

Article

Biomolecules with Antioxidant Capacity from the Seeds and Sprouts of 20 Varieties of *Chenopodium quinoa* Willd. (Quinoa)

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Abstract: Quinoa has acquired a great interest due to its high content of nutrients and biomolecules that have nutritional and medicinal properties. The aim of this study was to compare the total phenolic content (TPC), total flavonoids (TF), and the antioxidant capacity of 20 varieties of seeds and sprouts of quinoa extract. Quinoa seeds were germinated for 72 h and dried in an oven at 45 °C. The extracts were obtained by dynamic extraction using methanol. Phytochemical analysis with liquid chromatography coupled with mass spectrometry (LC-ESI-MS/MS), TPC, TF, and the antioxidant capacity was carried out and compared between both extracts. The TPC was determined with Folin-Ciocalteu reagent, TF with AlCl₃, and the antioxidant capacity was determined according to the DPPH and ABTS assays. Sprout extracts showed high values of TPC (31.28 ± 0.42 mg GAE/g; Pasankalla variety), TF (14.31 ± 0.50 mg EQ/g; black Coito variety), and antioxidant capacity (IC₅₀(DPPH): 12.69 ± 0.29 µg/mL and IC₅₀(ABTS): 3.51 ± 0.04 µg/mL; Pasankalla). The extracts of the Pasankalla variety revealed 93 and 90 phytochemical constituents in the seeds and sprouts, respectively, such as amino acids, phenolic acids, flavonoids, fatty acids, and triterpene saponins, among others. Quinoa sprouts showed a high content of TPC and TF, and high antioxidant capacity compared with seed extracts, especially the Pasankalla variety.

Keywords: Amaranthaceae; free radical; superfoods; phytochemical analysis; flavonoids; phenols; amino acids



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1. Introduction

Quinoa (*Chenopodium quinoa* Willd.) is a pseudocereal belonging to the Amaranthaceae family that is native to the Andean region in South America [1]. Peru is the leading quinoa-exporting country, exporting quinoa with a value of \$98.5 million dollars, followed by Bolivia, the Netherlands, the United States, Spain, Germany, Canada, France, Ecuador, and Belgium [2]. Quinoa seeds are known to have a high protein content ranging from 11% to 19%. The seeds are a source of amino acids (isoleucine, leucine, lysine, methionine, phenylalanine, threonine, tryptophan, valine, histidine, cysteine, tyrosine, glycine, arginine, proline, serine, glutamine, alanine, and aspartic acid), carbohydrates (49% to 68% dry weight), fat (2% to 9.5%), vitamins (thiamine, riboflavin, folic acid, and niacin), and minerals such as iron, zinc, magnesium, and copper (2.4% to 4.8%) [3]. Additionally, some phytochemical constituents such as saponins, phenolic compounds (ferulic, sinapinic and gallic acids, kaempferol, isorhamnetin, and rutin) [4], and peptides with therapeutic activity have been determined, making this crop very attractive for a wide range of food

products [5]. Quinoa has been traditionally used in tortillas, pasta, flour, cookies, bread, and soups, among others, and is considered to be a gluten-free superfood and a source of fiber dietary [6]. Thus, quinoa is considered to be an acceptable food worldwide and is highly recommended for vegetarians.

On the other hand, sprouts are obtained by germinating the seeds and provide multiple nutritional and therapeutic benefits to those who consume them in different ways, due to the increase in the availability of nutrients such as fatty acids and carbohydrates, as well as polyphenols and flavonoids, during the germination process, which improves their antioxidant capacity [7]. These changes are due to a multitude of biochemical processes, which generate alterations in the composition of primary and secondary metabolites, producing an intrinsic change in the phenolic compounds and antioxidant activity [8]. Sprouts can improve the nutritional quality of a grain by eliminating or inactivating some antinutritional factors and increasing the digestibility of proteins and starches [9]. During germination, the original composition of the seed changes: the nitrogen-containing proteins move towards smaller protein fractions, oligopeptides, and free amino acids (some increase; others decrease or are not altered). Consequently, the changes increase the biological protein value of the sprouts, and digestibility is higher than in seeds [10].

Studies have reported that quinoa sprouts have high levels of amino acids, peptides, vitamins, and minerals but also include antinutritional components such as tannin, lectin, trypsin inhibitor, and galactoside, although at lower values than in non-germinated seeds [11]. The main enzyme involved in the early phase during the sprouting of quinoa seeds seems to be α -amylase, which leads to the generation of new compounds [12]. Some biological studies in quinoa sprouts have reported hepatoprotective, antioxidant [13], and anti- α -amylase effects in vitro [14], and hypoglycemic effects in diabetic rats [15]. Currently, there are no studies on the antioxidant activity of a wide variety of quinoa sprouts grown in Peru. Thus, as the germination process is a strategy for obtaining sprouts and improving the antioxidant capacity, total phenols, and flavonoids, thereby increasing its nutraceutical value, the main aim of this study was to compare the total phenolic content, flavonoids, and antioxidant capacity of the seeds and sprouts of 20 varieties of quinoa and analyze the phytochemical constituents of varieties with major antioxidant capacity using liquid chromatography-mass spectrometry (LC-ESI-MS/MS). To carry out this study, four phases were developed:

- (a) Germinating 20 varieties of quinoa seeds under laboratory conditions and extracting their phytochemical constituents by maceration with methanol.
- (b) Determining the total phenolic content (TPC) and total flavonoids (TF) of the seeds and sprouts of quinoa.
- (c) Evaluating the antioxidant capacity of the seeds and sprouts of quinoa using the 2,2-diphenyl-1-picrylhydrazyl (DPPH) and 2,2'-azinobis-(3-ethylbenzothiazoline)-6-sulfonic acid (ABTS) methods.
- (d) Analyzing the phytochemical constituents of the seeds and sprouts of the quinoa variety with the best results obtained regarding the antioxidant capacity with liquid chromatography-mass spectrometry (LC-ESI-MS/MS).

2. Results

2.1. Germination Process

Sprouts were obtained in a time of 72 h, and measured between 1.7 and 2.3 cm in length for all varieties. However, the red variety achieved the greatest length among all varieties (2.1–2.3 cm). The other varieties had lengths as follows: White Junín Ayacucho, 1.7–1.9 cm; T-256, 1.8–1.9 cm; Pasankalla, 1.7–1.8 cm; Suano Puno, 1.7–1.9 cm; T-38, 1.8–2.0 cm; Yellow Sacaca, 1.9–2.0 cm; T-45, 1.7–1.9 cm; Santa Ana, 1.7–1.8 cm; T-61 Pomata, 1.8–1.9 cm; CQA-048, 1.8–2.0 cm; Black Collana, 1.7–1.9 cm; T-72 Huancayo, 1.8–1.9 cm; CQA-043, 1.8–1.9 cm; Salcedo, 1.8–2.0 cm; Ayacucho Compuesto, 1.7–1.9 cm; White Choclito, 1.7–1.9 cm; Yellow Maranganí, 1.9–2.1 cm; Black Coito, 1.7–1.9 cm; and Black, 1.8–2.0 cm. Figure 1 shows the

20 varieties of quinoa germinated under standard laboratory conditions of temperature, humidity, and time.

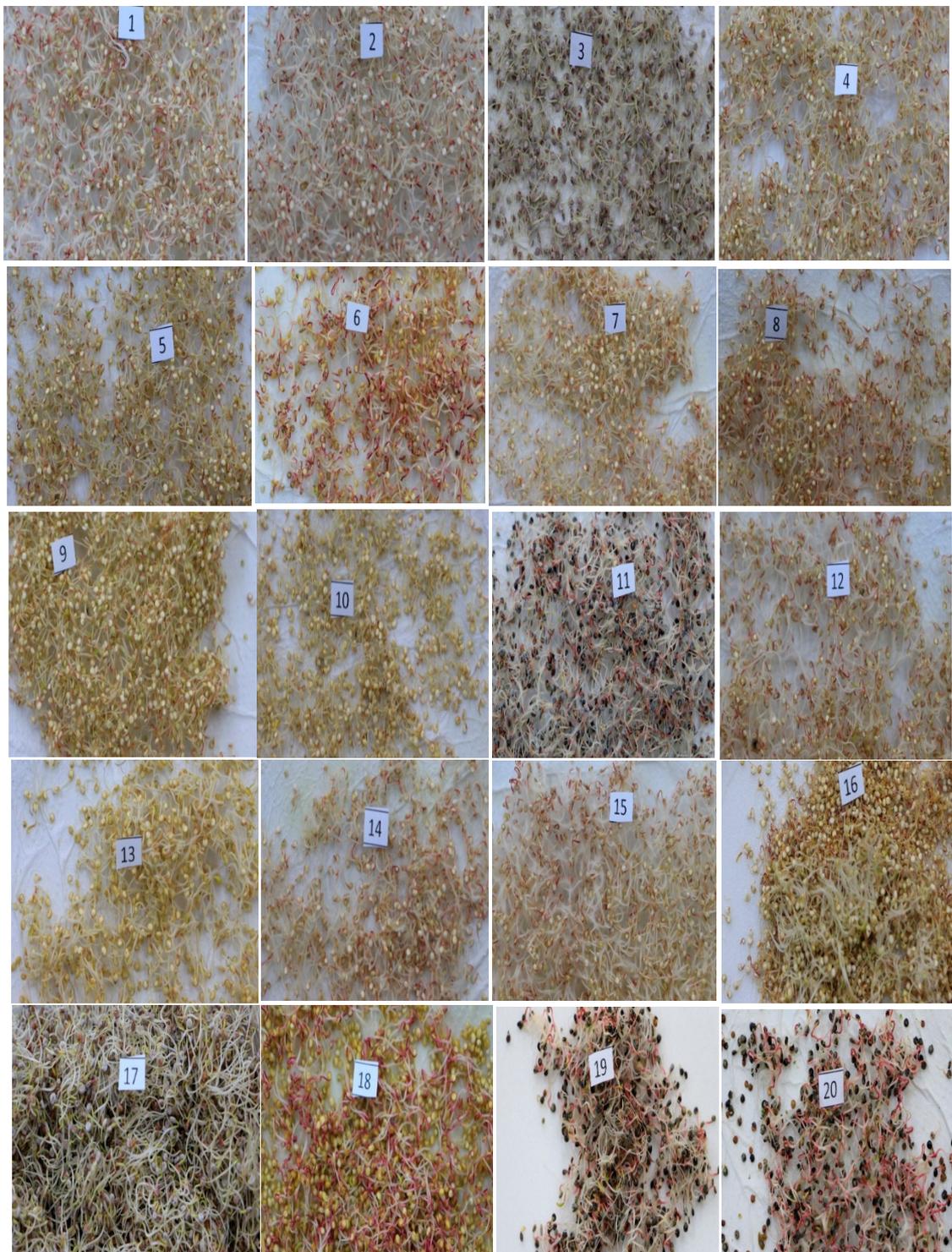


Figure 1. Twenty varieties of quinoa sprouts. (1), White Junín Ayacucho; (2), T-256; (3), Pasankalla; (4), Suano Puno; (5), T-38; (6), Yellow Sacaca; (7), T-45; (8), Santa Ana; (9), T-61 Pomata; (10), CQA-048; (11), Black Collana; (12), T-72 Huancayo; (13), CQA-043; (14), Salcedo; (15), Ayacucho Compuesto; (16), White Choclito; (17), Red; (18), Yellow Marangani; (19), Black Coito; (20), Black.

