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Article

Optimizing Protein-Rich Young Vegetative Quinoa (Chenopodium quinoa) Growth: Effects of Inter-Row Spacing and Genotype in Mediterranean Summer Cultivation

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Abstract: Young vegetative quinoa (YVQ) has gained attention as a high-protein leafy crop with potential for cultivation in Mediterranean and semiarid regions. We investigated the effects of interrow spacing and genotype on YVQ fresh and dry matter (DM) yield, protein content (PC), and protein yield during summer cultivation in northern Israel in two field experiments over two consecutive years (2020–2021). We hypothesized that row spacing and genotypic differences would significantly impact yield and PC. Inter-row spacing significantly affected plant density, ranging from 55–366 plants m⁻². Fresh and DM yields ranged from 4957–28,469 kg ha⁻¹ and 661–3662 kg DM ha-1, respectively. PC ranged from 20.5%–26.6% and was not significantly influenced by row spacing. Total protein yield ranged from 147-884 kg ha-1. Among genotypes, no significant differences were observed in fresh or dry biomass, PC, or protein yield (7477-17,776 kg ha-1, 1122-2199 kg DM ha-1, 21.2%-26.5%, 260-579 kg ha-1, respectively), suggesting that genetic variation among the tested accessions had minimal influence under the given growing conditions. Amino acid analysis confirmed the presence of all essential amino acids, fulfilling over 30% of the recommended daily intake per 100 g DM. These findings highlight YVQ as a promising, sustainable, and protein-rich leafy crop for Mediterranean agriculture. Further research should explore multiharvest potential, mechanical weeding, and optimized agronomic practices for commercial-scale production.

Keywords: amino acid; leaf; Mediterranean; semi-arid; superfood; sustainable

1. Introduction

In the last few years, there has been a notable increase in the demand for protein products. The worldwide protein ingredient market was estimated to be worth US\$38.5 billion in 2020, with an annual projected rise of about 10.5% between 2021 and 2028 [1]. Over the last century, animals (meat, eggs, fish, and dairy) have been the primary source of protein in western countries [2]. However, plant proteins may eventually take a share of the animal protein market due to lower production costs, and consumers' perception of them as healthier, more ethical, and environmentally friendly [3,4]. A well-known but underutilized protein-rich crop is the quinoa plant (*Chenopodium quinoa* Willd., Amaranthaceae) [5]. Originating in the Andean plateau of South America, this pseudo cereal is widely cultivated for grain production intended for human consumption [6,7]. Quinoa is regarded as a climate-resilient crop and has the remarkable ability to grow in harsh and unfavorable conditions [8,9]. Compared to the major cereal crops grown worldwide, such as barley, wheat, corn, and rice, quinoa grains' protein content (PC) is high [10,11]. Its high nutritional value is attributed to high levels of protein (11%–19%) that contain all of the essential amino acids (EAA) [12,13]. Individuals



with celiac disease may find it acceptable because it is gluten-free [14,15]. As a result, quinoa is commonly regarded as a "functional food" or "superfood," enhancing its commercial attractiveness [16]. Thus, quinoa production has increased dramatically in recent years, and the crop is now cultivated commercially in many countries outside of South America [17,18].

