Phenotypic Evaluation of Heritability, Agro-Morphological and Yield Characters of Sixteen *Amaranthus* Linn. Genotypes

Onuoha Samuel Ogechi and Olawuyi Odunayo Joseph

Genetics and Molecular Biology Unit, Department of Botany, University of Ibadan, Ibadan, Nigeria

Article history Received: 17-02-2017 Revised: 10-04-2017 Accepted: 20-07-2017

Corresponding Author: Onuoha Samuel Ogechi Genetics and Molecular Biology Unit, Department of Botany, University of Ibadan, Ibadan, Nigeria Email: sam4christ2005@gmail.com Abstract: The field experiment was conducted to evaluate the heritability, genetic variance, agro-morphological and yield characters of Sixteen Amaranthus genotypes. The seeds of the sixteen (16) genotypes of Amaranthus evaluated in this study were; NG/AA/MAY/09/027, NG/AO/11/08/042, NG/AO/11/08/039. NG/AA/03/11/010, NG/SA/DEC/07/0423, NG/SA/DEC/07/0412, NGB01667, NGB01601, NGB01283, NGB01271, NGB01276, NGB01259, NGB01644, NGB01234, NGB01613 and NGB01662. The results showed that there were variability performances in growth and yield characters of Amaranthus genotypes. NG/AA/MAY/09/027 and NG/AO/11/08/039 had the best growth characters while NG/AO/11/08/042 had best yield performance compared to other genotypes. The stem length recorded the best heritability estimate of 95.5% while weight of dry leaf, weight of fresh and dry inflorescent had least (47.7%). The plant height had a positive significant correlation with number of leaf (r = 0.53), leaf width (r = 0.57), number of branches (r =0.56) but a strong positive correlation with stem length (r = 0.97), stem girth (r = 0.75), number of inflorescent (r = 0.68), inflorescent length (0.64) and inflorescent width (r = 0.72). Prin. 1 accounted for the highest variation in growth and yield characters with proportion of 0.3376 and eigen value of 4.7269, while Prin. 14 was the least with proportion of 0.0003 and eigen value of 0.0038. Therefore, there could be genetic improvement of NG/AA/MAY/09/027 and NG/AO/11/08/039 genotypes for further improvement of Amaranthus.

Keywords: Amaranthus, Heritability, Agro-Morphological, Variance

Introduction

The genus Amaranthus of the order Caryophillalales comprises of more than 60 species C4 dicotyledonous herbaceous plants. Amaranthus species are cultivated in Central and South America, Africa and some parts of Asia as ornamentals, some are a source of highly nutritious pseudo-cereals and vegetables while others are notoriously weeds (Holm et al., 1997; Steckel, 2007). It has received attention due to its essential nutrients for the human diet (Tucker, 1986; Bressani et al., 1992). Amaranthus species are tolerant to infestation by herbivorous insects under field conditions and can grow successfully under varied soil and agro-climatic conditions such as bright sunlight, high temperatures and low moisture (Prakash and Pal, 1991; Brenner et al., 2010; Angel and Paulina, 2011). It can also tolerate a variety of unfavorable soil conditions such as high salinity, acidity, or alkalinity (Tucker, 1986). Besides

other crops, cultivation of this vegetable will not only increase food production but also provide balanced nutrition, food security, health security and poverty alleviation (Buragohain *et al.*, 2013).

Despite the perceived usefulness and untapped potentials of this vegetable, *Amaranthus* are underutilized making their potential economic value remaining "underexploited". It has also been reported that it has been neglected for many years by researchers, policy makers and funding agencies and thus currently threatened by extinction. Hence, improvement of this vegetable is highly needed to ensure maximum agronomic yield and high productivity of *Amaranthus* with a view to conserve the germplasm.

This study aimed at evaluating the heritability, genetic variance, agro-morphological and yield characters of Sixteen *Amaranthus* genotypes so as to improve the production of *Amaranthus spp.* for proper documentation of *Amaranthus* germplasm.



© 2017 Onuoha Samuel Ogechi and Olawuyi Odunayo Joseph. This open access article is distributed under a Creative Commons Attribution (CC-BY) 3.0 license.

Materials and Method

Experimental Site and Amaranthus Germplasm

This study was carried out at the Nursery Farm of the Department of Botany, University of Ibadan, Oyo state, Nigeria. The site lies between Latitude 7°02' 49" and 7°43' 21" N longitude 3°31' 58" and 4°08' 20" E with an altitude of 150 m in the valley at 275 m above sea level at moderate annual rainfall of 1,205 mm (Amanambu and Egbinola, 2013). The seeds of the sixteen Amaranthus spp. Genotypes were sourced from the National Centre for Genetic Resources and Biotechnology (NACGRAB), Moor plantation, Ibadan, Nigeria. The NG/AA/MAY/09/027, genotypes were NG/AA/03/11/010, NG/AO/11/08/042, NG/AO/11/08/039, NG/SA/DEC/07/0423, NG/SA/DEC/07/0412. NGB01667. NGB01601. NGB01283, NGB01271, NGB01276, NGB01259. NGB01644, NGB01234, NGB01613 and NGB01662.

Experimental Design, Plant Spacing and Planting Method

The Experiment was a complete Randomized Design with the polythene bags properly spaced at a distance of 65 cm between genotypes and 45 cm within genotypes. The young shoots were transplanted in pairs in each labeled polythene bags replicated four times. The seeds were first planted in the nursery through broadcasting for three weeks before transplanting in pairs to the well-labeled experimental polythene bags at the Nursery Farm of the Department of Botany, University of Ibadan, Oyo state, Nigeria. The cultivation was monitored and watered daily to resist drought.

