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PHENOTYPIC AND GENOTYPIC VARIANCE AND HERITABILITY OF STAY GREEN CHARACTER AMONG 22 ELITE SORGHUM (Sorghum bicolor (L.) Moench) GENOTYPES

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ABSTRACT

Delaying of leaf senescence (stay green) is an effective strategy for increasing of crop yield, particularly under water-limited conditions. Study of genotypic variance of stay green is very useful for breeding program of stay green character. Replicated field experiment was carried out in Farm Experimental Station, Faculty of Agriculture Padjadjaran University during dry season. The objectives of this study were: i) to estimate genotypic and phenotypic variances of stay green character; ii) to evaluate and investigate stay green performance of the 22 elite sorghum genotypes. High genotypic and phenotypic variances were observed for plant morphology (plant height, stem diameter), leaf morphology (leaf number, leaf width, leaf length and total leaf area) and yield component (panicle length) and were supported by high estimation of heritability value. Corresponding heritability ranged between 0.77 and 0.94. Genotypic variance of stay green characters was narrow and estimation of broad sense heritability was low. Stay green characters in sorghum might be a quantitative character that was controlled by many genes. Unpad 1.3, Unpad 1.1 and 1090 genotype showed good performance for stay green character. These genotypes can be recommended as stay green parental in breeding program.

Key words: Genotypic variance, Heritability, Phenotypic variance, Sorghum, Stay green

INTRODUCTION

Sorghum is not a new plant in Indonesia and is known already with different name in some region of Indonesia (Anas, 2007). However, sorghum has not been intensively planted in Indonesia. Sorghum could be used either as an alternative food or as bio-ethanol production in some countries (Hoshikawa, 1990; Jorapur & Rajvanshi, 1991). Sorghum is more tolerant to drought than other crop plants (Murty & Kumar, 1995; Metcalfe & Elkins, 1980) and sorghum is very suitable for Indonesia because growth rate of marginal land in Indonesia was very high every year (Anas *et al.*, 2007).

Delaying of senescence or stay green character is an important character because it can improve adaptation of plant to drought especially after flowering (Rosenow & Clark, 1981), lodging resistance and disease resistance to *Macrophominia phaseolina* (Tassi) Goid (Mughogho & Pande, 1984). Stay green is also associated with level of carbohydrate content in stem and increase of grain filling period of plant in water stress condition (Borrell & Douglas, 1996). Breeding programs for stay green character has been done in United States (Rosenow *et al.*,1983) and Australia (Henzell *et al.*, 1992). Stay green plants showed higher total biomass than those did not have stay green character. Stay green plants can store more food in their stem (Borrell *et al.*, 2000a). Stay green plants showed also high concentration of leaf nitrogen and were associated with a higher efficiency in transpiration (Borrell *et al.*, 2000b).

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Genotypic variance and heritability are very useful genetic parameters that determine effectiveness of selection process in plant breeding program (Sleper & Poehlman, 2006). Some of cereal crops show genetic variation of leaf senescence during grain filling period (Thomas & Smart, 1993). Sorghum show also variation for stay green character, and is strongly influenced by environment and genetic background of plant (Thomas & Howarth, 2000). There is considerable genetic diversity of sorghum (Anas & Yoshida, 2004, Murray *et al.*, 2009; Shehzad *et al.*, 2009) and gene control of leaf senescence (*See*) has been cloned successfully from several plant species. Homologous genes *See* also has been isolated from several plant fruits and part of flowers of some plants (Nam, 1997; Medina-Sua'rez *et al.*, 1997).

Broad genetic background of elite sorghum genotypes was used in field experiment to study variability and heritability of stay green character. These elite sorghum genotypes have been collected from diverse research center and had been bred from different genetic background of parents (Anas & Yoshida, 2000). Objectives of this study were: a) to estimate genotypic variance and heritability of stay green character, b) to evaluate performances of stay green character of some elite sorghum genotypes.

MATERIALS AND METHODS

Experimental design, Genetic materials and Observation

Field trial was arranged in Randomized Complete Block Design with two replications at the experimental farm station of Faculty of Agriculture, Padjadjaran University (Unpad), Jatinangor West Java. Treatments consist of 22 elite sorghum genotypes of Plant Breeding Lab. Faculty of Agriculture Unpad (Table 1). Planting space was 10 cm x 75 cm with a distance of 1 m between plots.

