

WWJMRD 2022; 8(03): 1-6 www.wwimrd.com International Journal Peer Reviewed Journal **Refereed** Journal Indexed Journal Impact Factor SJIF 2017: 5.182 2018: 5.51, (ISI) 2020-2021: 1.361 E-ISSN: 2454-6615 DOI: 10.17605/OSF.IO/K6V8H Clet Wandui Masiga^{1*}, Joseph Esimu¹, Brian Ssemugenze¹ Janet Nagasha¹ Emma Walimbwa¹, Daniel Mushikoma¹, Godfrey Kasiime¹, Demas Lukoye Kutosi¹Didas Mugisha1, Nipher Twikirize1, Solomon Malenje1, Eliakim Mwijuka1, Prima Akatwijuka1, Michael Kiboma1, Colophina Katemel, Stephen Okoth1 **Tropical Institute of Development Innovations** (TRIDI), Kampala, Uganda John Ndabagye Uganda Silk Producers Association, Kampala, Uganda. Ismail Baruhagara³, Geoffrey Sempiri³ Uganda National Council for Science and Technology, Kampala - Uganda Mugume Naboth Ngambe⁴ Sheema Local Government/Ankole Western University, Kabwohe, Uganda Sayed Mohammad Ali Mousavi⁵ Iran Agro Industrial Group, Kampala, Uganda Everlyn Nguku⁶ International Centre of Insect Physiology and Ecology (ICIPE), Nairobi, Kenya Emmanuel Omene⁷ Elioda Tumwesigye⁷ Ministry of Finance, Planning and Economic Development, Kampala Samuel Wangoda⁷ Ministry of Science, Technology & Innovation, Kampala, Uganda. Emmanuel Omene⁸ Elioda Tumwesigye⁸ Ministry of Finance, Planning and Economic Development, Kampala.

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Morphological Evaluation of mulberry genotypes across different agro-ecological conditions in Uganda

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Abstract

Mulberry (Morus spp.) is the chief source of food for the silkworm (Bombyx mori L.) which grows in diverse climatic conditions and is regarded as a unique plant on this earth due to its ability to be cultivated in different forms; multiple uses of leaf foliage and its positive impact in environment, bioremediation of polluted sites, conservation of water, prevention of soil erosion and improvement of air quality by carbon sequestering and pharmaceutical, food, cosmetic and health care industries has gained the attention of industrialists. Eight different mulberry genotypes were evaluated using eleven different traits related to agronomic and quality attributes as a feed for silkworm. The experiments were conducted in nine different districts across Uganda using a randomized complete block design. Analysis of variance was performed on quantitative characters to assess the morphological diversity. The results obtained revealed a high significant difference among the genotypes across all the locations. The traits that were significantly different across the mulberry accessions included lamina width and petiole length (P ≤ 0.01), petiole width and growth height (P \leq 0.05), internodes distance and number of branches (P \leq 0.001). Mysore Local and Thailand varieties had the highest plant height, Kanva2 had the highest number of branches, the longest bud length was observed in Local variety while the leaf yield was more in Thailand. The correlation coefficient showed a positive significant association of all the traits with leaf yield. The results obtained indicated that genotypes performed slightly differently in differently locations due to different factors studied. Overall, we therefore recommend farmers in these diverse ecological zones to grow these mulberry varieties, with good agronomic practices such as proper spacing, timely weeding and pest and disease management in order to obtain consistent high yields.

Keywords: Correlation coefficient, Genetic diversity, Mulberry, traits, Uganda.

Introduction

Mulberry (*Morus spp.*), the unique food source of domesticated silkworm (Bombyx mori L.), is a perennial plant propagated exclusively through stem cuttings (clones) and thus the acquired true-to-type plant population. The *morus spp* is a widely distributed crop in the temperate, sub-tropical and the tropical regions of the world and is highly adaptable to a wide range of climatic, topographical and soil conditions (Hoseini et al., 2018). Though it is believed that this species originated from the lower slopes of Himalayas in China and India, its current distribution across the regions reflects particularly its adaptability to a wide range of environmental conditions (Tikar and Vijayan., 2010). Plant species with a wide range of environmental adaptations often exhibit numerous morphological and physiological characteristics (Cordell et al., 1998) and this morphological variability in mulberry is believed to have contributed to its growth and survival under disruptive environmental conditions across the world (Gray, 1990). History states that this plant has been used in

sericulture industry throughout the world (Natic et al., 2015). In recent years, mulberry-based sericulture has been increasing in Rwanda, Kenya, Uganda and other East African countries.

