Selection of SSR Markers for drought resistant sugarcane in Thailand

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Abstract Drought is the most significant environmental stress factor affecting agriculture worldwide and improving yield under drought conditions is a primary goal of plant breeding. In Thailand, sugarcane productivity decreased by approximately 12.5 tons/hectare from 2014 to 2016 as a result of drought. Drought tolerance comprises the combination of many characteristics. Indirect screening methods based on physiological and genomic traits are used to select drought-tolerant varieties with increased growth efficiency. Physiological traits observed in this study were stomatal conductance (g_s) , PSII maximum quantum yield (F_v/F_m) , and leaf chlorophyll content (SPAD index). These were correlated with agronomic traits such as height, weight, and a number of stalks. Fourteen candidate SSR markers were selected as genomic data to determine the grouping relationship. Under drought conditions, our results indicated that maintaining higher photosynthetic efficiency (F_v/F_m) and chlorophyll content (SPAD index) showed the potential to achieve greater growth under water stress conditions. K93-219 had the highest values of physiological traits followed by KPS01-12, UT12, MPT10-52, and MPT03-320, respectively. A phylogenetic tree of 4 SSR markers gave an interesting pattern suggesting that K93-219 and UT12 were close neighbor groups followed by MPT10-54 and KPS01-12, respectively. Results can be used to model selective varieties of sugarcane in Thailand. This research demonstrates an alternative method for screening sugarcane varieties that can adapt and grow under water stress conditions, and offers opportunities to develop breeding approaches for crop improvement in Thailand.

Keywords: SSR marker, sugarcane, physiology, drought

Introduction

In Thailand, more than 75% of the sugarcane is produced in the northeastern region, with slightly more than half of the production located in rain-fed areas of mainly sandy soil (Ministry of Agriculture and Cooperatives, 2013). As

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a result, moisture content tends to decrease quickly along with fertile qualities. In addition, El Niño and La Niña events have recently been a natural global occurrence and adversely affected rainfall in Thailand. Climate change has produced variations in temperature, rain distribution, and precipitation frequency. As a result, all crops in Thailand now face varied growing conditions. One of the most pronounced changes in climatic conditions is drought.

Drought is one of the major abiotic stresses limiting global growth and development of crops including sugarcane (da Silva et al., 2012). Drought alters plant-water relationships via stomatal activity, carbon fixation and other plant metabolic functions which result in changes in growth and development. Stomatal closure during water stress reduces the flux of carbon dioxide into the leaf and photosynthesis declines (Flexas et al., 2006 and Lakshmanan and Robinson, 2014). In addition, water stress-induced leaf senescence reduces leaf area and thus water loss, thereby contributing to the maintenance of a favorable water balance at the whole plant level. As a result, photosynthesis of the whole canopy is lowered with reduced stomatal conductance (g_s) and green leaf area. From a crop production perspective, these responses negatively impact yield (Rivero et al., 2009). Physiological traits are directly or indirectly associated with crop growth and yields (Silva et al., 2007; Inman-bamber et al., 2012, and Wilkinson et al., 2012). Physiological adaptations to water stress at the whole plant level are highly complex due to the diverse responses elicited by distinct plant species to different levels of water stress and involve both deleterious and adaptive changes (Kramer and Boyer, 1995). The net effect of water deficit is largely determined by the dynamics, duration, and intensity of soil water depletion, atmospheric water demand and other prevailing environmental stresses including the stage of plant growth and phenology (Grant, 2012).

Drought mostly affects sugarcane during the summer season when the plant is at the tillering and elongation stage at around the 3rd to the 7th month after propagation. In response to drought, sugarcane displays pronounced morphological and growth responses that include a reduction in stalk and leaf elongation and accelerated leaf senescence. Together, these responses result in a decrease in biomass production and sucrose yield due to reduced rates of photosynthesis and crop growth as well as altered partitioning of assimilates between roots, leaves, structural stem material and stored sucrose (Singels *et al.*, 2000; Singels and Inman-bamber, 2002; Inman-bamber, 2004, and Singels *et al.*, 2010). Lost production yield associated with drought is estimated at 12.5 tons/hectare in the Thai sugarcane industry.