Data were recorded for 11 characters consisting of morphological and physiological characteristics. Total leaf area of sorghum was measured as a function of the sixth leaf using equation $TLA = 14.355 \, (SLA_{LW})^{0.857} \, (Chinnamuthu \, et \, al., \, 1989)$. SLA_{LW} is the sixth leaf area (maximum length multiplied by maximum width) of sorghum plants. Analysis of stay green character was estimated from component of leaf color (LC). Brightness and color of sixth leaf was measured using Digital Color Analyzer tool on three primary colors Red-Green-Blue. The data were then processed using the Color Picker (Adobe Photoshop CS.3 Version 10.1, 1990-2007). RGB values obtained were then plotted in 3D graphics using SPSS Ver. 16.0.1 (2007).

Table 1. Material genetic in field experiment

	Released in 2001, Maros			
2 UPCA-S1	Released iii 2001, Maios	12	Wray	Sweet sorghum, Jepang
	Released in 1972, Maros	13	Cantel	Local variety, Maros
3 Kawali	Released in 2001, Maros	14	II-24	Elite genotype, Unpad
4 Unpad 1.1	Variety, Unpad	15	10.1	Elite genotype, Unpad
5 B.100	Vaeriety, BATAN	16	10.2	Elite genotype, Unpad
6 4.1	Elite genotype, Unpad	17	2.1	Elite genotype, Unpad
7 1090	Elite genotype, Maros	18	2.2	Elite genotype, Unpad
8 ICR 3	Elite genotype, ICRISAT India	19	Keller	Sweet sorghum, Jepang
9 Super Sorgo	Sweet sorghum, Jepang	20	Batari	Variety, Maros
10 I-34	Elite genotype, Unpad	21	Unpad 1.3	Elite genotype, Unpad
11 Big S. Sugar	Sweet sorghum, Jepang	22	Taomitsu	Sweet sorghum, Jepang

Statistical Analysis

Differences among genotypes were analyzed by Scott Knott test (Gaspersz, 1995). Phenotypic (σ_f^2) and genotypic variance (σ_g^2) and heritability were estimated based on value of expected mean square according Singh and Chaudhary (1979) and Bos & Caligari (1995). Criteria of heritability were estimated according Stansfield (1991). Phenotypic and genotypic variance was then compared with twice of standard deviation values of genotypic and phenotypic variance. Standard deviation of the phenotypic variance and genotypic variance was calculated according Anderson and Bancroft (1952) cited Pinaria *et al.* (1997) and Wahdah *et al.* (1996).

RESULTS AND DISCUSSION

The data collected were subjected to ANOVA using SPSS for assessment of the stay green genotypes. Genotypic differences were highly significant for all characters observed except leaf color (G), leaf brightness and chlorophyll content. Number of leaf (NL), leaf length (LL), leaf width (LW) and total leaf area (TLA) showed wide genotypic and phenotypic variance (Table 2). Genotypic variance is one of the genetic parameters that determine effectiveness of selection process in plant breeding program (Sleper & Poehlman, 2006). Effectiveness of selection process of a character depends on environmental effect for these characters. Morphological characters of leaf showed high estimation of heritability values (Table 2). This suggested that the diversity of leaf morphology was mainly due to genetic factors and selection was more effective on these characters.

Sweet sorghum and white grain sorghum Unpad 1.1 showed higher number of leaf than Batari, Numbu, Cantel and B100 genotypes (Table 3). The NL character determined crop canopy (TLA) that was associated with evapotranspiration and photosynthesis activity of plants (Arkin et al., 1983). Improvement of TLA character will increase wide of crop canopy that will enhance absorption of sunlight by plants, which in turn will increase rate of transpiration and photosynthesis process. It will also reduce evapotranspiration from soil so will keep moisture of soil.