2.2. Total Phenolic Content

The TPC of sprouts was found to range from 19.15 ± 1.54 to 31.28 ± 0.42 mg GAE/g of methanolic extract, being highest in the Pasankalla variety, CQA-048, Black Collana, and Black Coito. On the other hand, in quinoa seed extracts, the variation was from 11.72 ± 0.32 to 28.32 ± 0.49 , being greater in the Pasankalla, Black Collana, and Black Coito varieties (Table 1). There was a significant difference between sprout and seed extracts for TPC, (paired sample t-test; $p < 0.05$), with TPC being higher in sprout extracts than in seed extracts, with an average of 24.57 ± 3.49 mg GAE/g in sprout extracts and 20.12 ± 4.37 mg GAE/g in seed extracts.

Table 1. Total phenolic content (TPC) and total flavonoids (TF) in the sprouts and seeds of 20 varieties of quinoa.

Variety	TPC mg EAG/g ME		TF mg EQ/g ME	
	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD
1. White Junín Ayacucho	23.32 ± 1.63	20.95 ± 0.79	11.52 ± 0.26	$8.77 \pm 0.26^*$
2. T-256	24.78 ± 0.21	$13.82 \pm 1.04^*$	11.23 ± 0.19	10.23 ± 0.95
3. Pasankalla	31.28 ± 0.42	$28.32 \pm 0.49^*$	13.48 ± 0.38	$11.52 \pm 0.92^*$
4. Suano Puno	19.62 ± 0.42	$17.25 \pm 0.66^*$	8.60 ± 0.48	8.56 ± 0.38
5. T-38	21.05 ± 0.40	21.75 ± 1.25	10.06 ± 0.57	9.81 ± 0.25
6. Yellow Sacaca	24.22 ± 0.31	23.58 ± 0.61	11.19 ± 0.38	$8.23 \pm 0.29^*$
7. T-45	21.02 ± 0.15	19.38 ± 2.06	11.06 ± 0.21	$8.39 \pm 0.38^*$
8. Santa Ana	23.02 ± 0.74	$18.23 \pm 1.01^*$	9.94 ± 0.63	$7.06 \pm 0.33^*$
9. T-61 Pomata	21.12 ± 1.50	$15.55 \pm 0.20^*$	10.94 ± 0.33	$8.73 \pm 0.31^*$
10. CQA-048	28.82 ± 0.67	$21.32 \pm 0.72^*$	7.44 ± 0.50	$6.23 \pm 0.26^*$
11. Black Collana	28.58 ± 1.21	$26.98 \pm 0.25^*$	13.44 ± 0.58	$8.73 \pm 0.14^*$
12. T-72 Huancayo	19.15 ± 1.54	18.58 ± 0.65	12.35 ± 0.48	$9.81 \pm 0.45^*$
13. CQA-043	26.05 ± 0.17	$11.72 \pm 0.32^*$	12.15 ± 0.08	11.31 ± 0.50
14. Salcedo	20.98 ± 1.99	$12.38 \pm 0.61^*$	11.94 ± 0.13	$9.81 \pm 0.45^*$
15. Ayacucho Compuesto	28.05 ± 0.53	$21.42 \pm 1.17^*$	11.19 ± 0.25	10.98 ± 0.40
16. White Choclito	24.02 ± 0.78	20.78 ± 1.86	11.52 ± 0.31	$9.90 \pm 0.26^*$
17. Red	26.05 ± 0.36	$20.45 \pm 0.44^*$	12.31 ± 0.50	$10.52 \pm 0.19^*$

Table 1. Cont.

Variety	TPC mg EAG/g ME		TF mg EQ/g ME	
	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD
18. Yellow Maranganí	27.98 \pm 0.70	22.82 \pm 1.12 *	13.52 \pm 0.44	10.98 \pm 0.52 *
19. Black Coito	28.18 \pm 0.35	24.42 \pm 0.75 *	14.31 \pm 0.50	9.94 \pm 0.13 *
20. Black	24.12 \pm 0.64	20.78 \pm 0.35 *	12.31 \pm 0.45	9.73 \pm 0.38 *
Total Average \pm SD	24.57 \pm 3.49	20.12 \pm 4.37 *	11.52 \pm 1.67	9.46 \pm 1.40 *

* $p < 0.05$ (paired sample *t*-test); mg GAE/g ME: mg equivalent to gallic acid per g of methanolic extract.; mg EQ/g ME: mg equivalent to quercetin per g of methanolic extract.

2.3. Total Flavonoids

In sprouts, the flavonoid content varied from 7.44 \pm 0.50 to 14.31 \pm 0.5 mg EQ/g of extract, being highest in the Black Coito, Yellow Maranganí, Pasankalla, and Black Collana varieties. In seed extract, the variation was from 6.23 \pm 0.26 to 11.52 \pm 0.92 EQ/g. The results showed a significant increase in the total flavonoids in sprouts compared with seed extracts, but in some varieties, the increase was not significant, such as in T-256, Suano Puno, T-38, CQA-043, and Ayacucho Compuesto (Table 1).

2.4. The Antioxidant Capacity Equivalent to Trolox

Table 2 shows the antioxidant capacity equivalent to Trolox (TEAC) of sprout and seed extracts, with the variation ranging, respectively, from 25.90 to 37.65 and from 25.03 to 29.60 μ mol ET/mg of extract for the radical DPPH and from 57.05 to 90.84 and from 45.80 to 67.04 μ mol ET/mg of extract for the radical ABTS. The sprouts with the highest antioxidant capacity for the radical DPPH were Pasankalla, White Junín Ayacucho, Yellow Sacaca, Black Collana, and Red; while for the radical ABTS, those with the highest antioxidant capacity were Black Collana, Black, Pasankalla, Suano Puno, Yellow Maranganí, Red, and Black Coito. Furthermore, it was found that in most of the varieties, significant differences appeared (Student's *t*-test; $p < 0.05$), with antioxidant capacity being greater in sprouts than in seed extracts.

Table 2. Antioxidant capacity equivalent to Trolox (TEAC) of the radical DPPH and ABTS of methanolic extracts of sprouts and seeds of 20 varieties of quinoa.

Variety	TEAC-DPPH μ mol TE/mg MS		TEAC-ABTS μ mol TE/mg MS	
	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD	Quinoa Sprouts Mean \pm SD	Quinoa Seeds Mean \pm SD
1. White Junín Ayacucho	31.26 \pm 0.56	28.47 \pm 1.44	64.78 \pm 1.63	54.68 \pm 0.48 *
2. T-256	28.38 \pm 0.27	25.88 \pm 0.72 *	62.84 \pm 1.65	61.95 \pm 0.96
3. Pasankalla	37.65 \pm 0.88	29.60 \pm 0.54 *	78.79 \pm 0.86	54.19 \pm 0.41 *
4. Suano Puno	25.90 \pm 0.36	25.24 \pm 0.22	78.66 \pm 2.02	53.10 \pm 1.03 *
5. T-38	27.67 \pm 0.30	25.03 \pm 0.18 *	59.96 \pm 5.62	48.88 \pm 1.52 *

Table 2. Cont.

Variety	TEAC-DPPH μmol TE/mg MS		TEAC-ABTS μmol TE/mg MS	
	Quinoa Sprouts Mean ± SD	Quinoa Seeds Mean ± SD	Quinoa Sprouts Mean ± SD	Quinoa Seeds Mean ± SD
6. Yellow Sacaca	30.54 ± 1.17	29.37 ± 0.82	63.21 ± 0.60	58.38 ± 2.14 *
7. T-45	25.94 ± 0.29	25.54 ± 0.17 *	60.55 ± 3.46	45.80 ± 0.37 *
8. Santa Ana	28.08 ± 0.07	26.93 ± 0.23 *	60.74 ± 1.06	53.66 ± 0.55 *
9. T-61 Pomata	25.92 ± 0.15	25.77 ± 0.15 *	65.40 ± 0.96	50.56 ± 2.29 *
10. CQA-048	26.32 ± 0.12	25.64 ± 0.17	62.29 ± 2.59	64.91 ± 5.06
11. Black Collana	29.26 ± 0.40	25.90 ± 0.15 *	90.84 ± 2.22	60.56 ± 4.28 *
12. T-72 Huancayo	26.97 ± 0.40	25.75 ± 0.20	68.67 ± 0.64	59.40 ± 0.09 *
13. CQA-043	26.17 ± 0.55	25.89 ± 0.23	57.05 ± 2.62	56.35 ± 0.34
14. Salcedo	26.21 ± 0.24	25.96 ± 0.23 *	64.95 ± 0.83	58.21 ± 0.19 *
15. Ayacucho Compuesto	26.45 ± 0.27	26.26 ± 0.20	68.91 ± 0.61	65.74 ± 0.25 *
16. White Choclito	27.30 ± 0.23	26.88 ± 0.31 *	58.84 ± 2.73	57.92 ± 0.75
17. Red	28.60 ± 0.20	26.93 ± 0.36 *	75.79 ± 1.26	67.04 ± 0.79 *
18. Yellow Marangani	27.51 ± 0.29	26.20 ± 0.12 *	78.11 ± 1.69	63.76 ± 0.70 *
19. Black Coito	28.04 ± 0.10	26.56 ± 0.16 *	69.41 ± 0.87	63.68 ± 0.93 *
20. Black	27.67 ± 0.25	26.09 ± 0.06 *	78.79 ± 2.36	56.43 ± 0.52 *
Total Average ± SD	28.09 ± 2.68	26.50 ± 1.30 *	68.43 ± 8.96	57.71 ± 5.83 *

* $p < 0.05$; paired sample t -test.

