnut fruit quality parameters such as weight, crude protein, phenolic, fatty acid, amino acid, organic acid and hormone contents to elucidate the physicochemical characteristics and antioxidant capacity of twenty-six walnut genotypes from Çoruh Valley, Turkey.

MATERIALS AND METHODS

Plant material. Walnut samples were collected from inside the native walnut population of Çoruh Valley, in Turkey. In the Valley, walnut has been widely cultivated since long years. Different districts

were visited. The walnut fruits were harvested randomly taken from the same trees, which were all between 20 and 40 years old, in September during 2019-2020. Walnut fruits were immediately transported to the laboratory and held in an oven 3 days at 30 °C, after harvest. Physical analyses were quickly determined. The fruits were stored in the shell, closed in plastic bags, and frozen to -20 °C, until the analyses. Chemical analyses were performed in three, physical analyses were performed in ten replications.

Analysis of fatty acids. Fatty acid composition of the air-dried seed walnut samples was determined using the modified fatty acid methyl ester (FAME) method [14]. The oil was extracted three times from 2 g samples by homogenizing with petroleum ether. For FAME, 1 ml of methylation reagent (80 ml of methanol + 0.5 g of sodium methoxide + 20 ml of isoktan) was added to 50 mg oil. The mixture was vortexed and left at room temperature for 24 hours; then 0.25 ml of isocene was added. Samples were centrifuged at 2400-x g for 5 minutes at 5 °C and the liquid was transferred to labeled Wheaton flasks and stored at 20 °C. FAME of the fatty acids (0.5 μ l) were analyzed with a gas chromatograph (Perkin Elmer Auto System XL, USA) equipped with a fused silica capillary column (MN FFAP (50 m x 0.32 mm i.d.; film thickness 0.25 μ m), flame ionizing detector (FID). Operated under the following conditions: oven temperature program, 120 °C for 1 min, then raised to 240 °C at a rate of 6 °C/min and kept at 240 °C for 15 min; injector and detector temperatures, 250 and 260 °C; respectively, carrier gas, helium at a flow rate of 14 psi; split ratio, 1/20 ml/min. The contents of palmitic (C16:0), stearic (C18:0), oleic (C18:1), linoleic (C18:2) and linolenic (C18:3) acids were determined with a computing Integrator.

Amino acid analysis. 0.1 N HCl was added to one gram of fresh sample, homogenized with ultraturraks, and incubated at 40 °C for 12 hours. Samples were then vortexed, centrifuged at 1200 rpm for 50 minutes, and the supernatants filtered through 0.22 µm filters (Millex Millipore). Then supernatants were transferred to glass vials and sent for HPLC analysis. Amino acids were extracted from the samples and were analyzed as described by Antoine et al. [15]. The amino acid derivatives were analyzed by HPLC on a Zorbax Eclipse-AAA 4.6 x150 mm, 3.5 µm columns (Agilent 1200 HPLC). The samples were analyzed by measuring the absorbance at 254 nm, and the amino acids were identified by comparison with standards. O-phthalaldehyde (OPA), fluorenylmethyl-chloroformate (FMOC) and 0.4 N Borate was used for derivation processes in an auto sampler. The following were used as the mobile phase in the chromatography system: mobile phase. A: 40 mM NaH₂PO₄ (pH 7.8) and mobile phase B: Acetonitrile/Methanol/Water (45/45/10, v/v/v) solutions. The flow rate of the mobile phase was 2 mL min⁻¹ and the column temperature was 40 °C. Aspartate, glutamate, asparagine, serine, glutamine, histidine, glycine, threonine, arginine, alanine, tyrosine, cysteine, valin, methionine, tryptophan, phenylalanine, isoleucine, leucine, lysine, hydroxyproline, sarcosine, proline quantities from PGPR samples were determined as pmol/µl after 26 minutes derivation process in HPLC.

Organic acid analysis. Deionized water (10 ml) was added to one gram of fresh, and homogenized with an ultra-turraks. The solution was centrifuged at 1200 rpm for 50 minutes, then the supernatants were filtered through 0.22 µm filters (Millex Millipore). The supernatants were transferred to glass vial and sent for HPLC analysis. The organic acids were analyzed by HPLC on Zorbax Eclipse-AAA 4.6 x 250 mm, 5 µm columns (Agilent 1200 HPLC) and absorbance of 220 nm in UV detector. Flow rate was 1 ml min⁻¹ and column temperature was 25 °C. Oxalic, propionic, tartaric, butyric, malonic, malic, lactic, citric, maleic, fumaric, and succinic acids were determined using 25 mM potassium phosphate (pH 2.5) as the mobile phase.

Hormone analysis. Extraction and purification processes were performed according to Battal and Tileklioglu [16]. First, methanole (%80) was cooled to -40 °C and added to fresh samples [17]. After the mixture was homogenized with ultra-turraks for 10 minutes and then incubated for 24 hours in the dark. The samples were filtered with (Whatman No:1) and then supernatants were filtered although 0.45 µm filters [18]. Supernatants were evaporated to dryness at 35 °C by evaporator pumps. Dried supernatants were solved using 0.1 M KH₂PO₄ (pH 8.0). Extracts were centrifuged at 5000 rpm for 1 hour at 4 °C to separating fatty acids [19]. Polvinilpolipirilidon (PVPP),

1 g, was prepared and added to supernatants to separate phenolic and colored matters [20]. Supernatants with PVPP were filtered with What-man No:1 filter paper to remove PVPP [21]. For further specific separation, a Sep-Pak C-18 (Waters) cartridge was used. Hormones adsorbed by the cartridge were transferred to vials using 80% methanol. The hormones were analyzed by HPLC on a Zorbax Eclipse-AAA C-18 column (Agilent 1200 HPLC) and absorbance measured at 265 nm in a UV detector. The flow rate was set to 1.2 ml/min and at column, temperature was 25 °C. Gibberellic acid, salicylic acid, indol-3-acetic acid (IAA), abscisic acid (ABA) was determined by using 13% acetonitrile (pH 4.98) as the mobile phase.

Element Analysis. The Kjeldahl method and a Vapodest 10 Rapid Kjeldahl Distillation Unit (Gerhardt, Königswinter, Germany) were used to determine total N [22] and Ca, Mg, K, P, Fe, Cu, Mn, Zn, B, Se contents of walnut. after wet digestion of dried and ground samples using a HNO₃-H₂O₂ acid mixture (2:3 v/v) with three steps (1st; 145 °C, 75% RF, 5 min; 2nd; 180 °C, 90% RF, 10 min and 3rd; 100 °C, 40% RF, 10 min) in a microwave oven (Bergof Speedwave Microwave Digestion Equipment MWS-2) [23]. The contents Ca, Mg, K, P, Fe, Cu, Mn, Zn, B, Se of types were determined using an Inductively Couple Plasma spectrometer (Perkin-Elmer, Optima 2100 DV, ICP/OES, Shelton, CT 06484-4794, USA) [24].

Statistical Analysis. Biplot analysis was made to determine similarities/ dissimilarities, performances and stabilities in genotypes and parameters. Genotypes are symbolized as GN in biplot analysis.