In response to this challenge, sugarcane breeders focus on conventional breeding of new varieties but the characteristics of drought tolerance are

complicated. Moreover, conventional breeding takes a long time with high investment to screen and select an elite variety. Sugarcane is an annual crop and 7-10 years are required to propagate a new variety (Juttupoornpong *et al.*, 2012). Indirect methods for the selection of new sugarcane cultivars offer the potential to select and screen elite varieties based on the use of phenotypic data related to drought characteristic traits.

The candidate varieties were planted in a greenhouse for a period of 8 months. SSR markers were selected using a phylogenic tree that related physiological data as stomatal conductance (g_s), PSII maximum quantum yield (F_v/F_m), leaf chlorophyll content (SPAD index), and agronomic traits such as height, number of stalks per pot, and weight.

The objective was to select SSR markers for drought-resistant sugarcane in Thailand. Conventional methods use sexual reproduction to breed many seedlings. Sugarcane was planted as single seeds in a row and the seedlings were transplanted into individual pots. Normally, potential sugarcane was selected by physiological data talking for several years. SSR marker can be indicated trait associate to reduce the time for selection characteristic of sugarcane at least 30 percent on a breeding program and remained varieties to have the constant characteristic.

Materials and Methods

Plant material and growth conditions

Sugarcane (*Saccharum officinarum* L. varieties K93-219, KPS01-12, KK3, PHIL66-07, MPT10-54, UT12, MPT03-320, SP50, TBY20-2248, and KPK98-40) were planted in a greenhouse at Mitr Phol Sugarcane Research Center Phu Khiao district, Chaiyaphum Province, Thailand. Each variety was planted by RCDB (Randomized complete block design) in three pots of 15-inch diameter and grown under two conditions drought (D) and irrigated (I). Fertilizer was applied three times, 1) on the planting day at ratio 312.5 kg of 16-16-8 NPK per hectare, 2) at 3 months with ratio 312.5 kg of 21-7-18 NPK per hectare, and 3) at 4.5 months with ratio 156.25 kg of 21-7-18 NPK per hectare.

Measurement of physiological and morphological parameters

Measurements of phenotypic data collected at the 3rd, 5th and 7th month stages of sugarcane growth were edited from Somkit *et al.* (2007); Sharma (2009), and da Silva *et al.* (2012). Each pot in the greenhouse was used to collect the data in each block. The germination rate of sugarcane was measured

at 45 days. Number of stalks per pot, the diameter of each stalk, number of internodes, and height of sugarcane were measured at the 3rd, 5th and 7th month stages of growth.

PSII maximum quantum yield (F_v/F_m): Chlorophyll fluorescence was measured by Handy PEA (Hansatech Instruments, UK) using at least four leaves per sample. The leaves were shielded from light for 15-30 min using leaf clips before measurement following the method adjusted from Kautsky and Hirsch (1931) for use in sugarcane.

Stomatal conductance (g_s) : Measurements were made using three Porometer AP-0 Instruments (Delta-T Devices, Cambridge, UK) and average values were calculated. Usability concerned daily temperature and humidity, with appropriate storage time between 8 and 14 hours. The method was adjusted from Stiles (1970) for use in sugarcane.

Leaf chlorophyll content (SPAD index): SPAD index was measured using SPAD-502 plus (Konica Minolta, Japan). Average values were measured in the middle portion of the four youngest fully expanded leaves (Dray *et al.*, 2012).

Molecular analysis

Fourteen SSR genetic markers were selected by a literature review that related to drought in sugarcane and 10 varieties in the greenhouse were extracted for DNA. PCR reaction was improved from the protocol of Hoisington (1992). Twenty microliters consisted of 5 µl of 10 ng/µl DNA sample, 1.0 µl of 25 mM MgCl₂, 1.2 µl of 2.5 mM dNTP Mix, 2.0 µl of 10X *Tag* buffer, 5.0 µl of 1.0 µM for forward and reverse primers, 2.0 µl of %100 glycerol, 0.2 µl of 5 U/µl *Tag* polymerase, and 3.6 µl of ddH₂O. PCR amplification was performed by T100 Thermal Cycle Bio-Rad (Bio-Rad, CA, USA) in a program of 94°C 3 min, 35 cycles of 94°C 1 min, annealing 2 min, 72°C 1 min, extension at 72°C 5 min and kept at 4°C. Annealing temperatures ranged between 50 and 56°C depending on each SSR primer. A pre-test was performed by agarose gel electrophoresis and then the PCR product was analyzed by the ZAG system (ZAG DNA Analyzer AATI, CA, USA).