2.5. The Half Inhibitory Concentration (IC_{50}) of the Methanolic Extracts of Sprouts and Seeds of Quinoa

The half inhibitory concentration (IC_{50}) (Table 3) represents the reduction to 50% of the initial absorbance of the DPPH and ABTS radicals, with the average variation for all varieties ranging from 12.69 to 18.45 mg/mL in sprout extracts and from 16.15 to 19.09 mg/mL in seed extracts using the DPPH assay. In the ABTS assay, the results ranged from 3.05 to 4.71 and from 4.13 to 6.04 mg/mL in sprout and seed extracts, respectively. There was a significant difference ($p < 0.05$) in the IC_{50} of the radicals DPPH and ABTS between the sprout and seed extracts, being lower in sprouts than in seed extracts.

Table 3. Half inhibitory concentration (IC₅₀) of the radicals DPPH and ABTS of methanolic extracts of sprouts and seeds of 20 varieties of quinoa.

Variety	IC ₅₀ (mg/mL)			
	DPPH		ABTS	
	Quinoa Sprouts Mean ± SD	Quinoa Seeds Mean ± SD	Quinoa Sprouts Mean ± SD	Quinoa Seeds Mean ± SD
1. White Junín Ayacucho	15.29 ± 0.27	16.81 ± 0.83	4.27 ± 0.11	5.06 ± 0.05 *
2. T-256	16.84 ± 0.16	18.47 ± 0.52 *	4.40 ± 0.12	4.47 ± 0.07
3. Pasankalla	12.69 ± 0.29	16.15 ± 0.30 *	3.51 ± 0.04	5.11 ± 0.04 *
4. Suano Puno	18.45 ± 0.26	18.93 ± 0.16	3.52 ± 0.10	5.21 ± 0.10 *
5. T-38	17.27 ± 0.19	19.10 ± 0.14 *	4.64 ± 0.45	5.66 ± 0.18 *
6. Yellow Sacaca	15.66 ± 0.60	16.28 ± 0.46	4.38 ± 0.04	4.74 ± 0.17 *
7. T-45	18.42 ± 0.20	18.71 ± 0.12 *	4.58 ± 0.26	6.04 ± 0.05 *
8. Santa Ana	17.02 ± 0.04	17.75 ± 0.14 *	4.55 ± 0.08	5.16 ± 0.06 *
9. T-61 Pomata	18.43 ± 0.08	18.54 ± 0.11 *	4.23 ± 0.06	5.48 ± 0.25 *
10. CQA-048	18.15 ± 0.09	18.63 ± 0.12	4.45 ± 0.18	4.28 ± 0.32
11. Black Collana	16.33 ± 0.22	18.45 ± 0.10 *	3.05 ± 0.08	4.58 ± 0.32 *
12. T-72 Huancayo	17.72 ± 0.27	18.55 ± 0.14	4.03 ± 0.04	4.66 ± 0.01 *
13. CQA-043	18.26 ± 0.39	18.46 ± 0.17	4.86 ± 0.22	4.91 ± 0.03
14. Salcedo	18.23 ± 0.17	18.40 ± 0.16 *	4.26 ± 0.06	4.75 ± 0.02 *
15. Ayacucho Compuesto	18.07 ± 0.18	18.20 ± 0.14	4.01 ± 0.04	4.21 ± 0.02 *
16. White Choclito	17.51 ± 0.15	17.34 ± 0.20	4.71 ± 0.21	4.78 ± 0.06
17. Red	16.71 ± 0.12	17.75 ± 0.23 *	3.95 ± 0.29	4.13 ± 0.05
18. Yellow Maranganí	17.37 ± 0.19	18.24 ± 0.09 *	3.54 ± 0.07	4.34 ± 0.05 *
19. Black Coito	17.04 ± 0.06	17.99 ± 0.11 *	3.99 ± 0.05	4.41 ± 0.06 *
20. Black	17.27 ± 0.15	18.31 ± 0.04 *	3.51 ± 0.11	4.90 ± 0.05
Total Average ± SD	17.14 ± 1.37	18.05 ± 0.84 *	4.12 ± 0.50	4.84 ± 0.51

* $p < 0.05$; paired sample t -test.

Table 4 shows positive correlations between antioxidant capacity and both TPC and total flavonoids, and a negative correlation with IC₅₀, with a significant difference in both cases ($p < 0.01$). This correlation indicates that while the concentration of TPC and TF increased in sprout extracts, their antioxidant capacity also increased and, inversely, as TPC and TF became higher, the IC₅₀ reduced.

Table 4. Pearson’s correlation coefficients among total phenols, total flavonoids, antioxidant capacity (TEAC), and the half inhibitory concentration (IC₅₀) of the radicals DPPH and ABTS in sprouts and seeds of quinoa.

Correlations		TEAC-DPPH	TEAC-ABTS	IC ₅₀ DPPH	IC ₅₀ ABTS
TPC of quinoa seeds	Pearson’s correlation	0.480 **	0.352 **	−0.477 **	−0.331 **
	<i>p</i> -value	<0.0001	0.006	<0.0001	0.010
TF of quinoa sprouts	Pearson’s correlation	0.372 **	0.407 **	−0.393 **	−0.404 **
	<i>p</i> -value	0.003	0.001	0.002	0.001
TPC of quinoa sprouts	Pearson’s correlation	0.436 **	0.106	−0.433 **	−0.087
	<i>p</i> -value	<0.0001	0.421	0.001	0.508
TF of quinoa seeds	Pearson’s correlation	0.092	0.202	−0.098	−0.214
	<i>p</i> -value	0.483	0.121	0.455	0.100

** The correlation is significant at the 0.01 level (bilateral).

2.6. Phytochemical Analysis of Methanolic Extracts of Sprouts and Seeds of *C. quinoa* (Pasankalla Variety)

Phytochemical analysis was carried out by LC-ESI-MS/MS for the Pasankalla variety due to its high TPC and TF values and antioxidant capacity, as shown in Tables 1–3. Our results indicated that the sprout extract had 90 phytochemical constituents, of which 45 were observed in ESI (−), 33 in ESI (+), and 12 in both modes. In the seed extract, 93 compounds were determined, of which 58 metabolites were observed in ESI (−), 28 in ESI (+), and 7 in both modes, as presented in Table 5. Figure 2 shows the ESI-positive and -negative chromatographic profiles for both sprouts and seeds of the Pasankalla variety.

Table 5. Number of annotated metabolites (via MS and MS/MS) in each extract according to the ESI (−) and ESI (+) ionization modes.

C. quinoa (Pasankalla Variety)	ESI (−)	ESI (+)	ESI (+/−)	Total
Seeds	58	28	7	93
Sprouts	45	33	12	90

The retention times (Rt), adductions, experimental, and theoretical m/z values, ppm error, MS/MS spectrum (m/z : absolute intensity), SMILES (simplified molecular input line entry system) string, InChIKey (IUPAC international chemical identifier), and tentative compounds are available in the Supplementary Table S1 and Supplementary Table S2.

