

Response of Cowpea Genotypes to Drought Stress in Uganda

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Abstract

Moisture stress is a challenge to cowpea production in the drought prone areas of eastern and north eastern Uganda, with yield losses of up to 50% reported. Genotypes grown by farmers are not drought tolerant. This study was therefore, undertaken at Makerere University Agricultural Research Institute Kabanyolo to identify cowpea genotypes tolerant to drought. Thirty cowpea accessions comprising of Ugandan landraces and released varieties, Brazilian lines, Makerere University breeding lines, elite IITA germplasm and seven IITA drought tolerant lines as checks were screened for drought tolerance at vegetative and reproductive stages. The experiment was designed as a 2 × 37 factorial and laid out in a split-plot arrangement, 37 genotypes of cowpea at two soil moisture stress levels (T1, no stress and T2, severe stress) with all factorial combinations replicated two times in a screen house. The genotypes showed considerable variability in tolerance to drought. Genotypes were significantly different for chlorophyll content ($P \leq 0.01$), efficiency of photosystem II ($P \leq 0.05$), non-photochemical quenching ($P \leq 0.05$), recovery ($P \leq 0.01$), delayed leaf senescence ($P \leq 0.01$), grain yield ($P \leq 0.01$), 100 seed weight ($P \leq 0.05$), number of pods per plant and number of seeds per pod ($P \leq 0.001$). There was a high significant positive correlation between chlorophyll content and efficiency of photosystem II ($r = 0.75$, $P \leq 0.001$) implying that chlorophyll content and efficiency of photosystem II could be used as efficient reference indicators in the selection of drought tolerant genotypes. Genotypes SECOW 5T, SECOW 3B, SECOW 4W, WC 30 and MU 24 C gave relatively high yields under stress and no stress conditions, maintained above mean chlorophyll content, efficiency of photosystem II and had good recovery scores from stress and thus were tolerant to drought stress induced at both vegetative

and reproductive stages.

Keywords

Drought Tolerance, Moisture Stress, Non-Photochemical Quenching, Split Plot

1. Introduction

Cowpea is one of the most important food legumes in the drier regions of the tropics and sub-tropics where drought is a major production constraint due to low and erratic rainfall [1]. It provides a cheap source of protein, vitamins and carbohydrates to small scale farmers in East Africa [2]. In Uganda, cowpea is the fourth most widely grown legume crop after common bean, groundnut and soy bean. The crop is intensively cultivated in the northern and eastern regions of Uganda [3] such that it provides an estimated 60% of the protein required by the population in these regions [4]. Despite its widespread cultivation in Uganda, the yields of cowpea in farmers' fields are still very low averaging 300 - 500 kg·ha⁻¹ against a yield potential of 1500 to 3000 kg·ha⁻¹ [5]. The low yields have been attributed to a number of biotic and abiotic stresses, low yielding local varieties, seed scarcity and poor soils [6]. Although cowpea is considered as being more drought tolerant than many other crops, its productivity is negatively affected by prolonged droughts and high temperatures [7] which are currently attributed to the effects of climate change. Climate models have predicted increased severity and frequency of droughts in Sub-Saharan Africa [8]. Development and adoption of drought tolerant varieties is one of the options to cope with the changing climate.

Cowpea is sensitive to soil moisture stress during the vegetative and reproductive growth stages [9]. Drought stress during flowering and pod filling impacts negatively on flower development, pollination, pod setting and grain filling leading to reduced number of pods per plant, reduced seed weight and low seed yield [10]. Plants respond and adapt to drought stress through the induction of various morphological and physiological responses [11]. Screening cowpea for tolerance to drought at either vegetative or reproductive stages elsewhere has focused mainly on morphological, biochemical, physiological and yield related indicators [12] [13] [14] [15] [16]. However, in Uganda such studies have not been reported, considering the multiplicity of factors involved in drought tolerance, much more robust screening techniques to characterize drought tolerance are needed. The study therefore sought to identify candidate drought tolerant lines from Uganda that can be used for future breeding applications using physiological and agro-morphological indicators.