Although mulberry has been in cultivation for centuries in different parts of the world, identification of the best performing varieties in Uganda is necessary. Proper and appropriate selection of the tropical cultivars based on the morphological characteristics/suitability is a necessity in availing reliable genetic resources for the farmers. This is because leaf is the economic product in mulberry cultivation unlike other agricultural crops. Mulberry leaf contributes 38.2% towards the production of a successful cocoon crop (Boraiah 1986).

Furthermore, Selection of suitable genotypes from gene pool therefore requires a thorough knowledge of morphological characteristics of different genotypes for farmer utilization (Yilmaz et al., 2012). Traditional methods for cultivar identification are based on the observation of phenotypic characteristics as they aid fast and simple evaluation of variability and; hence, are considered as an effective means of preliminary evaluation of assessing genetic diversity among morphologically distinguishable accessions (C'olic' et al., 2012). Furthermore, morphological characterization is the official method accepted for registration and protection of new cultivars (Ertan, 2007). Therefore, the main objective of this study was to assess the morphological and genetic variability of the mulberry cultivars available in Uganda and for sericulture use.

2. Materials and Methods

Eight mulberry genotypes (All foreign accessions as shown in Table 1) were evaluated in seven locations and different seasons. The locations were; Sheema, Mubende, Nakaseke, Bulambuli, Pallisa, Bukedea, Iganga, Bulambuli 2 and Zombo. These are located in eastern, north eastern and north western Uganda during 2019-2020. These locations represent the potential areas for sericulture in Uganda and are characterised by arid and semi-arid conditions. All the genotypes used in the study were obtained from China, India and Kenya (Table 1). In all locations, genotypes were planted in 4.0 m \times 5.0 m plots at a spacing of 1.0 M \times 1.0 M laid out in a Randomised Complete Block Design (RCBD) with each genotype replicated three times. Throughout the experimental period, the plots were kept free of weed by hoeing. No Fertilizers or supplementary water through irrigation was applied during the trials. Major pests and diseases (Leaf spot, leaf rust and powdery mildew) were controlled by 1-2 sprays (depending on pest pressure) using Dimethoate 40%EC and Rocket 44 EC (Profenofos 40% + Cypermethrin 4%) at recommended rates.

After one year of establishment in the field, data was recorded by following a 5-crop schedule. Data from the 5 plants were recorded on various yield attributing traits such as number of tillers (NT); plant height (HT), total shoot length (TSL), nodal distance (ND), number of branches per plant, weight of 100 leaves, leaf area, leaf yield/plant and leaf yield (LY) were recorded. These measurements used were adopted from Adolkar et al. (2007) and Food and Agriculture organization (FAO), (2007).

A combined analysis of variance to assess the significance

of genotype \times environment interactions was carried out using R version 4.0.1 (R Core Team, 2014). The mean values for all the agronomic traits were later used for correlation matrix

Table 1: Different mulberry varieties and their original	gin.
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S.NO	Genotype	Species	Country of origin
1	Thika	M. alba L.	Kenya
2	Kanva2	M. indica L.	India
3	Embu	M. alba L.	Kenya
4	Thailand	M. alba L.	Thailand
5	Chinese	M. alba L.	China
6	Mysore Local	M. indica L.	India
7	S.36	M. indica L.	India
8	S.54	M. indica L.	India

Adopted from Vijayan et al., 2003 and Wangari Peris and Ngode Lucas (2016).