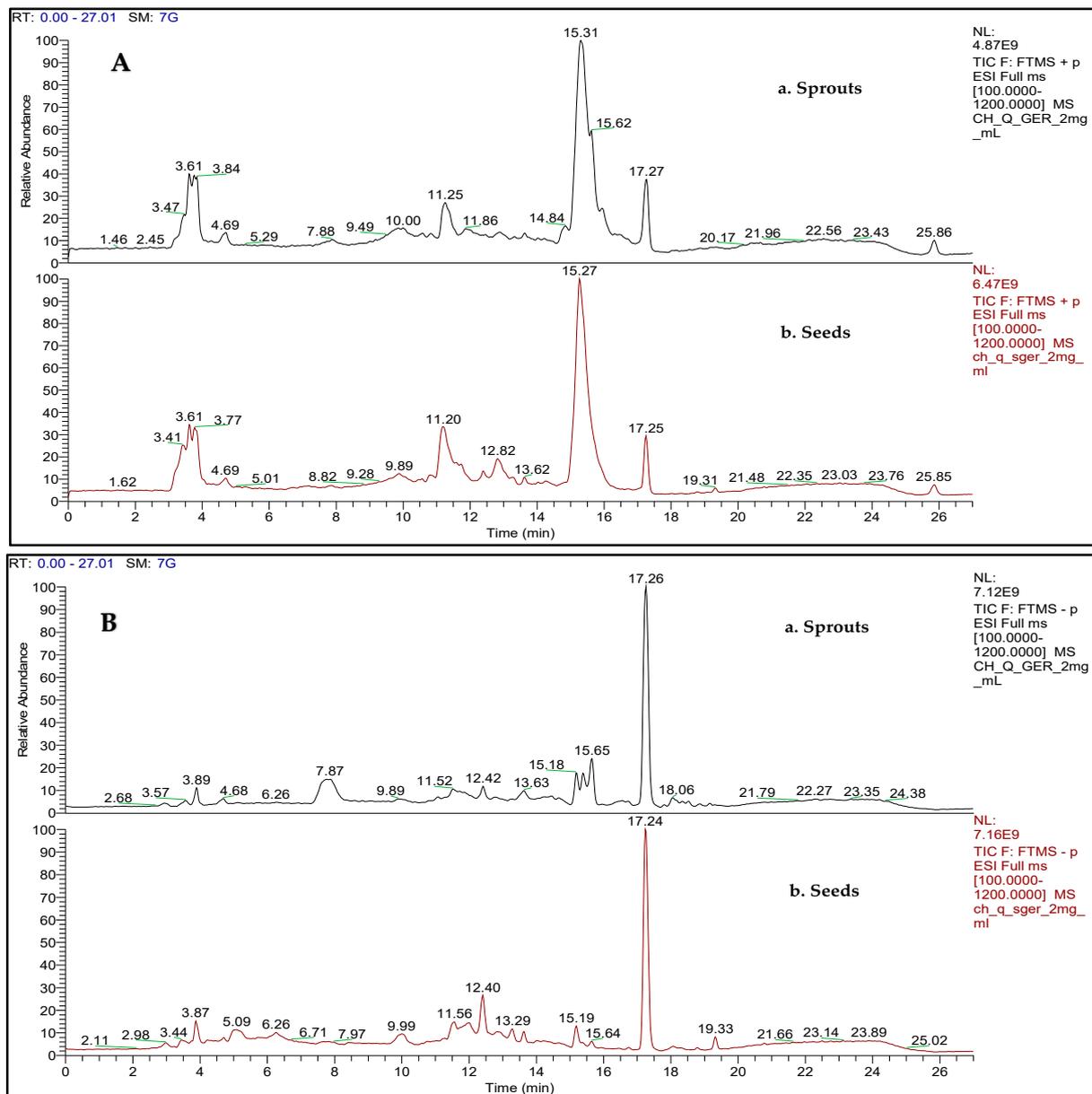


Figure 2. Chromatographic profile (LC-MS) of *C. quinoa* extracts (Pasankalla variety): **(A)**: ESI (+) ionization mode; **(B)**: ESI (−) ionization mode.

The phytochemical constituents determined in the extracts of Pasankalla sprouts (Table 6) were classified as (i) primary metabolites, such as amino acids and derivatives ($n = 23$), organic acids ($n = 14$), monosaccharide sugar acids and sugar alcohols ($n = 8$), disaccharides and oligosaccharides ($n = 7$), lipids ($n = 8$), and nucleobases/nucleosides ($n = 5$); and (ii) secondary metabolites, such as phenolic acids ($n = 2$), triterpenoids ($n = 4$), O-glycosyl compounds ($n = 4$), phenolic glycosides ($n = 2$), flavonoid-O-glycosides ($n = 2$), alkaloids and derivatives ($n = 1$), triterpene saponins ($n = 4$), coumarins ($n = 1$), and other compounds ($n = 13$).

Table 6. Phytochemical constituents of quinoa sprouts (Pasankalla variety) determined by LC-ESI-MS/MS.

#	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
1	3.85	340.1885884	C ₁₈ H ₂₈ O ₆	[5-acetyloxy-3-(hydroxymethyl)-2-oxo-6-propan-2-ylcyclohex-3-en-1-yl] 3-methylpentanoate	Menthane monoterpenoids
2	3.91	130.0266086	C ₅ H ₆ O ₄	Citraconic acid	Organic acids
3	4.30	313.131408	C ₁₈ H ₁₉ NO ₄	Feruloyl tyramine	Ferulic acid and derivatives
4	4.40	138.031694	C ₇ H ₆ O ₃	Salicylic acid	Salicylic acids
5	4.64	146.0579087	C ₆ H ₁₀ O ₄	2-Methylglutaric acid	Methyl-branched fatty acids
6	4.69	311.1157579	C ₁₈ H ₁₇ NO ₄	Feruloyl dehydrotyramine	Ferulic acid and derivatives
7	4.75	132.0422586	C ₅ H ₈ O ₄	Glutaric acid	Dicarboxylic acids and derivatives
8	5.12	118.0266086	C ₄ H ₆ O ₄	Succinic acid (Isomer I)	Dicarboxylic acids and derivatives
9	5.35	173.1051933	C ₈ H ₁₅ NO ₃	n-Acetyl-L-leucine (Isomer I)	Leucine and derivatives
10	5.77	118.0266086	C ₄ H ₆ O ₄	Succinic acid (Isomer II)	Dicarboxylic acids and derivatives
11	5.79	162.0528233	C ₆ H ₁₀ O ₅	β-hydroxy-β-methylglutaric acid (Isomer I)	Hydroxy fatty acids
12	5.87	123.0320284	C ₆ H ₅ NO ₂	Isonicotinic acid	Pyridinecarboxylic acids
13	6.03	154.0266086	C ₇ H ₆ O ₄	2,3-Dihydroxybenzoic acid	Salicylic acids
14	6.24	173.1051933	C ₈ H ₁₅ NO ₃	n-Acetyl-L-leucine (Isomer II)	Leucine and derivatives
15	6.31	162.0528233	C ₆ H ₁₀ O ₅	β-hydroxy-β-methylglutaric acid (Isomer II)	Hydroxy fatty acids
16	6.68	219.1106725	C ₉ H ₁₇ NO ₅	Pantothenic acid (Isomer I)	Vitamin B5
	6.69	219.1106725	C ₉ H ₁₇ NO ₅		
17	7.09	219.1106725	C ₉ H ₁₇ NO ₅	Pantothenic acid (Isomer II)	Vitamin B5
	7.12	219.1106725	C ₉ H ₁₇ NO ₅		
18	7.84	298.1568945	C ₁₉ H ₂₂ O ₃	Auraptin	Coumarins
19	7.86	129.042593	C ₅ H ₇ NO ₃	L-Pyroglutamic acid	Alpha amino acids and derivatives
20	10.07	480.3087035	C ₂₇ H ₄₄ O ₇	NCGC00168839-02![(2S,3R,5R,10R,13R,14S,17S)-2,3,14-trihydroxy-10,13-dimethyl-17-[(2R,3R)-2,3,6-trihydroxy-6-methylheptan-2-yl]-2,3,4,5,9,11,12,15,16,17-decahydro-1H-cyclopenta[a]phenanthren-6-one Syn. Phytoecdysteroids	Phytoecdysteroids
	10.08	480.3087035	C ₂₇ H ₄₄ O ₇		
21	10.52	648.3873477	C ₃₆ H ₅₆ O ₁₀	(2S,3S,4S,5R,6R)-6-[[[(3S,6aR,6bS,8aS,14bR)-8a-carboxy-4-(hydroxymethyl)-4,6a,6b,11,11,14b-hexamethyl-1,2,3,4a,5,6,7,8,9,10,12,12a,14,14a-tetradecahydricen-3-yl]oxy]-3,4,5-trihydroxyoxane-2-carboxylic acid Syn. NCGC00381031-01_C36H56O10_Olean-12-en-28-oic acid, 3-(beta-D-glucopyranuronosyloxy)-23-hydroxy-, (3beta,5xi,9xi,18xi)-	Triterpene saponins
22	10.91	372.1420321	C ₁₇ H ₂₄ O ₉	Syringin	Phenolic glycosides
23	11.06	356.110732	C ₁₆ H ₂₀ O ₉	NCGC00180844-02!(E)-3-[4-methoxy-2-[(2S,3R,4S,5S,6R)-3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxyphenyl]prop-2-enoic acid Syn. 2-O-Glucosyloxy-4-methoxycinnamic acid	Phenolic glycosides
24	11.50	244.069536	C ₉ H ₁₂ N ₂ O ₆	Uridine	Pyrimidine nucleosides
25	12.30	810.440171	C ₄₂ H ₆₆ O ₁₅	NCGC00347541-02_C42H66O15_beta-D-Glucopyranose, 1-O-[(3beta,5xi,9xi,18xi)-3-(beta-D-glucopyranuronosyloxy)-29-hydroxy-28-oxoolean-12-en-28-yl]-	Triterpene saponins
26	12.37	477.285539	C ₂₃ H ₄₄ NO ₇ P	Lysophosphatidylethanolamine LPE 18:2	Lipids
27	12.42	453.285539	C ₂₁ H ₄₄ NO ₇ P	Lysophosphatidylethanolamine LPE 16:0	Lipids
	12.43	453.285539	C ₂₁ H ₄₄ NO ₇ P		
28	12.44	152.0334253	C ₅ H ₄ N ₄ O ₂	Xanthine (Isomer I)	Xanthines
28	12.86	519.3324892	C ₂₆ H ₅₀ NO ₇ P	Lysophosphatidylcholine LPC 18:2	Lipids

Table 6. Cont.

