Article

Quantitative Assessment of Trace and Macro Element Compositions of Cassava (*Manihot esculenta*) Storage Roots Enriched with B-Carotene as Influenced by Genotypes and Growing Locations

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Abstract: Cassava's important mineral contents depends on some factors, including genetic and growing locational factors. The study aimed to evaluate the influence of genotype and growing locations on the mineral concentrations in yellow-fleshed cassava root genotypes. Twenty-five pipeline yellow-fleshed cassava genotypes and three white-fleshed varieties (check samples) were planted at five different experimental fields for two seasons, each representing the major agroecological zones in Nigeria. Standard laboratory protocols were employed in the sampling to ensure zero contamination, and the trace and macro elements were determined using the inductively coupled plasma optical emission spectroscopic method (ICPOES). The trace and macro elements identified in all the genotypes and varieties investigated were Fe, Mn, B, Cu, Mo, Co, Ni, Zn, and Al; Ca, Mg, Na, K, P, and S respectively. Genotype and growing location had a highly significant ($p < 0.05$) effect on all the trace elements except Ti and Cr. However, there was no interactive effect between genotype and growing location on all the trace elements except for Pb and Zn. Among the explanatory variables, the variable growing location was the most influential on macro and trace elements. Conclusively, genotypes 01/1442 and 01/1273 have outstanding trace and macro element concentrations.

Keywords: yellow-fleshed cassava; macronutrient; micronutrient; genotypic variations; growing locations

1. Introduction

Cassava has primarily been identified as one of the staple foods that is low in nutritional quality, and this due to the low protein content of its storage roots [1,2]. However, cassava contains Fe, Cu, Ca, Zn, Mg, and K, but most of these minerals are found more in the root peel than in the peeled root, a result of processing [3]. This makes the mineral content in cassava lower than sorghum and maize but comparable with many legumes except soybean [2]. Thus, cassava nutritional value can be considered lower than cereals, legumes, and some other roots and tubers. The importance of cassava as food in sub-Saharan Africa (SSA), especially Nigeria, has proved beyond doubt that it is an economic crop. Besides, it is an easily cultivated subsistence crop, a drought-tolerant, multiple-year crop that can be kept in the ground until needed; this qualifies cassava as a food security crop. Nutritionally, it provides an economical source of dietary energy in the form of carbohydrates [4]. The usefulness of cassava does not end with the roots; its leaves also are used in the preparation of relishes whose protein quality compares favorably with soybean with high levels of vitamins and minerals.

Hidden hunger, also known as micronutrient deficiencies, happens when the vitamin and mineral intake or absorption is very low to keep the good health and development in children and everyday physical and mental function in adults [5]. It is as a result of the unbalanced nature of diets consumed by people in most of the developing countries, especially Nigeria. Micronutrients deficiency is a challenge faced by the entire world. In most developing countries, deficiencies of Fe and Zn are part of the five micronutrients malnutrition that have been identified to be of public health significance. Biofortification is one of the approaches employed by plant breeders to increase the micronutrient content of crops using conventional breeding, such as cassava. There are different types of cassava varieties based on their peel and pulp colors. Generally, the peel appears cream, brown, light brown, and pinkish brown. At the same time, the pulp's appearance could be yellow, orange, cream, and white. The color usually indicates the beta-carotene and moisture content [6]. The white pulp has been the most cultivated in West Africa and especially, Nigeria. The yellow root has its origin traced to the Amazon basin, which over the years, has been cross-bred with the African white varieties using conventional breeding techniques to boost the carotenoid and micronutrient concentrations in cassava [7].

Minerals are essential elements that are not synthesized by the body but are essential for the body to function optimally [8]. These can be grouped into trace and macroelements. They function in overlap with other macronutrients and vitamins to ward off sicknesses and, most importantly, degenerative diseases. These minerals are actively involved in the development of biochemical processes, in which their essential roles are catalytic, functional, and structural [9]. These can be divided into trace and macro elements. Trace or microelements are usually needed in very minute quantities, while macro elements are needed in large quantities in the body [9]. Macro and trace elements have very varied functions, e.g., as electrolytes, as enzyme constituents, and as building materials, e.g., in bones and teeth. Microelements include Fe, Zn, Mn, B, Cu, Ni, Al, Ti, Cr, and Pb. Macro elements include Ca, Na, K, P, and S.

However, it is of importance to note that the concentration of these micronutrients contained in plant organs depends on several interacting factors, including soil type, plant type/variety, and the growing locations and its management [2,10]. As with all crops, the nutritional composition of roots and tubers varies from place to place depending on the climate, soil, crop variety, and other factors [11]. The mineral content of cassava, as stated earlier, depends on several factors, including genetic and growing locational factors. It could be inferred that there is a direct relationship between mineral concentrations in cassava roots and the elemental concentration present in the soil [12]. Mohan et al. [13] studied the mineral composition of nineteen cassava varieties. They discovered that K is the most abundant mineral present in cassava tubers with values ranging from 3270–10,870 mg/kg. In contrast, the second most abundant mineral is Ca in the range of 1083–1858 mg/kg the average Ca content reported was 1502 mg/kg, and Mg in cassava tuber showed a range of 540–895 mg/kg with a mean content of 740 mg/kg. The S content ranged from 1000–1290 mg/kg while Fe content varied from 68–248 mg/kg, Mn content ranged from 15.40–33.2 mg/kg. Zn content ranged from 15.0–28.2 mg/kg, Cu content ranged from 2.2–4.2 mg/kg, and B content ranged from 4.12–7.62 mg/kg. Efforts of several researchers to increase the mineral content of cassava tubers have assisted in improving the nutritional load of some varieties [14,15]. Biofortified crops, such as yellow-fleshed cassava roots could provide an excellent source of certain micronutrients and macronutrients for people that could not be reached by other interventions.

The research aimed to evaluate the impact of genotype and growing locations on the macro and trace elements in yellow-fleshed cassava root genotypes. Cassava breeders, processors, and dieticians will use the research findings from this study in identifying which cassava genotypes is suitable for cultivation in all agro-ecologies and contain optimum nutrients capable of combating hidden hunger.

2. Materials and Methods

2.1. Fresh Cassava Root Genotypes

A total of 25 yellow-fleshed cassava genotypes from unlimited yield trial (UYT) and three white roots varieties used as checks (30,572, TME 1, and 91/02324), were collected from the multi-growing locations field trial at five different research farms of the International Institute of Tropical Agriculture, Ibadan, Nigeria. The five trial growing locations (Ibadan, Ubiaja, Onne, Mokwa, and Zaria) were from different agroecological zones of Nigeria (Table 1). The trial was planted in a complete block design with four replicates and during the rainy season in July. There was no application of fertilizers or herbicides during the experiment. The weeding was done manually when necessary. The roots of all the yellow-fleshed genotypes and white-fleshed varieties were harvested at 12 months after planting (MAP). The two middle rows were harvested per plot, and only two replications were processed for analysis.