Statistical analysis

The greenhouse experiment was arranged in a randomized complete block design with three replications. Plants were grown under well-watered and water stress conditions. All phenotypic data were clarified and analyzed by SPSS (IBM, NY, USA) using the general linear model (GLM), ANOVA and Tukey's test at a 95% confidence interval. Then, the average of the three values was reported as the measured value with the standard deviation. All genetic data from ZAG were analyzed by ProSize software (AATI, CA, USA). Analyzed data were exported to Microsoft Excel® and GeneTools (Syngene, MD, USA) and then a phylogenetic tree was constructed by GeneDirectory (Syngene, MD, USA). The phylogenetic tree was compared with phenotypic data to select SSR marker groups related to drought characteristics. Highest score groups from the phenotypic data were likely to give a high yield of sugarcane in drought conditions, while lowest score groups were likely to give a low yield of sugarcane in drought conditions. SSR markers were combined and assessed to form a phylogenetic tree that gave the best pattern to explain both agronomic and physiological results.

Results

Growth and physiology in drought conditions

Sugarcane was planted in a greenhouse using three replications of (RCBD), and data were collected during the 3rd, 5th and 7th month of sugarcane growth. All data were analyzed by SPSS using the general linear model, ANOVA, and Tukey's test at a 95% confidence interval. Significant effects (P<0.05) were observed for different interaction on six physiological parameters as a number of stalks per plant, height, weight, stomatal conductance, photosynthesis efficiency and chlorophyll content (Table 1). A few sugarcane stalks showed significance at the 3rd and the 5th months but no significance at the 7th month. Varieties having the greatest characteristics of the stalk were KK3 at the 3rd month and MPT10-54 at the 5th month. The one-way interaction of variance components on height, photosynthesis efficient, and chlorophyll content were significant at the 3rd, 5th and 7th months of growth. PHIL66-07 was the main character in the height parameter for all months; however, some varieties recorded a high score in some months e.g. K93-219 and MPT10-54 in the 5th month, and KPS01-12 and MPT10-54 in the 7th month. For photosynthesis efficiency, K93-219, MPT03-320, SP50, and TBY20-2248 had the highest score in the 5th month, while MPT10-54 gave the highest score in the 7th month. UT12 had the highest score of chlorophyll content in the 7th month. The treatment characteristic was not significant for weight and stomatal conductance. The highest average score of stomatal conductance was recorded in KK3 and MPT10-54 in the 5th and 7th month. respectively. On the other hand, the highest average score of weight was recorded in SP50.

Genetic markers of drought tolerance with sugarcane

Fourteen SSR markers were tested for sugarcane genetic diversity and separated into five groups as ABA signaling pathway, protein signaling, electron transport hemoprotein, photosynthesis, and hydrogen peroxide removal (Table 2). The dendrogram of the SSR based analyses of 10 sugarcane cultivars that exhibited high diversity of DNA polymorphism clearly discriminated between genotypes (Figure 1). SSR markers amplify specific regions with different patterns between the groups. The dendrogram was constructed using four markers (SSR9, SSR80, A19, A70) that related to drought function that ABA signaling pathway and hydrogen peroxide removal. Four groups of genetic patterns were separated in the dendrogram. KK3 showed the most difference at 0.66 coefficient in the first group. The second group consisted of PHIL66-07 and MPT03-320 at 0.716 coefficient. TBY20-2248 and KPK98-40 showed similarities at 0.79 coefficient in the third group together with the SP50 cultivar. The last group consisted of K93-219, UT12, MPT10-54, and KPS01-12.