Many researchers used TLA character as indicator for studying of stay green character in sorghum. Stay green character in sorghum is often associated with tolerance to drought stress after flowering (post-flowering drought tolerance) (Xu et al., 2000). Slowing of senes-

Table 2. Genotypic and phenotypic variance and estimation of heritability of some characters

Character	$\sigma_{_{gen}}^{^{2}}$	Criteria	$\sigma_{_{phe}}^{^{2}}$	Criteria	Н	Criteria
Leaf number	0.99	Н	1.20	Н	0.83	Н
Leaf width (cm)	0.63	Н	0.95	Н	0.66	Н
Leaf length (cm)	81.18	Н	96.05	Н	0.84	Н
Total leaf area (cm2)	290727.00	Н	379725.80	Н	0.77	Н
Leaf color (green)	50.96	M	238.16	Н	0.21	M
Leaf brightness	2.84	M	26.21	Н	0.19	L
Chlorophyll content (unit)	36.98	M	162.20	Н	0.23	M
Plant height (cm)	2804.72	Н	2971.95	Н	0.94	Н
Stem diameter (cm)	0.05	Н	0.07	Н	0.69	Н
Panicle length (cm)	12.00	Н	14.23	Н	0.84	Н
Seed weight plant-1 (g)	146.44	L	368.27	Н	0.39	M

Note: σ_{gen}^2 = genotypic variance; σ_{fen}^2 = phenotypic variance; H= heritability; H= High; L= low; M= medium.

cence is one of effective strategies to increase yields of cereal crop in water-stress condition. There were association between stay green and grain yield that was affected by environment and genetic background of plant (Jordan *et al.*, 2012). Five sweet sorghums showed good TLA and not significantly different with superior grain sorghum genotypes Unpad 1.1 or Unpad 1.3 (Table 3). Super Sorgo showed the highest TLA character and followed by 2.1 and UPCA-S1 genotype. On the other hand, I-34, II-24, ICR3, Batari and Cantel cultivars showed low TLA character.

Low genotypic variance was exhibited by components of stay green character, such as green component of leaf color (LC). Heritability of green component of LC, chlorophyll content (CC) of leaf and leaf brightness (LB) were relatively low (Table 2). This suggested that effect of environment was high to components of stay green character and selection for these characters were less effective. Stay green characters in sorghum might be a quantitative character that was controlled by many genes and was strongly influenced by environmental factors. Some researchers reported that many genes scattered in several places in sorghum genome controlled stay green character.

Stay green character of some plants was controlled by *See* gene (Nam, 1997; Medina-Sua'rez *et al.*, 1997). However, a lot of research on genetic control of stay green character used QTLs approach in their study (Subudhi *et al.*, 2000; Borrel *et al.*, 2000a; Tao *et al.*, 2000). Kebede *et al.* (2001) have reported nine QTLs in seven-linkage group of sorghum genome associated with stay green character. While four QTLs (*Stg1*, *Stg2*, *Stg3* and *Stg4*) correlated with stay green character and *Stg2* QTL was most important in controlling stay green characters in sorghum (Harris *et al.*, 2006). Stay green character and some components of stay green character was controlled by additive gene (Borrel *et al.*, 2000a) and was associated with multiple locations of linkage group B and I of sorghum genome (Tao *et al.*, 2000).

Table 3. Performance of some characters of 22 sorghum genotypes

Genotype	L	.N	TL	Α.	L	C(G)	L	В	C	С	P	Н	SI)	P	L
Numbu	9	а	3179	b	130	а	52	а	55	а	187	С	1.97	b	20	а
UPCA-S1	11	b	3298	b	121	а	49	а	36	а	161	С	2.12	b	23	b
Kawali	11	b	2565	а	106	а	43	а	32	а	157	С	1.81	а	25	С
Unpad 1.1	12	b	3095	b	122	а	49	а	45	а	134	b	2.07	b	23	b
B.100	10	а	3243	b	130	а	52	а	39	а	170	С	1.86	а	21	b
4.1	9	а	2437	а	112	а	46	а	34	а	177	С	1.64	а	19	а
1090	11	b	3287	b	122	а	49	а	34	а	243	d	2.02	b	23	b
ICR 3	10	b	1936	а	118	а	48	а	41	а	112	а	1.69	а	22	b
Super Sorgo	11	b	3527	b	120	а	49	а	54	а	249	d	2.35	b	28	С
I-34	8	а	1563	а	125	а	52	а	34	а	125	а	1.38	а	16	а
Big Super S.	11	b	3202	b	117	а	48	а	55	а	254	d	2.14	b	29	С
Wray	11	b	3120	b	79	а	36	а	54	а	255	d	2.36	b	23	b
Cantel	10	а	2435	а	106	а	44	а	53	а	180	С	1.97	b	24	С
II-24	8	а	1769	а	121	а	49	а	55	а	117	а	1.72	а	25	С
10.1	11	b	2566	а	120	а	48	а	39	а	140	b	1.89	а	27	С
10.2	10	b	2450	а	118	а	48	а	40	а	132	b	1.76	а	22	b
2.1	9	а	3461	b	127	а	51	а	47	а	243	d	2.12	b	26	С
2.2	11	b	2814	b	114	а	47	а	58	а	131	b	1.90	а	26	С
Keller	11	b	3172	b	107	а	44	а	68	а	274	d	2.19	b	24	b
Batari	9	а	1997	а	116	а	47	а	37	а	224	d	1.60	а	33	С
Unpad 1.3	9	а	3267	b	136	а	54	а	37	а	219	d	1.83	а	22	b
Taomitsu	11	b	3042	b	100	а	48	а	43	а	248	d	2.07	b	29	С