Data Collection

After one week of transplanting, data taken weekly on growth characters included plant height (cm), number of leaves, stem length (cm), stem girth (cm) and leaf area (cm²). The plant height, stem length and leaf area were measured using a meter rule while stem girth was measured with a vernier caliper. Quantitative and qualitative data on flower characters were collected at maturity for six (6) weeks, this included number of inflorescence, inflorescence width (cm), inflorescence length (cm) and number of branches. The number of inflorescence and number of branches were done by counting, while inflorescence width and inflorescence length were determined by measurement using a metre rule. The inflorescence color and plant color were also determined by observation. After harvesting, the biomass of fresh and dry inflorescence and leaves were determined for each of the genotype using weighing balance. Heritability and genetic variance was also determined using these formulas:

$$Genotypic variance = \frac{Genotype \ MS - Error \ MS}{Replicate}$$

Phenotypic Variance : Genotypic Variance + Error MS

Statistical Analysis

The data were subjected to Analysis of Variance (ANOVA) and difference in means was separated using DMRT at 95% probability level (p<0.05). The relationship among the quantitative and qualitative traits were established using Pearson correlation coefficient and Principal Component Analysis (PCA). In addition, Heritability, Phenotypic Coefficient of Variance (PCV), Genotypic Coefficient of Variance (GCV) were also determined.

Results

Qualitative Traits in Amaranthus Genotypes

The qualitative traits observed in genotypes of Amaranthus are shown in Table 1. Genotypes NG/AA/MAY/09/027, NG/AA/03/11/010, NGB01601, NG/AO/11/08/039, NGB01271, NGB01276, NGB01259, NGB01644, NGB01234 and NGB01662 had a plant and spike/inflorescence color of green while NG/AO/11/08/042, NG/SA/DEC/07/0423, NG/SA/DEC/07/0412, NGB01667, NGB01283 and NGB01613 had plant and spike/inflorescence color of green but with a shade of purple. The spike/inflorescence colors were observed to vary from green to green with a shade of purple. The grain colors were observed to be TAN which is a light-brown color across all the genotype.

Mean Square Variance of Growth Characters in Amaranthus Genotypes

The result of the mean square variance of growth character for *Amaranthus* from Table 2 shows that the genotype and weeks after planting produced significant (p<0.01) effect on Plant height, Number of leaves, Stem length, Stem girth and Leaf width but non-significant on Leaf length for both genotype and week.

Mean Square Variance of Yield Characters in Amaranthus Genotypes

The result of the mean square variance of yield characters in Table 3 shows that the genotype and weeks after planting produced significant effect (p<0.01) on Number of Inflorescence, Inflorescence length, Inflorescence width, Number of branches, Fresh leaf biomass, Weight of dry leaf, Weight of fresh inflorescent and Weight of dry inflorescent.

S/N	Genotypes	Plant color	Inflorescence colour	Grain color
1	NG/AA/MAY/09/027	Green	Green	TAN
2	NG/AA/03/11/010	Green	Green	TAN
3	NG/AO/11/08/042	Green with a shade of purple	Green with some purple	TAN
4	NG/AO/11/08/039	Green	Green	TAN
5	NG/SA/DEC/07/0423	Green with a shade of purple	Green	TAN
6	NG/SA/DEC/07/0412	Green with a shade of purple	Purple	TAN
7	NGB01667	Green with a shade of purple	Green with some purple	TAN
8	NGB01601	Green	Green	TAN
9	NGB01283	Green with a shade of purple	Green	TAN
10	NGB01271	Green	Green	TAN
11	NGB01276	Green	Green	TAN
12	NGB01259	Green	Green	TAN
13	NGB01644	Green	Green	TAN
14	NGB01234	Green	Green	TAN
15	NGB01613	Green with a shade of purple	Green with some purple	TAN
16	NGB01662	Green	Green	TAN

Table 1. Qualitative traits in *Amaranthus* genotypes

Table 2. Mean square variance of growth characters at different stages in Amaranthus genotypes

Source of variation	Df	Plant height	Number of leaves	Stem length	Stem girth	Leaf length	Leaf width
Genotype	15	5388.43***	9750.21***	3929.57***	0.69***	130.70 ^{ns}	28.79***
Weeks	8	85011.33***	1480897***	50962.01***	2.53***	200.01 ^{ns}	43.57***
Replicates	3	3.97	4.14	2.67	0.03	1.08	0.17
Model	26	29266.49	10182.21	17947.99	1.18	137.07	30.04
Error	549	100.28	267.14	46.09	0.02	3.18	0.60
Corrected total	575						

* = Significant at p<0.05, ** = Highly significant at p<0.01, *** = Highly significant at p<0.001, ns = Non-significant, Df = Degree of freedom

	ld characters at different st	

							Weight of	Weight of	Weight of
Source of		Number of	Inflorescence	Inflorescence	Number	Fresh leaf	dry leaf	fresh	dry
Variation	Df	Inflorescence	length	width	of branches	biomass	biomass	inflorescent	inflorescent
Genotype	15	676.76***	317.62***	31.28***	72.69***	4.37***	0.44***	55.02***	10.65***
Weeks	8	13579.17***	12084.07***	718.93***	793.67***	614.38***	92.06***	6001.86***	1691.68***
Replicate	3	216.58 ^{ns}	14.14 ^{ns}	6.21 ^{ns}	8.05 ^{ns}	0.00^{ns}	0.00^{ns}	0.00^{ns}	0.00^{ns}
Model	26	4593.63	3903.05	239.97	287.07	191.56	28.58	1878.47	526.66
Error	549	116.32	44.87	4.52	5.95	0.96	0.09	12.03	
Corrected total	575								

* = Significant at p<0.05, ** = Highly significant at p<0.01, *** = Highly significant at p<0.001, ns = Non-significant, Df = Degree of freedom

Genotypic Effect of Growth Characters in Amaranthus Genotypes

The result of the mean performance of genotypic effect on growth character of *Amaranthus* reveals significant (p<0.01) effect on *Amaranthus* genotypes as shown in Table 4. NG/AA/MAY/09/027 was significantly higher for Plant height and Stem length compared to other genotypes. Also, leaf width produced significant effect for NGB01271 while Stem girth and Leaf length were significantly higher for NG/AO/11/08/039 but different from other genotypes. NGB01644 was significantly higher for Number of leaves than other genotypes.