2. Materials and Methods

2.1. Genetic Materials

Thirty medium maturing cowpea genotypes consisting of landraces from north-

ern, west central and eastern regions of Uganda, varieties released by the National Semi-Arid Resources Research Institute (NaSARRI), Brazilian lines, and Makerere University breeding lines were studied at vegetative and reproductive stages. The thirty medium maturing cowpea genotypes were selected based on their yield potential and disease resistance qualities. Seven drought tolerant genotypes from IITA-Kano, Nigeria, were used as checks.

2.2. Study Area

The study was conducted between May and December 2015 in a water proof screen house at the Makerere University Agricultural Research Institute-Kabanyolo (MUARIK), Uganda. MUARIK is located at an altitude of 1217 meters above sea level on coordinates 0.16°24'16 N and 32.5°27'34E, approximately 19 km in the northeast of Kampala at Kabanyolo in Lake Victoria Crescent [17]. MUARIK has deep ferrallitic soils with pH range of 5.2 to 6.0. Average temperature during experiment ranged from 25°C - 31°C while relative humidity ranged from 71% - 90%. The rainfall pattern of MUARIK is bimodal, with an annual average of 1160 mm, with the first rains from March to May, while the second rains are from September to November.

2.3. Experimental Design

The experiments were laid out in a randomized complete block design in a split plot arrangement with two replications, two watering regimes as main plots and the test genotypes as subplots. The two water regimes were no stress (pot capacity-50% maximum water capacity) and severe stress. Eight seeds per each of the 30 cowpea genotypes and 7 drought tolerant checks were planted in the screen house in 10 litre-plastic pot perforated at the base and filled with 10 kg top soil and later thinned to 4 seeds per pot 8 Days after planting (DAP). The soil used was composed of 66% sand, 16% clay, 18% silt, 0.15% total nitrogen, 0.53 Cmoles/kg of potassium and 4.24 parts per million (ppm) of phosphorus. NPK fertilizer was applied to soil at the rate of 6 g/pot, based on soil analysis in order to provide the recommended nutrient requirement for cowpea [18]. The NPK fertilizer was applied to each pot by incorporating it into the soil at planting to remove nutrient deficiency as a limiting factor [18].

2.4. Treatments

Water stress treatment at vegetative stage was imposed after 10 - 14 days of initial growth of the plants. The initial soil moisture level for the two treatments was maintained at field capacity (50% maximum water capacity). Plants in the well-watered treatment (no stress) were maintained at field capacity. A soil moisture meter (MO750 manufactured by Extech Instruments, USA) was used to determine the moisture level in the soil. Plants undergoing a severe stress did not receive water after the initial establishment for 30 days' interval [9] [19]. After 30 days of induced drought stress, watering was resumed. The presence or absence of new growth and whether the growth is from the apical or basal me-

ristem and recovery was recorded.

Water stress treatment at reproductive stage was imposed after the emergence of flower buds for of each of the genotype in a plot [20]. The drought stress treatment did not receive water for 20 days [21] after stress induction while the non-stress treatment received water twice a week until physiological maturity. After the elapse of the 20 days of drought stress induction, the drought stressed plants received water twice a week up to physiological maturity.

2.5. Data Collection

Data was collected on the following growth parameters on individual plant basis in line with the international plant genetic resources cowpea descriptors; wilting, delayed leaf senescence. Stem greenness was scored on a scale of 0 to 5, with 0 being completely yellow and 5 being completely green [22]. Wilting was scored using both the IBPGR (IB) [23] and [24] scales: total number of leaves per plant; number of leaves showing wilting signs per plant. Wilting was calculated weekly, from the first week to the final week of stress, as the ratio between leaves showing wilting signs and the total number of leaves per plant. Stem greenness and regrowth was scored as recovery parameters, after the two weeks of re-watering. Stem greenness was scored using a scale of 1 - 5, where 1 was yellow and 5 was completely green. Regrowth (recovery) was scored using three categories: *i.e.* 1 with no re-growth; 3 with regrowth from auxiliary buds; and 5 with re-growth from the apical meristems [25].