RESULTS

Fruit Quality Parameters. The fruit weight, width and length of the walnut genotypes are shown in Figure 2. The fruit weight of the genotypes varied within the wide range, from 7.33 to 13.14 g. The fruit weight was higher in genotype GN6 (13.14 g) followed by genotype GN18 (12.91g). The fruit width of the genotypes varied within the wide range, from 28.47 to 36.57 mm. The fruit width was highest in genotype GN18 (36.56 mm) followed by genotype GN19 (34.52 mm). The fruit size of the genotypes varied within the wide range, from 31.91 to 45.18 mm. The fruit size was highest in genotype GN2 (45.18 mm) followed by genotype GN18 (42.17 mm). Differences were observed among genotypes in the fruit weight, width, and size. The fruit ash of the genotypes varied within the wide range, from 1.64% to 3.32%. The fruit ash was highest in genotype GN5 (3.32%), followed by genotype GN26 (2.46%).

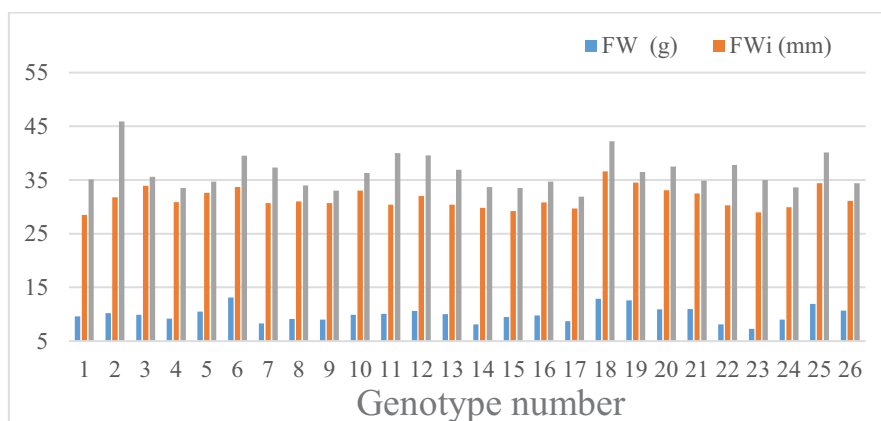


FIGURE 2
Fruit Weight (FW), Fruit Width (FWi), Fruit Length (FL) of walnut genotypes

TABLE 1
Fruit nutrient element contents of walnut genotypes

GN	N	Cp	P	K	Ca	Mg	Na	Zn	Cu	Mn	Fe	B	Se	A	
----	----	-%	-----					-----mg/kg							-%
1	1.9	12.1	3754	9215	3765	2214	242.2	29.7	15.6	24.5	32.6	22.3	86.6	1.9	
2	2.1	13.0	3812	9122	3789	2245	230.2	28.1	16.1	24.6	31.5	24.1	83.1	2.1	
3	2.2	13.5	3988	9134	3812	2234	222.2	29.9	15.4	25.4	34.5	23.4	92.5	2.2	
4	2.2	13.9	3874	9172	3887	2213	228.7	28.7	16.0	25.4	32.9	22.1	88.5	2.2	
5	3.3	20.7	3879	9233	3834	2265	229.8	27.7	15.2	26.3	33.1	21.1	95.4	3.3	
6	2.2	13.6	3912	9413	3712	2287	227.7	27.0	16.6	25.5	34.1	22.3	97.2	2.2	
7	1.7	10.7	3988	9230	3989	2130	224.5	28.1	17.0	24.3	34.2	21.0	90.9	1.7	
8	1.7	10.6	3856	9140	3944	2209	227.2	26.7	17.0	25.7	35.2	23.5	92.5	1.7	
9	1.6	10.2	3813	9187	3980	2167	229.8	25.0	18.1	25.1	32.9	22.3	89.7	1.6	
10	2.2	13.5	3987	9170	3945	2200	228.5	26.8	18.9	24.5	35.4	20.3	96.9	2.2	
11	2.3	14.5	4012	9121	4023	2214	231.1	26.1	17.3	25.0	36.1	21.1	88.7	2.3	
12	1.7	10.6	4094	8904	3972	2175	224.9	24.5	19.1	26.1	35.2	24.6	80.1	1.7	
13	1.7	10.8	4013	9234	3912	2190	231.5	25.0	16.7	25.9	37.1	21.0	79.5	1.7	
14	2.0	12.5	4015	9214	4009	2234	230.1	23.2	16.9	26.4	37.1	19.9	89.8	2.0	
15	2.1	13.0	4035	9087	4087	2176	229.8	26.5	18.1	26.0	38.2	22.3	80.7	2.1	
16	1.9	11.7	4123	9122	4092	2114	224.6	24.3	19.5	25.4	35.5	21.9	86.7	1.9	
17	2.0	12.5	3988	9100	4122	2186	231.9	20.6	19.2	26.1	35.1	19.5	92.6	2.0	
18	2.2	13.5	4098	9213	4005	2209	226.5	22.6	18.7	24.3	38.3	18.8	94.5	2.2	
19	2.0	12.6	3990	8977	4233	2261	224.9	23.4	18.1	25.5	36.1	17.7	78.6	2.0	
20	2.2	13.5	3870	8954	4198	2289	218.4	24.5	19.1	25.0	35.4	17.1	88.6	2.2	
21	2.0	12.6	3912	8844	4120	2241	216.9	20.1	16.6	27.7	33.1	18.4	92.7	2.0	
22	1.9	11.6	4122	8970	3987	2277	227.6	18.8	18.1	26.1	36.1	19.2	88.2	1.9	
23	2.2	13.8	4089	9032	4013	2018	225.1	22.3	19.2	25.3	38.8	15.7	94.6	2.2	
24	2.0	12.7	4032	8874	4093	2097	220.8	25.5	19.9	22.3	34.5	18.8	75.7	2.0	
25	2.2	13.8	4109	8942	4052	2133	218.9	21.4	19.0	24.5	35.4	20.1	69.4	2.2	
26	2.5	15.4	4187	8945	4129	2172	221.0	22.3	18.7	26.8	39.1	22.3	67.5	2.5	

GN: Genotype number, N: Nitrogen, Cp: Crude protein, P: Phosphorus, K: Potassium, Ca: Calcium, Mg: Magnesium, Na: Sodium, Zn: Zinc, Cu: Copper, Mn: Manganese, Fe: Iron, B: Boron, Se: Selenium, A: Ash