Genotypic Effect of Yield Characters in Amaranthus Genotypes

The result in Table 5 shows that the genotypic effect of Amaranthus yield related character was significant at P < 0.05. NG/SA/DEC/07/0412 was significantly higher for Number of inflorescence and weight of fresh inflorescent yield compared to other genotypes. For Number of inflorescence NG/SA/DEC/07/0412 did not differ statistically from genotype NG/AO/11/08/042, NG/AO/11/08/039. NG/AS/DEC/07/0423, NG/AS/DEC/07/0412, NGB01601 and NGB01276. For weight of fresh inflorescent NG/SA/DEC/07/0412 did not statistically from genotypes differ NG/AO/11/08/039, NGB01667 and NGB01234. Also,

Inflorescence length was higher for NG/AO/11/08/039 while Inflorescence width and Fresh leaf biomass were significantly higher for NG/AO/11/08/042 but different from other genotypes. However, it did not differ statistically from genotypes NG/SA/DEC/07/0412 and NGB01601. For Inflorescence width, NG/AO/11/08/042 did not differ statistically from genotypes NG/AO/11/08/039, NG/SA/DEC/07/0423, NG/SA/DEC/07/0412, NGB01667 and NGB01283. For Fresh leaf biomass, NG/AO/11/08/042 did not differ statistically from genotypes NG/AA/MAY/09/027, NG/AO/11/08/039 and NGB01601. NGB01601 had higher Number of branches than other genotypes while NGB01667 is significant for weight of dry leaf biomass and weight of dry inflorescent. However, NGB01667 did differ statistically from NG/AO/11/08/039, not NG/SA/DEC/07/0423, NG/SA/DEC/07/0412, NGB01283, NGB01271 and NGB01662 for the weight of dry leaf biomass. Also, for the weight of dry inflorescent NGB01667did not differ statistically from genotypes NG/AO/11/08/039, NG/SA/DEC/07/0423, NG/SA/DEC/07/0412, NGB01283, NGB01271, NGB01234 and NGB01662.

Heritability and Genotypic variance of Growth and Yield Traits of Amaranthus Genotype

The result of the component of variance for growth and yield traits in *Amaranthus* shown in Table 6 reveals that the phenotypic variance of both growth and yield characters were higher than the genotypic variance in all the characters evaluated. The values for the phenotypic and genotypic variance were highest at Number of leaves but least at weight of dry leaf. The stem length recorded the best heritability estimate of 95.5% while weight of dry leaf, weight of fresh and dry inflorescent had least (47.7%).

Table 4. Genotypic Effect of Growth Characters in Amaranthus genotypes

	Plant height	Number of	Stem length	Stem girth	Leaf length	Leaf width
Genotype	(cm)	leaves	(cm)	(cm)	(cm)	(cm)
NG/AA/MAY/09/027	98.64 ^a	37.28 ^{efg}	87.31 ^a	1.21 ^b	15.74 ^e	7.49 ^{de}
NG/AA/03/11/010	51.70^{f}	23.61 ⁱ	41.48 ⁱ	0.69 ⁱ	12.13 ^g	4.83 ^j
NG/AO/11/08/042	88.59 ^{bc}	48.14 ^d	72.68 ^{cd}	1.11 ^{def}	16.36 ^{cde}	7.11 ^{fg}
NG/AO/11/08/039	78.09 ^d	60.92 ^{bc}	68.77 ^{ef}	1.33 ^a	21.03 ^a	7.89 ^{bc}
NG/SA/DEC/07/0423	85.12 ^c	32.28 ^{gh}	67.09 ^{ef}	1.22 ^b	18.29 ^b	8.11 ^b
NG/SA/DEC/07/0412	91.69 ^b	44.86 ^{de}	78.09^{b}	1.08 ^{efg}	18.12 ^b	7.57 ^{cde}
NGB01667	85.45°	29.33 ^{ghi}	73.16 ^c	1.14 ^{cde}	16.70 ^{cde}	7.98 ^b
NGB01601	73.85 ^{de}	42.00 ^{def}	62.49 ^g	1.11 ^{def}	17.29 ^c	7.84 ^{bcd}
NGB01283	83.85 ^c	22.81 ⁱ	69.67 ^{cde}	1.10def	14.73 ^f	7.25 ^{ef}
NGB01271	85.29 ^c	27.89 ^{hi}	69.76 ^{cde}	1.14 ^{cde}	16.07 ^{de}	8.47 ^a
NGB01276	72.64 ^e	34.97 ^{fgh}	62.63 ^g	1.02 ^{gh}	16.28 ^{de}	7.07^{fg}
NGB01259	72.93 ^{de}	56.56 ^c	62.28 ^g	1.07^{fg}	16.42 ^{cde}	6.49 ^{hi}
NGB01644	77.94 ^d	80.83 ^a	69.25 ^{def}	1.19 ^{bc}	16.10 ^{de}	6.81 ^{gh}
NGB01234	78.03 ^d	66.33 ^b	69.99 ^{cde}	1.17 _{cde}	16.68 ^{cde}	6.71 ^{gh}
NGB01613	75.15 ^{de}	32.39 ^{gh}	65.88^{f}	1.06 ^{fg}	16.74 ^{cd}	6.72 ^{gh}
NGB01662	54.41 ^f	37.11 ^{efg}	50.09 ^h	0.99 ^h	14.46^{f}	6.14 ⁱ

Mean with the same letter in the same column are not significantly at p≥0.05 according to Duncan Multiple Range Test (DMRT)