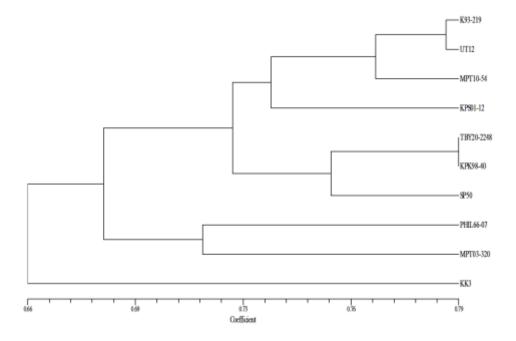


Figure 1. UPGMA dendrogram derived using Jaccard coefficient of similarity based on four SSR genetic distance data of 10 sugarcane varieties

Table 1. Sugarcane phenotypes in the greenhouse

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Treatment	Stalks per j	plant 5M	7M	Height 3M	5M	73.6	Weight 7M	Stomatal conductance		Photosynthesis efficiency 5M 7M		Chlorophyll content 7M	
K93-219-D	7±0.71 ^{ABC}	5±1.66 ABC	4±0.97 ^A	41.78±6.04 ^{BC}	89.89±12.71	7M 134.78±12. 96 ^{ABCD}	4.33±1.31 ^A	5M 2.48±2. 20 ^A	7M 3.01 ±2. 24 ^A	649.1±19.69 ^A	774.6±11. 08 ^{ABC}	35.51 ±6.55 ^{ABC}	
K93-219-I	6±1.17 ^{ABC} DE	5±1.50 ABC	$4\!\pm\!1.58^A$	$33.00\pm\!4.66^{CD}$	76.94±11.96 abc	146.44±19. 87 ^{ABC}	3.78±0.68 ^A	3.95 ±2. 74 ^A	3.54±2. 67 ^A	742.3±3.11 ^A	789.2±7.0 8 ^{AB}	31.23±5.14 ^{BC}	
KK3-D	9±4.00 ^A	5±2.51 ABC	4 ± 1.50^{A}	35.56±5.77 ^{CD}	$^{81.33\pm\!8.43^{A}}_{_{B}}$	105.67 ±12. 31 ^{CDEF}	3.67±0.44 ^A		2.11±1. 24 ^A	542.4 ± 18.93^{B}	777.2±13. 22 ^{ABC}	34.19±2.71 ^{BC}	
KK3-I	5 ± 2.06^{DE}	$\underset{ABC}{5\pm1.87}$	$4\!\pm\!1.48^A$	$_{F}^{28.22\pm5.52^{DE}}$	63.78±10.27 ABC	141.78±13. 65 ^{ABC}	4.22±1.13 ^A	3.85 ±3. 27 ^A	2.28±1. 32 ^A	$^{706.2\pm10.56^{A}}_{_{B}}$	792.7±11. 28 ^{AB}	35.44±6.71 ^{ABC}	
KPK98-40- D	5±1.33 ^{CDE}	4±1.73 ABC	3±1.09 ^A	39.11±7.85 ^{AB}	67.44±25.74 ABC	92.44±22.9 0 ^{DEF}	3.28±0.17 ^A	2.74±2. 35 ^A	2.78±1. 87 ^A	$^{648.2\pm\!11.38^{A}}_{_{B}}$	774.9±16. 91 ^{ABC}	33.91 ±4.66 ^{BC}	
KPK98-40-I	5±1.12 ^{CDE}	4±1.69 ABC	4±0.73 ^A	32.22 ± 7.21^{CD}	56.21 ±17.39 BCD	119.33±13. 13 ^{ABCDEF}	3.56±0.34 ^A	4.14±4. 02 ^A	3.48±2. 42 ^A	698.9±7.89 ^{AB}	780.9±15. 61 ^{ABC}	31.62±4.15 ^{BC}	
KPS01-12-D	$7\pm1.81^{ABC}_{DE}$	5±1.32 ABC	4±1.73 ^A	39.67 ±4.42 ^{AB}	90.56±6.88 ^A	119.11±28. 81 ^{ABCDEF}	4.33±0.75 ^A	4.23 ±2. 19 ^A	1.25±0. 74 ^A	623.6±12.32 ^A	776.8±17. 85 ^{ABC}	33.91 ±5.44 ^{BC}	
KPS01-12-I	7 ± 2.06^{ABC} DE	6±2.39 ABC	4±1.00 ^A	40.11±9.53 ^{AB}	70.04 <u>±2</u> 5.99 ABC	160.78±15. 65 ^A	4.00±0.47 ^A	3.30±1. 88 ^A	3.35±3. 04 ^A	570.8±22.09 ^A	776.7±12. 87 ^{ABC}	33.80 ± 2.56^{BC}	
MPT03-320- D	$_{\text{DE}}^{7\pm1.22^{ABC}}$	5±1.51 ABC	3±0.53 ^A	35.56±3.43 ^{CD}	69.22±9.77 ^A BC	91.22±20.3 6 ^{EF}	3.44±1.45 ^A	1.78±1. 60 ^A	1.68±1. 24 ^A	722.2±4.72 ^{AB}	780.2±8.6 7 ^{ABC}	33.30±2.72 ^{BC}	