Table 1. Characteristics of the five trial growing locations

Location	GPS	Agroecological Zone	Elevation	Annual Average Rainfall	Annual Average Temperatures	Soil Type
Ibadan	7°38' N, 3°89' E	Forest-savanna transition	227 masl	1312 mm	20.3–33.8 °C	Ferric Luvisols
Onne	4°41' N, 7°09' E	Humid forest	44 masl	4420 mm	22.3–33.4 °C	Dystic Nitosols
Ubiaja	6°65' N, 6°38' E	Subhumid forest	287 masl	1186 mm	19.9–32.6 °C	Dystic Nitosols
Mokwa	9°28' N, 5°05' E	Southern Guinea savanna	132 masl	1149 mm	18.1–37.3 °C	Dystic Nitosols
Zaria	11°16' N, 7°63' E	Northern Guinea savanna	687 masl	1076 mm	13.9–35.5 °C	Ferric Luvisols

2.2. Sampling Method

The sampling and sample preparation of the cassava roots for mineral analysis was done using the adapted method described by [16]. There was a selection of three storage roots (large—900–2300 g, medium—500–899 g, and small—200–499 g) and washed thoroughly with potable water to remove dirt and adhered sand particles before air-dried on a clean concrete floor [15]. The peeling of the storage roots was done manually using a stainless-steel knife, and the peeled roots were rinsed with deionized water to remove any contaminants. After peeling, the roots were cut longitudinally (from the proximal end to the distal end) into four equal parts. Two opposite sections from each root of each genotype were taken, combined, manually chopped into small pieces, and mixed thoroughly. A batch of samples from this lot was chosen for the mineral profiling analysis.

2.3. Determination of Trace Macroelement Content

The processed samples were placed in a petri dish and dried in an uncorroded conventional oven at 40 °C for 3 days. After drying, the samples were packed in labelled, mineral-free paper envelopes. The trace and the macro element content were determined using Inductively Coupled Optical Emission Spectrometry (ICP-OES) according to the validated method described by [17] and [15]. About 0.30 g of the dried sample was weighed to into 50 mL screw-cap polypropylene tubes and added 2.0 mL HNO₃ and 0.5 mL H₂O₂ to initiate the sample digestion. The samples were made to a final volume of 25 mL with 18 MΩ.cm water before injection. The sample flow rate was 2.0 mL min^{−1}, and the total analysis time per sample is approximately 2.5 min. Each sample was run in duplicate. Al concentrations of more than 5–10 mg/kg are frequently associated with contaminant Fe. Thus, the concentration of Al of less than 10 mg/kg was used to determine the quality of the data, and all samples had an Al level of

less than 10 mg/kg. The trace elements identified in all the genotypes and varieties investigated were Fe, Mn, B, Cu, Mo, Co, Ni, Zn, and Al, respectively. Besides, the macro elements identified in all the genotypes and varieties investigated were Ca, Mg, Na, K, P, and S, respectively.

2.4. Statistical Analysis

The data generated in this study were subjected to analysis of variance (ANOVA) using Linear Regression Model to establish the relationship between the independent variables and the dependent variables and descriptive statistics using the Statistical Analysis System Software (Cary, NC, USA) [18]. Means were separated using Fisher's protected least significance difference test at $p < 0.05$. Principal component analysis (PCA) and hierarchical cluster analysis (HCA) were also utilized.

3. Results and Discussion

3.1. Effect of Genotype and Growing Locations on the Trace Elements of Yellow-Fleshed Cassava Roots

Goodness-of-Fit Statistics for Minerals across the Five Growing Locations

The linear regression model used for the analysis of variance had a regression coefficient (R^2) for all the dependent variables (genotype and growing locations) that ranged from 0.5–0.9. It implies that 50–90% of the variability of the independent variables (trace elements) were explained by the three explanatory variables (genotype, growing locations, and two-way interaction between the genotype and growing locations). The mean square error (MSE) ranged from 0.1–30.7, and Mn had the highest while Cr had the lowest. The lower the MSE, the better the model in predicting the variability of the trace elements. The macro elements in the samples had a regression coefficient (R^2) for all dependent variables ranging from 0.2–0.7, implying that 20–70% variability of the independent variables for macro elements were explained by the three explanatory variables as described earlier. Five macroelements were recorded for the yellow-fleshed cassava genotypes studied. The mean square error (MSE) ranged from 4.1×10^3 – 1.47×10^7 , with the lowest for S and the highest for K. The high values for the MSE suggest the model may not be good enough in predicting the variability of macroelements.

Table 2 shows the mean square (MS) of the analysis of variance for the trace and macro elements of yellow-fleshed cassava roots from five growing locations. Genotype and growing locations had a highly significant ($p < 0.001$) effect on all the trace elements except Ti and Cr. However, there was no significant ($p > 0.05$) difference of genotype $\times$ growing locations interaction on all the trace elements, but a highly significant ($p < 0.001$) effect was observed for Pb and a slight significance ($p < 0.01$) for Zn. It implies that the growing locations and genotype affect the trace element contents of the yellow-fleshed cassava roots. Among the explanatory variables, the variable growing locations is the most influential for all the trace elements except B and Ti, where genotype is the most influential variable. Thus, the growing locations had a strong effect on the concentration of Fe, Zn, Mn, Cu, Ni, Al, B, and Pb, while genotype had a strong influence on B and Ti concentrations. This can be related to the type of soil used for planting. Acidic soils have been reported to be high in Al and Mn concentrations [19,20]. Genotype $\times$ growing locations interaction was highly significant for both Fe and Zn among 21 sweet potato genotypes from five growing locations in the eastern region of China [21]. Furthermore, another group of researchers evaluated genotype $\times$ growing locations interaction in recombinant inbred lines of wheat bread grains for Fe and Zn concentration. Genotype $\times$ growing locations interaction ($G \times E$) for Fe and Zn accounted for 29.46% and 23.24% of the total sum of squares, respectively, and the magnitude of $G \times E$ interaction was relatively high [22].

Table 2. Mean squares (MS) of analysis of variance (ANOVA) for trace and macroelements of yellow-fleshed cassava roots across five growing locations

Source	Parameters										
		Fe	Zn	Mn	B	Cu	Ni	Al	Ti	Cr	Pb
	DF	MS	MS	MS	MS	MS	MS	MS	MS	MS	MS
Genotype	27	60 ***	39.7 ***	146.5 ***	2.3 ***	4.9 ***	0.7 ***	30.1 **	0.4	0.1	0.8
Locations	4	285.4 ***	536.3 ***	1773.8 ***	1.33 *	54.7 ***	7.9 ***	85.2 **	0.1	0.2	13.4 ***
Genotype × Location	107	24.3	11.7 *	37.2	0.5	1.0 ***	0.3	14.0	0.3	0.2	0.8 ***
Source	DF	Ca MS	Mg MS	Na MS	K MS	P MS	S MS				
Genotype	27	2,378,351 ***	3,054,187 ***	133,832 **	63,007,980 ***	153,215 ***	23,015 ***				
Locations	4	22,331,768 ***	20,946,861 ***	589,581 ***	225,500,5890 ***	9,228,146 ***	200,022 ***				
Genotype × Location	107	633,863	896,698 ***	88,495 *	19,347,235 **	81,564 *	6228 ***				

DF—Degree of Freedom, ***, **, * Significant at $p \leq 0.001$, $p \leq 0.01$ and $p \leq 0.05$ respectively; ns—not significant $p > 0.05$.