Table 5. Genotypic Effect of Yield Characters in Amaranthus genotypes

	Number of	Inflorescence	Inflorescence	Number	Fresh leaf	Dry leaf	Weight of fresh	Weight of dry
Genotype	Inflorescence	length (cm)	width (cm)	of branches	biomass (g)	biomass (g)	inflorescent (g)	inflorescent (g)
NG/AA/MAY/09/027	8.14 ^d	4.50 ^e	1.53 ^g	3.06 ^{bcde}	1.38 ^{abc}	0.36 ^{de}	2.33 ^{cde}	1.51 ^{bcd}
NG/AA/03/11/010	9.19 ^{cd}	5.51 ^e	2.26^{efg}	1.64 ^{fg}	0.73 ^d	0.16 ^f	1.69 ^e	0.67^{e}
NG/AO/11/08/042	20.31 ^a	11.24 ^{bc}	4.74 ^a	3.11 ^{bcde}	1.83 ^a	0.37 ^{de}	3.83 ^{bcd}	1.51 ^{cde}
NG/AO/11/08/039	20.69 ^a	15.61 ^a	4.14 ^{abc}	4.08^{ab}	1.56 ^{ab}	0.43 ^{abcd}	3.98 ^{abc}	1.81 ^{abcd}
NG/SA/DEC/07/0423	13.31 ^{bcd}	9.76 ^{dc}	3.78 ^{abcd}	1.06 ^{gh}	0.89 ^{cd}	0.56^{ab}	3.29 ^{cde}	2.52 ^a
NG/SA/DEC/07/0412	21.06 ^a	12.65 ^{abc}	4.43 ^{ab}	2.72 ^{cdef}	0.87^{cd}	0.43 ^{abcd}	5.76 ^a	2.15 ^{abc}
NGB01667	12.08 ^{bcd}	10.44 ^c	3.96 ^{abc}	0.39 ^h	0.65 ^d	0.58^{a}	4.00 ^{abc}	2.58 ^a
NGB01601	19.97 ^a	13.92 ^{ab}	3.19 ^{cde}	5.00 ^a	1.37 ^{abc}	0.36 ^{de}	2.99 ^{cde}	1.02^{de}
NGB01283	12.83 ^{bcd}	11.35 ^{bc}	3.81 ^{abcd}	0.00^{h}	0.73 ^d	0.44 ^{abcd}	2.96 ^{cde}	2.14 ^{abc}
NGB01271	14.00 ^{bcd}	9.76 ^{dc}	3.41 ^{bcd}	1.19 ^{gh}	1.02 ^{cd}	0.54 ^{abc}	2.61 ^{cde}	2.31 ^{ab}
NGB01276	15.86 ^{ab}	10.25 ^c	3.14 ^{cde}	2.50 ^{def}	1.13 ^{bcd}	0.24 ^{ef}	2.67 ^{cde}	1.01 ^{de}
NGB01259	12.50 ^{bcd}	9.36d ^c	2.80 ^{def}	3.97 ^{abc}	1.11 ^{bcd}	0.40 ^{bcde}	2.07 ^{de}	1.66 ^{bcd}
NGB01644	11.47 ^{bcd}	9.19d ^c	2.77 ^{def}	3.64 ^{bcd}	0.61 ^d	0.34 ^{de}	4.19 ^{abc}	1.63 ^{bcd}
NGB01234	8.14 ^d	6.50d ^e	1.78^{fg}	3.83 ^{abc}	1.02 ^{cd}	0.38 ^{cde}	5.49 ^{ab}	1.79 ^{abcd}
NGB01613	14.14 ^{bc}	9.34d ^c	3.18 ^{cde}	2.03 ^{efg}	0.88 ^{cd}	0.33 ^{de}	1.94 ^{de}	1.35 ^{cde}
NGB01662	12.33 ^{bcd}	6.59d ^e	2.23 ^{efg}	2.92^{bcde}	0.75 ^d	0.46^{abcd}	1.84 ^e	1.76^{abcd}

Mean with the same letter in the same column are not significantly at $p \ge 0.05$ according to Duncan Multiple Range Test (DMRT)

