

EFFECT OF LOCATION, GENOTYPE AND THEIR INTERACTIONS FOR ANTHOCYANINS AND ANTIOXIDANT ACTIVITIES OF PURPLE WAXY CORN COBS

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ABSTRACT

Selection of purple waxy corn genotypes for high and stable anthocyanin content in corn cobs is important for breeding programs and anthocyanin production. The objectives of this study were to evaluate the effects of location, genotype and their interaction on anthocyanin content and antioxidant activities and to identify purple waxy corn genotypes with high and stable anthocyanin content. Five purple waxy corn and a white waxy corn genotypes were evaluated in a randomized complete block design with three replications at four locations with different elevations in Thailand. Location (L), genotype (G) and GxL interaction were significant for all characters. Waxy corn grown in Nakhon Ratchasima had the highest total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G), DPPH radical scavenging activity of phenolics (DPPH) and Trolox equivalent antioxidant potential (TEAC). Genotype KNDM4 had the highest TAC, Pg3G, Pn3G and DPPH. Its regression coefficient (b_i) was close to one but it had the highest Sd^2 , indicating specific adaptation to favorable environments. KNDM4 genotype performed better than other genotypes at unfavorable environments for all studied traits. This information is useful for breeding programs and anthocyanin production from purple waxy corn.

Keywords: Anthocyanin production, Different elevations, Favorable environments, Regression coefficient, Specific adaptation

INTRODUCTION

Waxy corn (*Zea mays* L. var. *ceratina*) is an important vegetable crop in East and Southeast Asia including China, Vietnam, Cambodia, Taiwan, Korea and Thailand. The world production of vegetable waxy corn is low compared to field corn and sweet corn. It is popular among people in these countries because of its glutinous starch in endosperm due to high amylopectin. Unlike field corn, waxy corn has received little interest for genetic improvement through breeding. The increase in demand for waxy corn and market potential arouses more attention from breeders to improve waxy corn. There are numerous special cultivars that contain colored pigments and give rise to numerous varieties of black and purple corn. Purple corn produced the anthocyanin pigment throughout the plant, especially high in the husk and cob regions,

although anthocyanin levels varied significantly among different plant parts (Li et al., 2008).

Anthocyanins have health benefits in reducing the incidence of cardiovascular disease, cancer, hyperlipidemias and other chronic diseases (de Pascual-Teresa and Sanchez-Ballesta, 2008; Wang and Stoner, 2008). Purple waxy corn genotypes had the highest total anthocyanin and antioxidant activity in seed at both immature and harvesting stages in comparison with white waxy corn, super sweet corn and field corn (Khampas et al., 2013). Recently, a study showed that purple waxy corn is a source of anthocyanins with a protective property against diabetic cataract (Thiraphatthanavong et al., 2014). Kernels and cobs of purple corn possess an excellent antioxidant activity, and the application of these natural food colorants by the food industry would be increased

considerably (Yang and Zhai, 2010). Corn cob contributed the most in phenolic compounds and antioxidant capacity, and can be considered as excellent novel sources of natural antioxidants for the functional food and dietary supplement markets (Cevallos-Casals and Cisneros-Zevallos, 2003). This makes it more attractive for nutraceutical and functional food market.

High and stable yield across a range of environments is the primary goal of plant breeding programs. Recently, breeding for high anthocyanins is a priority of waxy corn breeding program at Khon Kaen University as this character defines its possible use as colorants and functional food. Most previous studies on stability and genotype by environment interaction in agronomic traits of cereal crops have focused on yield and the studies on quality traits such as anthocyanin content and antioxidant activity are still lacking (Altay, 2012; Mitrović et al., 2012; Mohammadi et al., 2012). In wheat, cultivars Kate A1 and Mufitbey were found to be the most stable genotypes for all the environments whereas cultivar Gerek-79 was found to be the best performer for under poor soil and weather conditions (Altay, 2012), and significant genotype×environment interactions were observed among wheat genotypes (Altay, 2012; Mohammadi et al., 2012). In maize, environment accounted for the largest portion (77.83%) of total variation in grain yield, whereas genotype and genotype by environment accounted for 30% of total variation (Mitrović et al., 2012).