#	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
29	12.89	495.3324892	C ₂₄ H ₅₀ NO ₇ P	Lysophosphatidylcholine LPC 16:0	Lipids
	12.91	495.3324892	C ₂₄ H ₅₀ NO ₇ P		
30	12.93	131.0946286	C ₆ H ₁₃ NO ₂	Alanine betaine	Alanine and derivatives
31	12.93	517.3168391	C ₂₆ H ₄₈ NO ₇ P	Lysophosphatidylcholine LPC 18:3	Lipids
	12.94	517.3168391	C ₂₆ H ₄₈ NO ₇ P		
32	12.99	956.4980797	C ₄₈ H ₇₆ O ₁₉	6-[[[(3S,6aR,6bS,8aS,14bR)-4,4,6a,6b,11,11,14b-Heptamethyl-8a-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxycarbonyl-1,2,3,4a,5,6,7,8,9,10,12,12a,14,14a-tetradecahydropicen-3-yl]oxy]-3,5-dihydroxy-4-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxyoxane-2-carboxylic acid Syn. NCGC00385168-01_C48H76O19_Hexopyranose, 1-O-[(3beta,5xi,9xi,18xi)-3-[(3-O-hexopyranosylhexopyranuronosyl)oxy]-28-oxoolean-12-en-28-yl]-	Triterpene saponins
33	13.20	315.2773439	C ₁₈ H ₃₇ NO ₃	Dehydrophytosphingosine	Lipids
34	13.39	456.3603452	C ₃₀ H ₄₈ O ₃	Ursolic acid Syn. Isomer I	Triterpenoids
35	13.40	956.4980797	C ₄₈ H ₇₆ O ₁₉	6-[[[(3S,6aR,6bS,8aS,14bR)-4,4,6a,6b,11,11,14b-Heptamethyl-8a-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxycarbonyl-1,2,3,4a,5,6,7,8,9,10,12,12a,14,14a-tetradecahydropicen-3-yl]oxy]-3,5-dihydroxy-4-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxyoxane-2-carboxylic acid Syn. NCGC00385168-01_C48H76O19_Hexopyranose, 1-O-[(3beta,5xi,9xi,18xi)-3-[(3-O-hexopyranosylhexopyranuronosyl)oxy]-28-oxoolean-12-en-28-yl]-	Triterpene saponins
36	13.49	152.0684734	C ₅ H ₁₂ O ₅	Xylitol (Isomer I)	Sugar alcohols
37	13.60	291.0954163	C ₁₁ H ₁₇ NO ₈	N-fructosyl pyroglutamate	N-fructosyl amino acids
38	13.63	284.075684	C ₁₀ H ₁₂ N ₄ O ₆	Xanthosine	Purine nucleosides
	13.64	284.075684	C ₁₀ H ₁₂ N ₄ O ₆		
39	13.64	152.0334253	C ₅ H ₄ N ₄ O ₂	Xanthine (Isomer II)	Xanthines
40	13.72	152.0684734	C ₅ H ₁₂ O ₅	Xylitol (Isomer II)	Sugar alcohols
41	13.80	454.3446952	C ₃₀ H ₄₆ O ₃	NCGC00380944-01_C30H46O3_(3beta,5xi,9xi,13alpha,17alpha,18xi)-3-Hydroxy-13,28-epoxyurs-11-en-28-one Syn. 3-Hydroxy-11-ursen-28,13-olide	Triterpenoids
42	13.98	456.3603452	C ₃₀ H ₄₈ O ₃	Ursolic acid (Isomer II)	Triterpenoids
43	14.34	267.0967538	C ₁₀ H ₁₃ N ₅ O ₄	Adenosine	Purine nucleosides
44	14.45	180.063388	C ₆ H ₁₂ O ₆	Psicose	Monosaccharides
45	14.69	456.3603452	C ₃₀ H ₄₈ O ₃	Ursolic acid (Isomer III)	Triterpenoids
46	14.80	120.0575148	C ₈ H ₈ O	Phenylacetaldehyde	Phenylacetaldehydes
47	14.80	165.0789785	C ₉ H ₁₁ NO ₂	Phenylalanine	Amino acids
48	14.92	204.0898776	C ₁₁ H ₁₂ N ₂ O ₂	Tryptophan	Amino acids
49	15.03	131.0946286	C ₆ H ₁₃ NO ₂	Isoleucine	Isoleucine and derivatives
50	15.11	756.1901639	C ₃₆ H ₃₆ O ₁₈	NCGC00381212-01![[6-[2-(3,4-dihydroxyphenyl)-5,7-dihydroxy-4-oxochromen-3-yl]oxy-4,5-dihydroxy-2-[(3,4,5-trihydroxy-6-methyloxan-2-yl)oxymethyl]oxan-3-yl]](E)-3-(4-hydroxyphenyl)prop-2-enoate	Flavonoid-O-glycosides
51	15.14	756.2112932	C ₃₃ H ₄₀ O ₂₀	2-(3,4-Dihydroxyphenyl)-3-[(2S,3R,4S,5S,6R)-4,5-dihydroxy-3-[(2R,3R,4R,5R,6S)-3,4,5-trihydroxy-6-methyloxan-2-yl]oxy-6-[(2R,3R,4R,5R,6S)-3,4,5-trihydroxy-6-methyloxan-2-yl]oxymethyl]oxan-2-yl]oxy-5,7-dihydroxychromen-4-one Syn. Quercetin 3-O-rutinoside-(1-2)-O-rhamnoside	Flavonoid-O-glycosides
52	15.21	182.079038	C ₆ H ₁₄ O ₆	D-sorbitol	Sugar alcohols
53	15.31	117.0789785	C ₅ H ₁₁ NO ₂	Betaine	Alpha amino acids
54	15.39	180.063388	C ₆ H ₁₂ O ₆	Mannose (Isomer I)	Hexoses
55	15.41	136.0371732	C ₄ H ₈ O ₅	Threonic acid (Isomer I)	Sugar acids and derivatives

Table 6. Cont.

#	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
56	15.41	150.0528233	C ₅ H ₁₀ O ₅	Xylose	Pentoses
57	15.57	104.107539	C ₅ H ₁₄ NO	Choline	Cholines
58	15.61	136.0371732	C ₄ H ₈ O ₅	Threonic acid (Isomer II)	Sugar acids and derivatives
59	15.63	180.063388	C ₆ H ₁₂ O ₆	Mannose (Isomer II)	Hexoses
60	15.70	196.0583026	C ₆ H ₁₂ O ₇	D-gluconic acid (Isomer I)	Medium-chain hydroxy acids and derivatives
61	15.71	137.0476784	C ₇ H ₇ NO ₂	Trigonelline	Alkaloids and derivatives
	15.82	137.0476784	C ₇ H ₇ NO ₂		
62	15.87	145.0851265	C ₅ H ₁₁ N ₃ O ₂	4-Guanidinobutyric acid	Gamma amino acids and derivatives
63	15.98	196.0583026	C ₆ H ₁₂ O ₇	D-gluconic acid (Isomer II)	Medium-chain hydroxy acids and derivatives
64	15.96	181.0738931	C ₉ H ₁₁ NO ₃	Tyrosine	Tyrosine and derivatives
65	16.15	165.0789785	C ₉ H ₁₁ NO ₂	Phenylalanine	Phenylalanine and derivatives
66	16.16	212.0896027	C ₇ H ₁₆ O ₇	Volemitol	Sugar alcohols
67	16.45	293.1474519	C ₁₂ H ₂₃ NO ₇	N-fructosyl isoleucine	N-fructosyl amino acids
	16.48	293.1474519	C ₁₂ H ₂₃ NO ₇		
	16.54	293.1474519	C ₁₂ H ₂₃ NO ₇		
68	16.62	103.0633285	C ₄ H ₉ NO ₂	4-Aminobutyric acid Syn. 4-Aminobutanoic acid/GABA	Gamma amino acids and derivatives
69	17.15	147.0531577	C ₅ H ₉ NO ₄	L-Glutamic acid (Isomer I)	Glutamic acid and derivatives
70	17.18	342.1162113	C ₁₂ H ₂₂ O ₁₁	Melibiose (Isomer I)	O-glycosyl compounds
71	17.25	342.1162113	C ₁₂ H ₂₂ O ₁₁	Isomaltulose	O-glycosyl compounds
72	17.26	342.1162113	C ₁₂ H ₂₂ O ₁₁	Trehalose	Disaccharide
73	17.27	129.042593	C ₅ H ₇ NO ₃	Pyroglutamic acid	Alpha amino acids and derivatives
74	17.29	147.0531577	C ₅ H ₉ NO ₄	L-glutamic acid (Isomer II)	Glutamic acid and derivatives
75	17.38	119.0582431	C ₄ H ₉ NO ₃	Threonine (Isomer I)	L-alpha-amino acids
76	17.46	119.0582431	C ₄ H ₉ NO ₃	Threonine (Isomer II)	L-alpha-amino acids
77	18.02	165.0459638	C ₅ H ₁₁ NO ₃ S	Methioninesulfoxide	Alpha amino acids
78	18.02	342.1162113	C ₁₂ H ₂₂ O ₁₁	Melibiose (Isomer II)	O-glycosyl compounds
79	18.10	146.0691421	C ₅ H ₁₀ N ₂ O ₃	Glutamine	D-alpha-amino acids
	18.14	146.0691421	C ₅ H ₁₀ N ₂ O ₃		
80	18.12	105.042593	C ₃ H ₇ NO ₃	Serine	Serine and derivatives
	18.16	105.042593	C ₃ H ₇ NO ₃		
81	18.31	344.1318613	C ₁₂ H ₂₄ O ₁₁	Maltitol	Fatty acyl glycosides of mono- and disaccharides
82	18.40	132.053492	C ₄ H ₈ N ₂ O ₃	Asparagine	Asparagine and derivatives
	18.41	132.053492	C ₄ H ₈ N ₂ O ₃		
83	18.53	342.1162113	C ₁₂ H ₂₂ O ₁₁	Melibiose (Isomer III)	O-glycosyl compounds
84	18.84	504.1690346	C ₁₈ H ₃₂ O ₁₆	Melezitose (Isomer I)	Oligosaccharides
	18.85	504.1690346	C ₁₈ H ₃₂ O ₁₆		
	18.86	504.1690346	C ₁₈ H ₃₂ O ₁₆		
85	19.13	504.1690346	C ₁₈ H ₃₂ O ₁₆	Melezitose (Isomer II)	Oligosaccharides

Table 6. Cont.

#	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
86	19.14	504.1690346	C ₁₈ H ₃₂ O ₁₆	Maltotriose (Isomer I)	Oligosaccharides
87	19.33	504.1690346	C ₁₈ H ₃₂ O ₁₆	Maltotriose (Isomer II)	Oligosaccharides
88	19.34	504.1690346	C ₁₈ H ₃₂ O ₁₆	Raffinose	Oligosaccharides
89	21.88	155.0694765	C ₆ H ₉ N ₃ O ₂	L-Histidine	Histidine and derivatives
90	22.28	174.1116756	C ₆ H ₁₄ N ₄ O ₂	L-Arginine	L-alpha-amino acids

In the seeds (Table 7), the phytochemical constituents were classified as: (i) primary metabolites, such as amino acids and derivatives (n = 11), organic acids (n = 16), monosaccharides sugar acids and sugar alcohols (n = 8), disaccharides and oligosaccharides (n = 7), lipids (n = 14), and nucleobases/nucleosides (n = 5); and (ii) secondary metabolites, such as triterpenoids (n = 3), catechols (n = 3), phenolic glycosides (n = 2), flavonoids (n = 5), alkaloids and derivatives (n = 1), triterpene saponins (n = 1), and other compounds (n = 17).

Table 7. Phytochemical constituents of quinoa seeds (Pasankalla variety) determined by LC-ESI-MS/MS.