Source of v	ariation			G	enotypic	variance	(O^2g)	I	Phenotyp	ic varianc	$e(O^2p)$]	Heritabil	ity (%)		
Plant heigh	t			13	1322.040					1422.320				92.9		
Number of								2	2637.910			89.9				
Stem length					70.868				1016.970			95.5				
Stem girth	L			,	0.169			1	0.184				91.8			
												-				
Leaf length					31.879				35.064				90.9			
Leaf width					7.048				7.649				92.1			
Number of		ent		1	40.111				256.426			5	54.6			
nflorescen	length				68.186				113.060)		6	50.3			
nflorescen	t width				6.689				11.207			4	59.7			
Number of	hranches	2			16.683				22.638				73.7			
Weight of f					0.854				1.809				39.4			
Weight of c					0.0854				0.181				17.2			
	n y leal															
Wet yield	~				10.749				22.776				17.2			
Weight of c	ry infloi	rescent			2.081				4.409			2	17.2			
Table 7. Principal Characters	Components A Prin. 1	Analysis (PC. Prin. 2	/	and Yield Cha Prin. 4	racters of Ger Prin. 5	otypes of Amar Prin. 6	anthus spp. Prin. 7	Prin. 8	Prin. 9	Prin. 10	Prin. 11	Prin. 12	Prin. 13	Prin. 14		
PH	-0.2519	0.3319			0.4813	-0.2049	0.2745	-0.1748	0.043	-0.1351	-0.1294	-0.0597	0.603	-0.0037		
NL	-0.2319	0.3319			0.4813	0.6235	-0.0817	-0.1748	-0.1436	0.1331	0.1294	0.1214	0.003	0.0072		
SL	-0.2922	0.3213			0.4099	-0.1621	0.0811	-0.0191	-0.0523	0.0349	0.1219	0.1214	-0.7278	-0.0012		
SG	-0.1766	0.3794			-0.1888	-0.1767	-0.7272	0.1763	0.1235	-0.2963	-0.1586	-0.0441	0.0695	-0.0063		
LL	-0.1506	0.3805	-0.0073	0.0298	-0.5711	0.2148	0.4703	-0.1988	0.0316	-0.4241	-0.0607	-0.0051	-0.1188	-0.0039		
В	-0.2769	0.3092			-0.3689	-0.0291	0.0325	0.1959	-0.0882	0.7038	0.2215	-0.0612	0.1579	0.0095		
II	0.0187	0.2209		-0.0789	0.08	-0.0233	-0.0095	0.4288	-0.6027	-0.2084	0.0034	0.0117	0.0266	0.0059		
L	0.2836	0.2651		-0.0206	-0.0707	-0.1949	-0.2414	-0.7397	-0.2199	0.2231	-0.1318	0.0521	-0.0534	0.0115		
B	0.0173	0.2181		-0.2649	0.1395	0.3079	-0.1209	0.0414	0.6692	0.029	0.0841	0.0005	-0.0529	-0.0106		
NB Weight of	0.1374 0.3964	0.1679 0.1669		0.6181 -0.0233	-0.0439 -0.0132	-0.5166 -0.1627	0.273 -0.0099	0.2163	0.2964 0.0341	0.1835 -0.2185	-0.1041 0.8057	-0.0144 0.1976	-0.0217 0.1415	0.0106 0.1083		
resh leaf	0.3904	0.1009	-0.1371	-0.0233	-0.0132	-0.1027	-0.0099	-0.0000	0.0341	-0.2185	0.8037	0.1970	0.1415	0.1085		
Weight of Iry leaf	0.3974	0.1883	-0.1959	-0.1415	0.0103	0.0813	0.0702	0.1814	0.0107	0.0716	-0.2432	0.3997	0.0239	-0.692		
Weight of 0.38	410.2196	-0.171	-0.0271	0.1214	0.1177	0.0467	0.0682	-0.0428	0.0203	0.0147	-0.8392	-0.1314	-0.1201			
resh infloresco Weight of	ent 0.3895	0.1929	-0.2071	-0.1554	0.0372	0.1332	0.0747	0.1947	0.0163	0.0865	-0.3626	0.218	-0.0222	0.7029		
ry inflorescent	4 7260	2 0002	1.0577	1 2221	0.0770	0.55(2	0 4175	0 2240	0.2511	0.2107	0.1610	0.1054	0.0927	0.0020		
Eigenvalue Proportion	4.7269 0.3376	2.9902 0.2136		1.3231 0.0945	0.8779 0.0627	0.5563 0.0397	0.4175 0.0298	0.3249 0.0232	0.2511 0.0179	0.2197 0.0157	0.1618 0.0116	0.1054 0.0075	0.0837 0.006	0.0038		
H: Plant Height lumber of Branc able 8. Correlat	hes		,	U ,	,		th, LB: Leaf	Width, NI: N	Number of In	florescent, IL:	Inflorescent 1	Length, IB:	Inflorescent	Width, N		
										Weight of fresh leaf	Weight of dry leaf	Weight fresh dr		type		
Ή	PH	NL	SL	SG I	LL L	B NI	IL	IB	NB	inflorescent	inflorescent			Wee		
VL SL	0.5331* 0.9746**	0.5748*														
SL SG	0.9746** 0.7485**	0.5748* 0.5755*	0.7909**													

LL	0.2207	0.2027	0.2502	0.4257											
LB	0.5678*	0.2423	0.6090**	0.6603**	0.6210**										
NI	0.6840**	0.3955	0.6464**	0.4721	0.163	0.3416									
IL	0.6420**	0.3929	0.5913*	0.4965	0.1064	0.1994	0.7503**								
IB	0.7217**	0.3809	0.6752**	0.4808	0.1565	0.3681	0.8671**	0.7537**							
NB	0.5567*	0.5712*	0.5391*	0.4837	0.1152	0.1129	0.6351**	0.7012**	0.5449*						
Weight of	0.3688	0.2085	0.3267	0.3000	0-0.0442	0-0.0585	0.3766	0.7165**	0.3871	0.5296*					
fresh leaf															
Weight of	0.3889	0.2215	0.3443	0.3027	0-0.0254	0-0.002	0.3653	0.6918**	0.4023	0.4548	0.8973**				
dry leaf															
Weight of	0.3957	0.3137	0.3535	0.3116	0-0.0224	0-0.0342	0.374	0.6964**	0.4008	0.4927	0.8856**	0.9210**			
fresh inflorescent															
Weight of	0.3928	0.2279	0.348	0.3029	0-0.0229	0.0035	0.3561	0.6765**	0.4017	0.432	0.8722**	0.9940**	0.9321**		
dry inflorescent															
Genotype	0-0.1166	0.1491	-0.0882	-0.0121	-0.0595	0-0.1119	0-0.0466	0-0.0221	0-0.0641	0.0371	0-0.0475	0.0058	0-0.0083	0.0036	
Week	0.8919**	0.5257*	0.8741**	0.6648**	0.0642	0.3816	0.7279**	0.7595**	0.7530**	0.6756**	0.5175*	0.5271*	0.5097*	0.5208*	0
Replicate	0-0.0033	0-0.0035	0-0.0023	0-0.0207	0-0.0155	0-0.0032	0.0548	0-0.0133	0.0174	0-0.002	0.00000	0.0000	0.00000	0.0000	
* - Cionificant a		* _ Hiahh	Ciquificant	at a < 0.01.	DIL Dlout 1	Laight ML	Number of	Lagrage SI	Ctom Lo	noth CC.	Stom Cinth II	Loof Lonoth	ID: Loof Wid	th NL No.	mhor of

* = Significant at p<0.0 5, ** = Highly Significant at p<0.01; PH: Plant Height, NL: Number of leaves, SL: Stem Length, SG: Stem Girth, LL: Leaf Length, LB: Leaf Width, NI: Number of Inflorescent, IL: Inflorescent Length, IB: Inflorescent Width, NB: Number of Branches

Principal Components Analysis (PCA) of Growth and Yield Characters of Genotypes of Amaranthus spp.