Numerous studies have demonstrated that quinoa can be grown outside its native region in various agroecological settings, including semiarid and desert climates [6,18]. Other studies have investigated the influence of agrotechnical factors—such as plant density, row spacing, irrigation, and fertilization—on quinoa grain production [19-22]. Numerous recent studies have demonstrated the versatility of quinoa beyond its use for grain production. For instance, it has been suggested that the straw left over from the grain harvest might be fed to cattle because it is highly digestible and has comparatively high protein and mineral contents [23-25]. Others have indicated the significant potential of using the high nutritional quality and biomass of quinoa hay and silage as a high-quality feed for ruminants [23,26,27]. Young vegetative quinoa (YVQ)—an unconventional leafy green crop—has been recently investigated [28,29]. Similar to spinach, the soft stalks and green leaves of the quinoa plant have high nutritional value and can be eaten either fresh or cooked [30,31]. YVQ can be cultivated year-round in fields, greenhouses, and high tunnels [32]. According to several studies, quinoa leaves contain all of the EAA and have high PC (percentage in dry matter [DM]) that can reach 25%–37%, higher than quinoa grains (9.1%–15.7%) [30,33–35]. Quinoa leaves have a higher PC than spinach, chard (mangold), and broccoli [33]. In addition, quinoa leaves have high quantities of Cu, Mn, and K and moderate salt, Ca, P, and Zn levels [34]. In evaluating the nutraceutical potential of YVQ, Gawlik-Dziki et al. (2013) discovered that quinoa leaf extract includes notable concentrations of several bioactive polyphenols associated with a suppressive effect on the proliferation of prostate cancer cells [36]. Furthermore, YVQ has lower concentrations of saponins—bitter antinutritive metabolites found in quinoa—than the grains [37,38]. YVQ is usually harvested 30-62 days after sowing (DAS) [29,32], whereas detectable quantities of sapogenins—a triterpenoid aglycone that makes up around 50% of the saponin molecule—are first detected 82 DAS [39]. Moreover, quinoa shoots were found to have the highest saponin concentration at 100 DAS and the lowest 60 DAS [40].

A study conducted in Poland showed that YVQ should be harvested when it reaches a height of 20–30 cm. In spring and summer sowing, YVQ fresh weight (FW) biomass averaged 317–1389 and 245–777 g m⁻², respectively [31]. In Egypt, the fresh yield of YVQ harvested 45 DAS was higher, ranging from 2.05–4.14 kg m⁻² [41]. In another study conducted in Israel over the course of 2 years, five quinoa accessions were sown on three winter sowing dates (November, December, and January). When the plants reached 10% DM, they were harvested, and DM yield ranged from 574 to 1982 kg ha⁻¹ [29]. These results suggest the economic potential for cultivated YVQ, with Mediterranean and semiarid regions benefiting from the use of aerial vegetative sections of young quinoa as a novel, sustainable, and high-protein leafy crop.

Since there is little information available about growing YVQ in Mediterranean countries, our goal was to further research the crop's yield and PC to assess its potential as a novel, high-protein leafy crop in Israel. We investigated the impact of different genotypes and row spacings on biomass yield, PC, and protein production of YVQ during summer cultivation. We hypothesized that narrower row spacing would enhance biomass and protein yield due to increased plant density, and that genotypic differences would significantly impact these parameters.

2. Materials and Methods

2.1. Experimental Site

The experiments were conducted in northern Israel (The Gadash research farm, altitude 71 masl, 33°17'95" N lat; 35°58'32" E long) from June 15 to July 27, 2020, and from June 21 to August 3, 2021. The climate is Mediterranean and the rainy season generally lasts from October to May, with a mean annual rainfall of 500–550 mm (Israel Meteorological Service). There was no rainfall in the experimental plots during the experiment. During the experimental period of 2020, air temperature

ranged from 18.2–39.8°C in 2020, and from 17.2–40.2°C in 2021 (Figure 1). The soil at the experimental site is composed of 37% clay, 40% silt, and 23% sand with a pH of 7.6.

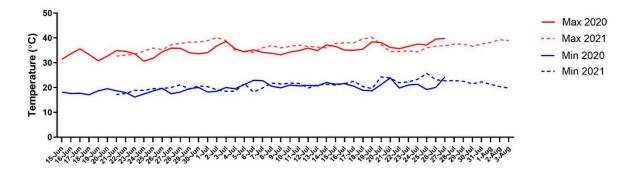


Figure 1. Daily minimum (Min) and maximum (Max) air temperatures at the Gadash research farm during the experiments of 2020 and 2021 (Source: Israel Meteorological Service).