MPT03-320- I	6±1.33 ^{ABC} DE	5±2.29 ABC	4±0.73 ^A	30.94±4.13 ^{CD} EF	54.22±12.81 BCD	113.00±22. 37 ^{BCDEF}	3.56±0.54 ^A	4.78±4. 18 ^A	3.91±2. 89 ^A	730.6±1.86 ^A	778.3±9.5 5 ^{ABC}	31.41±6.60 ^{BC}	
MPT10-54- D	8±1.22 ^{ABC}	6±1.13 ABC	5±1.39 ^A	37.67 ±4.12 ^{BC}	90.78±8.70 ^A	114.00±34. 84 ^{BCDEF}	4.78±1.48 ^A	3.41±1. 91 ^A	2.51±2. 20 ^A	641.2±12.52 ^A	782.1±21. 85 ^{ABC}	33.83 ±4.66 ^{BC}	
MPT10-54-I	7±2.69 ^{ABC}	7±3.77	4±1.13 ^A	33.78±5.52 ^{CD}	61.72±18.67 ABCD	159.22±15. 58 ^A	4.44±0.25 ^A	5.13±5. 02 ^A	4.46±3. 82 ^A	680.4±9.51 ^{AB}	796.9±4.4 6 ^A	29.60±3.73 ^C	
PHIL66-07- D	5 ± 2.05^{DE}	4±0.71 BC	4±2.12 ^A	46.89±9.06 ^{AB}	92.00±13.01	134.11±46. 95 ^{ABCDE}	4.00±0.42 ^A	3.39±2. 96 ^A	2.02±0. 82 ^A	667.9±9.62 ^{AB}	769.9±21. 99 ^{BC}	33.49±6.28 ^{BC}	
PHIL66-07- I	4±1.74 ^E	5±2.65 ABC	4±1.32 ^A	50.22±15.47 ^A	79.92±41.33 AB	158.78±12. 81 ^A	3.67±0.50 ^A	1.80±1. 74 ^A	2.35±1. 26 ^A	698.2±10.40 ^A	791.1±8.8 1 ^{AB}	32.86±3.14 ^{BC}	
SP50-D	9±1.83 ^{AB}	7±2.35 AB	4±0.50 ^A	40.00±9.45 ^{AB}	81.33±17.36 AB	112.89±29. 81 ^{BCDEF}	3.67±0.42 ^A	4.97 ±4. 33 ^A	3.64±3. 41 ^A	682.7±10.28 ^A	776.7±14. 55 ^{ABC}	34.31±3.22 ^{BC}	
SP50-I	6±1.20 ^{BCD}	6±1.36 AB	5±1.22 ^A	37.67 ±4.33 ^{BC}	63.08 ±20.90 ABC	148.89±30. 18 ^{AB}	5.33±1.49 ^A	4.14±3. 37 ^A	2.09±1. 58 ^A	744.3±2.76 ^A	785.6±12. 44 ^{ABC}	36.39±3.19 ^{ABC}	
TBY20- 2248-D	6±3.12 ^{BCD}	2±2.00	4±0.94 ^A	24.33 ±4.23 ^F	30.00±26.41	86.00±26.9 1 ^F	3.50±0.31 ^A	5.70±2. 62 ^A	1.74±0. 89 ^A	684.8±15.02 ^A	779.0±15. 40 ^{ABC}	36.07±3.89 ^{ABC}	
TBY20- 2248-I	6±2.74 ^{BCD} E	6±3.35 AB	4±1.50 ^A	26.33±3.67 ^{EF}	45.89 ±24.89 CD	93.47±51.5 8 ^{DEF}	4.42±0.48 ^A	3.50±2. 43 ^A	1.58±0. 78 ^A	746.8±4.91 ^A	782.2±11. 61 ^{ABC}	34.32±5.16 ^{BC}	
UT12-D	7±0.93 ^{ABC}	6±1.01 ABC	5±1.54 ^A	33.67 ±5.07 ^{CD}	70.22±10.79 ABC	115.56±23. 10 ^{BCDEF}	4.89±1.86 ^A	2.90±2. 17 ^A	1.28±0. 61 ^A	601.4±6.34 ^{AB}	766.2±14. 54 ^C	38.31±3.04 ^{AB}	
UT12-I	7±0.87 ^{ABC} DE	6±1.41 AB	5±1.33 ^A	34.89±5.44 ^{CD} EF	60.56±15.95 ABCD	135.56±26. 14 ^{ABCD}	4.56±0.95 ^A	2.18±1. 34 ^A	3.05 ±2. 61 ^A	727.0±6.34 ^{AB}	779.3±5.7 4 ^{ABC}	42.20±4.57 ^A	
SD	0.8876	0.9816	0.5995	3.1422	9.0312	12.174	0.4302	1.6705	1.0286	5.27	6.4398	2.1819	
CV	3.1443	3.4773	2.1239	11.132	31.994	43.129	1.5239	5.9181	3.6439	18.67	22.814	7.7296	
Significant	*	*	NS	*	*	*	NS	NS	NS	*	*	*	