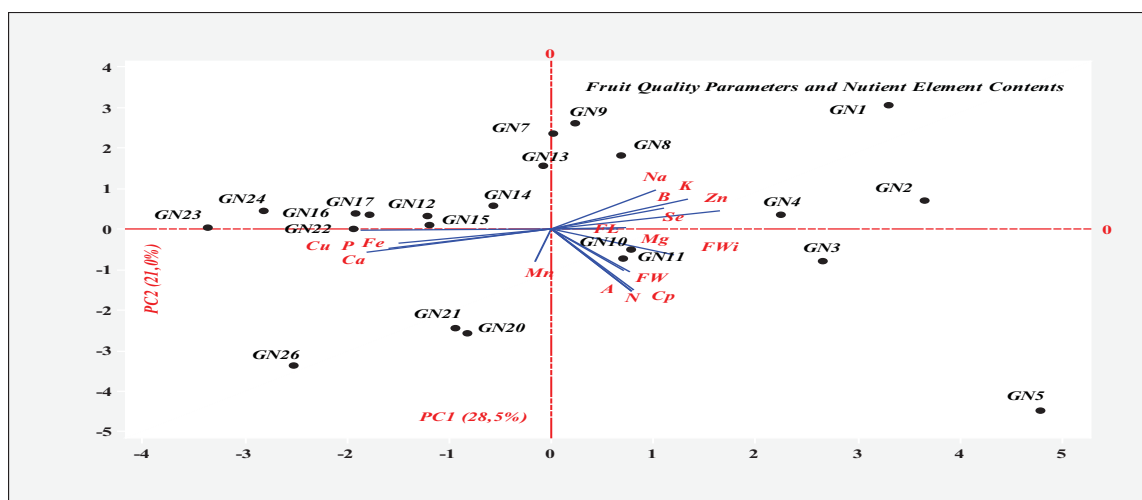


FIGURE 3

Fruit quality parameters and nutrient element contents of walnut genotypes

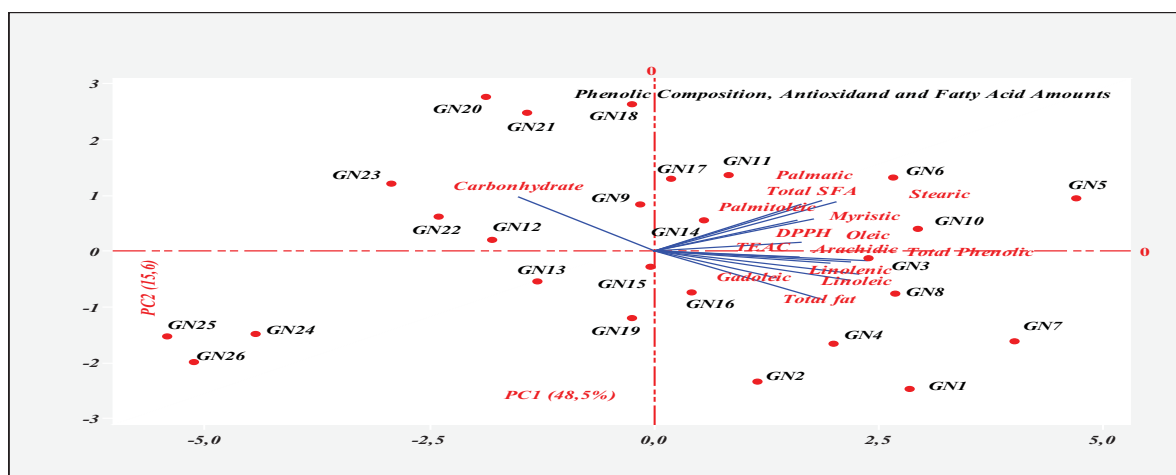


FIGURE 4

Phenolic composition, antioxidant and fatty acid amounts of walnut genotypes

Biplot analysis of fruit weight, fruit width, fruit length and minerals examined in the study are given in Figure 3. Fruit weight, fruit length, fruit width, ash amount and protein ratio were grouped under one group. In this case, it is seen that walnut genotypes do not show excessive variation in terms of these properties and exhibit similar trends. Three groups occurred for fruit quality parameters. In first group, GN26, GN21 and GN20; in second group, GN1, GN2, GN3, GN4, GN10 and GN11; and the third group had the other genotypes. Fruit weight, fruit length, fruit width, ash content and crude protein were stable and showed a high amount in terms of genotypes. As for genotypes, GN2, GN3, GN4, GN7, GN8, GN9, GN10 and GN11 genotypes were determined as rich and stable genotypes in terms of fruit characteristics.

Macro and Micro Nutrient Contents. The crude protein and mineral concentrations of the walnut genotypes are shown in Table 1. The crude protein content of the genotypes varied within the wide

range, from 10.21% to 20.71%. The crude protein content was highest in genotype GN5 (20.71%), followed by genotype GN26 (15.36%). The phosphorus content of the genotypes varied within the wide range, from 3754 mg/kg to 4187 mg/kg. The phosphorus content was highest in genotype GN26 (4187 mg/kg), followed by genotype GN16 (4123 mg/kg), and genotype GN22 (4122 mg/kg). The potassium content of the genotypes varied within the wide range, from 8844 mg/kg to 9413 mg/kg. The potassium content was highest in genotype GN6 (9413 mg/kg), followed by genotype GN13 (9234 mg/kg). The calcium content of the genotypes varied within the wide range, from 3712 mg/kg to 4233 mg/kg. The calcium content was highest in genotype GN19 (4233 mg/kg), followed by genotype GN20 (4198 mg/kg), and the same differences were observed among genotypes in the other nutrient elements etc. Mg, Na, Zn, Cu, Mn, Fe, B, and Se contents of walnut fruit. In biplot analysis (Figure 3), the parameters analyzed consisted of two groups. While Cu, Fe, Ca and P form one group, other minerals form another

TABLE 2
Total fat, carbohydrate, phenolic composition and antioxidant amounts of walnut genotypes

GN	Total fat	Carbohydrate	Total Phenolic	TEAC	DPPH
		%	mg GAE/100g FW	$\mu\text{molTE/g FW}$	
1	69.14	15.05	1698.18	113.212	83.613
2	66.71	16.49	1505.05	100.336	76.993
3	63.45	19.11	1367.88	111.191	80.050
4	67.86	14.31	1530.92	102.060	94.428
5	71.43	18.99	1825.72	121.714	86.167
6	63.15	19.41	1551.02	108.401	97.449
7	71.42	14.66	1754.09	116.939	90.449
8	66.29	19.77	1628.10	112.539	96.839
9	60.71	25.57	1491.12	109.407	88.109
10	64.58	18.18	1585.97	105.731	72.399
11	60.45	20.81	1303.20	99.879	95.562
12	58.74	27.52	1360.13	90.675	82.451
13	60.43	25.58	1484.13	98.941	86.567
14	63.45	20.50	1558.21	103.880	77.210
15	63.30	19.99	1389.79	118.652	91.237
16	66.87	18.12	1642.27	109.484	85.802
17	62.88	21.06	1544.44	102.962	81.618
18	57.48	25.30	1490.14	97.942	88.542
19	64.89	19.03	1593.76	106.250	87.344
20	53.75	29.00	1320.20	88.013	80.732
21	59.17	24.65	1453.19	96.879	81.908
22	60.03	24.12	1474.35	88.290	70.518
23	61.74	20.71	1269.33	80.622	70.843
24	62.02	21.70	1275.18	85.012	68.707
25	60.15	22.36	1236.73	82.448	70.712
26	60.74	19.98	1272.82	84.854	72.650

group. Genotypes are grouped under 3 groups. GN26, GN21 and GN20 created one group, while GN1, GN2, GN3, GN4, GN10 and GN11, created another group. The remaining genotypes formed a separate group.

Phenolic Composition, Antioxidant and Fatty Acids Amounts. The phenolic composition and antioxidant amount of the walnut genotypes are presented in Table 2. Considerable differences were observed among walnut genotypes in terms of phenolic composition, and antioxidant amounts.