2.2. Plant Material and Experimental Design

Two independent randomized plot experiments were conducted to examine the effect of (a) inter-row spacing and (b) genotype on quinoa parameters. Each experiment was repeated twice, once a year, for two consecutive years. For the inter-row spacing experiment, seeds of accession Mint Vanilla were sown in two, four, or six rows (80, 26, or 16 cm between rows, respectively) at an intrarow density of 50–60 seeds m⁻¹ using a manual seeder. Five quinoa accessions were used for the genotype experiment: 'Red Head', 'Mint Vanilla', 'Ivory', 'Oro de Valle', and 'Peppermint'. These accessions were chosen because they had performed well in previous experiments conducted in the experimental region [20,23,29,42]. Seeds were sown in six rows (16 cm between rows) at an intra-row density of 50–60 seeds m⁻¹ using a manual seeder. Seeds of all accessions were obtained from Wild Garden Seed (Philomath, OR). Quinoa seeds were sown in June 2020 and 2021 in all experiments in four 9.8-m² (5 m long x 1.96 m wide) plots (repeats). The flowerbeds were prepared after cultivation and rolling. The plots were fertilized with 92 kg N ha⁻¹ as urea (46% N). Sprinkler irrigation was applied during the growing period at a rate of 1500 m³ ha⁻¹. All plots were manually weeded once during cultivation.

2.3. Plant Density at Harvest and Yield Measurements

To determine plant density at harvest (plants m⁻²), plants were counted in a representative 1-m² area in each plot. The percentage of DM was determined from 30 DAS: five plants per plot with similar external characteristics were collected every 3 days and weighed before and after drying at 60°C for 48 h, and %DM was calculated. When these plants reached 10%–12% DM, the aerial parts of all plants in each experimental plot were manually harvested ~5 cm above the ground and weighed for fresh matter yield in each plot (42 and 43 DAS in 2020 and 2021, respectively). Representative samples of each plot were collected and weighed before and after drying at 60°C for 48 h to determine plant %DM. Plot DM yield was calculated as fresh matter yield x plant %DM.

2.4. Plant PC, Protein Yield, and Amino Acid Composition

The Kjeldahl method (AOAC, 2019) was used to determine each quinoa sample's N content (AOAC, 2019) at the Milouda and Migal laboratories (Kiryat Shmona, Israel), where PC (% of DM) was also estimated. The conversion factor used to transform N into protein was 6.25. Plot protein yield was calculated as DM yield x PC. The amino acid composition was determined at the Merieux NutriScience laboratories (Resana, Italy). Briefly, the samples were subjected to hot enzymatic hydrolysis of the proteins. The solution obtained from hydrolysis was diluted with deionized water and methanol and analyzed by high-pressure liquid chromatography (HPLC).

2.5. Statistical Analysis

Results were subjected to one-way ANOVA, and the significance between treatments was checked at *P* < 0.05 using JMP software version 11.0.0 (SAS Institute, Cary, NC). In case of significance, Tukey's HSD test was performed to compare all pairs and to examine the differences between treatments. Results were also subjected to a two-tailed Pearson correlation matrix using the 'corrplot' package in RStudio (Boston, MA), in the programming language R.

3. Results

3.1. Effect of Row Spacing on YVQ

3.1.1. Plant Density and DM

Plant density at harvest in 2020 and 2021 for the different row spacings is shown in Table 1. In both years, there were significant differences in plant density at harvest among all inter-row spacings tested. In 2020, DM at harvest ranged between 11.4% and 13.4%, with no significant difference between treatments. In 2021, DM at harvest ranged between 13.3% and 14.4% and was significantly higher in plots sown with 16 cm between rows compared to those sown at 26 and 80 cm between rows.

Table 1. Effect of row spacing on young vegetative quinoa (YVQ) yield parameters and protein content (PC) in the 2020 and 2021 experiments.