Chlorophyll content, efficiency of photosystem II and non-photochemical quenching were measured from 10:00 am to 12:00 pm in the middle three leaflet of the trifoliolate leaf from the plant base, which were mature but not senescent leaves using Photosynq device [26] (TerAvest D, May 2015, Photosynq coordinator personal communication). At maturity, the plants were harvested to determine yield components (number of pods per plant, number of seeds per pod, 100 seed weight, and seed yield per plant).

2.6. Data Analysis

Data was analyzed using Analysis of Variance with a split plot model in GENSTAT 12 to determine the significance of water regime, genotypes & genotypes by water regimes on yield & yield component parameters. The Linear Mathematical Model for Split plot experimental Design used was:

$$X_{ijk} = Y\cdots + M_i + B_j + d_{ij} + S_k + (MS)_{jk} + e_{ijk},$$

where X_{ijk} = mean observations, $Y\cdots$ = the experiment mean, M_i = the main plot treatment effect, B_j = replication or block effect, d_{ij} = the main plot error (error a), S_k = the subplot treatment effect, $(MS)_{jk}$ = the main plot and subplot treatment interaction effect, e_{ijk} = the subplot error (error b). i = a particular main plot treatment, j = a particular block, k = a particular subplot treatment. Genetic correlation was calculated from the components of variance and covariance for the genotype term using multivariate analysis of variance in Genstat (12th edition). The correlation coefficient and coefficient of determination values were

based on 37 Genotypes (n = 37).

3. Results and Discussion

3.1. Genetic Variability for Drought Tolerance in Cowpea Genotypes

The results from analysis of variance for chlorophyll content, efficiency of photosystem II, non-photochemical quenching (Table 1), drought recovery, leaf wilting index (LWI) (Table 2), seed yield and its components (Table 3) showed that Genotypes were significantly different, indicating the presence of genetic variation among the cowpea genotypes for these traits that is essential for improvement of the crop for drought tolerance.

Table 1. Mean squares of physiological drought stress indicators of cowpea genotypes at vegetative stage.

Source of variation	df	Chl C	Phi2	NPQ
Replication	1	559.91	0.00	1.53
water Levels	1	3706.00	1.56	200.02
Main Plot Error	1	218.59	0.07	2.08
Genotypes	36	72.12***	0.02*	2.60*
Genotypes × Water Levels	36	25.27	0.02*	2.40
Sub Plot Error	72	26.71	0.01	1.58

*, ***, significant at $P \leq 0.05$, $P \leq 0.001$ respectively, df: degrees freedom, ChlC: Chlorophyll content, Phi2: Efficiency of photosystem II, NPQ: Non-photochemical quenching.

Table 2. Mean squares of morphological drought stress indicators at vegetative stage.

Source of variation	df	Recovery	Stem Greenness	Wilting Index
Replication	1	0.2	5	0
Genotypes	36	3.56**	1.93ns	0.02**
Error	36	1.43	1.21	0.01

** significant at $P \leq 0.01$, ns: non-significant, df: degrees of freedom.

Table 3. Mean squares of seed yield and its components under drought stress at reproductive stage.

Source of variation	DF	NPP	NSP	100 SWT (g)	SY (g)
Replication	1	2.55	4.80	9.08	4.41
water Levels	1	30.50*	70.53	0.68	6.08
Main Plot Error	1	0.05	6.53	3.01	1.88
Genotypes	29	7.64***	12.20***	1.00*	2.59**
Genotypes × Water Levels	29	2.51	4.72*	1.18**	1.28
Sub Plot Error	58	2.22	2.74	0.51	0.90

*, **, *** significant at $P \leq 0.05$, $P \leq 0.01$, $P \leq 0.001$ respectively, df = degree of freedom, NPP = Number of pods per plant, NSP = Number of seeds per pod, SWT = Seed weight, SY = Seed yield, g = grams.