Several factors such as genetic, agronomic, pH, temperature, light intensity, type of light, and metallic ions, processing and storage affect anthocyanin stability (Bordignon-Luiz et al., 2007; de Rosso and Mercadante 2007; Mollov et al., 2007; Veigas et al., 2007; de Pascual-Teresa and Sanchez-Ballesta, 2008). Anthocyanin was mainly affected by pH and temperature (Cevallos-Casals and Cisneros-Zevallos, 2004). However, Zhao et al. (2008) found that anthocyanin in five Chinese purple corn hybrids was rather stable across a wide range of temperatures and times. Soil potassium concentration and population density did not significantly affect anthocyanin concentration in cobs, but growing location affected anthocyanin levels and the percentage of anthocyanins to total phenolics (Jing et al., 2007).

Table 2. The testing locations used in this study.

Locations	Geographical coordinates	Elevations (masl)	Planting date
Saraburi	N 14° 43' E 100° 47'	120	Dec 2012
Khon Kaen	N 16° 28' E 102° 49'	200	Nov 2012
Nakhon Ratchasima	N 14° 30' E 101° 30'	356	Dec 2012
Chiang Rai	N 20° 24' E 99° 53'	380	Feb 2013

The genotypes were grown in four-row plots with 5 m long and spacing of 0.80 m between rows and 0.25 m between plants within rows. Recommended cultural practices were used for all locations. The crop at all location was well-irrigated for optimum growth and yield. Sprinkler irrigation was available at the locations in Khon

However, effects of various factors such as genotypes, environments and their interaction on purple waxy corn anthocyanin have not been adequately studied. The objectives of this study were to evaluate the effects of location, genotype and their interaction on anthocyanin content and antioxidant activities and to identify purple waxy corn genotypes with high and stable anthocyanin content. The information obtained from this study is important for breeding of waxy corn for high anthocyanin content and antioxidant activity and selection of locations for production of waxy corn for use as raw material for functional food products.

MATERIALS AND METHODS

Plant Materials

Five purple waxy corn genotypes (KGW1, KGW2, KGW3, KNDM4 and Fancy 111) and one white waxy corn genotype (Fancy 121) were used as plant material in the study (Table 1). Most of them are F₁ hybrids except for KNDM4, which is an open-pollinated variety (Hussanun et al., 2014).

Table 1. The waxy corn varieties used in this study.

Genotypes	Seed color	Cob color	Types
1. KGW1	Purple	Purple	Hybrid
2. KGW2	Purple	Purple	Hybrid
3. KGW3	Purple	Purple	Hybrid
4. KNDM4	Purple	Purple	Opened-pollinated
5. FANCY 111 ^{1/}	Purple	Purple	Hybrid
6. FANCY 121 ^{1/}	White	White	Hybrid

^{1/} Commercial variety

Field Experiment

Six genotypes of waxy corn were evaluated in a randomized complete block design with three replications at four locations (Saraburi, Khon Kaen, Nakhon Ratchasima and Chiang Rai) with differences in elevations from 120 to 380 meters above sea level in the dry season 2012/13. These locations represented waxy corn growing areas in Thailand from the central plain to the North of Thailand. The details of locations were presented in Table 2.

Caen, Nakhon Ratchasima and Chiang Rai, whereas furrow irrigation was performed in Saraburi.

15-15-15 Fertilizer of N-P-K as basal dose at the rate of 171 kg ha⁻¹ was incorporated into the soil during soil preparation. Two splits of the fertilizer at the rate of 93.75 kg ha⁻¹ plus urea (46-0-0) at the rate of 93.75 kg ha⁻¹ for

first split and the fertilizer at the rate of 125 kg ha⁻¹ plus urea at the rate of 62.5 kg ha⁻¹ for second split were applied to the crop at 14 days after planting (DAP) and 30 DAP, respectively. At flowering stage, 13-13-21 fertilizer was applied at the rate of 156.25 kg ha⁻¹. Therefore, total

dose of fertilizers consisted of 150.65 kg ha⁻¹ nitrogen, 78.78 kg ha⁻¹ phosphorus and 91.27 kg ha⁻¹ potassium, respectively. Weather and soil data for all locations were presented in Table 3 and Table 4, respectively.