Note: Value followed with same alphabet in a column show not different based on *Scott-Knott* test at 5% level. LN=leaf number; TLA=total leaf area (cm2); LC(G)= green component of leaf color; LB=leaf brightness (unit); CC=chlorophyll content (unit); PH=plant height (cm); SD=stem diameter (cm); PL=panicle length (cm).

Green components of leaf color showed no significant differences among genotypes. Differences of stay green character among genotypes could not be distinguished during field trial test. High frequency of rainfall even dry season caused small difference of LC among genotypes. Stay green character will be expressed and are more pronounced among plants in water stress conditions. Plants that have no stay green character will rapidly senescence (Thomas & Howarth, 2000). In general, elite grain sorghum genotypes of Plant Breeding Lab. Padjadjaran University showed high average of LC and LB (Table 3). Otherwise, generally sweet sorghum and released cultivars showed high chlorophyll content (CC)

Using digital color analyzer, leaf of I-34 genotype showed dark color and contribution of green color component was low (Table 3). This suggested that leaves have started to dry up and was in line with early maturity character of this genotype. Judging from percentage contribution of green color components in overall RGB system, 1090 genotype, Unpad 1.3 and Unpad 1.1 genotype showed high percentage of green color components. This suggested that these genotypes could be developed as parent in stay green breeding programs.

Percentage of green component and brightness level of Taomitsu and Wray leaves were very low. In general, sweet sorghum genotypes showed lower level of brightness than grain sorghum genotype. Sorghum leaves that were starting to dry will have lower brightness level than green leaf. However, Wray and Keller have been used as parent in breeding program for high sugar content (Murray *et al.*, 2009).

Genotypic and phenotypic variance of plant morphology, such as plant height (PH), stem diameter (SD), number of node (NN) and some yield components characters showed wide variability (Table 2). This was also supported by high heritability of these characters. Genotypic and phenotypic variances that were sharing value equally and well supported by high heritability value will greatly important for plant development program.

PH and SD character of grain sorghum were shorter and smaller significantly than sweet sorghum genotypes. According Tsuchihashi & Goto (2005) difference of sorghum plant height was a result of increasing number of node (25%) and 75% was result of internode elongation. According Sabadin *et al.* (2012) there were not associations of stay green QTL with differences in flowering time/plant height. Grain sorghum Unpad 1.1 has been bred for short plant height to facilitate harvesting process. Meanwhile, stem diameter greatly affected biomass of yield stem. According Tsuchihashi & Goto (2004), proper cultivation techniques, spacing and planting density per unit area greatly affected process of thickening of sweet sorghum stem that determined weight of stem. Size of sweet sorghum stems was greatly influenced by four factors i.e. climate, soil type, fertilization and planting density. Additionally stem weight was strongly influenced by characteristics of stem and water availability during the growing process (Tsuchihashi & Goto, 2005).

CONCLUSION

Generally, morphological characters of sorghum showed high genotypic and phenotypic variance and was supported by high estimation of broad sense heritability value. Genotypic variance of stay green characters was narrow and estimation of broad sense heritability was low. Unpad 1.3, Unpad 1.1 and 1090 genotype could be developed as stay green sorghum and can be used as parent in breeding program of stay green character.

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