3. Results and Discussion

3.1 Effect of environment and genotypes on the growth characteristics of different mulberry varieties under study

Morphological analysis is considered as a first approach towards the assessment of genetic diversity in a plant species (Boubaya et al., 2009). In this study, an experiment was conducted to evaluate the agronomic performance of different mulberry accessions across different locations in Uganda and to understand their genetic variability. The results presented in Table 2 show the effect of environment and genotypes on the growth characteristics of different mulberry varieties in different districts in Uganda. It is evident from the morphological traits and the regions of cultivation that the varieties used in this study showed considerable variations across the different agro-ecological zones in Uganda. Analysis of variance showed a significant variation between the genotypes \times location (P<0.05). There was also significant variation among the parameters in all genotypes across all the agro-climatic regions except for the number of tillers which was not significant (P=0.721).

Means of different parameters across genotypes revealed that plant height, lamina length, lamina width, internode distance, number of branches, petiole length and bud length varied significantly ($p \le 0.05$) within the accessions while the number of tillers was not statistically significant in all the accessions. Thailand, S-36, Embu and Local variety had the largest leaves while Chinese variety had the smallest leaves. Local, Thika and Thailand were the highest plant height among the accessions respectively whereas the shortest was Chinese. Peris et al (2013) studied the morphological variations in different accessions in two different locations in Kenya and the results obtained concur with the results in this study that location (environment) significantly influences the performance of mulberry.

Analysis of variance for genotype × location showed that this interaction had an influence on the morphological traits revealing that lamina length, lamina width, petiole length, plant height, bud length, leaf area, internode distance, total plant biomass, number of branches, 100 leaf weight and diseased leaves were highly different ($p \le 0.05$) across the locations. On contrary, the number of tillers was not statistically significant across all the locations. Reaction to pests and diseases by all the genotypes across the locations was also highly significant

Table 2: Means of different phenotypic traits of 5 mulberry accessions evaluated across different environments in Uganda.