Table 3. Weather data at four locations.

Days after planting	Average Temperature (°C)	Solar radiation (MJ m ⁻² day ⁻¹)	Total rainfall (mm)	Relative humidity (%)
Saraburi				
0-30	28.5	15.3	49	68.7
31-60	27.1	17.8	0	62.7
61-90	29.7	18.9	3	66.8
mean	28.4	17.3		66.1
Khon Kaen				
0-30	26.9	17.1	23	60.0
31-60	23.7	17.6	0	52
61-90	26.7	17.1	6	52
mean	25.8	17.3		78.6
Nakhon Ratchasima				
0-30	25.1	16.9	5	68.4
31-60	25.0	18.2	46	62.4
61-90	26.7	19.7	42	71.1
mean	25.6	18.3		67.3
Chiang Rai				
0-30	25.2	16.9	53	74.3
31-60	27.1	19.8	8	71.4
61-90	28.8	20.8	84	62.3
mean	27.0	19.2		69.4

Table 4. Soil physical and chemical properties.

Locations	Soil types	pH (1:1 H ₂ O)	EC (1:5 H ₂ O) (dS/m)	Organic Matter (%)	Total nitrogen (%)	Available Phosphorus (mg/kg)	Available potassium (mg/kg)	Available calcium (mg/kg)
Saraburi	Sandy loam	7.4	0.09	1.22	0.06	234	155	2,432
Khon Kaen	Sandy	4.9	0.01	0.73	0.03	307	139	54
Nakhon Ratchasima	Loamy sand	7.1	0.13	2.49	0.13	52	162	3,137
Chiang Rai	Loamy sand	5.6	0.08	3.40	0.09	45	149	593

Corn cobs were collected at harvest maturity stage (R6) or 90 days after planting and stored at room temperature for analysis of anthocyanin and antioxidant activity. Data were recorded for total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G), DPPH radical scavenging activity of phenolics (DPPH) and the Trolox equivalent anti-oxidant potential (TEAC) as described by Yang et al. (2008).

Sample extraction

The portions of 0.5 g cob powder samples were weighted and pounded to fine powder using mortar and pestle. The extraction method used for TAC and antioxidant activity analysis was modified from the method reported previously (Yang et al., 2008). The samples were extracted in 25 ml of methanol containing 1% 1 M citric acid in falcon tubes. The mixtures were

vortexed and stored at 4 °C for 24 hours. The homogenates were centrifuged at 5000 rpm at 4 °C for 15 minutes. After centrifugation, the mixtures were acquired by filtration through Whatman No.1 filter papers, adjusted to 25 ml and stored in dark room at -20 °C until analysis for TAC and antioxidant activity.

Analysis of total anthocyanin content (TAC)

TAC was quantified by pH differential method (Giusti and Wrolstad, 2001). The absorbance of diluted extracts was measured to detect anthocyanins level at 510 nm and 700 nm with UV-Visible Spectrophotometer (model Specord 250 plus, Analytik jena, Germany) against a blank cell filled with distilled water. The absorbance was calculated as $Abs = (Abs_{510} - A_{700})_{pH 1.0} - (Abs_{510} - A_{700})_{pH 4.5}$. Then monomeric anthocyanin pigment in samples was calculated as;

Monomeric anthocyanin pigment = (Abs x MW x DF X 1000)/(ϵ x 1),

Where MW is the molecular weight of cyanidin 3-glucoside (MW of 449.2 g/mol), DF is dilution factor of the sample and ϵ is the molar extinction coefficient (Equal to 26900 M⁻¹ cm⁻¹).