The total oil of the genotypes varied within the wide range, from 53.8% to 71.4%. The total oil rate was highest in genotype GN5 (71.4%), followed by genotype GN7 (71.4%). The carbohydrate amounts of the genotypes varied within the wide range, from 14.3% to 29.0%. The carbohydrate amount was highest in genotype GN20 (29.0%), followed by genotype GN12 (27.5%). Similarly, the total phenolic composition of the genotypes varied within the wide range; from 1236 mg GAE/100 g FW to 1825 mg GAE/100 g FW. The total phenolic composition was highest in genotype GN5 (1825 mg GAE/100 g FW), followed by genotype GN7 (1754 mg GAE/100 g FW). TEAC, DPPH, palmitic acid,

myristic acid, stearic acid, arachidic acid, total SFA, palmitoleic acid, oleic acid, gadoleic acid, linoleic acid, and linolenic acid of the genotypes varied within the wide range. The highest TEAC, myristic acid, arachidic acid, total SFA, palmitoleic acid value in genotype GN5 (121 $\mu\text{molTE/g FW}$; 1.01%; 0.176%; 7.85%, and 2.32% respectively) (Table 2, Table 3). The highest DPPH, and palmitic acid values were found in genotype GN6 (97 $\mu\text{molTE/g FW}$, 5.40%, respectively). The highest stearic acid value was observed in genotype GN7 (1.40%). The highest oleic acid, linoleic acid, and linolenic acid values were in genotype GN3 (25.58%, 88.22%, and 11.03%, respectively). Moreover, the highest gadoleic acid was found in genotype GN2 (0.131%). Considering the biplot analysis in Figure 4, showing the phenolic composition, antioxidant and fatty acid amounts, the investigated properties formed 2 groups. While carbohydrates constitute a group alone, other features examined are grouped under one group. Genotypes are grouped under 4 groups. While GN5 created a single group, GN1, GN7 and GN8 created a second group. While GN20, GN21, GN23, GN24 and GN26 constitute the third group, other genotypes are grouped under the fourth group.

TABLE 3
Fatty acid amounts of walnut genotypes

GN	Fatty Acid (%)									
	Pal-matic	Myristic	Stea-ric	Ara-chidic	SFA	Pal-mitoleic	Oleic	Gado-leic	Lino-leic	Lino-lenic
1	4.81	0.60	1.24	0.169	6.81	1.93	22.34	0.129	87.51	10.94
2	4.62	0.72	1.02	0.166	6.53	2.09	20.18	0.131	83.00	10.37
3	5.14	0.71	1.32	0.117	7.29	2.11	25.58	0.126	88.22	11.03
4	4.92	0.77	1.03	0.090	6.81	2.13	23.12	0.083	84.71	10.59
5	5.30	1.01	1.36	0.176	7.85	2.32	19.54	0.028	81.70	10.21
6	5.40	0.75	1.39	0.097	7.64	2.18	22.15	0.068	79.78	9.97
7	5.05	0.69	1.40	0.172	7.31	1.71	23.55	0.125	83.03	10.38
8	5.14	0.61	1.32	0.173	7.24	1.69	22.45	0.127	78.89	9.86
9	4.99	0.60	1.33	0.091	7.00	1.64	19.26	0.029	73.79	9.22
10	5.38	0.75	1.38	0.147	7.66	2.16	25.34	0.127	79.07	9.88
11	4.93	0.80	1.27	0.127	7.13	2.32	23.98	0.099	57.42	7.18
12	4.45	0.67	1.04	0.116	6.28	1.92	25.10	0.027	53.84	8.73
13	4.42	0.60	1.14	0.163	6.32	1.73	18.66	0.126	54.92	6.86
14	4.99	0.69	1.28	0.137	7.10	2.12	21.50	0.058	50.87	8.36
15	4.49	0.72	1.15	0.064	6.42	2.08	25.30	0.028	58.28	7.28
16	4.82	0.65	1.24	0.069	6.77	1.90	20.66	0.128	53.42	6.68
17	5.14	0.69	1.32	0.093	7.25	2.00	22.98	0.031	44.28	7.53
18	5.25	0.75	1.25	0.138	7.39	2.17	20.13	0.029	49.73	6.22
19	4.37	0.70	1.12	0.092	6.28	2.01	21.56	0.123	51.44	6.43
20	4.92	0.75	1.27	0.070	7.01	2.17	19.66	0.093	53.86	6.73
21	5.15	0.70	1.32	0.074	7.25	2.02	18.34	0.022	39.72	5.97
22	4.90	0.64	1.13	0.070	6.74	1.86	19.20	0.127	37.09	5.64
23	5.26	0.58	1.35	0.075	7.26	1.21	17.45	0.025	44.12	5.52
24	4.21	0.56	1.08	0.060	5.90	1.04	19.23	0.016	50.27	6.28
25	3.86	0.60	0.99	0.055	5.50	1.21	18.42	0.014	42.21	5.28
26	3.75	0.59	0.96	0.054	5.35	1.46	17.12	0.016	44.05	5.51

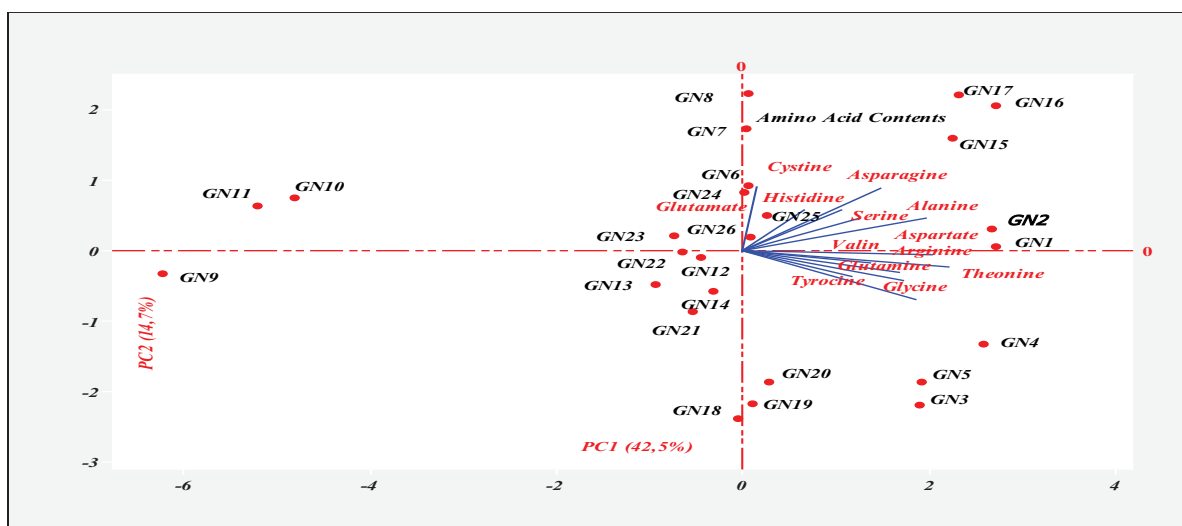


FIGURE 5
Amino acid contents of walnut genotypes

Amino Acids Contents. The amino acid contents of the walnut genotypes are presented in Table 4. Considerable differences were observed among walnut genotypes in terms of amino acid compositions. Overall, aspartate, glutamate, isoleucine, leucine and lycine were the most abundant amino acids, whereas all genotypes deficient in cystine, valin, and,

tryptophan. In general, the highest amino acid concentrations were detected in genotype GN16. On the other hand, most of the amino acids were also high in genotype GN15, genotype GN1, genotype GN4, and genotype GN17. Cystine, valin, tryptophan, lysine, sarcosine, serine, histidine and arginine which are deficient amino acids in walnut, changed