	LOCATION									VARIETIES									c	
TRAITS	Sheem a	Muben de	Nakase ke	Iganga	Bulam buli	Bulam buli 2	Bukede a	Pallisa	Zombo	S-36	S-54	THAIL AND	KAMU LI	KANV A-2	CHINE SE	THIK A	LOCA L	EMBU	s e	v (%)
bud length(m m)	4.6 85 ab	4.1 23 b	4.9 93 a	4.2 36 b	4.3 19 b	4.1 94 b	4.0 8b	4.9 93 ab	4.5 83a b	4.2 4bc	4.7 34b	4.8 02a b	4.1 23b c	3.9 25c d	3.4 56 c	4.5 55b c	5.3 9a	4.3 33 bc	1. 0 0 4	1. 4 1
lamina length(cm)	19. 45 06 2a b	18. 51 72 8b	20. 18 47 2a	20. 15 97 2a	19. 45 27 8a b	19. 55 27 8ab	20. 20 97 2a	18. 73 47 2b	20. 28 75 0a	19. 997 53b c	20. 179 01b c	24. 091 36a	19. 139 51c	19. 459 26c	13. 65 18 5d	19. 267 90c	21. 01 60 5b	19. 57 22 2c	2. 5 4 4	1 3. 9 5
lamina width(cm)	14. 71 6b	14. 71 3b	15. 53 8a b	15. 87 9a	14. 93 4a b	14. 60 1b	15. 80 2a	14. 69 7b	15. 45 8ab	16. 92b	15. 359 cd	18. 882 a	14. 698 de	15. 065 cde	10. 24 07 f	14. 040 7e	15. 66 7c d	16. 13 8b c	2. 1 1 1	1 0. 9 2
leaf area(sq.c m)	28 9.7 77 b	28 7.6 59 b	32 7.7 51 a	32 7.8 45 a	30 5.4 75 ab	29 9.4 93a b	32 8.2 91 a	29 9.4 93 b	32 7.8 51a	341 .64 1b	313 .66 6bc	463 .37 4a	286 .86 cd	296 .09 9cd	15 1.8 74 1e	274 .68 27d	33 4.3 97 b	33 3.9 92 b	1 0. 7 6	1 4. 3 9
petiole length(cm)	3.9 79 cde	4.2 62 bc	4.7 33 a	4.1 05 bc de	3.7 44 e	4.1 05 bcd e	4.4 13 ab	3.8 44 de	4.2 06 bcd	3.9 209 b	4.4 22a	4.7 17a	3.6 88b	3.7 555 b	3.7 25 b	4.4 37a	4.4 20 9a	4.8 44 a	0. 8 4 8	9. 7 9
internode distance(c m)	4.7 55 bc	4.7 27 bc	5.5 08 a	4.6 08 c	4.7 9b c	4.9 38 bc	5.0 72 b	4.5 37 c	5.0 85 b	4.8 64b	4.7 08b	5.1 96a b	4.9 03a b	4.9 87a b	3.8 c	5.4 01a	5.1 83 ab	5.1 27 ab	0. 9 8 1	1 0. 2 7
plant height(cm)	20 9.6 56 8 c	19 2.5 74 1 cd	28 6.3 61 1 a	27 4.0 06 9 a	16 0.8 90 3 e	18 2.0 95 8 de	24 6.5 69 4 b	16 3.5 43 1 e	20 5.0 95 8 cd	199 .02 1c	195 .53 70c	237 .76 54a b	214 .08 02b c	192 .38 40c d	16 8.3 65 4d	244 .77 78a	25 7.0 09 9a	19 4.0 55 6c d	1 0. 7 5	1 7. 4 6
Total Plant biomass (Kg)	84 6.5 06 2c	81 3.0 24 7c	17 00. 13 89 a	16 32. 76 39 a	80 2.4 72 2c	98 0.0 13 9c	13 11. 55 56 b	71 6.7 5c	15 48. 12 5ab	132 8.7 284 abc	109 8.3 951 abc	119 0.1 235 abc	104 1.2 346 bc	109 4.9 136 abc	61 7.0 12 3d	137 7.4 691 ab	14 20. 97 53 a	97 5.3 33 3c	1 0. 8 7	7. 5 9
Leaf yield/plant (kg)	21 1.7 35 1d	21 1.2 10 9d	32 9.6 36 4c	44 5.4 02 8a	39 7.3 61 1a b	41 5.6 42 8ab	45 6.1 90 3a	36 6.3 33 3	45 7.9 95 6a	404 .76 22b	385 .25 98b	467 .28 14a	363 .67 9bc	368 .22 28b c	17 8.1 67 2d	340 .60 88b c	39 4.2 56 b	31 7.4 35 6c	1 0. 4 8 8	2 0. 0 5
Number of branches	16. 1a	9.1 7b	11. 4b	6.0 9c	10. 6b	4.4 c	4.5 c	4.6 c	10. 36 b	7.8 bcd	8.5 6bc	4d	12. 703 a	11. 2ab	8.7 2b c	9.1 bc	7.0 1c d	6.6 d	7. 0 6 2	4. 5 3
number of tillers	1.9 2d	1.8 3d	1.7 7d	2.1 2c d	2.4 3b cd	2.5 2bc d	3.0 4a b	2.8 ab c	3.6 1a	2.3 8a	2.8 1a	1.9 a	2.4 5a	2.5 8a	2.5 4a	2.5 8a	2.3 5a	2.2 7a	1. 9 0 3	4. 4 2
number of diseased leaves	57. 8b	57. 2b	11 7.2 a	14. 6c	5.6 2c	10. 3c	8.4 8c	3.0 1c	7.0 5c	38. 5ab	32. 5ab	21. 18c	29. 7ab	25. 1b	22. 1c	35. 91a b	47. 5a	46. 3a	1. 2 9	5. 2 6
number of leaves destroyed by pests	26. 19 b	26. 1b	68. 12 a	5.2 7c	7.2 2c	7.2 2c	7.8 75 c	8.1 2c	7.9 7c	24. 1a	19. 7ab	14. 02a b	16. 8ab	16. 6ab	8.8 5b	22. 3a	21. 8a	24. 8a	1. 1 3 6	5. 5 5 3

Means in the same row having different letters are not significantly different at α =0.01 level of significance according to the Duncan's Multiple Range Test (DMRT).