Anthocyanin levels were presented in mg of cyanidin 3-glucoside equivalents (CGE) per 100 g of dry weight (DW).

HPLC analysis of anthocyanin components

Total anthocyanin content was determined by the method described by Kim et al. (2007) with minor modification, and individual anthocyanin components were quantified with high-performance liquid chromatography (HPLC), Shimadzu LC-20AC pushes (Shimadzu Co., Kyoto, Japan), SPD-M20A with a photodiode array detector (Waters, USA). A XselectCHS C-18 line (4.6 mm x 250 mm, i.d. 5 μ m). The extracts of cob samples were filtered through a 0.22 μ m filter at 30 °C. The anthocyanins were eluted at 1 mL/min using gradient system consisting of two solvents: (A) acidified methanol (methanol 0.1% HCl, 85:15, v/v) and (B) 10% formic acid. The gradient elutions of solvent B were performed as follows: for 0.0-0.5 min (80%), 0.5-9.5 min (15%), 9.5-10.0 min (5%), 10.0-15.0 min (80%), and 15.0-20.0 min (80%). The divided solutions were consequently detected and identified at 520 nm for chromatographic retention times. Three purified anthocyanins consist of cyanidin 3-glucoside, pelargonidin 3-glucoside, and peonidin 3-glucoside and one anthocyanidin (cyanidin chloride) (Sigma, USA) were used for quantification.

The purified anthocyanins extractions were utilized to aid the identification of individual anthocyanins. The standard solutions were prepared in acidified methanol (pH 1) by weighing exactly 200-300 μ g in 200 μ L; 20 μ L of standard solutions and then diluted to 500 μ L to prepare working standard. The standard anthocyanins revealed a linear relationship with a HPLC optimum peak area with a concentration range of 0.0-1.0 μ g. The coefficient of dedication (r^2) varied from 0.9933 to 0.9998 for a mixture of purified anthocyanins, which were divided by HPLC using 5, 10, and 15 μ L of operating standard solution.

Analysis of antioxidant activity

DPPH radical scavenging activity of methanolic extracts was determined according to Xu et al. (2010). Briefly, 0.5 mL of sample extract was added and mixed with 4.5 mL of 60 μ M DPPH demolished in methanol. The mixture was shaken thoroughly and allowed to stand for half an hour in the dark room. The absorbance was detected at 517 nm against the solvent blank. Trolox solution (100-1000 μ M) was used as a reference conventional. The outcomes were provided in micromoles Trolox equivalents (TE) per g of DW.

Analysis of Trolox equivalent antioxidant capacity (TEAC) measured by the reduction in radical cation of

ABTS by anti-oxidants was carried out as previously described by Jemai et al. (2009). Briefly, ABTS+ radical cation was conducted with a result of 7 mmol/L ABTS and 2.45 mmol/L potassium per sulfate. The combination was allowed to stay in dark room at ambient temperature 16-24 hours prior to use and should be used within 2 days. The ABTS solution was diluted with methanol for absorbance of 0.700 \pm 0.050 at 734 nm. All samples were diluted appropriately to provide 20-80% inhibition of the blank absorbance. 50 μ L of the diluted extract was taken by a pipette and combined with 1.9 mL of diluted ABTS solution. The analysis of the combination was performed in three replications. The combination was allowed to stand for 6 min at ambient temperature and the absorbance was instantly measured at 734 nm. Trolox solution (100-1000 μ M) was used as a reference point. The outcomes were indicated as micromoles Trolox equivalents (TE) per g of DW.

Statistical Analysis

Data for each location were analyzed separately and error variances were tested for variance homogeneity. As variances were homogeneous and the differences of error variances were not larger than 3 folds, then combined analysis of variance was performed across the locations. Linear regression was carried out to study stability of corn genotypes by plotting cultivar's means against environmental means for each genotype as described by Eberhart and Russell (1966). Means were compared statistically using least significant difference (LSD) at $P \leq 0.05$.