M = month, NS = not significant, D = drought condition, I = irrigated condition

Table 2. SSR markers related to drought response in sugarcane

No	Marker Name		Primer sequence (5'-3')	Range	Target	Significance	Reference	Sequence repeat*
1	SSR9	(F)	AAGAAAAGGAGGCCAAAAA	204-320	Proteinase phosphatase	Part of the ABA	(Sharma, 2009)	
		(R)	GCCAGGCAAGAGGATAAAA		2c homolog (ABA) Cysteine protease	signaling pathway	•	
2	SSR80	(F)	GTTCCCACCGCTGTCATC	229	component of protease	Part of the ABA	(Sharma, 2009)	gtc, cgc
		(R)	TACGAGCACGTGTCCAACTC		inhibitor complex (ABA)	signaling pathway		
3	SSR230	(F)	TTGTGCTGATGTTTCCTGCT	200-380	Patatin like protein		(Sharma, 2009)	ta, at
	33K230	(R)	CAAGAGAAGATGCCATTAGCC	200-380	r atatiii like proteili		(Sharma, 2009)	ta, at
4	GGD024	(F)	CCGAGTGTCCTCATCGCAGAAC	200 200	Auxin-independent	Part of the ABA	(Sharma, 2009)	cgc,
	SSR924	(R)	CTCTAGTCTCTTCATAACCTCTC	200-300	growth promoter (ABA)	signaling pathway		cctcgc
5	SCM7	(F)	ACGGTGCTCTTCACTGCT	157-169			(Lu et al., 2015)	taca
	SCIVI7	(R)	GGGCATACTTCCTCCTCTAC	137-109			(Lu et al., 2013)	tgcg
6	ESTB100	(F)	CCACGGGCGAGGACGAGTA	268	22 kDa drought- inducible protein	Protein signaling	(Oliveira et al., 2009)	ta, cgg
	E31B100	(R)	GGGTCCTTCTTCGCCTCGTG	208				
7	EGED 120	(F)	GCCCAGGTAATTATCCAGACTC	124	Putative auxin response		(01: 1 2000)	
	ESTB130	(R)	GCTGTTGCTCACTGGTTCC	124	factor 7a		(Oliveira et al., 2009)	gca
8	ESTC82	(F)	GGCGGCGGCTGGCTGGAT	149	Cytochrome b5	Electron	(Oliveira et al., 2009)	gcc, gca, caac
	E31C82	(R)	GATTTGTGGCTGGCGGAAGTGGAC	149		transport hemoproteins		
9	ESTC83	(F)	ATTTGTGGCTGGCGAGGTGGAC	149	Cytochrome b5	Electron	(Oliveira <i>et al.</i> , 2009)	caac, gcgt, cta
	ESICOS	(R)	GGCTGGCGGCTGGAT	149		transport hemoproteins	(Onvena et al., 2009)	
10	ESTC110	(F)	ACATGATCGCCGTCCTCTG	138	Putative cytochrome		(Oliveira <i>et al.</i> , 2009)	
		(R)	GCAAAGGCAGAAAAAGGTGTT	136	p450		(Onvena et at., 2007)	
11	ESTC117	(F)	GGGAGCGACGAACTGACG	295	Chlorophyll a/b-binding	Photosynthesis	(Oliveira et al., 2009)	acc, ccgt
		(R)	GATCCCGTCGCCAACAAC	275	protein	Thotosynthesis	(Onvena et at., 2007)	
12	ESTA19	(F)	CGCACCCGTTGACGAAGCAGT	191	Peroxidase 7	Hydrogen peroxide	(Oliveira et al., 2009)	ct, gc
		(R)	GTTCCTCGCGCTCCTCTGCT			removal	(0, 2, 2,	
13	ESTA70	(F)	GATGGAACCTGAAGATGAAGAGCA	175	Peroxidase	Hydrogen peroxide	(Oliveira et al., 2009)	tc total ga
		(R)	CCGGCCGGAGCACAACA			removal		
14	ESTB82	(F) (R)	CGTCGATCGAGATGAAGAAGG GAAGCAGTCGTGGAAGTGGAG	263	Putative peroxidase		(Oliveira <i>et al.</i> , 2009)	tatg, gc, cgt, agct, ctag, acg