Year	Row spacing (cm)	Plant density at harvest (plant m ⁻²)	Dry matter (%)	Fresh yield (kg ha ⁻¹)	Dry yield (kg DM ha ⁻	Protein content (%)	Protein yield (kg ha ⁻¹)
	16	$336 \pm 6 \text{ a}$	13.1 ± 0.6	28469 ± 639 a	3737 ± 243 a	23.9 ± 1.4	$884 \pm 24 \text{ a}$
2020	26	$193 \pm 27 \text{ b}$	13.4 ± 0.7	$27262 \pm 251 a$	$3662 \pm 176 a$	23.5 ± 1.1	$865 \pm 76 a$
	80	$55 \pm 4 c$	11.4 ± 0.2	20986 ± 555 b	$2396 \pm 54 \text{ b}$	26.6 ± 2.2	$633 \pm 39 \text{ b}$
2021	16	$345 \pm 37 \text{ a}$	14.4 ± 0.1 a	7423 ± 209	1072 ± 41	20.5 ± 0.4	220 ± 13
	26	$166 \pm 157 \mathrm{b}$	$13.4 \pm 0.1 \text{ b}$	6747 ± 1609	902 ± 212	23.8 ± 1.3	209 ± 43
	80	$65 \pm 3 \text{ c}$	$13.3 \pm 0.4 b$	4957 ± 775	661 ± 105	22.3 ± 2	147 ± 28

Note: Results are presented as mean value \pm SE of four 9.8-m² sample plots per treatment. DM, dry matter. Values in a column marked with different letters differ significantly by Tukey HSD test (P < 0.05).

3.1.2. Fresh and DM Yield

In 2020, the quinoa fresh yield was significantly higher in plots sown at 26 and 16 cm between rows than in those sown at 80 cm between rows. In 2021, the quinoa fresh yield was generally lower than that in 2020, with no significant difference between treatments (Table 1). In 2020, quinoa DM yield was significantly higher in plots sown at 26 and 16 cm between rows than in those sown at 80 cm between rows. In 2021, the DM yield was generally lower, with no significant difference between treatments (Table 1).

3.1.3. PC and Protein Yield

In plots sown in 2020 and 2021, the PC (% of DM) of YVQ showed no significant difference between treatments (Table 1). In 2020, protein yield was significantly higher in plots sown with 26 and 16 cm between rows compared to plots sown with 80 cm between rows. In 2021, the protein yield of YVQ showed no significant differences between treatments (Table 1).

3.1.4. Correlations Between Traits

Of the 21 correlation coefficients evaluated, 7 exhibited significant values (P < 0.05; Figure 2). Row spacing was negatively correlated to plant density and % DM. No significant correlations were found between this parameter and yield or quality parameters. The opposite trend was found for the correlations with plant density at harvest, which was positively correlated to % DM. The latter was negatively correlated to PC. Fresh yield was positively correlated to DM yield, and both were positively correlated with protein yield.

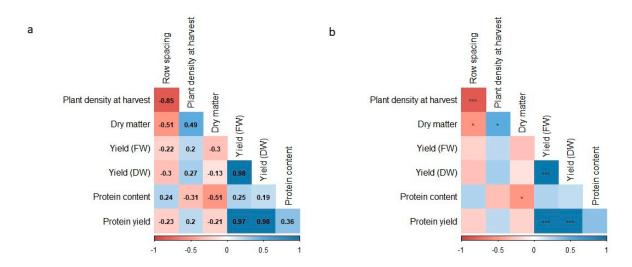


Figure 2. Pearson correlation coefficients among plant density, dry matter, yield (fresh weight [FW] and dry weight [DW]), protein content, and protein yield of YVQ. Blue and red colors represent positive and negative correlations, respectively. (a) Pearson correlation values. (b) Significance of Pearson correlation values (*P < 0.05, ***P < 0.001). Colors reflect correlation values between every two variables. Each value represents measurements from plots for all three row spacings during the 2 years of the experiment.

3.2. Effect of Genotypes on YVQ

In the second experiment, we examined the effect of different genotypes (accessions) on yield parameters and PC of YVQ during summer cultivation.

3.2.1. Plant Density and DM

In 2020, plant density at harvest ranged between 111 and 207 plants m⁻². Accession Oro de Valle plant density was significantly lower than that of all other accessions (Table 2). In 2021, plant density at harvest ranged between 163 and 279 plants m⁻². Accession Red Head plant density was significantly higher than that of all other accessions (Table 2). In the first year, DM at harvest ranged between 12.2% and 13.6% and in the second year, between 14% and 15.1%. There were no significant differences between the different accessions in either year (Table 2).

Table 2. Effect of genotype on YVQ yield parameters and PC in the 2020 and 2021 experiments.