#	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
1	3.19	340.1885884	C ₁₈ H ₂₈ O ₆	[5-Acetyloxy-3-(hydroxymethyl)-2-oxo-6-propan-2-ylcyclohex-3-en-1-yl] 3-methylpentanoate	Menthane monoterpenoids
2	3.72	145.0527638	C ₉ H ₇ NO	2-Hydroxyquinoline	Hydroquinolones
3	3.74	122.0367794	C ₇ H ₆ O ₂	3-Hydroxybenzaldehyde	Phenolic compounds
4	3.78	206.0579087	C ₁₁ H ₁₀ O ₄	Isoeugenitol	Chromones
5	3.80	152.047344	C ₈ H ₈ O ₃	4-Hydroxyphenylacetic acid	1-Hydroxy-2-unsubstituted benzenoids
6	3.85	168.0422586	C ₈ H ₈ O ₄	3,4-Dihydroxyphenylacetate (Isomer I) Syn. Homoprotocatechuic acid	Catechols
7	3.89	154.0266086	C ₇ H ₆ O ₄	Pyrocatechuic acid (Isomer I)	Salicylic acids
8	3.96	130.0266086	C ₅ H ₆ O ₄	Citraconic acid	Methyl-branched fatty acids
9	4.13	164.047344	C ₉ H ₈ O ₃	3-Hydroxycinnamic acid	Hydroxycinnamic acids
10	4.20	132.0786442	C ₆ H ₁₂ O ₃	2-Hydroxyisocaproic acid	Hydroxy fatty acids
11	4.20	160.0735588	C ₇ H ₁₂ O ₄	3-Methyladipic acid	Medium-chain fatty acids
12	4.22	168.0422586	C ₈ H ₈ O ₄	3,4-Dihydroxyphenylacetate (Isomer II) Syn. Homoprotocatechuic acid	Catechols
13	4.29	154.0266086	C ₇ H ₆ O ₄	Pyrocatechuic acid (Isomer II)	Salicylic acids
14	4.43	138.031694	C ₇ H ₆ O ₃	Salicylic acid	Salicylic acids
15	4.66	146.0579087	C ₆ H ₁₀ O ₄	2-Methylglutaric acid	Methyl-branched fatty acids
16	4.75	132.0422586	C ₅ H ₈ O ₄	Glutaric acid (Isomer I)	Dicarboxylic acids and derivatives
17	4.75	194.0579087	C ₁₀ H ₁₀ O ₄	<i>trans</i> -4-Hydroxy-3-methoxycinnamate (Isomer I)	Hydroxycinnamic acids
18	4.98	134.0215232	C ₄ H ₆ O ₅	Malic acid (Isomer I)	Beta hydroxy acids and derivatives
19	5.07	164.047344	C ₉ H ₈ O ₃	3-Hydroxycinnamic acid (Isomer I)	Hydroxycinnamic acids
20	5.14	118.0266086	C ₄ H ₆ O ₄	Succinic acid (Isomer I)	Dicarboxylic acids and derivatives
21	5.45	132.0422586	C ₅ H ₈ O ₄	Glutaric acid (Isomer II)	Dicarboxylic acids and derivatives

Table 7. Cont.

	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group	
	22	5.75	118.0266086	C ₄ H ₆ O ₄	Succinic acid (Isomer II)	Dicarboxylic acids and derivatives
	23	5.82	194.0579087	C ₁₀ H ₁₀ O ₄	<i>trans</i> -4-Hydroxy-3-methoxycinnamate (Isomer II)	Hydroxycinnamic acids
	24	5.83	123.0320284	C ₆ H ₅ NO ₂	Isonicotinic acid	Pyridinecarboxylic acids
	25	5.98	110.0367794	C ₆ H ₆ O ₂	Catechol	Catechols
	26	6.26	134.0215232	C ₄ H ₆ O ₅	Malic acid (Isomer II)	Beta hydroxy acids and derivatives
	27	6.26	164.047344	C ₉ H ₈ O ₃	3-Hydroxycinnamic acid (Isomer II)	Hydroxycinnamic acids
	28	7.05	219.1106725	C ₉ H ₁₇ NO ₅	Pantothenic acid	Secondary alcohols
		7.11	219.1106725	C ₉ H ₁₇ NO ₅		
	29	7.27	516.3298328	C ₂₇ H ₄₈ O ₉	MGMG 18:2	Lipids
	30	7.40	122.0480128	C ₆ H ₆ N ₂ O	Nicotinamide	Nicotinamides
	31	7.82	264.1110069	C ₁₃ H ₁₆ N ₂ O ₄	Phenylacetylglutamine	<i>N</i> -acyl- α amino acids
	32	7.83	129.042593	C ₅ H ₇ NO ₃	5-Oxo-D-proline Syn. D-Pyroglutamic acid	Proline and derivatives
	33	8.38	514.3141828	C ₂₇ H ₄₆ O ₉	NCGC00380867-01_C27H46O9_9,12,15-Octadecatrienoic acid, 3-(hexopyranosyloxy)-2-hydroxypropyl ester, (9Z,12Z,15Z)-	Glycosylmonoacylglycerols
	34	9.53	494.3243536	C ₂₈ H ₄₆ O ₇	NCGC00169545-02[(2S,3R,5R,10R,13R,14S,17S)-2,3,14-trihydroxy-10,13-dimethyl-17-[(2R,3R,5R)-2,3,6-trihydroxy-5,6-dimethylheptan-2-yl]-2,3,4,5,9,11,12,15,16,17-decahydro-1H-cyclopenta[a]phenanthren-6-one Syn. Makisterone A	Phytoecdysteroids
	35	10.03	480.3087035	C ₂₇ H ₄₄ O ₇	NCGC00168839-02[(2S,3R,5R,10R,13R,14S,17S)-2,3,14-trihydroxy-10,13-dimethyl-17-[(2R,3R)-2,3,6-trihydroxy-6-methylheptan-2-yl]-2,3,4,5,9,11,12,15,16,17-decahydro-1H-cyclopenta[a]phenanthren-6-one Syn. Ecdysterone	Phytoecdysteroids
		10.04	480.3087035	C ₂₇ H ₄₄ O ₇		
	40	10.06	722.5097847	C ₃₈ H ₇₅ O ₁₀ P	[3-[[2,3-Dihydroxypropoxy]-hydroxyphosphoryl]oxy-2-hexadecanoyloxypropyl] hexadecanoate Syn. Dipalmitoylphosphatidylglycerol	Phosphatidylglycerol
	41	10.90	372.1420321	C ₁₇ H ₂₄ O ₉	Syringin	Phenolic glycosides
	42	10.96	478.0747403	C ₂₁ H ₁₈ O ₁₃	Quercetin-3-glucuronide	Flavonoid-O-glucuronides
	43	11.49	244.069536	C ₉ H ₁₂ N ₂ O ₆	Uridine	Pyrimidine nucleosides
	44	12.03	281.1124038	C ₁₁ H ₁₅ N ₅ O ₄	2'-O-methyladenosine	Purine nucleosides
	45	12.36	477.285539	C ₂₃ H ₄₄ NO ₇ P	Lysophosphatidylethanolamine LPE 18:2	Lipids
	46	12.40	453.285539	C ₂₁ H ₄₄ NO ₇ P	Lysophosphatidylethanolamine LPE 16:0	Lipids
	47	12.59	131.0946286	C ₆ H ₁₃ NO ₂	Alanine betaine	Alanine and derivatives
	48	12.76	521.3481393	C ₂₆ H ₅₂ NO ₇ P	Lysophosphatidylcholine LPC 18:1	Lipids
		12.78	521.3481393	C ₂₆ H ₅₂ NO ₇ P		
	49	12.85	519.3324892	C ₂₆ H ₅₀ NO ₇ P	Lysophosphatidylcholine LPC 18:2	Lipids
	50	12.88	152.0684734	C ₅ H ₁₂ O ₅	L-arabitol (Isomer I)	Sugar alcohols
	51	12.89	495.3324892	C ₂₄ H ₅₀ NO ₇ P	Lysophosphatidylcholine LPC 16:0	Lipids
		12.90	495.3324892	C ₂₄ H ₅₀ NO ₇ P		
	52	12.91	517.3168391	C ₂₆ H ₄₈ NO ₇ P	Lysophosphatidylcholine LPC 18:3	Lipids
		12.95	517.3168391	C ₂₆ H ₄₈ NO ₇ P		
	53	13.11	639.3383623	C ₂₉ H ₅₄ NO ₁₂ P	Hexosyl LPE 18:2	Lipids
	54	13.16	315.2773439	C ₁₈ H ₃₇ NO ₃	Dehydrophytosphingosine Syn. 4-Hydroxy-8-sphingenine	Lipids
	55	13.43	956.4980797	C ₄₈ H ₇₆ O ₁₉	6-[[[(3S,6aR,6bS,8aS,14bR)-4,4,6a,6b,11,11,14b-heptamethyl-8a-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxycarbonyl-1,2,3,4a,5,6,7,8,9,10,12,12a,14,14a-tetradecahydricen-3-yl]oxy]-3,5-dihydroxy-4-[3,4,5-trihydroxy-6-(hydroxymethyl)oxan-2-yl]oxyoxane-2-carboxylic acid Syn. NCGC00385168-01_C48H76O19_Hexopyranose, 1-O-[(3beta,5xi,9xi,18xi)-3-[(3-O-hexopyranosyl)hexopyranuronosyl]oxy]-28-oxoolean-12-en-28-yl]-	Triterpene saponins
	56	13.61	291.0954163	C ₁₁ H ₁₇ NO ₈	<i>N</i> -fructosyl pyroglutamate	<i>N</i> -fructosyl amino acids

Table 7. Cont.