Discussion

Normally, sugarcane is harvested once per year, with the drought season from March to April (tillering stage) and May to August (elongation stage) (Thai Encyclopedia for Youth, 1980). Lacking water at both stages significantly impacted sugarcane yield. On the other hand, drought stress of sugarcane from November to February stimulates increased soluble sucrose to crush in the sugar factory (Juttupoornpong et al., 2012). Similar to the greenhouse experiment, most varieties showed that lack of water activated the pathway of the drought stress response. The main characteristics of sugarcane under drought stress comprise a complex response consisting of both agronomic and physiological traits to adapt to drought conditions (Khonghintaisong, 2017). Drought stress reduced the stalk diameter and biomass (Jangpromma et al., 2012) with reduced numbers of stalks (Robertson et al., 1999). In the absence of water, sugarcane color intensity value (SPAD) relates to the amount of chlorophyll in the leaves (Jangpromma et al., 2010) which promotes the creation of food through photosynthesis. Sugarcane varieties that are not resistant to water deficiency have low SPAD values. Moreover, relative water content, respiration rate and value of stomatal conductance in leaves of droughtresistant varieties have a better response to drought than weak varieties (Graca et al., 2010).

Three parameters (number of stalks per plant, height, and weight) were related to drought response and these were measured during the 3rd, 5th and 7th months of sugarcane grown in the greenhouse (Table 1). These agronomic traits have an important impact on the total yield of sugarcane in harvesting season (Chaves et al., 2002). After the elongation stage, sugarcane growth enters the sucrose accumulation stage when growth rate decreases and total yield becomes invariable. MPT10-54 showed good potential to survive in drought conditions with a higher number of stalks under drought treatment than irrigation treatment in more than 50% of all treatment varieties. Moreover, MPT10-54 in drought conditions had the second highest weight parameter after UT12. MPT10-54 in drought conditions showed decreased growth rate (height) in the 7th month, similar to all treatment varieties but higher than the irrigation condition in the 3rd and 5th months as K93-219, KK3, KPK98-40, MPT03-320, and SP50. On the other hand, K93-219 showed some interesting patterns as the high value of height in drought conditions more than in irrigated conditions in the drought period of the 3rd to the 5th months, then growth rate reduced after irrigation during the 7th month, similar to KPS01-12. Drought varieties should remain yield or increase growth rate in drought condition comparing with the normal condition, so the number of stalks per plant, height, and weight can be criteria to select drought characteristic (Ribeiro et al., 2013). Thus, KPS01-12, K93219, MPT10-54, and UT12 showed drought agronomic traits with higher yield in drought conditions than in irrigation conditions.