The growing locations had highly significant ($p < 0.001$) effects on all macroelements. In comparison, genotype had a highly significant ($p < 0.001$) effect on Ca, K, and S, a significant effect ($p < 0.05$) on P, and a slightly significant effect on Na. Genotype and growing locations bring significant ($p < 0.05$) information to explain the variability of the dependent variable Ca, Mg, Na, K, P, and S. However, among the factors, the growing location is the most influential for the macroelements. Ref. [23] reported that cassava thrives best on soils with adequate macronutrients—especially Ca, P, and K—coupled with the ability to extract high concentrations of nutrients from the soil. K plays an essential role in the transport of starch and sugar from the above-ground parts of the plant and, thus, their accumulation in tubers, the reason why it is the macroelement with the highest concentration across all locations and genotypes [24]. The leaves dropping from cassava plants as maturity progresses usually act as mulch for the plant, and traditional farm practices like mulching increases soil P and K, which in turn prevents a decline in soil Ca and Mg [19,25]. However, not all soil types have these macro elements in required quantities to support cassava growth because there seems to exist a direct relationship between the concentration of macroelements in cassava roots and their abundance in the soil [24]. Ultimately, a plant's ability to obtain adequate amounts of macronutrients depends critically on the availability of these minerals in the soil concerning factors such as precipitation, temperature, wind, soil type, and soil pH [26].

3.2. Trace and Macroelement Content of Yellow-Fleshed Cassava Roots by Genotype

Tables 3 and 4 show the trace and macroelement content of yellow-fleshed cassava roots by genotype across the five growing locations. The concentrations of Fe, Zn, and Mn were found to be higher than B, Cu, Ni, Al, Ti, Cr, and Pb, and implies that cassava roots could be a good source of Fe, Zn, and Mn which are highly essential micronutrients. The Fe content ranged from 2.6–4.9 mg/kg (mean 10.4 ± 5.5 mg/kg), Zn from 3.4–30.9 mg/kg (mean 11.1 ± 4.4 mg/kg), and Mn from 1.6–57.8 mg/kg (mean 14.7 ± 8.3 mg/kg). Also, B had a mean value of 2.0 ± 0.8 mg/kg, Cu 3.2 ± 1.4 mg/kg, Ni 1.3 ± 0.6 mg/kg, Al 5.8 ± 4.1 mg/kg, Ti 0.5 ± 0.5 mg/kg, Cr 0.4 ± 0.3 mg/kg, and Pb 1.3 ± 0.9 mg/kg. On the other hand, among the macroelements, K has the highest concentrations ranging from 2100–43,000 mg/kg (mean $10,637 \pm 7172$), Ca from 410–7600 mg/kg (mean 1604 ± 1057), Mg from 4600–8100 mg/kg (mean 1576 ± 1077), P from 450–3100 mg/kg (mean 967 ± 458), S, the lowest, from 138–700 mg/kg (mean 297 ± 94), while the element with the widest concentration range is Na, 4–4100 mg/kg (mean 167 ± 297). Chavez et al. [27] reported the average concentrations of five

trace elements (Fe, Zn, Mn, B, Cu) and six macroelements (Ca, Mg, Na, P, K, S) for 20 yellow-fleshed cassava genotypes. For Fe, the average concentration was 9.6 mg/kg, Zn 6.4 mg/kg, Mn 1.2 mg/kg, B 2.4 mg/kg, and Cu 2.2 mg/kg, which are low compared with values in this study except for boron. The concentration reported for the selected macroelements are Ca 590 mg/kg, Mg 1153 mg/kg, Na 66.4 mg/kg, P 1284 mg/kg, K 8903 mg/kg, and S 273 mg/kg. All the values fall within the range, as analyzed in this study.

Table 3. Mean value of trace elements of yellow-fleshed cassava roots by genotypes and across five growing locations