	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
57	13.61	610.1533845	C ₂₇ H ₃₀ O ₁₆	Rutoid Syn. Rutin	Flavonoid-O-glycosides
58	13.61	284.075684	C ₁₀ H ₁₂ N ₄ O ₆	Xanthosine	Purine nucleosides
	13.62	284.075684	C ₁₀ H ₁₂ N ₄ O ₆		
59	13.62	152.0334253	C ₅ H ₄ N ₄ O ₂	Xanthine	Xanthines
60	13.73	152.0684734	C ₅ H ₁₂ O ₅	L-arabitol (Isomer II)	Sugar alcohols
61	13.79	454.3446952	C ₃₀ H ₄₆ O ₃	NCGC00380944-01_C30H46O3_(3beta,5xi,9xi,13alpha,17alpha,18xi)-3-Hydroxy-13,28-epoxyurs-11-en-28-one Syn. 3-Hydroxy-11-Ursen-28,13-Olide	Triterpenoids
62	13.97	456.3603452	C ₃₀ H ₄₈ O ₃	Ursolic acid (Isomer I)	Triterpenoids
63	14.10	770.2269433	C ₃₄ H ₄₂ O ₂₀	7-Methylquercetin-3-galactoside-6''-rhamnoside-3'''-rhamnoside	Flavonoid -O-glycosides
64	14.21	221.0899371	C ₈ H ₁₅ NO ₆	N-acetylmannosamine	N-acyl-alpha-hexosamines
65	14.29	180.063388	C ₆ H ₁₂ O ₆	Psicose	Monosaccharides
66	14.69	456.3603452	C ₃₀ H ₄₈ O ₃	Ursolic acid (Isomer II)	Triterpenoids
67	14.76	165.0789785	C ₉ H ₁₁ NO ₂	Phenylalanine	Amino acids
68	14.90	204.0898776	C ₁₁ H ₁₂ N ₂ O ₂	Tryptophan	Indolyl carboxylic acids and derivatives
69	14.93	283.0916684	C ₁₀ H ₁₃ N ₅ O ₅	Guanosine	Purine nucleosides
70	14.93	742.1956431	C ₃₂ H ₃₈ O ₂₀	NCGC00180410-02!3-[(2S,3R,4S,5S,6R)-6-[[[(2R,3R,4R,5R,6S)-3-[(2S,3R,4R)-3,4-dihydroxy-4-(hydroxymethyl)oxolan-2-yl]oxy-4,5-dihydroxy-6-methyloxan-2-yl]oxymethyl]-3,4,5-trihydroxyoxan-2-yl]oxy-2-(3,4-dihydroxyphenyl)-5,7-dihydroxychromen-4-one]Syn. Quercetin 3-(2R-apsiosylrutinoside)	Flavonoid-O-glycosides
71	15.04	131.0946286	C ₆ H ₁₃ NO ₂	Isoleucine	Isoleucine and derivatives
72	15.11	104.107539	C ₅ H ₁₄ NO	Choline (Isomer I)	Cholines
73	15.14	756.2112932	C ₃₃ H ₄₀ O ₂₀	2-(3,4-Dihydroxyphenyl)-3-[(2S,3R,4S,5S,6R)-4,5-dihydroxy-3-[(2R,3R,4R,5R,6S)-3,4,5-trihydroxy-6-methyloxan-2-yl]oxy-6-[[[(2R,3R,4R,5R,6S)-3,4,5-trihydroxy-6-methyloxan-2-yl]oxymethyl]oxan-2-yl]oxy-5,7-dihydroxychromen-4-one Syn. Quercetin 3-O-rutinoside-(1-2)-O-rhamnoside	Flavonoid-O-glycosides
74	15.19	182.079038	C ₆ H ₁₄ O ₆	D-sorbitol	Sugar alcohols
75	15.27	117.0789785	C ₅ H ₁₁ NO ₂	Betaine	Alpha amino acids
76	15.39	135.0544952	C ₅ H ₅ N ₅	Adenine	6-Aminopurines
77	15.59	104.107539	C ₅ H ₁₄ NO	Choline (Isomer II)	Cholines
78	15.64	180.063388	C ₆ H ₁₂ O ₆	Mannose	Hexoses
79	15.78	212.0896027	C ₇ H ₁₆ O ₇	Perseitol	Sugar alcohols
80	15.79	137.0476784	C ₇ H ₇ NO ₂	Trigonelline	Alkaloids and derivatives
81	15.84	145.0851265	C ₅ H ₁₁ N ₃ O ₂	4-Guanidinobutanoic acid	Gamma amino acids and derivatives
82	16.49	293.1474519	C ₁₂ H ₂₃ NO ₇	N-fructosyl isoleucine	N-fructosyl amino acids
	16.52	293.1474519	C ₁₂ H ₂₃ NO ₇		
83	17.24	342.1162113	C ₁₂ H ₂₂ O ₁₁	Trehalose	Disaccharides
84	17.25	342.1162113	C ₁₂ H ₂₂ O ₁₁	Maltose	Oligosaccharides
85	17.25	342.1162113	C ₁₂ H ₂₂ O ₁₁	Isomaltulose	Oligosaccharides
86	17.25	342.1162113	C ₁₂ H ₂₂ O ₁₁	Melibiose	Oligosaccharides
87	17.29	147.0531577	C ₅ H ₉ NO ₄	L-glutamic acid	Glutamic acid and derivatives
88	18.13	146.0691421	C ₅ H ₁₀ N ₂ O ₃	Glutamine	D-alpha-amino acids
89	18.30	344.1318613	C ₁₂ H ₂₄ O ₁₁	Maltitol Syn. 4-O-alpha-D-Glucopyranosyl-D-glucitol	Hexoses
90	19.31	504.1690346	C ₁₈ H ₃₂ O ₁₆	Maltotriose	Oligosaccharides

Table 7. Cont.

	Rt (min)	Theoretical Mass (Neutral Form)	Molecular Formula (Neutral Form)	Predicted Metabolite	Chemical Group
91	19.33	504.1690346	C ₁₈ H ₃₂ O ₁₆	Raffinose	Oligosaccharides
92	20.64	179.0793724	C ₆ H ₁₃ NO ₅	D-mannosamine	Hexoses
93	20.79	666.2218579	C ₂₄ H ₄₂ O ₂₁	Tetrasaccharides (Hex-Hex-Hex-Hex)	Oligosaccharides

3. Discussion

Polyphenolic compounds are secondary metabolites present in plants, which are divided into flavonoids and non-flavonoids, the first being responsible for the antioxidant capacity, exerting this through various mechanisms such as transition metal chelators, free radical scavengers, and enzyme inhibitors [16]. The antioxidant properties of secondary metabolites are related to vasodilatory, lipid-lowering, antiaging, and anti-inflammatory, modulating apoptosis processes in the vascular endothelium, but these molecules could also be influenced by factors such as the number and position of the phenolic hydroxyl groups, steric effects, and molecular properties [17]. In our results, the content of total phenols and flavonoids found in quinoa sprouts presented differences in each variety analyzed, being influenced by the type of seed, the cultivation site, maturity, storage, and germination conditions, as the flavonoids play an important role in pigmentation [18]. It is known that the phenolic compounds present in plants are formed during their development and under stress conditions; these include simple phenols, phenolic acids, coumarins, flavonoids, stilbenes, hydrolysable and condensed tannins, lignans, and lignins [19]. Additionally, these polyphenols could be altered during the germination process, increasing their content and the antioxidant capacity [20].

In our study, the variation in TPC and TF differed from the studies of Valencia et al., in which the TPC varied from 0.783 to 3437 mg GAE/g in quinoa seeds [21], and that of Carciochi et al. [22], with values of TPC of 39.3 ± 0.9 mg GAE/100 g and TF of 11.06 mg of quercetin/100 g in sprouts. These were higher in our study due to the type of solvent used in the maceration process. In the same way, when the antioxidant activity of the content of polyphenols and flavonoids was evaluated in the red and yellow varieties of quinoa, there was a significant increase after 9 days of germination. In a similar study, the antioxidant capacity in germinated seeds was greater compared with seeds of *C. quinoa*, increasing up to twofold, similar to the increase in phenolic compounds and antioxidant capacity observed after 72 h of germination [13]. In our study, a wide range of values were observed for phenolic compounds and flavonoids, as well as for the antioxidant activity in each variety of quinoa studied, which can be explained by the characteristics of each seed, variation in the availability of nutrients, and activation of the antioxidant machinery during germination.

Several studies have shown nutritional improvements in quinoa sprouts, such as in crude quinoa flour (CQF) and germinated quinoa flour (GQF), where the CQF/GQF ratio increased the nutritional quality of pasta. Chemical analysis indicated an increase in the proportion of proteins by 37% and a decrease in phytic acid by 77%, which means that the germination process is an effective method to minimize phytic acid content in seeds. Pasta with a high CQF/GQF ratio had an increased content of Ca, K, Fe, Mn, Mg, P, and Zn, and thus using GQF is recommended in the production of bread, cakes, and cookies to take advantage of their nutritional properties, which provide a high content of proteins, minerals, TPC, and amino acids, and a low amount of phytic acid [23]. During germination, quinoa seeds undergo relevant physical and chemical changes; the maximum intensity of macromolecular modification occurs at 48 h. The germinated material contains micronutrients with improved bioavailability. This has a great impact on quinoa, as it improves the technological properties of quinoa, as well as some of its nutritional characteristics, enhancing the use of quinoa sprout flour as an ingredient in food formulation [12]. The germination process of quinoa seeds is an effective technique to enhance the content of total

phenols and total flavonoids and to improve the antioxidant capacity, as was demonstrated in quinoa (*C. quinoa*) and kiwicha (*Amaranthus caudatus*) [24], where the sprouts had enhanced content of coumaric acid and kaempferol tri-glycoside in quinoa and caffeoylquinic acid in kiwicha. Additionally, a significant increase was observed in the phenolic content and the antioxidant capacity through malting quinoa sprouts [25] and *Amaranthus caudatus* sprouts [26].

4. Materials and Methods

4.1. Collection of Quinoa Seeds

Fifteen certified varieties were provided by the Agrarian Research Institute (INIA, Ayacucho, Peru) and five varieties were collected between November and December 2019 in the districts of Huamanguilla and Acocro of the province of Huamanga. These are registered with the following names: White Junín Ayacucho, T-256, Pasankalla, Suano Puno, T-38, Yellow Sacaca, T-45, Santa Ana, T-61 Pomata, CQA-048, Black Collana, T-72 Huancayo, CQA-043, Salcedo, Ayacucho Compuesto, White Choclito, Red, Yellow Marangani, Black Coito, and Black.