The result from Table 7 delineates the *Amaranthus* genotype into fourteen principal component axes; Prin.

1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13 and 14. Prin. 1 which constituted the highest accounted for 0.3376 of the total proportion with eigen value of 4.7269, while Prin. 14 was the least with proportion of 0.0003 and eigenvalue of 0.0038. Weight of dry leaf from Prin. 1

had the highest eigen vector of 0.3973 while Number of leaves was the least with (-0.0793). Also Prin. 2 produced the highest eigen vector for Leaf length at 0.3805 while weight of fresh leaf biomass had the least at 0.1669. Prin. 3 produced the highest eigen vector at 0.5883 for number of inflorescent while Leaf length produced the least at (-0.0073). Prin. 4 produced the highest eigen vector at 0.6181 for Number of branches while Inflorescent length had the least at (-0.0206). Prin. 5 produced the highest eigen vector at 0.4813 for Plant height while weight of fresh leaf biomass had the least at (-0.0132). Prin. 6 produced the highest eigen vector at 0.6235 for Number of leaves while Number of Inflorescent had the least at (-0.0233). Prin. 7 produced the highest eigen vector at 0.4703 for leaf length while Weight of fresh leaf biomass had the least at (-0.0095). Prin. 8 produced the highest eigen vector at 0.4288 for number of inflorescent while Weight of fresh leaf had the least at (-0.0066). Prin. 9 produced the highest eigen vector at 0.6692 for inflorescent width while Weight of fresh inflorescent biomass had the least at (-0.0428). Prin. 10 produced the highest eigen vector at 0.7038 for Leaf width while Plant height has the least at (-0.1351). Prin. 11 produced the highest eigen vector at 0.8057 for Weight of fresh leaf biomass while leaf length has the least at (-0.0607). Prin. 12 produced the highest eigen vector at 0.3997 for Weight of dry leaf biomass while leaf length has the least at (-0.0051). Prin. 13 produced the highest eigen vector at 0.6030 for plant height while number of Branches has the least at (-0.0217). Prin. 14 produced the highest eigen vector at 0.7029 for Weight of dry inflorescent biomass while Stem length has the least at (-0.0012).

Correlation Co-efficient among Characters in Genotype of Amaranthus spp.

The correlation result is shown in Table 8. The plant height had a positive significant correlation with Number of leaf (r = 0.5331), Leaf width (r = 0.5678), Number of branches (r = 0.5567) and a strong positive correlation with Stem length (r = 0.9746), Stem girth (r = 0.7485), Number of inflorescent (r = 0.6840), Inflorescent length (0.6420), Inflorescent width (r = 0.7217) and Week (r = 0.8919). Number of leaf had a positive correlation with Stem length (r = 0.5748), Stem girth (r = 0.5755), Number of branches (r = 0.5712) and Week (r = 0.5257). In addition, Stem length showed a strong positive correlation with Stem girth (r = 0.7909), Leaf width (r =0.6090), Number of inflorescent (r = 0.6464), Inflorescent width (r = 0.6752) and Week (r = 0.8741) and a positive correlation with Inflorescent length (0.5913) and Number of branches (r = 0.5391). Moreover, Stem girth had a strong positive correlation with Leaf width (r = 0.6603) and Week (r = 0.6648). Leaf length had a strong positive correlation with Leaf width (r = 0.6210). Number of inflorescent had a strong positive correlation with Inflorescent length (r = 0.7503), Inflorescent width (r = 0.8671), Number of branches (r =

0.6351) and Week (r = 0.7279). Inflorescent length had a strong positive correlation with Inflorescent width (r =0.7537), Number of branches (r = 0.7012), Weight of fresh leaf (r = 0.7165), Weight of dry leaf (r = 0.6918), Weight of fresh inflorescent (r = 0.6964), Weight of dry inflorescent (r = 0.6765) and Week (r = 0.7595). Inflorescent width had a strong positive correlation with Week (r = 0.6756) and a positive correlation with Number of branches (r = 0.5449). Number of branches had a strong positive correlation with Week (r = 0.6756) and a positive correlation with Weight of fresh leaf (r = 0.5296). Weight of fresh leaf had a strong positive correlation with Weight of dry leaf (r = 0.8973), Weight of fresh inflorescent (r = 0.8856), Weight of dry inflorescent (r = 0.8722) and a positive correlation with Week (r = 0.5175). Weight of dry leaf had a strong positive correlation with Weight of fresh inflorescent (r = 0.9210), Weight of dry inflorescent (r =(0.9940) and a positive correlation with Week (r = 0.5271). Weight of fresh inflorescent had a strong positive correlation with Weight of dry inflorescent (r = 0.9321) and a positive correlation with Week (r = 0.5097). While Weight of dry inflorescent had a positive correlation with Week (r = 0.5208).

Dendogram Showing the Relationship of Yield Characters Among the Amaranthus Genotypes

The dendogram showing the relationship of Yield Characters among the *Amaranthus* Genotypes is shown in Fig. 1. There are two major clusters sub-divided into four (4) groups. Genotype NG/SA/DEC/07/0423 and NGB01271 is closely related but different from genotype NGB01613 while genotype NGB01667 and NGB01283 is similar than genotype NGB01276. Also, genotype NGB01259 and NGB01644 are closely related compared to genotype NGB01662. Again, genotype NG/AA/03/11/010 and NG/AA/MAY/09/027 are closely related than genotype NGB01234 while genotype NG/AO/11/08/042 and NG/SA/DEC/07/0412 are similar as also observed in genotype NG/AO/11/08/039 and NGB01601.