Genotype	Parameters (mg/kg)									
	Fe	Zn	Mn	B	Cu	Ni	Al	Ti	Cr	Pb
01/1442	14.9 ab	15.7 ab	25.8 a	3.6 a	4.7 ab	1.5 abc	9.4 a	1.1 a	0.4 a	1.6 ab
01/1413	14.7 ab	12.4 abcd	17.5 abc	2.7 abcd	3.4 bcdef	1.8 ab	8.6 a	0.6 a	0.7 a	1.6 ab
01/1663	13.9 ab	12.5 abcd	17.2 abc	2.6 abcd	3.1 cdefg	1.2 abc	7.9 a	0.9 a	0.5 a	2.3 a
01/1115	15.6 a	15.9 a	17.0 abc	2.4 bcde	4.0 abcd	1.9 a	5.8 a	0.5 a	0.4 a	1.4 ab
01/1273	12.5 ab	13.7 abc	16.4 abc	2.8 abc	4.4 abc	1.3 abc	9.0 a	0.7 a	0.3 a	1.6 ab
01/1335	11.1 ab	12.6 abcd	21.5 ab	2.6 abcd	3.4 bcdef	1.5 abc	8.4 a	0.7 a	0.4 a	1.2 ab
94/0330	12.3 ab	11.8 abcd	14.7 bc	2.3 bcde	3.0 defg	1.6 abc	6.0 a	0.7 a	0.5 a	1.3 ab
01/1610	12.9 ab	12.1 abcd	18.2 abc	2.2 bcde	3.8 abcde	1.4 abc	7.1 a	0.5 a	0.3 a	1.5 ab
01/1331	10.4 ab	12.1 abcd	19.3 abc	2.8 ab	3.1 cdefg	1.1 abc	7.6 a	0.7 a	0.4 a	1.2 ab
95/0379	11.2 ab	12.5 abcd	9.2 c	2.1 bcde	3.3 cdefg	1.3 abc	7.2 a	0.5 a	0.5 a	1.2 ab
01/1371	9.4 ab	10.3 cd	14.7 bc	2.1 bcde	3.5 bcdef	1.5 abc	6.1 a	0.5 a	0.4 a	1.5 ab
01/1277	10.2 ab	11.3 abcd	16.0 bc	2.3 bcde	2.6 efg	1.4 abc	6.8 a	0.7 a	0.3 a	1.2 ab
01/1412	11.5 ab	11.4 abcd	14.7 bc	2.7 abcd	2.6 efg	0.9 c	5.1 a	0.6 a	0.3 a	1.4 ab
01/1649	11.2 ab	9.8 cd	14.7 bc	2.1 bcde	3.0 cdefg	1.3 abc	5.6 a	0.4 a	0.3 a	1.3 ab
01/1380	8.4 ab	10.5 cd	10.3 c	2.2 bcde	3.5 bcdef	1.2 abc	5.0 a	0.4 a	0.3 a	1.3 ab
01/1235	12.1 ab	8.9 cd	11.4 c	2.0 bcde	2.0 g	1.0 bc	5.2 a	0.7 a	0.5 a	1.1 ab
98/2132	8.0 ab	10.9 bcd	15.1 bc	2.1 bcde	3.3 cdefg	1.4 abc	4.2 a	0.4 a	0.3 a	1.1 ab
01/1206	8.4 ab	12.5 abcd	16.1 bc	1.8 bcde	2.3 fg	1.0 bc	5.6 a	0.6 a	0.3 a	1.2 ab
01/1404	8.0 ab	10.9 bcd	15.6 bc	2.1 bcde	2.9 defg	1.1 abc	4.7 a	0.4 a	0.3 a	1.6 ab
01/1368	10.1 ab	9.9 cd	11.2 c	1.8 bcde	3.1 cdefg	1.2 abc	4.3 a	0.3 a	0.3 a	1.7 ab
01/1224	9.7 ab	9.7 cd	14.3 bc	1.8 bcde	3.1 cdefg	1.1 abc	5.1 a	0.4 a	0.3 a	1.1 ab
01/1646	7.9 ab	10.0 cd	15.4 bc	1.9 bcde	2.5 efg	1.6 abc	4.5 a	0.4 a	0.3 a	1.2 ab
94/0006	8.1 ab	9.6 cd	8.9 c	1.3 e	3.2 cdefg	1.6 abc	5.0 a	0.5 a	0.3 a	1.0 b
90/01554	7.6 b	8.4 d	13.4 bc	2.0 bcde	2.4 fg	1.1 abc	4.2 a	0.4 a	0.3 a	1.1 ab
96/1089A	7.6 b	10.5 cd	11.3 c	1.5 de	2.3 fg	1.1 abc	3.5 a	0.3 a	0.3 a	1.2 ab
30,572(chk)	9.4 ab	9.1 cd	9.5 c	1.6 de	2.7 defg	1.0 bc	4.2 a	0.4 a	0.4 a	1.2 ab
TME 1(chk)	7.7 ab	7.6 d	13.2 bc	1.6 cde	2.7 defg	1.1 bc	3.4 a	0.2 a	0.3 a	1.6 ab
91/02324(chk)	8.2 ab	9.0 cd	10.3 c	1.8 bcde	4.9 a	1.6 abc	3.7 a	0.5 a	0.4 a	1.1 ab
Minimum	2.6	3.4	1.6	0.6	0.9	0.5	0.2	0.1	0.3	1
Maximum	48.9	30.9	57.8	7.1	8	5.3	25	5.5	4.4	9.2
Mean	10.4	11.1	14.7	2.2	3.2	1.3	5.8	0.5	0.4	1.3
Standard deviation	5.5	4.4	8.3	0.8	1.4	0.6	4.1	0.5	0.3	0.9
Pr > F(Genotype)	***	***	***	***	***	***	**	ns	ns	ns

***, **, Significant at $p \leq 0.001$ and $p \leq 0.01$ respectively; ns -not significant $p > 0.05$. Means with different same letters within the same column are significantly different.

The Fe and Zn content of the yellow-fleshed cassava is comparable with some raw vegetables and even higher for some genotypes [2]. Interestingly, macro element concentrations for K, Ca, and Na in some yellow-fleshed cassava genotypes surpass values reported for popular cereals and soybean, a tested and trusted legume [2]. Fe and Zn are essential micronutrients for children and adults. Iron helps the human body in the creation of new red blood cells, and together with Cu, boosts red blood cells, thereby preventing anaemia [3]. Zn, on the other hand, strengthens the immune system, assists in protein synthesis, boosts gene expression, aids wound healing, among many more benefits. In Nigeria, about 75 percent of the preschool children and 67 percent of pregnant women are reported to be anemic, and 20 percent of children below five years suffer from zinc deficiency [28]. K plays an important role in the tuberization of cassava because it uses it as a primary osmolyte to osmotically adjust and

positively respond to water stress [29]. Therefore, this accounts for its abundance in cassava roots. K is a vasodilator due to its ability to reduce strain on the cardiovascular system. Also, it maintains fluid balance in the body for efficient metabolic activities. From the results, all the yellow-fleshed cassava genotypes will convincingly combat diseases brought about by deficiencies of these essential minerals, which are very low in the white varieties commonly consumed.

Table 4. Mean value of macro elements of yellow-fleshed cassava roots by genotypes and across five growing locations

Genotype	Ca (mg/kg)	Mg (mg/kg)	Na (mg/kg)	K (mg/kg)	P (mg/kg)	S (mg/kg)
01/1442	2060.0 abcde	2517.0 abc	410.4 ab	12,690.0 abcde	1121.0 ab	384.0 ab
01/1115	1665.0 abcdef	2236.0 abcd	285.0 ab	14,850.0 abc	959.0 ab	412.0 a
01/1610	1328.0 abcdef	2612.0 ab	138.4 ab	10,240.0 abcde	1097.0 ab	384.0 ab
01/1273	2388.0 ab	2972.0 a	567.7 a	15,780.0 a	873.0 b	236.0 de
01/1331	1795.9 abcdef	1798.6 abcde	100.3 b	15,006.9 abc	1407.4 a	297.9 abcde
01/1371	2591.0 a	1921.0 abcde	356.0 ab	11,720.0 abcde	849.0 b	289.2 bcde
01/1663	2176.0 abcde	2254.0 abcd	112.4 ab	15,610.0 ab	903.0 b	319.0 abcd
01/1335	2246.0 abcd	1230.0 cde	102.9 b	14,230.0 abcd	1109.0 ab	331.0 abcd
01/1404	1538.0 abcdef	1749.0 abcde	221.5 ab	11,390.0 abcde	991.0 ab	285.2 bcde
01/1413	1519.0 abcdef	1870.0 abcde	213.4 ab	9910.0 abcde	990.0 ab	318.0 abcd
01/1649	1096.0 bcdef	1473.0 bcde	90.3 b	11,510.0 abcde	964.0 ab	331.0 abcd
01/1224	2379.0 abc	1185.0 de	125.2 ab	8850.0 cde	790.0 b	332.0 abcd
01/1380	1019.0 def	1686.0 abcde	134.8 ab	9390.0 abcde	945.0 b	320.0 abcd
90/01554	1909.0 abcdef	1498.0 bcde	164.5 ab	8320.0 de	920.0 b	271.6 cde
01/1277	1684.0 abcdef	1433.0 bcde	203.7 ab	11,530.0 abcde	862.0 b	231.2 de
95/0379	1704.0 abcdef	1425.0 bcde	119.2 ab	10,100.0 abcde	948.0 ab	253.7 cde
94/0330	1550.0 abcdef	1521.0 bcde	87.3 b	7660.0 e	1106.0 ab	281.0 bcde
01/1412	1936.0 abcdef	1498.0 bcde	100.1 b	12,580.0 abcde	835.0 b	253.5 cde
01/1368	1175.0 bcdef	1500.0 bcde	187.9 ab	8660.0 cde	883.0 b	280.4 bcde
98/2132	1257.0 bcdef	1182.0 de	174.8 ab	9200.0 bcde	962.0 ab	275.1 cde
96/1089A	1642.0 abcdef	1028.0 de	119.7 ab	8670.0 cde	911.0 b	280.2 bcde
01/1646	1743.0 abcdef	1355.0 bcde	57.4 b	10,060.0 abcde	881.0 b	249.8 cde
01/1235	1076.0 cdef	1571.0 bcde	81.1 b	10,040.0 abcde	783.0 b	288.8 bcde
01/1206	1726.0 abcdef	1065.0 de	102.8 b	9980.0 abcde	881.0 b	208.5 e
94/0006	938.0 ef	1045.0 de	59.8 b	8220.0 de	1018.0 ab	304.4 abcde
91/02324(chk)	1111.0 bcdef	763.0 e	106.6 ab	9600.0 abcde	1094.0 ab	350.0 abc
30572(chk)	1094.0 bcdef	874.0 e	131.6 ab	6670.0 e	952.0 ab	287.6 bcde
TME 1(chk)	723.0 f	796.0 e	69.0 b	7750.0 de	1072.0 ab	254.9 cde
Minimum	410	460	4	2100	450	138
Maximum	7600	8100	4100	43,000	3100	700
Mean	1604	1576	167	10,637	967	297
Std. deviation	1057	1077	297	7172	458	94
Pr > F(Genotype)	***	***	*	***	**	***