TABLE 4
Amino acid contents of walnut genotypes

GN	Aspar- tate	Gluta- mate	Aspara- gine	Ser- ine	Gluta- mine	Histi- dine	Glycine	The- onine	Argi- nine	Alanine	Tyro- cine	Cys- tine	Valin
pmol/μg													
1	2733	5340	6137	3971	3467	2290	3181	2909	4440	4750	1261	556	940
2	2793	5180	5937	4286	3340	2123	3156	2985	4377	4769	1234	614	956
3	2690	5088	5460	4161	3429	1894	3471	2781	4321	4388	1376	547	976
4	2687	5412	5739	4094	3588	1974	3425	2864	4334	4449	1380	590	877
5	2743	5292	5499	4075	3500	1823	3336	2877	4274	4434	1332	577	912
6	2579	4916	5679	4135	3339	2123	3012	2653	3992	5058	1190	627	944
7	2639	5368	5868	4035	3244	2067	3059	2665	4061	5022	1217	670	844
8	2636	5268	5867	4211	3249	2100	2883	2741	3946	5131	1119	663	890
9	2287	4572	5083	3792	3114	1976	2641	2509	3232	2963	1097	551	807
10	2273	5360	5318	3934	3228	1870	2613	2610	3247	2939	1021	606	891
11	2204	5012	5413	4120	3266	1876	2747	2561	3185	2912	1053	585	754
12	2492	5564	5828	4064	3363	1966	3169	2794	3792	4619	1107	487	796
13	2504	5736	5558	3594	3509	2050	3109	2692	3680	4520	1182	534	821
14	2542	5492	5727	3978	3434	1894	3257	2891	3744	4665	1063	517	743
15	2507	5840	6285	4313	3514	2098	3229	2710	4224	5160	1129	570	952
16	2687	5980	6421	4465	3467	2055	3180	2748	4197	5199	1084	552	925
17	2719	5680	6227	4388	3424	2150	3100	2798	4178	5233	1058	606	925
18	2561	4876	5305	4161	3334	1929	3359	2793	4187	4294	1056	434	945
19	2617	4944	5085	4222	3393	2055	3297	2738	4256	4237	1085	445	930
20	2614	5080	5344	4075	3370	1948	3225	2890	4140	4339	1012	476	970
21	2605	4612	5729	4030	3452	2146	3036	2686	4041	4552	1134	459	817
22	2671	5032	5490	3971	3468	2203	2921	2635	3959	4597	1061	509	863
23	2662	4896	5671	4106	3362	2198	2908	2730	3987	4559	1064	495	800
24	2739	4940	5727	4167	3423	2324	3007	2618	4053	4655	1096	534	821
25	2664	5048	5426	4291	3384	2330	3075	2692	4123	4588	1109	551	866
26	2514	5372	5423	4109	3412	2309	3124	2714	4134	4500	1124	540	885

TABLE 4 continue
Amino acid contents of walnut genotypes

GN	Methionine	Trvptophan	Phenylalanine	Isoluecine	Leucine	Lvsine	Hydroxvproline	Sarcosine	Proline
nmol/μg									
1	2072	989	1044	1388	2850	2574	552	4326	59
2	1991	940	1051	1392	2662	2738	625	4294	58
3	1882	952	1076	1295	2573	2817	552	5232	55
4	2026	912	1005	1204	2453	2906	581	5192	56
5	1767	959	1113	1250	2665	2909	514	5361	53
6	1768	806	1151	1290	2750	2458	581	4269	49
7	1927	838	1235	1358	2629	2376	545	4369	51
8	1645	831	1226	1239	2563	2524	624	4166	47
9	1923	700	1059	1131	2606	2265	494	4276	50
10	1861	724	1094	1311	2752	2164	499	4339	55
11	1808	736	1096	1230	2698	2356	456	4310	48
12	1707	837	1095	1230	2263	2432	586	4329	55
13	1814	813	1122	1292	2312	2679	537	4580	65
14	1592	873	1100	1276	2489	2523	554	4430	53
15	1940	890	1178	1330	2535	2438	535	4487	59
16	1892	929	1196	1319	2447	2673	507	4340	54
17	1817	958	1102	1279	2434	2610	532	4384	59
18	2114	977	1123	1122	2517	2537	544	4249	53
19	2204	912	1031	1230	2588	2475	562	4153	55
20	1999	985	1160	1198	2610	2574	545	4057	51
21	1788	929	1213	1317	2247	2547	535	4372	59
22	1620	979	1120	1233	2070	2673	562	4457	58
23	1838	930	1049	1334	2220	2605	518	4269	57
24	1920	948	1114	1318	2262	2669	502	4376	56
25	1876	941	1096	1350	2403	2571	542	4309	58
26	1912	954	1176	1412	2322	2645	554	4312	63

considerably among genotypes. The range of cysteine, valin, tryptophan, lysine, sarcosine and proline contents of walnut genotypes were from 434 (GN 18) to 670 (GN 7) pmol μl^{-1} , 743 (GN 14) to 976 (GN 3) pmol μl^{-1} , 700 (GN 9) to 989 (GN 1) pmol μl^{-1} , 2164 (GN 10) to 2909 (GN 5) pmol μl^{-1} , 4057 (GN 20) to 5361 (GN 5) pmol μl^{-1} and 47 (GN 8) to 65 (GN 13) pmol μl^{-1} , respectively.

In terms of biplot analysis (Figure 5), when looking at the amino acid of the studied walnut genotypes, all the amino acids examined constitute a single group, while the genotypes are grouped under 4 groups. While GN9, GN10 and GN11 form a group, GN18, GN19 and GN20 form the second group. Again, GN3, GN4 and GN5 constitute the third group, while the remaining genotypes are grouped under the fourth group.

Organic Acid Contents. As seen from the Table 5, succinic, oxalic, lactic, malic, citric, and propionic acids were the most abundant, whereas tartaric, butyric, maleic and fumaric acids were the scarcest, respectively. Organic acid content of the genotypes

varied within the wide range. Generally, most of the organic acids were high in genotype GN11, GN14, GN22, GN23 and GN26, and low in genotype GN2, GN4, GN5, and GN6. The highest total organic acid contents were determined in genotype GN11 (48.80 ng μl^{-1}). The range of oxalic acid contents of walnut genotypes were from 5.51 to 7.56 ng μl^{-1} . The highest oxalic acid contents were determined in genotype GN8 (7.56 ng μl^{-1}). Similarly, the range of succinic acid contents of walnut genotypes were from 5.67 to 9.64 ng μl^{-1} . The highest succinic acid contents were determined in genotype GN26 (9.66 ng μl^{-1}).