3.2 Correlation coefficient matrix

The association of different agronomical traits were analysed through Pearson simple correlation matrix. Correlation coefficient analysis was used to depict the degree and extent of relationship among important plant traits being the main basis for crop selection. The results in this study revealed that different traits were associated directly or indirectly with each other.

Phenotypic correlation coefficient was highly significant for all the important characters like yield and yield related characters (Table 3). Lamina length showed significant positive correlation with lamina width. Leaf area positively correlated with both lamina length and lamina width. Number of branches per plant, number of seeds per pod, pod length and test weight exhibited positive and significant association with seed yield per plant. Bud length expressed positive significant correlation with laminar length, lamina width and leaf area. Petiole length showed significant positive correlation with laminar length, lamina width, leaf area and bud length. Internode distance showed significant positive correlation with lamina width, leaf area, bud length and petiole length but non-significantly correlated with lamina length. Number of branches per plant showed a negative significant correlation with lamina length, lamina width, and leaf area and showed a positive significant correlation with bud length. Number of tillers per plant revealed a negative insignificant association with all the traits. Plant height expressed positive significant association with lamina length, lamina width, leaf area, bud length and internode distance while petiole length and number of branches were positively correlated though not significant. Plant height was negatively correlated with the number of tillers. The 100-leaf weight exhibited positive and significant correlation with lamina length, lamina width, leaf area, petiole length, internode distance and plant height while it had a negative association with bud length and the number of branches. Number of diseased leaves expressed a positive significant association with bud length, petiole length, internode distance, number of branches and plant height but lamina length, lamina width, leaf area positively correlated though not significant while number of tillers negatively correlated. Number of leaves destroyed by pests showed positive significant association with lamina width, leaf area, bud length, number of branches, plant height and internode distance while lamina length and number of diseased leaves were positively correlated though not significant. Leaf yield expressed positive significant association with all the traits under study namely lamina length, lamina width, number of primary branches per plant, plant height, total shoot length, internodal distance and single leaf weight. Other traits are associated with each other and formed a complex relationship among them.

Mulberry is one of the most important commercial crops grown extensively as sole food plant for silkworm (*Bombyx mori*). Mulberry (*Morus* spp.) is a perennial and high biomass producing plant which also plays an important role in keeping the mother nature under constant phase by reducing the global warming through uptake of carbon dioxide from atmosphere and in return produce and release oxygen into the environment; which purifies the air and provide life to animals and other organisms (Rohela et al., 2020). Katul and Novick, 2009 stated that mulberry plants also contribute to soil health; retains water in soil subsurface and cools overheated urban areas through its evapotranspiration mechanism of water cycle. It continues to be grown throughout the year in tropical and sub-tropical regions of the world (Vijayan et al., 2004). A large number of mulberry varieties are grown in several geographical regions of the world both for their edible fruit and for the silk industry.

The results in this study showed significant differences among the mulberry genotypes for some morphological traits studied. The study further revealed that the interaction between varieties and environment was slightly significant suggesting that each of these varieties can perform relatively the same way regardless of the growing environment/location. However, in these nine environments, the results for morphological traits showed better performance at Zombo station compared to other stations. This can be attributed to the differences in terms of soil type, rainfall patterns, temperatures, humidity and other important aspects that affect plant growth and yield in Uganda. The results were in agreement with Chambel et al. (2004) who stated that the environmental changes are not only responsible for changing the population phenotypic structure based on its genetic makeup to survive in the new environment and the level of each individual to alter its phenotype according to its environment. Not only does phenotypic plasticity make a genotype behave differently in different environments but also similarly in a particular environment (Bradshaw, 1965). Several authors (Peris et al., 2013, Gray, 1990, Karst and Lechowics (2007) reported different modifications in mulberry as a means for adaptation to environment. Leaf variations in size, stomatal size and leaf polymorphism were as a result of phenotypic plasticity (Kitajima et al., 1997).