RESULTS AND DISCUSSION

Variations in anthocyanin and antioxidant activity

Location (L), genotype (G) and GxL interaction were significant ($P \leq 0.01$) for total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G), DPPH radical scavenging activity of phenolics (DPPH) and the Trolox equivalent anti-oxidant potential (TEAC) (Table 5). Location contributed to large portions of total variations in TAC (44%), C3G (56%), Pg3G (60%), Pn3G (62%), DPPH (52%) and TEAC (57%).

Genotype contributed to relatively large portion of total variation in TAC (42%), but genotype contributed to rather small portions of total variations in C3G (18%), Pg3G (20%), Pn3G (2%), DPPH (24%) and TEAC (11%). G x L interactions contributed to medium to small portions of total variations in TAC (13%), C3G (25%), Pg3G (20%), Pn3G (15%), DPPH (22%) and TEAC (32%).

Anthocyanins are phytochemicals that commonly occur in many cereal and fruit crops, and the trait is quantitatively inherited (Vizzotto et al., 2007). In corn, anthocyanins are present in all tissues and found at high concentrations in kernel skin and cobs (Moreno et al., 2005). In fruit crops such as grape, grapes may be either white or colored, ranging from the lightest pink to the darkest purple tones according to the amount of

anthocyanin accumulated in the berry skin, which is a crucial trait for both wine quality and human nutrition (Fournier-Level et al., 2009). In peach, anthocyanin content, phenolic content, and antioxidant activity were higher in red-flesh than in light-colored flesh peaches (Vizzotto et al., 2007). Genetic variation in anthocyanins has been reported in maize (Chander et al., 2008), grape

(Fournier-Level et al., 2009) and wheat (Knievel et al., 2009). The results indicated that selection for high TAC would be possible among purple waxy corn genotypes. Variation among locations also indicated the importance of selection for the best location for anthocyanin production in corn.

Table 5. Mean squares for total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G), DPPH radical scavenging activity of phenolics (DPPH) and the Trolox equivalent anti-oxidant potential (TEAC) of six waxy corn lines/cultivars.

SOV	df	TAC	C3G	Pg3G	Pn3G	DPPH	TEAC
		mg/100gDW	mg/100gDW	mg/100gDW	mg/100gDW	μmol/gDW	μmol/gDW
Location (L)	3	1,004,062**	60,464**	6,320**	1,564**	179.4**	86.1**
		(44)	(56)	(60)	(62)	(52)	(57)
Replication (R)/L (a)	8	432	50	6	2	1.4	0.1
Genotype (G)	4	15,975**	14,582**	1,542**	411**	61.4**	12.3**
		(42)	(18)	(20)	(2)	(24)	(11)
G x L	12	75,107**	6,596**	512**	96**	18.5**	12.0**
		(13)	(25)	(20)	(15)	(22)	(32)
Error (b)	32	2,726	67	5	3	0.4	0.1
C.V. (%) (a)		4.1	15.6	14.0	12.5	8.0	7.5
C.V. (%) (b)		10.3	18.1	12.0	15.2	4.1	5.4

** Highly significant difference at $p \leq 0.01$

Values in parenthesis are percentages of sum squares.

Effects of location and genotype on anthocyanins and antioxidant activity

Location was the main source of variation in TAC, C3G, Pg3G, Pn3G, DPPH and TEAC of five purple waxy corns and one white waxy corn. Nakhon Ratchasima had the highest TAC of 730 mg/100 gDW followed by 418 mg/100 gDW in Saraburi, whereas Khon Kaen and Chiang Rai had rather low TAC, accounting for 275 and 271 mg/100 gDW, respectively (Table 6). Similar patterns were also observed for C3G, Pg3G, Pn3G and DPPH. For these parameters, Nakhon Ratchasima was still highest followed by Saraburi, whereas Khon Kaen and Chiang Rai were similar and lower than Nakhon Ratchasima and Saraburi. Nakhon Ratchasima was also highest for TEAC followed by Khon Kaen, Saraburi and Chiang Rai, respectively.