Dendogram Showing the Relationship of Growth Characters Among the Amaranthus Genotypes

The relationship of Growth Characters among the Amaranthus Genotypes is depicted in the dendogram as shown in Fig. 2. There are three major clusters subdivided into five (5) groups. Genotype NG/AO/11/08/042 and NG/SA/DEC/07/0412 are similar compared to genotype NG/AA/MAY/09/027 while Genotype NGB01667 and NGB01271 are similar but different from NG/SA/DEC/07/0423 and NGB01283. Also, genotype NGB01276 and NGB01613 are closely related compared NGB01601. to genotype Again, Genotype NG/AO/11/08/039 and NGB01234 are similar but different from genotype NGB01259 and NGB01644 while Genotype NG/AA/03/11/010 and NGB01662 are similar as seen in Fig. 2.

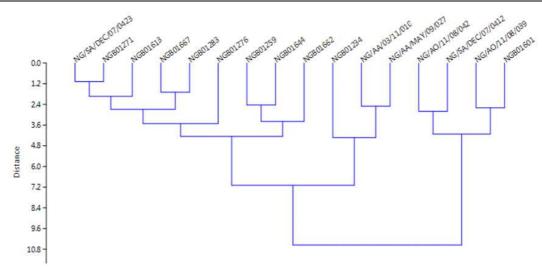


Fig. 1. Showing the relationship of yield characters among the Amaranthus genotypes

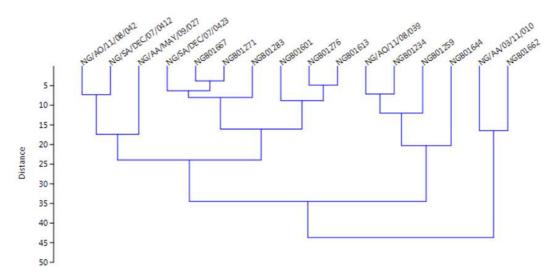


Fig. 2. Showing the relationship of growth characters among the Amaranthus Genotypes



Fig. 3. NG/AA/MAY/09/027 genotype showing the best growth character



Fig. 4. NG/AO/11/08/042 genotype showing the best yield attribute

Discussion

The findings from this study showed that there are variations in the performance of growth and yield characters studied among the *Amaranthus* genotypes. This is in accordance with the reports of Nwangburuka *et al.* (2012) and Olawuyi *et al.* (2014) who considered genetic variability as essential in crop breeding. The genotypic effect also had significant expression on the traits evaluated in *Amaranthus*. Variability in performance of genotypes also affected the growth performance of *Amaranthus*. The variations shown by the characters were due to high genetic diversity, differences of growing type and differences on the type of adaptation (Kulakow, 1987; Mujica and Jacobsen, 2003).

The best performance of growth and yield characters exhibited by NG/AA/MAY/09/027, NG/AO/11/08/039 and NG/AO/11/08/042 genotypes (Fig. 3 and Fig. 4) could be due to genetic variation of these genotypes. Selections based on this characters and the genetic diversity inherent in the plants could thus improve productivity considerably. These performances shown by *Amaranthus* also suggest hybridization breeding procedure for crop improvement with desired traits in the parents line.

The findings from correlation coefficient shows that the characters were mostly positively related as similarly observed by Olawuyi *et al.* (2012). The correlation between the characters implies that selection based on plant height will favour all growth and yield characters. This will enhance the rate of productivity and yield.

Prin 1 accounted for the highest variation as previously observed by Olowe *et al.* (2013) and Olawuyi *et al.* (2015). The results of the Principal components analysis reveals the pattern of variation

among the characters studied and the characters that accounted most for variation within a group of entries (Ogunbodede and Omueti, 1997; Aremu *et al.*, 2007). It implies that the Principal Component Analysis (PCA) can be quantified from the contribution of the different variable to each principal component as revealed by the eigen vector (Lezzoni and Pritts, 1991).

This phenotypic expressions might be due to environmental influences; exacerbating this problem is the presence of considerable morphological variation within cultivated populations (Sauer, 1967; Espitia, 1992). This shows that their genotypes and species genetic make-up played a huge role in the phenotypes expressed in this studies.

The phenotypic variance of both growth and yield characters were higher than the genotypic variance in all the characters studied. Heritability of growth traits were higher than yield traits in *Amaranthus* genotypes. This shows that the proportion of genotypic effect to phenotypic effect was higher at growth than maturity. This conforms to the report of Palaniappan *et al.* (1999) who observed an improvement in general crop performance. This supported the findings of Chadha and Paul, 1984; Gautam and Srinivas, 1992; Prasad *et al.*, 2004; Singh and Kumar, 2005; Babu and Patil, 2005 who reported high heritability and genetic advance for yield characters for *Solanum melongena.*

Conclusion and Recommendation

The variations in the genotypes could be sufficient basis for crop improvement. NG/AA/MAY/09/027, NG/AO/11/08/039 and NG/AO/11/08/042 are promising genotypes that could be selected and explored for future breeding in improvement of *Amaranthus* vegetable. This will further enhance proper documentation and conservation of *Amaranthus* germplasm.

Acknowledgement

Special thanks to Department of Botany, University of Ibadan, Oyo state, Nigeria for making their facilities accessible for the research work. Also, unreserved appreciation goes to National Centre for Genetic Resources and Biotechnology (NACGRAB), Moor plantation, Ibadan, Nigeria for providing the seeds.

Funding Information

This work was supported and funded by both authors.

Author's Contributions

This work was carried out in collaboration among the authors. Both authors participated actively and synergistically in all works, analysis and the entire process of the writing, editing and approval of the manuscript.