3.2. Physiological and Morphological Response of Cowpea Genotypes to Drought Stress at Vegetative Stage

Although the interaction between genotypes and water levels when averaged across the two water levels was found to be non-significant, there were detectable differences in the performance of genotypes under drought stress. Drought stress consistently and significantly reduced the efficiency of photosystem II, Chlorophyll content of the tested genotypes (**Table 4**) though the effect varied in both improved and local genotypes. The reduction in efficiency of photosystem II was supported by an increase in non-photochemical quenching (NPQ) as shown by a strong negative correlation ($r = -0.76$, $P \leq 0.001$), suggesting that a greater portion of the energy was thermally dissipated (**Figure 1** and **Table 5**). [27] reported that an increase in NPQ protects the plant against photo damage and this reduces the quantum yield of photosystem II. NPQ plays a key role in the protection of PSII from photodamage. NPQ is considered as an indicator of excess excitation energy [28]. The maximum quantum efficiency of photosystem II provides a measure of the rate of linear electron transport, hence, an indication of overall photosynthetic capacity [29].

Genotypes SECOW 4W, SECOW 5T, WC 67B, NE 6 maintained higher levels of chlorophyll content in their leaves similar to drought tolerant checks such as Dan ILA, IT 89KD-288 and IT 93K-452-1 under water stress (**Table 4**). The average chlorophyll content (SPAD) measurement under no stress and severe

Table 4. Mean of chlorophyll characteristics and drought recovery.

Genotype	No Water Stress			Water Stress			Recovery
	SPAD	Phi2	NPQ	SPAD	Phi2	NPQ	
SECOW 4W	57.32	0.56	1.02	50.87	0.49	1.43	5
SECOW 2W	54.30	0.59	1.27	46.97	0.37	5.61	3
182	58.17	0.63	0.92	42.58	0.28	4.01	5
SECOW 1T	50.60	0.56	1.30	42.03	0.29	5.28	5
IT 91	58.83	0.55	0.89	43.62	0.32	4.54	5
3306	56.45	0.59	0.87	46.33	0.28	3.34	5
SECOW 5T	53.43	0.58	0.72	51.82	0.48	1.99	5
WC27	54.97	0.66	0.58	35.45	0.30	3.01	5
NE 45	54.32	0.62	0.78	46.25	0.30	3.98	5
WC64	55.22	0.68	0.65	34.77	0.30	1.59	1
SECOW 3B	53.62	0.62	0.66	48.70	0.51	2.50	5
SECOW 5 X SECOW1	56.25	0.62	0.48	47.03	0.42	1.83	5
NE 20	56.17	0.59	1.01	46.77	0.37	3.35	5
NE50	58.03	0.62	0.95	43.15	0.16	5.14	3
MU17	55.77	0.64	0.59	45.57	0.46	1.94	3
MU15	58.20	0.65	0.71	41.50	0.39	4.48	5

Continued

WC8	46.10	0.37	0.44	41.92	0.26	4.40	3
SECOW 5 X NE39	50.17	0.57	2.18	45.23	0.33	5.83	5
NE 44	55.13	0.63	0.77	33.27	0.26	4.57	1
WC30	55.62	0.56	1.57	48.87	0.54	1.55	5
NE 70	54.70	0.60	0.90	42.38	0.23	3.07	1
IT 84	56.02	0.61	0.87	45.43	0.34	4.81	5
BRAZIL5	57.42	0.56	0.87	34.15	0.24	4.53	1
NE 67	56.27	0.52	1.13	40.77	0.26	5.65	3
WC67B	59.48	0.61	1.45	53.32	0.48	2.43	5
MU 24C	58.12	0.70	0.65	48.03	0.31	4.39	5
NE 5	50.28	0.56	0.87	47.75	0.47	1.70	3
NE30	56.47	0.54	1.03	46.97	0.36	4.90	5
WC 17	55.20	0.58	0.88	45.28	0.48	4.12	5
NE6	56.55	0.60	0.92	52.53	0.51	1.64	3
Checks							
DANILA	62.22	0.51	1.31	54.95	0.65	0.78	5
IT89KD-288	62.28	0.61	1.11	54.52	0.46	1.98	5
IT93K-452-1	55.53	0.55	1.35	54.27	0.59	0.74	5
IT96D-610	65.93	0.58	1.08	60.63	0.60	0.86	5
IT97K-499-38	61.63	0.63	0.85	52.43	0.51	1.67	5
IT97K-819-118	57.37	0.70	0.60	41.82	0.45	3.35	5
IT98K-128-3	59.65	0.59	1.01	50.55	0.31	4.30	5
Grand Mean	56.32	0.59	0.95	46.31	0.39	3.28	4.19
LSD(5%)	7.70	0.16	0.82	12.67	0.25	3.51	2.42
CV%	7	13	42	14	32	53	30