Another three parameters as SPAD, stomatal conductance, and photosynthesis efficiency were measured during the 5th and 7th months in the greenhouse. Some varieties had the potential to survive in drought conditions such as K93-219, MPT10-54, and MPT03-320 by growth rate because their value of stomatal conductance in drought conditions was less than in irrigation conditions. The stomatal conductance parameter affects stoma opening in the sugarcane leaves (Pirasteh-Anosheh et al., 2016). A high value of stomatal conductance parameter indicates high water loss to the environment which is the worst scenario in drought conditions (Basnayake et al., 2012). However, when sugarcane grows continuously it requires water for photosynthesis (Dodd, 2003). The photosynthesis efficiency parameter determines whether drought stress sugarcane can survive (Wang et al., 2018). KPS01-12 and MPT03-320 had high photosynthesis rates and they produced biomass as height. It has been reported that increasing chlorophyll content promotes sugarcane growth in drought conditions (Tripathi et al., 2019). K93-219, KPK98-40, KPS01-12, MPT03-320, MPT10-54, PHIL66-07, and TBY20-2248 had higher chlorophyll contents in drought conditions than irrigation conditions, while K93-219, MPT10-54, KPS01-12, and MPT03-320 had drought physiological response related with phenotype to give higher yield in drought conditions than in irrigation conditions.

Four SSR markers as SSR9, SSR80, A19, and A70 were analyzed by the Jaccard coefficient and UPGMA clustering method. Separate sugarcane varieties were grouped into patterns related to physiological data. Kanagaraj et al. (2010) reported that using more SSR marker-related trait responses improved the accuracy of predicting candidate varieties. Phylogenetic tree constructed by SSR marker analysis can relate with trait form result of cluster analysis. Each variety may have same or like the relationship between them and presence of specific effect within the group which could be due to population structure effect to closely or so far (Tabkhkar et al., 2018). Thus, this can support the hypothesis that the marker-trait association is independent of population structure. Physiological traits showed that each sugarcane variety had different performance in each period. KK3 as the control treatment gave a total performance in drought conditions less than MPT10-54, K93-219, MPT03-320, and KPK98-40, while K93-219 and MPT10-54 had the highest agronomical traits. Conversely, MPT03-320, TBY20-2248, and KPK98-40 had the highest physiological traits focusing on specific traits. For genetic data, results showed that K93-219 and UT12 had a close specific relationship, followed by MPT 10-54 and KPS01-12. This group showed drought efficiency for adaptation to drought stress. The second group as KK3, MPT 03-320, and PHIL66-07 were insensitive to drought stress, while moderate drought stress was shown by the group of TBY20-2248, KPK98-40, and SP50. Four markers were related with drought function via the ABA signaling pathway and hydrogen peroxide removal. Drought is one of the most severe abiotic factors restricting plant growth and yield. Numerous gene functions in drought response are regulated by abscisic acid (ABA) dependent and independent pathways (Liu et al., 2018). Hydrogen peroxide (H₂O₂) functions as a signal molecule in plants under abiotic and biotic stress to improving drought tolerance (Guler and Pehlivan, 2016). Changes in H₂O₂ response to environmental conditions, in parallel with changes in abscisic acid (ABA) and oxidative stress markers, together with lignin accumulation, xylem, and sclerenchyma differentiation, and leaf area were also investigated (Jubany-Mar í et al., 2009). In Thailand, drought affects many regions and reduces sugarcane yield by at least 2 ton/rai. Effective sugarcane candidate varieties for drought adaptation may be able to resolve this problem and offer increased benefits to farmers. Our results suggested that four efficient SSR markers can be rearranged in Thai sugarcane varieties to relate with physiological data.

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Compliance with Ethical Standards

Conflict of interest: All authors of this research paper declare no conflicts of interest.

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