Ethics

The authors declares that there is no conflict of interest and this manuscript conforms to the ethical standards specified by the American Journal of Agricultural and Biological Sciences.

References

- Amanambu, C.A. and C.N. Egbinola, 2013. Climate variation assessment based on rainfall and temperature in Ibadan, South-Western. Nig. J. Environ. Earth Sci., 3: 2224-3216.
- Angel, H.O. and B.R. Paulina, 2011. Amaranth: A pseudo-cereal with nutraceutical properties. Curr. Nutriti. Food Sci., 7: 1-9.
- Aremu, C.O., M.A. Adebayo, M. Oyegunle and J.O. Ariyo, 2007. The relative discriminatory abilities measuring Genotype by environment interaction in soybean (Glycine max). Agric. J., 2: 210-215
- Babu, S.R. and R.V. Patil, 2005. Genetic variability and correlation studies inegg plant (*Solanummelongena L*.). Madras J. Aric. Res., 95: 18-23.
- Brenner, D.M., D.D. Baltensperger, P.A. Kulakow, J.W. Lehmann and R.L. Myers *et al.*, 2010. Genetic Resources and Breeding of *Amaranthus*. In: Plant Breeding Reviews, Jules J. (Ed.), John Wiley and Sons, Hoboken, ISBN-10: 0470650087, pp: 227-285.
- Bressani, R., A. Sanchez-Marroquin and E. Morales, 1992. Chemical composition of grain amaranth cultivars and effects of processing on their nutritional quality. Food Rev. Int., 8: 23-49.
- Buragohain, J., V.B. Singh, B.C. Deka, A.K. Jha and K. Wanshnong *et al.*, 2013. Collection and evaluation of some underutilized leafy vegetables of Meghalaya Indian. J. Hill Farming, 26: 111-115.

- Chadha, M.L. and B. Paul, 1984. Genetic variability and correlation studies in eggplant (*Solanummelongena L*.). Indian J. Hort., 41: 101-107.
- Espitia, E., 1992. Amaranth germplasm development and agronomic studies in Mexico. Food Rev. Int., 8: 71-86.
- Gautam, B. and T. Srinivas, 1992. Study on heritability, genetic advance and characters association in brinjal (*Solanummelongena L.*). SouthIndian Hort, 40: 316-318.
- Holm, L., J. Doll, E. Holm, J. Pancho and J. Herberger, 1997. World weeds: Natural histories and distribution. 1st Edn., John Wiley and Sons, Toronto.
- Kulakow, P.A., 1987. Genetics of grain amaranths. J. Hered. 78: 293-297.
- Lezzoni, A.F. and M.P. Pritts, 1991. Application of principal component analysis to horticultural research. Hort. Sci., 26: 334-338.
- Mujica, A. and S.E. Jacobsen, 2003. The genetic resources of Andean grain amaranths (*Amaranthus* caudatus L., A. cruentus L. and A. hypochondriacus L.) in America. Plant Genetic Resources Newsletter, 133: 41-44.
- Nwangburuka, C.C., O.A. Denton, O.B. Kehinde, D.K. Ojo and A.R. Popoola, 2012. Genetic variability and heritability in cultivated okra (Abelmoschusesculentus[L.] moench). Spanish J. Agric. Res., 10: 123-129.
- Ogunbodede, B.A. and O. Omueti, 1997. Regional research project on maize and cassava report on research activities on maize breeding and utilization. Sponsored Eleven Coastal West African Countries.
- Olawuyi, O.J., D.T. Ezekiel-Adewoyin, A.C. Odebode, D.A. Aina and G.E. Esenbamen, 2012. Effect of arbuscularmycorrhizal fungi (Glomusclarum) and organomineral fertilizer on growth and yield performance of Okra (Abelmoschusesculentus). African J. Plant Sci., 6: 84-88.
- Olawuyi, O.J., S.G. Jonathan, F.E. Babatunde, B.J. Babalola and O.S. Yaya *et al.*, 2014. Accession x treatment interaction, variability and correlation studies of pepper (*Capsicum spp.*) under the influence of ArbuscularMycorrhiza Fungus (Glomusclarum) and Cow Dung. Am. J. Plant Sci., 5: 683-690.
- Olawuyi, O.J., O.B. Bello, C.V. Ntube and A.O. Akanmu, 2015. Progress from selection of some maize cultivars' response to drought in the derived savanna of Nigeria. AGRIVITA J. Agric. Sci., 37: 8-17.
- Olowe, O.M., A.C. Odebode, O.J. Olawuyi and A.O. Akanmu, 2013. Correlation, principal component analysis and tolerance of maizegenotypes to drought and diseases in relation to growth traits. Am. Eurasian J. Agric. Environ. Sci., 13: 1554-1561.
- Palaniappan, S.P., A. Jeyabal and S. Chelliah, 1999. Evaluation of integrated nutrient management in summer sesame (*Sesamum indicum L*.). Sesame Saffl. Newslett.

- Prakash, D. and M. Pal, 1991. Nutritional and anti nutritional composition of vegetable and grain amaranth leaves. J. Sci. Food Agric., 57: 573-583.
- Prasad, M., N. Mehta, S.N. Diokshit and S.S. Nishal, 2004. Genetic variability, genetic advance and heritability in brinjal (*Solanummelongena L.*). Orissa J. Hort., 32: 26-29.
- Sauer, J.D., 1967. The grain amaranths and their relatives: A revised taxonomic and geographic survey. Annals Missouri Botanical Garden, 54: 103-137.
- Singh, O. and J. Kumar, 2005. Variability, heritability and genetic advance in brinjal. Indian J. Hort., 62: 265-267.
- Steckel, L.E., 2007. The dioecious *Amaranthus spp.*: Here to stay. Weed Technol., 21: 567-570.
- Tucker, J.B., 1986. Amaranth: The once and future crop. Bioscience, 36: 9-13.