***, **, * Significant at $p \leq 0.001$, $p \leq 0.01$ and $p \leq 0.05$ respectively; ns—not significant $p > 0.05$. Means with different same letters within the same column are significantly different.

Most of the yellow-fleshed genotypes had higher values of the trace elements than the three white-fleshed genotypes used as check samples (Table 4). This observation shows that the yellow-fleshed cassava roots compared favorably with the white-fleshed cassava roots in terms of microelement content, and it confers an additional nutritional benefit to the yellow-fleshed roots. There were no significant ($p > 0.05$) differences among the genotypes for Al, Ti, Cr, and Pb. It bolsters the earlier observation that genotype did not play a significant influence on these trace elements. However, they are all < 10.0 mg/100 g, unlike the Fe, Zn, and Mn that had higher values. The respective genotypes had the highest Fe (01/1442), Zn (01/1442), Mn (01/1442), B (01/1442), Cu (01/1442), Ni (01/1115), Al (01/1442), Ti (01/1442), Cr (01/1413), and Pb (01/1163) content, while the respective genotypes with the lowest values were: Fe (90/01554, 96/1089A), Zn (90/01554), Mn (94/006), B (94/006), Cu (01/1235), Ni (01/1412), Al (96/1089A), Ti (01/1368), and Pb (94/006). Genotype 01/1442 was outstanding because it showed

the highest values for almost all the minerals studied. However, genotypes 90/01554 and 90/1089A were the poor ones because they had the lowest amounts of Fe and Zn, the essential micronutrient in terms of their nutritional importance. Genotype 01/1442 is a potential pipeline genotype that could be advanced for release as a high micronutrient cassava variety. It will help to address the micronutrient deficiencies in the cassava growing and consuming communities of Nigeria.

The macro element content of yellow-fleshed cassava roots by genotype shows a general increase in values as compared to white-fleshed cassava roots. There were no significant differences among the genotypes for all the elements at a 5% significance level. However, these genotypes have the highest Ca (01/1371), Mg (01/1273), Na (01/1273), K (01/1273), P (01/1331), S (01/1115), while the respective genotypes exhibit the lowest Ca (94/0006), Mg (96/1089A), Na (01/1646), K (94/0330), P (01/1235), and S (01/1206). Genotype 01/1273 showed the highest values for half of the elements under observation, while no genotype can be qualified as inferior because none has low value across all elements. Ref. [30] reported there was no difference in concentrations of Fe and Zn in 10 cassava genotypes planted in two growing locations, namely, Nampula and Luipo. However, differences were significant at another growing location known as Umbeluzi with values for Fe ($p < 0.001$) and Zn ($p = 0.0162$) ranging from 8–24 mg/kg and from 8–19 mg/kg, respectively. These reported values are comparable with results in this research, although high values such as the ones reported were not recorded. This follows the same trend as observed that the growing locations have a highly significant effect ($p < 0.05$) in the trace element concentration of yellow-fleshed cassava (Table 5). Akin-Idowu et al. [31] reported a similar trend where eight yam genotypes were planted in two growing locations (Ibadan and Onne).

Table 5. Mean value of trace and macro elements of yellow-fleshed cassava roots by growing locations

Growing Locations	Parameters (mg/kg)									
	Fe	Zn	Mn	B	Cu	Ni	Al	Ti	Cr	Pb
Mokwa	12.8 a	11.3 b	22.1 a	2.3 a	2.1 d	1.6 a	6.4 a	0.6 a	0.4 a	1.5 b
Zaria	10.7 ab	15.6 a	19.2 a	2.3 ab	2.7 c	1.6 a	6.2 a	0.4 a	0.3 a	2.1 a
Ibadan	12.5 a	10.6 b	11.0 bc	2.2 ab	3.7 b	1.5 a	7.2 a	0.5 a	0.4 a	1.0 c
Ubiaja	8.7 bc	6.8 c	12.5 b	2.1 ab	2.8 c	1.0 b	5.4 ab	0.5 a	0.4 a	1.1 c
Onne	7.6 c	11.2 b	8.9 c	1.9 b	4.6 a	0.8 b	3.9 b	0.5 a	0.3 a	1.0 c
Pr > F (Locations)	***	***	***	**	***	***	**	ns	ns	***

Growing Locations	Parameters (mg/kg)					
	Ca	Mg	Na	K	P	S
Onne	1220.9 b	1816.1 b	247.7 a	4323.2 c	1668.2 a	388.7 a
Mokwa	2345.4 a	1945.7 ab	182.3 ab	14,408.9 b	840.2 b	221.0 c
Ubiaja	1059.1 b	2257.5 a	276.0 a	3769.6 c	924.5 b	292.3 b
Zaria	2258.0 a	967.8 c	34.5 c	18,199.4 a	612.6 c	288.5 b
Ibadan	1164.6 b	880.4 c	85.1 bc	12,908.9 b	795.0 b	293.4 b
Pr > F (Growing locations)	***	***	***	***	***	***

***, ** Significant at $p \leq 0.001$ and $p \leq 0.01$ respectively; ns—not significant $p > 0.05$. Means with different same letters within the same column are significantly different.