The biplot analysis of the organic acid amounts of the walnut genotypes showed that all organic acids examined constitute a single group, while the genotypes are grouped under 4 groups. While GN11 created a stand-alone group, GN2, GN4, GN5, GN6 and GN17 formed the second group. Again, GN1, GN3, GN7, GN8, GN9 and GN16 constitute the third group, while the remaining other genotypes are grouped under the fourth group (Figure 6).

TABLE 5
Organic acid contents of walnut genotypes

GN	Oxalic	Propionic	Tartaric	Butyric	Malonic	Malic	Lactic	Citric	Maleic	Fumaric	Succinic
Ng/ μl											
1	6.44	5.46	1.20	1.87	3.74	5.44	4.32	4.55	0.30	1.13	7.71
2	6.51	5.10	1.06	1.68	3.51	5.26	3.87	4.38	0.28	1.07	7.87
3	6.74	5.63	1.12	1.77	3.87	5.18	4.57	4.16	0.34	1.04	7.83
4	6.23	5.20	1.23	1.62	3.38	5.69	3.59	4.27	0.43	1.02	5.67
5	5.51	4.42	1.36	1.54	3.59	5.87	3.25	4.10	0.41	1.19	6.21
6	6.22	5.37	1.26	1.47	3.63	5.43	3.87	4.28	0.40	1.25	7.55
7	6.88	5.84	1.45	1.97	4.02	4.95	4.71	3.88	0.28	1.16	7.19
8	7.56	5.64	1.73	1.78	3.94	5.11	4.64	4.23	0.28	1.15	6.70
9	7.17	6.01	1.80	1.88	4.09	4.69	4.95	3.49	0.32	1.07	7.31
10	7.14	6.07	1.45	1.73	4.65	6.06	4.79	4.80	0.66	1.39	9.04
11	6.42	5.29	1.58	1.65	4.87	6.96	4.45	5.83	0.61	1.56	9.58
12	6.14	5.49	1.67	1.95	4.41	6.59	3.93	4.54	0.63	1.02	8.62
13	6.80	5.91	1.85	1.69	4.20	5.80	4.11	5.36	0.23	1.28	7.67
14	6.72	6.34	1.73	1.82	4.41	6.78	4.72	5.61	0.31	1.47	8.74
15	6.80	6.08	1.97	1.73	4.01	6.27	4.64	4.96	0.30	1.19	8.63
16	6.58	4.85	1.63	1.93	3.53	5.74	4.31	4.27	0.35	1.06	7.88
17	6.24	4.26	1.68	1.82	3.70	6.59	4.64	4.70	0.31	1.09	8.15
18	6.10	4.91	1.78	1.67	4.67	6.17	4.40	4.57	0.33	1.30	8.01
19	6.63	5.52	1.46	1.89	4.12	5.35	4.43	4.38	0.36	1.36	9.51
20	6.31	6.57	1.89	1.92	4.33	5.95	4.53	4.32	0.38	1.24	8.99
21	6.92	6.66	1.66	1.80	4.09	4.99	4.67	4.11	0.39	1.28	9.64
22	6.71	6.07	1.72	1.88	3.74	7.25	5.06	4.30	0.53	1.34	9.45
23	6.54	6.45	1.85	1.78	3.95	7.42	4.72	4.51	0.48	1.43	9.10
24	6.23	6.33	1.66	1.61	4.01	6.42	5.69	4.70	0.46	1.42	8.79
25	6.51	6.73	1.73	1.68	4.09	6.22	5.24	4.45	0.55	1.38	9.23
26	6.38	6.54	1.77	1.83	4.19	6.09	5.60	4.84	0.56	1.27	9.66

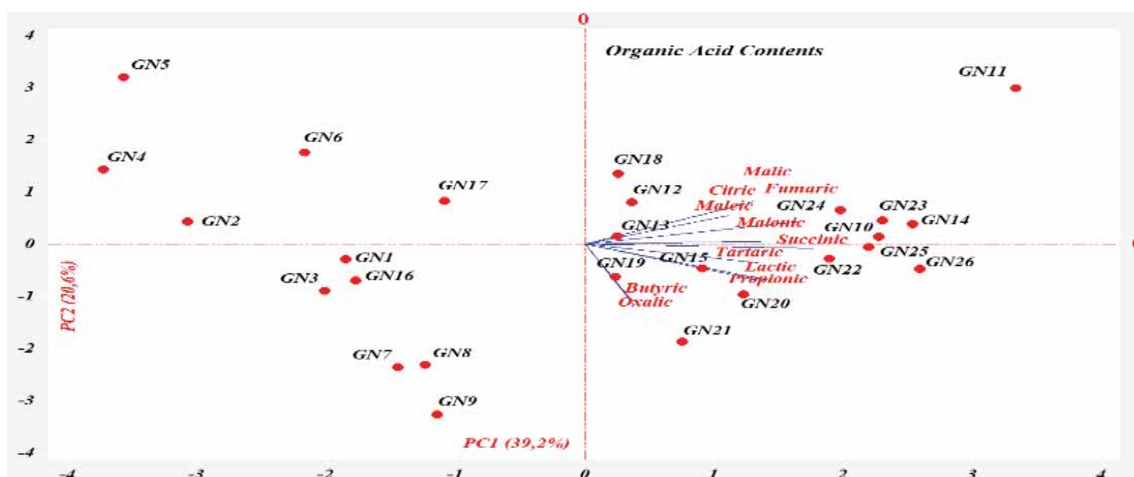


FIGURE 6
Organic acid contents of common walnut genotypes

TABLE 6
Hormone contents of walnut genotypes

GN	Gibberellic acid	Salicylic acid	Absisic acid	Indole acetic acid
	Ng/μl			
1	291.4	55.0	0.14	10.8
2	290.9	47.6	0.13	11.4
3	291.0	57.7	0.12	12.6
4	271.8	63.6	0.13	13.5
5	262.7	57.6	0.11	14.4
6	318.5	57.8	0.12	14.0
7	323.1	61.1	0.10	13.5
8	312.5	54.8	0.10	12.9
9	328.6	65.5	0.13	14.3
10	346.3	72.1	0.14	15.6
11	325.6	65.6	0.11	16.1
12	336.6	57.6	0.13	13.5
13	341.7	60.1	0.12	14.3
14	338.1	54.6	0.14	13.4
15	330.6	72.8	0.10	12.8
16	324.8	75.2	0.10	13.9
17	302.3	81.3	0.14	14.6
18	280.6	68.2	0.11	12.1
19	275.8	68.8	0.13	13.4
20	288.7	73.4	0.13	13.9
21	298.2	72.4	0.14	13.2
22	267.1	68.7	0.14	14.8
23	298.8	73.5	0.11	14.1
24	293.5	71.3	0.13	15.3
25	304.9	74.5	0.13	14.6
26	290.8	70.9	0.15	15.5