Year	Accession	Plant density at harvest (plant m ⁻²)	Dry matter (%)	Fresh yield (kg ha-1)	Dry yield (kg DM ha ⁻¹)	Protein content (%)	Protein yield (kg ha ⁻¹)
Re	ed Head	176 ± 6 a	12.2 ± 0.6	17776 ± 794	2175 ± 196	24.4 ± 0.8	533 ± 56
M	Iint Vanilla	$183 \pm 16 a$	13.1 ± 0.3	16852 ± 389	2199 ± 54	26.3 ± 1.1	579 ± 25
2020Iv	vory	$207 \pm 14 a$	13.6 ± 0.6	15969 ± 752	2182 ± 173	24.6 ± 2.1	539 ± 69
O	ro de Valle	$111 \pm 4 \text{ b}$	12.3 ± 0.4	14923 ± 1056	1852 ± 182	26.5 ± 1.7	486 ± 40
Pe	eppermint	$186 \pm 6 a$	12.7 ± 0.7	16531 ± 1609	2136 ± 298	25.9 ± 1.4	556 ± 85
R	ed Head	279 ± 7 a	14 ± 0.4	9342 ± 1620	1292 ± 209	22.7 ± 1	290 ± 44

Mint Vanilla	$168 \pm 13 \text{ b}$	14.5 ± 0.3 83	$37 \pm 686 120$	$99 \pm 98 21.9 \pm 0.5$	265 ± 19
2021Ivory	$181 \pm 14 \text{ b}$	14.5 ± 0.4 982	21 ± 1241 141	7 ± 177 21.5 ± 1	300 ± 29
Oro de Valle	$163 \pm 6 \text{ b}$	15.1 ± 0.2 747	77 ± 1387 112	$2 \pm 198 23.4 \pm 0.7$	260 ± 42
Peppermint	$213 \pm 20 \text{ b}$	15.1 ± 0.5 98	$47 \pm 793 149$	$4 \pm 153 21.2 \pm 1.3$	315 ± 33

Note: Results are presented as mean value \pm SE of four 9.8-m² sample plots per treatment. Values in a column marked with different letters differ significantly by Tukey HSD test (P < 0.05).

3.2.2. Fresh and DM Yield

Quinoa fresh yield was generally lower in 2021 than in 2020 (Table 2). There were no significant differences between the different accessions in either year. Similarly, DM yield in 2021 was generally lower than that in 2020, with no significant differences between the different accessions in either year (Table 2).

3.2.3. PC and Protein Yield

In 2020, YVQ PC ranged between 24.4% and 26.5% and in 2021, between 21.2% and 23.4%. There were no significant differences in PC between the accessions in either year (Table 2). YVQ protein yield in 2021 was generally lower than that in 2020, with no significant differences between the different accessions in either year (Table 2).

3.3. Amino acid Composition of YVQ

Accession Mint Vanilla had relatively high protein yield values and is currently the main quinoa accession cultivated in Israel. It was therefore chosen for amino acid composition analysis. The YVQ contained all of the EAA (Table 3). For comparison, Table 3 also shows the recommended daily EAA intake for humans (g per 70 kg body weight). Histidine, isoleucine, leucine, lysine, methionine, phenylalanine (with tyrosine), threonine, tryptophan, and valine levels in the YVQ were 33%, 39%, 37%, 24%, 31%, 58%, 60%, 21%, and 35% of the recommended daily intake, respectively.

Table 3. Amino acid composition of accession Mint Vanilla from the 2021 growing season.