4.2. Germination Process

The seeds were washed with hypochlorite 0.02% (*w/v*) for 20 min, rinsed several times with distilled water, and placed on absorbent paper moistened with distilled water in Technopor containers covered with paper towels and incubated at room temperature (between 18 and 22 °C) for 72 h until good sprouts had been obtained. The sprouts were harvested, dried at 45 °C for 48 h, then crushed and stored under refrigeration [27].

4.3. Preparation of the Methanolic Extract

Ten grams of each sample of sprouts and seeds was subjected to dynamic extraction with 100 mL of methanol (1:10), using a magnetic stirrer for 4 h at room temperature, then filtered with Whatman No. 1 paper and concentrated on a rotary evaporator until dry. Each extract was refrigerated until further use at 4 °C.

4.4. Determination of Total Phenolic Content (TPC)

In total, 50 µL of the methanolic extract (10 mg/mL) was mixed with 1 mL of distilled water, 0.5 mL of 0.2 N Folin-Ciocalteu reagent, and 2.5 mL of 5% sodium carbonate, then the sample was allowed to react in the darkness for 40 min at room temperature (20°C). The absorbance was read at 725 nm using a UV-Vis Genesys 150 Thermo Scientific spectrophotometer. A standard curve was made with a gallic acid solution (50 µg/mL) at concentrations of 10, 20, 30, 40, and 50 µg/mL. The results are presented in mg equivalent to gallic acid per g of methanolic extract (mg GAE/g of extract) [28].

4.5. Determination of Total Flavonoids

In total, 0.5 mL of the extract (10 mg/mL) was mixed with 1 mL with distilled water and 0.15 mL of 5% sodium nitrite; 5 min later, 0.15 mL of 10% aluminum chloride was added, then at 6 min, 2 mL of 4% sodium hydroxide was added. The sample was made up to 5 mL with distilled water, mixed, and allowed to react in the darkness for 15 min at room temperature. The absorbance was read at 510 nm against a blank. A standard curve was made with quercetin (200 µg/mL) at concentrations of 40, 80, 120, 160, and 200 µg/mL. The flavonoid content is presented as mg equivalent to quercetin per g of dry methanolic extract (mg QE/g of extract) [29].

4.6. Determination of the Antioxidant Capacity by the Free Radical Sequestration Method with 2,2-Diphenyl-1-picrylhydrazyl

For this assay, 150 µL of extract (10 mg/mL) was mixed with 2850 µL of a methanolic solution of DPPH radicals (20 mg/L) with the absorbance adjusted to 0.6 ± 0.02 nm. After mixing, the sample was incubated in the dark for 30 min and the absorbance was

read at 515 nm. The standard curve was elaborated with Trolox at concentrations of 0 to 800 $\mu\text{mol}/\text{mL}$ [30]. The antioxidant capacity equivalent to Trolox (TEAC) was calculated with the following formula:

$$TEAC \frac{\mu\text{molTROLOX}}{\text{mg ME}} = IC_{50TROLOX} \left(\frac{\mu\text{mol}}{\text{mL}} \right) / IC_{50sample} \left(\frac{\text{mg}}{\text{mL}} \right)$$

To calculate the half inhibitory concentration (IC_{50}), the percentage of inhibition of the DPPH radical was determined at concentrations of 5, 10, and 20 mg/mL of methanolic extract according to the following equation:

$$\% \text{ inhibition of the DPPH radical} = \frac{abs_{control} - abs_{sample}}{abs_{control}} \times 100$$

where $abs_{control}$ is the absorbance of the control without the sample at $t = 0$ min, and abs_{sample} is the absorbance of the sample at $t = 30$ min.

4.7. Determination of the Antioxidant Capacity by the Sequestration Method with the Radical Cation of the 2,2'-Azinobis-(3-ethylbenzothiazoline)-6-sulfonic acid

A standard solution (ST) was prepared by mixing 10 mL of ABTS (4.06 mg/mL) with 10 mL of potassium persulfate (0.7 mg/mL) and reacted for 12 h. The working solution (ST) was prepared with 1 mL of each extract and 60 mL of methanol. The absorbance was adjusted to 0.7 ± 0.02 with methanol at a wavelength of 734 nm, then 150 μL of the extract (5 mg/mL) was mixed with 2850 μL of the extract solution and incubated in the dark for 7 min, followed by reading the absorbances at 734 nm [31]. The standard curve was made with Trolox at 0–400 $\mu\text{mol}/\text{mL}$. The antioxidant capacity equivalent to Trolox (TEAC) was expressed as $\mu\text{mol ET}/\text{mg}$ of the extract.

$$TEAC \frac{\mu\text{molTROLOX}}{\text{mg ME}} = IC_{50TROLOX} \left(\frac{\mu\text{mol}}{\text{mL}} \right) / IC_{50sample} \left(\frac{\text{mg}}{\text{mL}} \right)$$

To calculate the half inhibitory concentrations (IC_{50}), the percentage of inhibition of the ABTS radical was determined at concentrations of 1, 5, and 10 mg/mL as follows:

$$\% \text{ inhibition of the ABTS radical} = \frac{abs_{control} - abs_{sample}}{abs_{control}} \times 100$$

where $abs_{control}$ is the absorbance of the control without the sample at $t = 0$ min and abs_{sample} is the absorbance of the sample at $t = 7$ min.

4.8. Phytochemical Analysis by LC-ESI-MS/MS of the Main Constituents of Methanolic Extracts of the Sprouts and Seeds of *C. quinoa* (Pasankalla Variety)

4.8.1. Preparation of the Sample

The methanolic extracts of the sprouts and seeds of *C. quinoa* were weighed and diluted with methanol until a final concentration of 2 mg/mL had been obtained. Next, each sample was vortexed for 1 min and subsequently centrifuged for 10 min at 10,000 rpm. Finally, 800 μL of the 1 mg/mL solution supernatant (methanol:water, 1:1) was removed in vials for LC-MS analysis in a Dionex UltiMate 3000 liquid chromatograph (Thermo Fisher Scientific, San José, CA, USA) coupled to a Thermo QExactiveTM Plus Orbitrap mass spectrometer (Thermo Fisher Scientific, Bremen, Germany) with an electrospray ionization source.

4.8.2. Chromatographic Conditions

This analysis used a chromatographic column XBridge[®] Amide BEH water (150 mm \times 4.6 mm \times 3.5 μm). Solvent A was 0.1% formic acid in water and Solvent B was 0.1% formic acid in ACN. The gradient elution of the method was as follows: 0–2 min, B 95%;

2–17.0 min, B 50%; 17–20.0 min, B 50%; 20.0–21.0 min, B 95%; 21.0–27.0 min, B 95%. The flow rate was 500 $\mu\text{L min}^{-1}$ with injection of 8 μL and a column oven temperature of 40 $^{\circ}\text{C}$.

4.8.3. Mass Spectrometry Conditions

A full scan experiment combined with a fragmentation experiment (MS/MS) was performed for both electrospray ionization modes (ESI + and –). The ESI source parameters were as follows: spraying voltage: 3.9 kV (+) and 3.6 kV (–); envelope gas flow rate: 50 (arbitrary values); auxiliary gas flow: 10 (arbitrary values); tube lens voltage: 50 V; probe heater temperature: 400 $^{\circ}\text{C}$; capillary temperature: 300 $^{\circ}\text{C}$.

1. (. ESI +) mode: full MS mode parameters: 35,000 resolution; ACG target (automatic gain control): 5e5; maximum IT (injection time): 100 ms; scan range: 100–1200 m/z .

Dd-MS² (data-dependent acquisition experiment, DDA) mode parameters: 17,500 resolution; ACG objective: 1e5; maximum IT: 50 ms; loop count, 3; isolation window: 1–2 m/z ; topN, 3; NCE (stepped normalized collision energy): 15, 30, and 40.

2. (. ESI –) mode: full MS mode parameters: 35,000 resolution; ACG objective: 5e5; maximum IT: 100 ms; range, 100–1200 m/z .

Dd-MS² (data-dependent acquisition experiment, DDA) mode parameters: 17,500 resolution; ACG objective: 1e5; maximum IT: 50 ms; loop count, 3; isolation window: 1–2 m/z ; topN: 3; NCE: 15, 20, and 40.

Data acquisition and processing were performed with Thermo XcaliburTM software version 3.0 (Thermo Fisher Scientific Inc., Waltham, MA, USA) with the Qual Browser, and metabolite annotations were performed with MS-Dial software version 4.70 (Riken, Osaka University, Suita City, Japan) using the MS-Dial metabolomics MPS spectral kit library (available at: <http://prime.psc.riken.jp/compms/msdial/main.html>; last updated on 13 April 2021).

4.9. Data Analysis

The results are presented as the means plus standard deviation of three repetitions. The differences between the means were analyzed using paired sample t-test for total phenols, flavonoids, antioxidant capacity, and the half inhibitory concentration (IC_{50}), using SPSS software. Pearson's correlation coefficient was determined to establish the relationships among total phenols and flavonoids, antioxidant capacity (TEAC), and the half inhibitory concentration (IC_{50}), with a p -value less than 0.05 being significant.

5. Conclusions

Based on our results, we concluded that quinoa sprouts germinated for 72 h had higher total phenolic content and total flavonoids compared with seed extracts, and these correlated with its high antioxidant capacity. Furthermore, sprout extracts had better IC_{50} and TEAC values in the DPPH and ABTS assays. The best variety of quinoa was Pasankalla, which showed a high antioxidant capacity and also contained 90 and 93 phytochemical constituents in the sprout and seed extract, respectively. Some chemical groups highlighted were amino acids, organic acids, phenolic acids, flavonoids, fatty acids, lipids, saponins, and sugars, with a greater diversity of essential amino acids found in sprouts than in seeds.

Supplementary Materials: The following are available online at <https://www.mdpi.com/article/10.3390/plants10112417/s1>, Table S1: List of compounds putatively identified by LC-HRMS/MS in the extract of *Chenopodium quinoa* sprouts. Table S2: List of compounds putatively identified by LC-HRMS/MS in the extract of *Chenopodium quinoa* seeds.

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Data Availability Statement: The data that support the findings of this study are available from the corresponding author upon reasonable request.

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