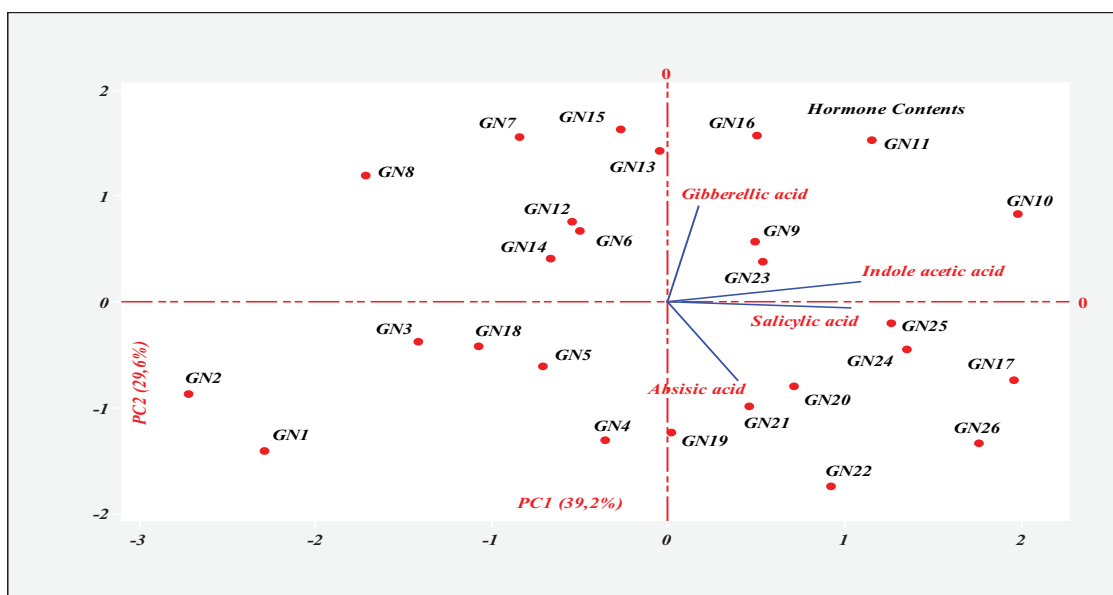


FIGURE 7
Hormone contents of common walnut genotypes

Hormone Contents. Walnut genotypes showed differences in terms of hormone content (Table 6). Gibberellic acid content of the genotypes ranged between 262.73 ng μl^{-1} to 346.28 ng μl^{-1} . The highest gibberellic acid contents were determined in genotypes GN10, GN12, and GN13, whereas genotypes GN5, GN19, and GN22 had the lowest gibberellic acid contents. The amount of salicylic acid, which was the second abundant hormone following gibberellic acid, varied among walnut genotypes. Genotype number 17 (81.32 ng μl^{-1}) had higher salicylic acid content than those of the other genotypes. The indole acetic acid content of the genotypes also varied within the wide range, from 10.78 ng μl^{-1} to 16.09 ng μl^{-1} . The amount of indole acetic acid was high in genotype GN 10, 11, 24, and 26, and low in genotype GN 1, 2, 3, 8, 15, and 18. Abscisic acid content of the genotypes ranged between 0.10 ng μl^{-1} (genotype GN 7, 8, 15, and 16) to 0.15 ng μl^{-1} (genotype GN 26). If we look at the hormones in Figure 7, the parameters examined consisted of 3 groups. While gibberellic acid 1 group constituted another group of abscisic acid, indole formed a separate group in acetic and salicylic acid. In terms of genotypes, genotypes were determined to form 4 groups. While GN1, GN2, GN3, GN4, GN5 and GN18 are creating a group, GN6, GN7, GN8, GN12, GN13, GN14 and GN15 have created the second group. The third group included GN9, GN10, GN11, GN16 and GN23, while the fourth group included GN17, GN19, GN20, GN21, GN22, GN24, GN25 and GN26 (Figure 7).

DISCUSSIONS

Fruit Quality Parameters. Turkey has an important walnut production rate due to the favorable

climatic and soil characteristics. Turkey is rich in walnuts population and is one of the leading countries in the world in walnut production until the end of the year. Walnut culture in Turkey as well, each made because the seeds come from very ancient times have different characteristics from each other and adapted to the ecological conditions of the region in the form of a wide genetic variation. In this genetic variation, it is extremely important that the types which are highly qualified are protected from reproduction. Turkey's having such a rich genetic variation makes it possible to achieve success in breeding work as soon as possible. Determination of fruit quality characteristics and fatty acid composition of these Turkish walnut varieties and genotypes is very important for commercial culture and oil industry [25]. This study results showed that fruit weight and fruit quality parameters significantly varied among plants. Akca and Sen [26] showed nut length as 39.97 mm, nut diameter as 33.59 mm of the promising walnut genotype. Khatkhat et al. [27] determined the nut weight (1.90-12.30 g), for the walnut cultivars. Our results are higher than that shown by Ozkan and Koyuncu [28], who studied the walnut cultivars and found nut weights between 8.43-11.09 g. This could be due to the differences in the ecological and genetic properties of walnut cultivars growing in Turkey. The results of the nut properties showed that fruits of walnut cultivars have superior physical and chemical properties.

Mineral Nutrient Contents: Similar to the results reported in the literature [29, 30], our findings also showed that walnuts contain high levels of potassium (390-700 mg 100 g⁻¹), phosphorus (310-510 mg 100 g⁻¹) and magnesium (90-140 mg 100 g⁻¹), and low sodium (1-15 mg 100 g⁻¹). The calcium content in walnuts genotypes ranged from 58-91 mg

100 g⁻¹ [31], 640-1180 mg/kg [32]. Calcium values found in our study were considerably higher than those reported by Lavedrine et al. [31]. Magnesium and sodium contents of this study were generally higher than those reported by many studies in different walnut varieties. Copper, manganese and zinc contents were similar to values reported by other researchers [31, 32]. Content of elemental composition of walnuts can be influenced by genotype, cultivar, different ecology and different soil. While the levels of Cu, Ca, F and P are insufficient on the basis of genotypes, genotypes have a higher amount of minerals in terms of the minerals that make up the third group. Although all minerals except Mn have a stable course in genotypes, genotypes have a richer structure in terms of Na, K, B, Zn, Se, Mg and N. This shows that genotypes have given a higher quality product in terms of these minerals throughout the Valley. Genotypes of GN2, GN3, GN4, GN10 and GN11 are both stable and richer genotypes in terms of mineral substance.

Phenolic Composition and Antioxidant Amounts. Our results concerning phenolic composition and fatty acid were in agreement with the results obtained from walnut cultivars [33] but higher than those of Ozcan et al. [35]. Linoleic acid and linolenic acid contents were similar to the results of Zwartz et al. [33] but higher than the results of Çaglarırmak [36]. There were some differences in the fatty acid composition between our findings and the literature. It is thought that this difference may come from many factors such as the region where it is cultivated, the harvesting time, the year of harvest, and the cultural processes applied, such as irrigation [7]. Although they contain very different nutrients, the most important nutrient provided by walnuts is the dietary fat. The walnut oil is rich in essential fatty acids and the fatty acid composition is mainly composed of oleic, linoleic and linolenic acids. The walnut genotypes with high fatty acid, especially linoleic and linolenic acids, have beneficial effects on human health especially for cardio vascular system [37]. The ratios of oleic, linoleic and linolenic acids to each other are important to the economic and nutritional value of the nuts [25]. Therefore, fatty acid compositions are an important criterion for identification of walnut genotypes in different regions. The consumption of walnut is increasing as consumers are showing an interest in their valuable fatty acid content. It is reported that polyunsaturated fatty acids have a very important role in human nutrition. Polyunsaturated fatty acids found in walnut play an important role in the prevention of cardiovascular diseases and prevention of vascular occlusion [38]. Pereira et al. [37] determined fats (68.83%-72.14%), carbohydrates (3.75-7.16%), and ash (3.31-4.26%) for the six walnut cultivars grown in Portugal. In Italy, internal fat content (54.2-72.2%) linoleic (46.9-68.6%), oleic (10.0-25.1%), linolenic (6.9-17.6%),