Amino acids		Recommended daily intake
(g 100 g DM ⁻¹)		(g per 70 kg body weight)
Essential		
Histidine	0.23 ± 0.01	0.7
Isoleucine	0.55 ± 0.04	1.4
Leucine	1.01 ± 0.05	2.73
Lysine	0.51 ± 0.04	2.1
Methionine	0.22 ± 0.01	0.7
Phenylalanine ^b	0.65 ± 0.04	1.75a
Threonine	0.63 ± 0.03	1.05
Tryptophan	0.0588 ± 0.0069	0.28
Valine	0.64 ± 0.05	1.82
Non-essential		
Alanine	0.74 ± 0.05	
Arginine	0.64 ± 0.04	
Aspartic acid	1.24 ± 0.06	
Cystine + Cysteine	0.20 ± 0.01	
Glutamic acid	1.54 ± 0.5	
Glycine	0.75 ± 0.03	
Proline	0.63 ± 0.03	
Serine	0.65 ± 0.03	
Tyrosine	0.36 ± 0.04	

4. Discussion

Our previous study demonstrated significant potential for cultivating YVQ in Israel and other Mediterranean and semiarid regions during the winter [29]. In general, YVQ yield is higher in summer vs. winter cultivation, whereas PC tends to be lower in summer than in winter. Ultimately, protein yield is higher during summer cultivation. For example, in this study, the protein yield of accession Mint Vanilla in June 2020 reached 884 kg ha⁻¹ in plots sown with 16-cm row spacing, whereas in December 2020, it was only 359 kg ha⁻¹ with the same spacing (Rubinovich et al., 2023). However, it is important to note that these two experiments were conducted in different locations, which may have influenced the results. In addition, YVQ yield was generally lower in this experiment's second year than in the first. Because both experiments in this study were conducted in the same location, environmental conditions, such as air temperature, may have influenced the outcomes. Notably, maximum temperatures were higher in 2021 than in 2020, exceeding 40°C (Figure 1). Although quinoa is known for its stress tolerance, heat stress can adversely impact the growth and development of quinoa and other plants [25,44–46]. The effects of prolonged heat exposure on YVQ growth and quality therefore warrant further investigation, especially given the increasing frequency of extreme heat events due to global warming [47].

4.1. Effect of Row Spacing on Yield Parameters

Row spacing was significantly negatively correlated with plant density at harvest (Table 1, Figure 2). This was expected, as plots with broader row spacing have fewer seeds sown per unit area than those with narrower row spacing. Similar results were obtained in a previous study that examined the same row spacings in quinoa [20]. Moreover, a similar effect of row spacing on YVQ yield was observed in both years—it was generally, albeit not significantly, negatively correlated with fresh and dry yields (Table 1, Figure 2). However, no significant differences in yield parameters were found between the 16 cm and 26 cm row-spacing treatments in either year. In contrast, in 2020, the yield in the 80 cm row-spacing treatment was significantly lower than in the 16 cm treatment (Table 1). These findings hold practical significance for commercial growers, because reducing the number of seeds per unit area could help lower crop production costs. However, since wider row spacing and lower plant density may lead to thicker quinoa stems [20,48], potentially reducing the plants' palatability, the 16-cm row spacing may be the most advantageous for producing high-quality quinoa for leafy green use. Future studies should explore this aspect further.

Data on the impact of row spacing on YVQ are limited, but a study in India found that quinoa foliage yields generally increase with wider row spacing, peaking at 25 cm compared to 15 or 20 cm spacings [49]. Conversely, studies investigating quinoa grown for grain production have reported higher grain yields with narrower row spacing. For instance, Sief et al. (2015) observed the highest quinoa grain yield at a 20-cm inter-row spacing, outperforming 30 and 40 cm spacings. Another study found that a 30-cm row spacing results in greater quinoa grain yield than 40 or 50 cm spacings [51]. Similar trends have been observed in other plant species, where reduced row spacing leads to increased yields. For example, intermediate wheatgrass hay and straw yields were higher when planted at 15- or 30-cm row spacing compared to 61 cm [52]. Similarly, Hungarian vetch (Vicia pannonica Crantz) demonstrated significantly greater forage yield at 17.5 vs. 35 cm spacing (Albayrak et al., 2011). On the other hand, a study in Brazil reported no significant change in quinoa grain yield with increasing plant density (Spehar and Rocha, 2009); and a study conducted in Denmark noted that quinoa plots with 25- or 12.5-cm row spacing yield less than those with a 50 cm spacing [54]. This discrepancy may be attributed to the fact that in the latter study, only the plots with the highest planting density were manually weeded, reducing competition for resources [54]. Because our experimental results were derived from small plots and primarily relied on manual labor, further research is needed to determine optimal row spacing for YVQ production in commercial fields. Future studies should consider additional agronomic factors, such as mechanical weeding and harvesting methods.