Checks: IITA drought tolerant lines were included for screening at vegetative stage, NE & WC: Local Genotypes, SECOW: Serere released varieties, MU: Makerere breeding lines.

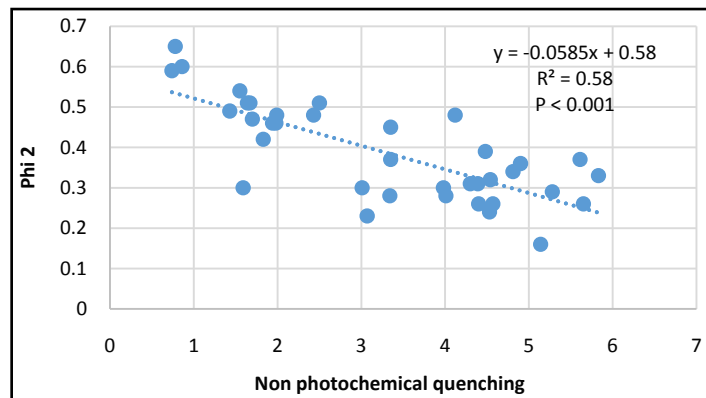


Figure 1. Regression analysis of efficiency of photosystem II and non-photochemical quenching under soil moisture stress.

Table 5. Correlation analysis between chlorophyll characteristics and drought recovery of cowpea genotypes.

	SPAD	NPQ	Phi2	Recovery
SPAD	1			
NPQ	-0.55***	1		
Phi2	0.75***	-0.76***	1	
Recovery	0.56***	-0.12	0.43**	1

** , *** significant at $P \leq 0.01$, $P \leq 0.001$ respectively.

stress conditions were 56 and 46 respectively (**Table 4**). Drought stress imposed at the vegetative stage, significantly ($P \leq 0.05$) decreased the relative chlorophyll content. The results are consistent with [30]. [31] reported that severe drought stress on wheat significantly decreased the chlorophyll content of the leaves. The decrease in chlorophyll under drought stress is mainly the result of damage to chloroplasts caused by active oxygen species [32]. The predictability of the efficiency of photosystem II from the chlorophyll content (**Figure 2**) as shown by the coefficient of determination of 56% indicated a high association between efficiency of photosystem II and chlorophyll content ($r = 0.75$, $P \leq 0.001$). Similar findings were reported by [15]. Furthermore, there were strong and moderate significant positive correlation between chlorophyll content with drought recovery and efficiency of photosystem II with drought recovery respectively (**Table 5**). Suggesting that Genotypes that maintained a higher chlorophyll content and efficiency of photosystem II such as SECOW 4W, SECOW 5T, WC 30, WC 67B had higher chances of recovering from drought. As such Chlorophyll content and Phi2 could be used as reliable and efficient reference indicators in the selection of drought tolerant genotypes. Similar findings were reported by [15]. The low predictability of non-photochemical quenching from chlorophyll content as indicated by the coefficient of determination of 30% implied a low degree of association between NPQ and chlorophyll content. This could be due to damaged membranes which reduced the mobility of the electron transport chain carriers and reduced proton gradient across thylakoid membrane [33].

Some cowpea genotypes showed high leaf wilting index scores (LWI) after the first week of stress, for example genotypes WC 64 and BRAZIL 5. On the other hand, the following genotypes showed lower wilting index scores during the whole stress period; IT98K-205-8, IT97K-499-38, IT98K-503-1, NE 67, SECOW 1T, SECOW 3B, WC 27, NE 20, IT98K-1283, MU 15, IT 84, 182, IT97K-819-118, DAN ILA. The results were consistent with previous studies in cowpea, which also identified leaf wilting within the first week of water stress [34]. Although some genotypes had higher wilting index scores but they were able to fully recover after rehydration. This was attributed to Type 2 drought tolerance mechanism that most genotypes had [34]. These genotypes derived water from their lower canopies, in order to support apical meristematic growth during water stress. Such genotypes show relatively high values of LWI and yet fully recovered after re-watering [25].