palmitic (3.9-11.4%) and stearic fatty acids (1.1-5.2%) were determined for 190 wild walnut types [38]. The results of this study are in good agreement with this notion except for the percent carbohydrates and ash contents. Al-Bachir [39] determined the protein (22.85%), fat (67.35%) and ash (1.26%) content. The results of the fat content and ash content are in agreement with the results of this study but there is variation in term of protein content. These differences and variations can be attributed to environmental conditions, horticulture procedures and composition of walnut fruits. The fact that other studied properties except carbohydrates show similar stability in genotypes reveals that the genotypes sampled throughout the Valley do not show much variation in terms of these substances. Therefore, these substances show almost similar variations on the basis of genotypes. Genotypes showed a rich and stable structure in terms of all phenolic and antioxidant compounds except carbohydrates. Regarding the genotypes, GN1, GN2, GN3, GN4, GN8, GN10, GN11, GN14 and GN16 have been determined as rich and stable varieties in terms of phenolic and antioxidant compounds.

Amino Acid Contents. Previous studies showed that amino acids may contribute to yield, growth and nutrient element uptake from soil in different plant species under stress conditions for different crops. Proline, alanine, serine, and asparagine also delayed wilting of maize under stress conditions, proline, glycine, alanine, leucine, threonine, lysine, arginine, tryptophan and phenylalanine inhibited stomatal opening while histidine, methionine, aspartic acid, glutamic acid, asparagine and glutamine promoted stomatal opening of *Vicia faba*, histidine, proline, glutamine, methionine and glycine promoted calcium uptake in phaseolus seedlings, proline relieved salt toxicity in barley plant lets by changing salt transport from root to shoot and increasing proline content increased K⁺ content and alleviated salt stress effects on growth of *Vigna radiate* cultures [40]. In the literature, the walnut plant was reported to have a very rich amino acids content. We also found that although the amounts in the plant varied, almost all the genotypes examined throughout the Valley showed quite stable amino acid levels and no major variations were observed in this change (Figure 3). In terms of amino acid levels and stability, GN1, GN2, GN15, GN24 and GN25 genotypes were determined to have a rich and stable structure in terms of amino acids.

Organic Acid Contents. Organic acids make osmotic adjustment and provide a greater cation balance in the plant. Acetic, abscisic, glycolic, malonic, oxalic, and formic acid play a very important role in nutrient uptake (P, Fe and Mn) of plants grown in low nutrient soils. These acids differ from plant species in response to nutrients starvation [41]. Similarly,

these acid anions form complexes with Ca, Al and Fe present in soil as in-soluble phosphates of calcium, iron and aluminium, and liberate P for uptake by roots [42]. Additionally, these acids can desorb P from sesquioxide surfaces by anion exchange [43], and also maintain sulphate mobility in rhizosphere soil through competitive displacement from adsorption sites. The concentrations of fumaric, malic and citric acids can also chelate Fe and Mn in iron and manganese oxides (i.e. Fe₂O₃ and MnO₂), thus making them available for uptake by the plant [42]. Organic acids are the main inputs for many biochemical events in walnuts, as in all plants. Organic acids in genotypes examined in Çoruh Valley were found to be similar in terms of both quantity and change. In this case, the effect of both environmental and genotypic characteristics in the whole genotypes throughout the Valley is almost similar and the organic acid levels in walnuts showed very close changes. All the organic acids examined were determined in high amounts and stable on the basis of genotypes, while the fruit characteristics, protein ratio and ash were found to be not stable. Briefly, GN10, GN11, GN12, GN13, GN14, GN15, GN20, GN22, GN23, GN24, GN25 and GN26 were determined as high and stable varieties in terms of organic acid (Figure 3).

Hormone Contents. The effects of indol-3-acetic acid (IAA), a main auxin in plants, on plants are as follows: to promote root formation of steels, to provide parthenocarpic fruit, to provide adventitious root formation, to prevent fruit and leaf breakdown, to increase fruit attitude, to provide early flowering of buds and to prevent weed growth. In plant cells, The IAA is produced from tryptophan (TRP) in a variety of semi-independent and tryptophan-independent ways. Synthesis of auxin is controlled developmentally and environmentally. Indole-3-acetic acids are known for their effects such as cell division, root development, expansion of root surface area, helping the plant at the entrance of nutrients [44]. Cytokinins are the hormones that initiate the cell division. It is especially synthesized in the root meristems and then transported to the green parts of the plant through xylem. Cytokinins stimulate plant cell division, control root meristem differentiation, inhibits primary root elongation and lateral root formation, but can promote root hair development. The most obvious effect of gibberellins is to increase the prolongation of cells. Gibberellins enhance particularly stem tissue, promote root elongation and lateral root extension. In biplot analysis, although genotypes differed in terms of hormones, IAA and salicylic acid levels in genotypes were determined to be more stable and higher. In other words, the samples examined in the Valley in terms of these two hormones did not change much and they were rich in these two hormones. Salicylic and indole acetic acid have been determined as hormones that are stable on the basis of genotypes and contain high

amounts of genotypes. Again, GN9, GN10, GN17, GN20, GN23, GN24 and GN25 have been determined as stable varieties with high hormone content. In addition, it has been demonstrated that genotypes have different hormone levels throughout the Valley in terms of hormone levels and were not stable in this distribution (Figure 6).

CONCLUSIONS

The present study showed that cultivars affected walnut fruit quality parameters, weight, nutrient contents, crude protein, phenolic components, fatty acid, amino acid, organic acid and hormone contents of walnut in Çoruh Valley, Turkey. The selection of cultivars can improve the economic yield and directly influence the yield components. It can reasonably be concluded that the results of this study showed that cultivars of walnut had a significant effect on fruit quality parameters, weight, nutrient contents, crude protein, phenolic components, fatty acid, amino acid, organic acid and hormone contents of walnut. Different cultivars conditions were caused by a variation in phenolic component, crude protein, fatty acid and nutrient contents. The nut weight of walnut was affected by other components of yield parameters. However, probability in the variations of nut weight and yield parameters was affected by amino acid, organic acid and hormone contents of walnut. Genotypes have a rich and stable structure in terms of other fruit quality parameters, minerals, organic acids, phenolic composition, antioxidant and fatty acids amounts and amino acids. In other words, these characteristics did not show much variation in genotypes grown in the Valley. However, there was a great variation in hormone levels among genotypes. This means that although there may be variations in hormone content in walnut cultivation along the Çoruh Valley, walnut genotypes are expected to display a rich and stable feature across the Valley in terms of other parameters. In conclusion, GN2, GN10, GN15, GN24, GN25 and GN26 genotypes were determined as rich and stable genotypes.

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