4.2. Effect of Genotype on Yield Parameters

No significant differences were observed in yield or quality parameters among the different accessions grown under the same row spacing (Table 2). These findings align with our previous study, which showed minimal variation in YVQ yield and quality parameters among the same accessions during winter cultivation [29]. In contrast, Pathan et al. (2023) reported significant differences in YVQ yield among three different quinoa accessions, with accession Ames 13724 consistently producing the highest yield of leafy greens over 3 years.

In our study, the absence of a genotype effect could be attributed to the genetic similarity among the selected accessions or the short growth period, which may have limited the expression of genetic differences. Nevertheless, in both years of our experiment, accession Peppermint generally exhibited higher protein yield than most of the other accessions, whereas accession Oro de Valle consistently showed lower yield parameters. Notably, these trends were also observed in winter cultivation, where 'Peppermint' tended to yield more and 'Oro de Valle' less than other accessions [29]. These results suggest that the accession Peppermint holds strong commercial potential under Mediterranean growing conditions. However, the poor performance of 'Oro de Valle' may not necessarily indicate incompatibility with the local environment; rather, it could be due to its low plant density in both years of the study, likely resulting from a low germination rate. Because this study did not assess germination rate, further research is needed to investigate this factor. Our data suggest that under uniform management, all five genotypes are comparably well adapted to Mediterranean summer conditions, and factors such as sowing density and year-to-year climate variations generally influenced yield more than genetic differences. Thus, quinoa's broad adaptability might result in agronomic practices being more critical to success than the choice of cultivar – provided the cultivars are all reasonably well adapted to the region.

'Mint Vanilla' is currently the main quinoa accession cultivated in Israel. It also recorded the highest DM and protein yield in the first year of the experiment, leading to its selection for the amino acid composition analysis. In most cases, 100 g of DM from 'Mint Vanilla' YVQ provided over 30% of the recommended daily intake of EAA for a 70 kg adult [55] (Table 3). This highlights the high nutritional quality of YVQ protein, which is primarily determined by its content of EAA, which humans cannot synthesize [34]. Given that the daily protein requirement for an adult is 0.66 g kg body weight-1 [55], consuming 200 g DM from YVQ could meet the protein and most EAA needs for a 70 kg individual. This has significant economic and social implications, because ensuring adequate intake of high-quality protein remains challenging for certain populations, particularly those who have reduced or eliminated animal proteins in their diets and rely on cereals, grains, and legumes as primary protein sources [10].

Optimizing quinoa cultivation for young vegetative biomass can therefore benefit Mediterranean agriculture by providing a resilient, water-efficient source of high-quality protein for human and potentially animal consumption. This study focused on a single harvest and did not explore the potential for multiple harvests or regrowth, which could further enhance cumulative yield. Moreover, some agronomic practices (e.g., fertilizer rate, irrigation volume) were held constant, limiting our ability to generalize the results to other management systems. Thus, YVQ could be effectively integrated into sustainable cropping systems by further experimenting and refining agronomic practices, and addressing climate change and resource-scarcity challenges.

5. Conclusions

The present study demonstrated that inter-row spacing affects YVQ biomass and protein yield, whereas genotypic differences among the tested accessions were relatively minor. YVQ grown in the Mediterranean summer yielded substantial biomass with high PC despite the high temperatures and limited irrigation. Our results highlight YVQ's potential as a sustainable, protein-rich crop for Mediterranean climates characterized by hot and dry summers. These conditions, typical of many Mediterranean regions, underscore YVQ's potential to contribute to food security under climate

change. The consistently high PC and quality observed in the YVQ further emphasize its value as a high-nutrition crop in regions aiming to improve local protein sources.

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Abbreviations

The following abbreviations are used in this manuscript:

DAS days after sowing

DM dry matter FW fresh weight

EAA essential amino acids PC protein content

YVQ young vegetative quinoa

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