Yield-wise, no single variety was consistently superior in terms of agronomic performance across all the agroecological zones as performance varied in different locations. The above observation therefore gave a clear impression that yield traits in mulberry were variable and their response was not only dependent on genetic influence but also on the environment. In the combined analysis, Thailand variety exhibited the largest leaf sizes, short internode distance and fairly high branching attributes in all environments, thereby having the best traits that contributed to higher leaf yields compared to other varieties studied. These slight variations in performance observed amongst these genotypes may have been due to differences in genetic characteristics and environmental adaptations of the individual genotypes, which influence their Nutrient absorption capacity, photosynthetic area accumulation, water use efficiency (WUE) among others. Similar results were reported by Gray, (1990) that leaf characters like lamina length, width, petiole length, width and height changes as the environment changes. Similar observations on association of different traits with leaf yield was also reported by Sarkar et al., (1987), Tikader and Roy (2001), Tikader and Dandin (2005) and Vijayan et al. (1997).

Table 3: Pearson's correlation coefficient matrix for the different quantitative traits across the different environments.

	lami na lengt h	lami na widt h	leaf are a	bud len gth	petiole length	intern ode distan ce	number of branch es	numb er of tillers	plan t heig ht	Number of diseased leaves	number of leaves destroyed by pests	100le af weig ht	total plant biomas s
lamina length													
lamina width	0.88 1***												
leaf area	0.95	0.95											

	1***	9***											
bud length	0.09 6***	0.09 1***	0.0 9**										
petiole length	0.49 33** *	0.41 4***	0.4 584 ***	0.219 ***	8								
internode distance	0.39 8	0.31 6***	0.3 361 ***	0.1 624 ***	0.281	2***							
number of branches	- 0.16 3***	- 0.17 81	- 0.2 006	0.2 312 ***	-0.056	-0.031							
number of tillers	- 0.09 45	- 0.09 33	- 0.1 049	0.2 402	- 0.0446	- 0.0625	- 0.0139						
plant height	0.44 03** *	0.37 ***	0.4 032 ***	0.2 751 ***	0.4526	0.2584 ***	0.0829	- 0.148 7					
Number of diseased leaves	0.01 74	0.04 4	0.0 336	0.6 073 ***	0.1465 ***	0.0914 ***	0.2838 ***	- 0.156 5	0.3	3948***			
number of leaves destroyed by pests	0.07 25	0.10 6***	0.0 949 ***	0.5 274 ***	0.1883 ***	0.1657 ***	0.2641 ***	0.123 3	0.41 03* **	0.0853			
100leaf weight	0.58 05** *	0.56 71** *	0.6 01* **	- 0.3 435	0.3114 ***	0.1767 ***	- 0.1861	0.045 7***	0.38 24* **	-0.1097	-0.0117		
Leaf yield/plant	0.33 74** *	0.31 97** *	0.3 252 ***	0.0 577	0.3261 ***	0.2065 ***	0.2644 ***	0.027 2***	0.67 44* **	0.3691** *	0.4755***	0.45	99***

Conclusion

The results obtained in this study revealed that there was a slight difference in the agronomic performance of these mulberry varieties. According to the findings above, it can be concluded that Kanva2 which produced the highest number of branches and Thailand with highest leaf yield can be recommended for selection as good cultivars for farmer utilization in different geographical regions of Uganda since the main goal of mulberry production is high yield of leaves for silk worm production. This study has helped bridge the knowledge gap regarding different mulberry accessions and their performance in different parts of the country and this forms basis of proper and appropriate selection of the tropical cultivars based on the morphological traits for the farmers in Uganda. But it has been revealed that pests and diseases, genetic diversity directly affected leaf yield of mulberry therefore further research should be conducted to study pests and diseases of mulberry and their management, genetic diversity of different mulberry accessions and their contribution to adaptation to different agro-ecological environments. Future experiments should also consider evaluating these varieties over more locations to be able to determine the best adapted mulberry varieties with acceptable performance regardless of the season of cultivation.

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