There was a significant ($p < 0.05$) difference between the mineral content of the samples between the growing locations. It can be observed that Mokwa, Zaria, and Ibadan growing locations have higher concentrations for all trace elements except for Onne that records the highest concentration of Cu (Table 5). In the case of the macro elements, Ubiaja and Onne locations had higher values for all the elements except K. The variations of the mineral content further established that mineral content could fluctuate greatly depending on genetic and climatic factors, agricultural procedures, the composition of the soil, and ripeness of the harvested crops, among other factors. This applies to both macro and trace elements. However, changes in the mineral content usually occur also in the processing of raw

materials, e.g., in thermal processes and material separation, which determine the available mineral content in the cassava products.

3.3. Principal Component (PC) and Cluster Analysis of the Trace and Macroelements of Yellow-Fleshed Cassava Roots Across Five Growing Locations

Table 6 showed the eigenvector of the first three axes of the Principal components (PCs) of the trace elements of yellow-fleshed cassava roots across the genotypes and growing locations. Four PCs (PC1 to 4) explained 75.2% of the data, but PC1 and PC2 accounted for above 50%, and PC1 had the highest eigenvalue of 3.5, while PC2 and PC3 had only 1.5 and 1.4, respectively. The minerals with common characteristics were loaded on each of the PCs. Thus, Fe, Mn, B, and Al were on PC1, Ti, Pb, and Mn on PC2, while Zn, Cu, and Cr were loaded on PC3. There was a positive and significant ($p < 0.05$) correlation among all the minerals on PC1, a positive and significant ($p < 0.05$) correlation between Mn and Pb. At the same time, both have a significant ($p < 0.05$) negative correlation with Ti on PC2. On PC3, Zn and Cu correlate negatively with Cr. In a recent study, Mn and B; and K and P were reported to be positively correlated [15]. There is a similar observation for Mn and B, but a negative correlation was observed for K and P in this study.

Table 6. Eigenvectors of the first three axes of the principal components (PC) analysis of the trace and macro elements of yellow-fleshed cassava roots

Parameters	PC1	PC2	PC3
Fe	0.4 ***	—	—
Zn	—	—	−0.4 ***
Mn	0.4 ***	0.4 ***	—
B	0.4 ***	—	—
Cu	—	—	−0.5 ***
Ni	—	—	—
Al	0.4 ***	—	—
Ti	—	−0.4 ***	—
Cr	—	—	0.5 ***
Pb	—	0.6 ***	—
Eigenvalue	3.5	1.5	1.4
Cumulative %	35.2	50.5	64.1
Parameters	PC1	PC2	PC3
Ca	—	0.6 ***	0.4
Mg	0.4 ***	0.5 ***	—
Na	0.4 ***	0.5 **	—
K	−0.5 ***	—	0.5 **
P	0.5 ***	—	0.4 **
S	0.4	—	0.6
Eigenvalue	2	1.7	1.1
Cumulative %	34.1	61.9	80.3

Cut-off point of eigenvector = 0.4. **, *** Significant at $p \leq 0.01$, and $p \leq 0.001$, respectively.

Furthermore, the principal component analysis for macroelements (PC1 to 3) explained the 80.3% of the data with PC1, having the highest eigenvalue of 2.0, PC2 1.7, and PC3 1.1. The macro elements contributing to PC1 are Mg, Na, K, P, and S, with K having a significant negative correlation with the others. Macro elements contributing to PC2 are Ca, Mg, and Na, all having significant positive correlations with one another. Ca, K, P, and S are loadings on PC3, having only K and P correlating significantly, while Ca and S have no significance. A similar result of negative correlation between Na and K in yellow-fleshed cassava root was reported by [15]. Thus, the macro elements in their respective loadings are related together, which will help in breeding for suitable agronomic traits with the desired macro element(s) in the yellow-fleshed cassava genotypes under consideration. It has been established

that macro elements have been absorbed more than trace elements by the stem and root crops [19,32], which is evident from their principal components. However, across the five growing locations, a highly significant difference is still observed for both trace and macro elements, which proves beyond a reasonable doubt that a breeder will have clarity in making decisions involving trace and macro elements as regards growing locations.

3.4. Multivariate Cluster Analysis

Table 7 and Figure 1 show a cluster of the yellow-fleshed cassava roots genotypes based on their dissimilarity for the 16 quantitative mineral traits investigated. This produced three groups of genotypes across all growing locations. Using the Euclidean distance for dissimilarity and Ward's method as the agglomeration method, show specific portions of variance explained by each of the trait (Figure 1). Cluster 1 had 12 yellow-fleshed genotypes with two white-fleshed varieties (TME1 and 91/02324) used as checks. It implies that these genotypes compared well with the two check varieties. Cluster 2 had eight yellow-fleshed genotypes, and Cluster 3 had five yellow-fleshed genotypes and one white-fleshed check variety (30,572). It was observed that the genotypes identified with the highest trace element concentration (01/1442) and macroelement concentration (01/1273) clustered in the same group (2) with genotype 01/1273 on top of the group while genotype 01/1442 is at the bottom. The genotypes identified as the poor ones (90/01554) and (96/1089A) with the lowest Fe and Zn are in groups 3 and 1, respectively. This strongly suggests that cassava genotypes in group 2 contain high concentrations of essential trace and macroelements.

Table 7. Cluster analysis of yellow-fleshed cassava roots across all growing locations using all parameters

Cluster	1	2	3
	01/1224	01/1273	90/01554
	94/0006	01/1404	01/1277
	94/0330	01/1412	01/1368
	96/1089A	01/1335	01/1380
	01/1663	01/1610	01/1235
	01/1115	01/1371	30572
	01/1413	01/1206	
	98/2132	01/1442	
	95/0379		
	01/1646		
	91/02324		
	TME 1		
	01/1649		
	01/1331		

Furthermore, none of the check samples, which are white varieties, fell into group 2, two of them were in group 1 and the other one in group 3. The check samples are typical of the genotypes identified as deficient in minerals compared with pipeline genotypes that are typically referred to as 'improved'. According to the dendrogram, there are several noticeable dissimilarities/differences among genotypes in groups 1, 2, and 3 (Figure 1).