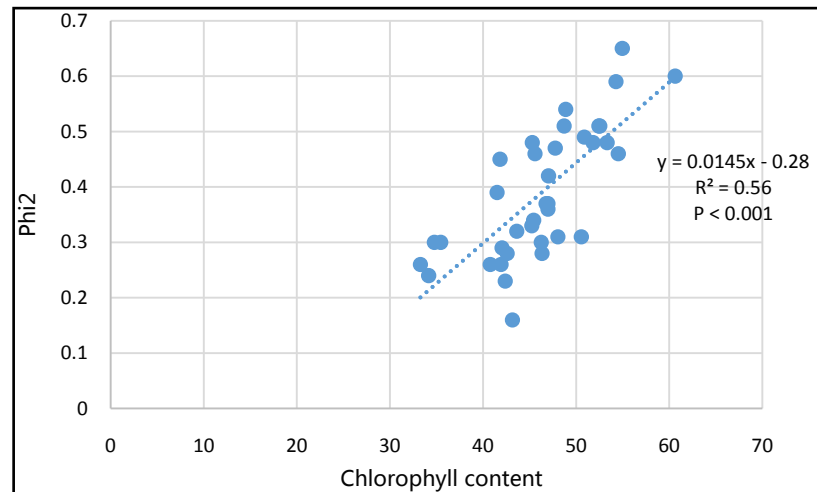


Figure 2. Regression analysis of efficiency of photosystem II and chlorophyll content under soil moisture stress.

Recovery after stress is a process that involves the rearrangement of many metabolic pathways to repair drought-induced damage and to resume plant growth. There was variability in drought recovery among genotypes. Similar findings were reported by [35]. Genotypes that maintained high chlorophyll content under drought stress showed a greater ability to recover from stress. Particularly genotypes SECOW 5T, SECOW 3B, SECOW 4W, WC 67B, NE 6, MU 24C and WC 30 maintained above mean chlorophyll content, efficiency of photosystem II and had a greater ability to recover from stress (Table 5).

3.3. Yield and Its Components Indicators of Drought Tolerance at Reproductive Stage

Grain yield under drought is the primary trait for selection in crop improvement programs [1]. Different morphological and physiological characters contribute to grain yield. Therefore, the knowledge of genetic relationship between grain yield and its components can improve the efficiency of selection in breeding programs. The mean seed yield performance of the thirty (30) cowpea genotypes ranged between 2.7 to 6.0 grams/plant under no soil moisture stress and 0.7 to 5.7 grams/plant under soil moisture stress (Table 5). Genotype NE 30 and WC 8 gave the highest seed yield under no stress and stress respectively. Genotype WC 17 produced the highest number of pods both under no soil moisture stress and soil moisture stress conditions while IT 84 produced the highest number of seeds per pod both under no soil moisture stress and soil moisture stress conditions (Table 6). Drought stress reduced the seed yield and its components, but, improved genotypes and landraces responded differently to the stress. Brazil 5, NE 45 and NE 50 experienced a seed yield reduction of 83% and 36% respectively under soil moisture stress while WC 8, MU24C, Secow 5T, Secow 3B, MU15, IT84 and Secow 4w gave relatively higher yields. [21] reported that the reduction in seed yield under water stress was associated with the decrease in yield components such as number of pods per plant and number of seeds per pod. Cowpea

has been reported to be very sensitive to water stress at flowering and pod-filling stages [36]. Drought stress caused flower abortion, pod dropping and reduced seed filling [37]. Therefore, the reduction in number of pods per plant and number of seeds per pod in this study was attributed to abscission of reproductive structures and limitation of dry matter partitioning to reproductive sink. Genotypes that maintain higher yields under drought stress are desirable as such, genotypes SECOW 5T, SECOW 3B, SECOW 4W, WC 8, MU 24C, MU15, and IT 84 were classified as tolerant because they produced above mean yields and had minimal yield reduction under water stress. The presence of a strong positive significant genetic correlation among seed yield, pods per plant and seeds per pod implied that yield gains under drought stress in the tested genotypes could be made by simultaneously selecting for pods per plant and seeds per pod (Table 7).

Table 6. Genotype means for yield and its components.

Genotypes	Seed yield		Number of pods per plant		Number of seeds per pod	
	no stress	Stressed	no stress	Stressed	no stress	Stressed
SECOW 4W	4.7	5.0	5.8	6.3	16.7	14.7
SECOW 2W	4	5.3	7.5	6.8	16.3	17.7
182	2.7	2.7	4.3	6.5	10.7	13.3
SECOW 1T	5.3	4.3	5.2	5.5	15	19.7
IT 91	3.7	3.7	4	2.3	16	16.0
3306	2.7	3.0	6.5	4.8	15.3	13.7
SECOW 5T	4.3	5.3	5.2	6.2	15	17.3
WC27	4.7	4.0	6.7	7.5	15.3	15.3
NE 45	3.7	2.3	5.7	3.5	16.3	17.0
WC64	5.3	4.0	8.5	8.8	16.7	19.3
SECOW 3B	4.3	5.3	6.8	7.8	15	14.0
SECOW 5X SECOW 1	3.7	2.7	4	3.3	16.7	13.3
NE 20	5.7	4.7	6.7	7.8	14	16.7
NE50	4.7	3.0	3.7	4.3	16	17.7
MU17	5	4.7	6	6.8	16.7	16.0
MU15	4.3	5.3	4.8	4.7	14.3	15.7
WC8	5.7	5.7	7.2	6.8	16.7	14.7
SECOW 5 X NE 39	3.3	3.3	6.5	6.8	14.7	13.3
NE 44	4.3	5.0	8.2	8.5	14.3	15.3
WC30	5	3.7	6.2	3.0	16.3	15.3
NE 70	5.3	4.3	9.3	9.2	13.3	19.7
IT 84	4.3	5.0	3.7	4.5	19.7	17.0
BRAZIL5	4	0.7	5	4.3	11.7	9.7

Continued

NE 67	5	4.0	6.2	4.0	19	15.7
WC67B	5	3.3	6.5	4.8	14.7	16.0
MU 24C	4	5.0	6.8	4.8	15.7	12.0
NE 5	4	4.7	6.5	5.0	18	16.7
NE30	6	5.0	7	6.3	17	17.0
WC 17	5	3.3	6.3	5.3	17.3	20.3
NE6	3.3	2.3	4.5	2.8	8.7	13.3
MEAN	4.4	4.0	6.0	5.7	15.4	15.8
LSD (5%)	0.70	0.70	1.18	1.18	2.09	2.09
CV (%)	19.23	29.27	23.32	32.55	14.58	15.15

NPP = Number of pods per plant, NSP = Number of seeds per pod, SWT = Seed weight in grams, GY = Grain yield in grams, NE & WC: Landraces, SECOW: Serere released varieties, MU: Makerere breeding lines.

Table 7. Genetic correlation among seed yield, number of pods per plant and number of seeds per pod under intermediate stress of 30 cowpea genotypes.

	Number of pods per plant	Number of seeds per pod
Seed Yield	0.57**	0.66***

** , *** significant at $P \leq 0.01$, $P \leq 0.001$ respectively.

4. Conclusion

Considerable variability was observed in morphological, physiological and yield related indicators among the tested cowpea genotypes under drought stress and no stress conditions. Mean yield under stress and non-stress, maintenance of above average chlorophyll content and efficiency of photosystem II, their ability to quickly recover from stress were useful in identifying drought tolerant genotypes. However, the use of wilting scores alone to identify drought tolerant genotypes was not conclusive. Genotypes SECOW 5T, SECOW 3B, SECOW 4W, WC 30 and MU 24 C were recommended as sources of drought tolerance and can be utilized as parents for breeding programs.

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