3.5. Pearson Correlation Coefficients of Trace and Macroelements of Yellow-Fleshed Cassava Roots Genotypes Across Five Growing Locations

Table 8 shows the Pearson correlation coefficients of trace elements of yellow-fleshed cassava roots genotypes across five growing locations. It can be deduced that Fe, Al, Cr, Ni, Mn, Zn, B, and Ti, have strong significant ($p < 0.001$) correlations across the growing locations while Zn, Pb, and Cu have weak ($p < 0.05$) correlation across the growing locations. However, some also exhibit negative correlations, which can be a basis for predicting fertilizer formulation if the need for application arises. The increased concentration of a trace element could lead to the depletion of another; thus, care is needed during a

breeding program to ensure the purpose of biofortification is not forfeited. In consideration of the macroelements, there are mild to strong correlations across the growing locations, but several negative correlations are observed. S has a significant ($p < 0.05$) negative correlation with K and Ca ($r = -0.154$, -0.119), respectively, but a significant positive ($p < 0.001$) correlation with P ($r = 0.505$). This correlation shows the possibility of breeding yellow-root cassava varieties with high K and P mineral content. Also, K has a significant ($p < 0.05$ and $p < 0.01$) negative correlation with Mg and Na ($r = -0.170$ and -0.216). However, the negative correlations observed may not be detrimental in real-life situations because of research reports that micronutrients are stable across growing locations. Moreover, it implies that it is feasible to combine the high micronutrient content with high yield in breeding, unlike protein content and yield that are negatively correlated [33].

Figure 1. Dendrogram (Ward's minimum variance clusters) showing the relationship among different cassava roots genotypes across growing locations.

Table 8. Pearson correlation coefficients of trace and macro elements of yellow-fleshed cassava roots across genotypes and growing locations

Parameters	Ni	Cr	Fe	Al	Ti	B	Mn	Zn	Pb	Cu
Ni	1									
Cr	0.316 ***	1								
Fe	0.401 ***	0.636 ***	1							
Al	0.174 **	0.158 **	0.663 ***	1						
Ti	0.09	0.165 **	0.527 ***	0.728 ***	1					
B	0.054	0.186 **	0.426 ***	0.449 ***	0.317 ***	1				
Mn	0.350 ***	0.111	0.404 ***	0.394 ***	0.234 ***	0.512 ***	1			
Zn	0.287 ***	0.015	0.334 ***	0.310 ***	0.153 *	0.466 ***	0.503 ***	1		
Pb	0.169 **	−0.016	0.123 *	0.116	0.001	0.048	0.426 ***	0.3	1	
Cu	−0.027	−0.066	−0.038	0.006	0.068	0.147 *	−0.215 **	0.210 **	−0.217 **	1
Parameters	S	P	K	Ca	Mg	Na				
S	1									
P	0.505 ***	1								
K	−0.154 *	−0.327 ***	1							
Ca	−0.119 *	−0.156 *	0.531 ***	1						
Mg	0.086	0.263 ***	−0.170 *	0.289 ***	1					
Na	0.045	0.139 *	−0.216 **	0.124 *	0.597 ***	1				

*, **, *** Significant at $p \leq 0.05$, $p \leq 0.01$ and $p \leq 0.001$ respectively.

4. Conclusions

This study provides information on the concentration of trace and macroelements of yellow-fleshed cassava genotypes based on the interaction between their genetic make-up and the growing locations in which they are planted. It could be concluded that among the explanatory variables, the variable growing locations was the most influential on macro and trace elements. However, genotype had a strong influence on B and Ti. Most of the yellow-fleshed genotypes had higher values of the trace elements than the three white-fleshed genotypes used as check samples. The trace elements that positioned themselves to be the most abundant in yellow-fleshed cassava regardless of the genotype are Fe, Mn, and B. At the same time, Mg, Na, K, P, and S showed up as the most abundant for macroelements. The genotypes 01/1273, 01/1404, 01/1412, 01/1335, 01/1610, 01/1371, 01/1206, and 01/1442 were clustered together and identified to be different from the white-fleshed varieties used as checks and were promising pipelines. However, from the group, genotypes 01/1442 and 01/1273 have outstanding trace and macroelement concentrations, respectively. They could be advanced in the breeding cycle as new varieties to combat hidden hunger in Nigeria. However, all genotypes of the yellow-fleshed cassava can be predicted to perform well in any of the growing locations in terms of their trace and macroelement concentrations. The research findings from this study supported the claim that yellow-fleshed cassava roots could be a good source of certain micronutrients and macronutrients capable of combating hidden hunger in most of the cassava growing and consuming areas of the developing countries. However, there is a need for the investigation of the effect of traditional processing and bioavailability of the most important minerals of the identified yellow-fleshed cassava genotypes.

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Data Availability: All data are deposited into CKAN, an IITA data repository, and can be accessed <http://data.iita.org/>. Also, the data could be made available on request from the corresponding author.

References

1. Alamu, E.O.; Maziya-Dixon, B.; Dixon, A.G. Evaluation of proximate composition and pasting properties of high-quality cassava flour (HQCF) from cassava genotypes (*Manihot esculenta* Crantz) of β -carotene-enriched roots. *LWT* **2017**, *86*, 501–506. [[CrossRef](#)]
2. Montagnac, J.A.; Davis, C.R.; Tanumihardjo, S.A. Nutritional value of cassava for use as a staple food and recent advances for improvement. *Compr. Rev. Food Sci. Food Saf.* **2009**, *8*, 181–194. [[CrossRef](#)]
3. Temesgen, Z.T.; Bakalo, B.; Tamirat, T. Medicinal, Nutritional and Anti-Nutritional Properties of Cassava (*Manihot esculenta*). *Acad. J. Nutr.* **2019**, *8*, 34–46. [[CrossRef](#)]
4. Chandrasekara, A.; Joseph Kumar, T. Roots, and tuber crops as functional foods: A review on phytochemical constituents and their potential health benefits. *Int. J. Food Sci.* **2016**, 3631647. [[CrossRef](#)] [[PubMed](#)]
5. Von Grebmer, K.; Saltzman, A.; Birol, E.; Wiesmann, D.; Prasai, N.; Yin, S.; Yohannes, Y.; Menon, P. 2014 *Global Hunger Index: The Challenge of Hidden Hunger*; International Food Policy Research Institute: Bonn, Germany; Washington, DC, USA; Dublin, Ireland, 2014.
6. Ayetigbo, O.; Latif, S.; Abass, A.; Müller, J. Comparing characteristics of root, flour, and starch of biofortified yellow-flesh and white-flesh cassava variants, and sustainability considerations: A review. *Sustainability* **2018**, *10*, 3089. [[CrossRef](#)]

7. Talsma, E.F.; Brouwer, I.D.; Verhoef, H.; Mbera, G.N.K.; Mwangi, A.M.; Demir, A.Y.; Maziya-Dixon, B.; Boy, E.; Zimmermann, M.B.; Melse-Boonstra, A. Biofortified yellow cassava and vitamin A status of Kenyan children: A randomized controlled trial. *Am. J. Clin. Nutr.* **2016**, *103*, 258–267. [CrossRef] [PubMed]
8. Belitz, H.D.; Grosch, W.; Schieberle, P. *Food Chemistry*, 4th ed.; Springer: Berlin/Heidelberg, Germany, 2009; p. 1116. [CrossRef]
9. CÎRÎÎNĂ, P.D.; CĂPĂÎÎNĂ, C. Considerations on the Role of Mineral Substances in the Organism. *Synthesis* **2018**, *4*, 322–326.
10. Gregory, P.J.; Wahbi, A.; Adu-Gyamfi, J.; Heiling, M.; Gruber, R.; Joy, E.J.; Broadley, M.R. Approaches to reduce zinc and iron deficits in food systems. *Glob. Food Secur.* **2017**, *15*, 1–10. [CrossRef]
11. Woolfe, J.A.; Woolfe, W.J.; Poats, S.V. *The Potato in the Human Diet*; Cambridge University Press: Cambridge, UK, 1987; p. 248. ISBN 978-0521112994.
12. Odoom, E. The Potassium Status of Soils and Its Absorption by Cassava in Selected Farms in the Central Region. Ph.D. Thesis, University of Cape Coast, Cape Coast, Ghana, 2017. Available online: <http://hdl.handle.net/123456789/3436> (accessed on 15 March 2020).
13. Mohan, G.; Raju, J.; Shiny, R.; Abhilash, P.V.; Soumya, S. Biochemical, mineral, and proximate composition of Indian cassava varieties. *IJCS* **2019**, *7*, 1059–1065.
14. Ghandilyan, A.; Vreugdenhil, D.; Aarts, M.G. Progress in the genetic understanding of plant iron and zinc nutrition. *Physiol. Plant.* **2006**, *126*, 407–417. [CrossRef]
15. Alamu, E.O.; Maziya-Dixon, B.; Parkes, E.; Dixon, A.G. Variations of Macro-and Microelements in Yellow-Fleshed Cassava (*Manihot esculenta* Crantz) Genotypes as a Function of Storage Root Portion, Harvesting Time, and Sampling Method. *Appl. Sci.* **2020**, *10*, 5396. [CrossRef]
16. Maziya-Dixon, B.; Alamu, E.O.; Dixon, A.G. Variation in the evaluation of cis-and trans-carotene in yellow-fleshed cassava (*Manihot esculenta* Crantz) varieties as a function of the storage root portion and sampling method. *LWT* **2016**, *70*, 296–301. [CrossRef]
17. Wheal, M.S.; Fowles, T.O.; Palmer, L.T. A cost-effective acid digestion method using closed polypropylene tubes for inductively coupled plasma optical emission spectrometry (ICP-OES) analysis of plant essential elements. *Anal. Methods* **2011**, *3*, 2854–2863. [CrossRef]
18. SAS. *Statistical Analysis System Software, Version 9.4*; SAS Institute Inc.: Cary, NC, USA, 2012.
19. Howeler, R.H. Cassava mineral nutrition and fertilization. In *Cassava, Biology, Production and Utilization*; Hillocks, R.J., Thresh, J.M., Bellotti, A.C., Eds.; CABI Publishing: Wallingford, UK, 2002; pp. 115–147.
20. Vanlauwe, B.; Descheemaeker, K.; Giller, K.E.; Huising, J.; Merckx, R.; Nziguheba, G.; Wendt, J.; Zingore, S. Integrated soil fertility management in sub-Saharan Africa: Unravelling local adaptation. *Soil* **2015**, *1*, 491–508. [CrossRef]
21. Chen, J.; Lu, G.; Wang, G. Effects of genotype and growing location on iron and zinc contents in sweetpotato (*Ipomoea batatas* L.). *Crop Pasture Sci.* **2009**, *60*, 684–690. [CrossRef]
22. Gopalareddy, K.; Singh, A.M.; Ahlawat, A.K.; Singh, G.P.; Jaiswal, J.P. Genotype-growing locations interaction for grain iron and zinc concentration in recombinant inbred lines of a bread wheat (*Triticum aestivum* L.) cross. *Indian J. Genet.* **2015**, *75*, 307–313.
23. Ande, O.T. Soil suitability evaluation and management for cassava production in the derived Savanna Area of Southwestern Nigeria. *Int. J. Soil Sci.* **2011**, *6*, 142. [CrossRef]
24. Wang, Z.H.; Li, S.X.; Malhi, S. Effects of fertilization and other agronomic measures on nutritional quality of crops. *J. Sci. Food Agric.* **2008**, *88*, 7–23. [CrossRef]
25. Kirnak, H.; Kaya, C.; Higgs, D.; Gercek, S. A long-term experiment to study the role of mulches in the physiology and macro-nutrition of strawberry grown under water stress. *Aust. J. Agric. Res.* **2001**, *52*, 937–943. [CrossRef]
26. Maathuis, F.J. Physiological functions of mineral macronutrients. *Curr. Opin. Plant Biol.* **2009**, *12*, 250–258. [CrossRef]
27. Chávez, A.L.; Bedoya, J.M.; Sánchez, T.; Iglesias, C.; Ceballos, H.; Roca, W. Iron, carotene, and ascorbic acid in cassava roots and leaves. *Food Nutr. Bull.* **2000**, *21*, 410–413. [CrossRef]
28. Narayanan, N.; Beyene, G.; Chauhan, R.D.; Gaitán-Solís, E.; Gehan, J.; Butts, P.; Siritunga, D.; Okwuonu, I.; Woll, A.; Jiménez-Aguilar, D.M.; et al. Biofortification of field-grown cassava by engineering expression of an iron transporter and ferritin. *Nat. Biotechnol.* **2019**, *37*, 144–151. [CrossRef] [PubMed]

29. Omondi, J.O.; Lazarovitch, N.; Rachmilevitch, S.; Kukew, T.; Yermiyahu, U.; Yasuor, H. Potassium, and storage root development: Focusing on photosynthesis, metabolites, and soluble carbohydrates in cassava. *Physiol. Plant.* **2020**, *169*, 169–178. [[CrossRef](#)]
30. Burns, A.E.; Gleadow, R.M.; Zacarias, A.M.; Cuambe, C.E.; Miller, R.E.; Cavagnaro, T.R. Variations in the chemical composition of cassava (*Manihot esculenta* Crantz) leaves and roots as affected by genotypic and environmental variation. *J. Agric. Food Chem.* **2012**, *60*, 4946–4956. [[CrossRef](#)] [[PubMed](#)]
31. Akin-Idowu, P.E.; Asiedu, R.; Maziya-Dixon, B.; Odunola, A.; Uwaifo, A. Effects of two processing methods on some nutrients and anti-nutritional factors in yellow yam (*Dioscorea cayenensis*). *Afr. J. Food Sci.* **2009**, *3*, 086–093.
32. Ismail, F.; Anjum, M.R.; Mamon, A.N.; Kazi, T.G. Trace metal contents of vegetables and fruits of Hyderabad retail market. *Pak. J. Nutr.* **2011**, *10*, 365–372. [[CrossRef](#)]
33. Gregorio, G.B. Progress in breeding for trace minerals in staple crops. *J. Nutr.* **2002**, *132*, 500S–502S. [[CrossRef](#)]

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