In general, the highest TAC and anthocyanin components in corn cobs were found in Nakhon Ratchasima compared to the nearest locations in Khon Kaen and Saraburi. Nakhon Ratchasima is located at higher elevation and had lower temperature during growing season and lower phosphorus. Low temperature and low phosphorus might be the cause of the high TAC and its components.

Low temperature induced anthocyanin synthesis in maize seedlings (Christie et al., 1994). Anthocyanin concentration increased rapidly during low temperature in apple and peach shoots (Leng et al., 2000). Anthocyanins and phenolics contents were enhanced with higher light

intensity and lower temperature in potato tubers (Reyes et al., 2004). Visible and UVB radiation, cold temperatures and water stress has been shown to induce anthocyanin synthesis in plant tissue (Chalker-Scott, 1999).

However, low temperatures in the absence of either visible light or UVB prevent anthocyanin biosynthesis (Janda et al., 1996; Oren-Shamir and Levi-Nissim, 1997). Wang and Zheng (2001) also found that high temperature (30/22 day/night °C) condition enhanced phenolic acid, flavonols, and anthocyanins in strawberry.

Phosphorus deficiency induces anthocyanins accumulation in maize (Cobbina and Miller, 1987). Anthocyanin content was increased and growth was decreased on nitrogen and phosphorus deficiency in Cruciferous (Hodges and Nozzolillo 1996). However, Chiang Rai also had low phosphorus but it had the lowest anthocyanins possibly due to high temperature and high solar radiation during reproductive periods. It should be note that stresses of low temperature and low phosphorus were the factors enhancing anthocyanin accumulation.

Nakhon Ratchasima (low temperature and low phosphorus) also had the highest antioxidant activities (DPPH and TEAC) but Chiang Rai (high temperature and low phosphorus) had the lowest highest antioxidant activities. These results were in agreement with previous study of Lachman et al. (2008) who reported that low temperature during vegetative period showed high antioxidant activity in potatoes.

Table 6. Means for total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G), DPPH radical scavenging activity of phenolics (DPPH) and the Trolox equivalent anti-oxidant potential (TEAC) of six waxy corn lines/cultivars evaluated in four locations 2012/13.

	Saraburi	Khon Kaen	Nakhon Ratchasima	Chiang Rai
TAC (mg/100gDW)				
KGW1	247 ^l	671 ^m	609 ^e	129 ^{kl}
KGW2	449 ^g	341 ^h	982 ^c	373 ^h
KGW3	328 ^{hi}	263 ^{ij}	547 ^{ef}	330 ^{hi}
KNDM4	1,132 ^{ab}	826 ^d	1,166 ^a	490 ^{fg}
FANCY111	346 ^h	148 ^k	1,076 ^b	302 ^{hij}
FANCY121	4 ^m	6 ^m	3 ^m	2 ^m
Means	418 ^B	275 ^C	730 ^A	271 ^C
C3G (mg/100g DW)				
KGW1	5 ^{fg}	4 ^{fg}	37 ^d	1 ^g
KGW2	27 ^{de}	9 ^{fg}	246 ^a	2 ^g
KGW3	16 ^{ef}	7 ^{fg}	93 ^b	3 ^g
KNDM4	96 ^b	23 ^e	246 ^a	5 ^{fg}
FANCY111	9 ^f	2 ^g	72 ^c	1 ^g
FANCY121	1 ^g	1 ^g	1 ^g	1 ^g
Means	26 ^B	8 ^C	116 ^A	2 ^D
Pg3G (mg/100g DW)				
KGW1	3 ^{hi}	3 ^{hi}	9 ^l	2 ^{hi}
KGW2	11 ^f	5 ^{gh}	54 ^c	3 ^{hi}
KGW3	10 ^f	5 ^{ghi}	29 ^d	3 ^{hi}
KNDM4	30 ^d	15 ^e	86 ^a	7 ^{fg}
FANCY111	16 ^e	5 ^{gh}	63 ^b	3 ^{hi}
FANCY121	2 ^{hi}	2 ⁱ	2 ^{hi}	2 ^{hi}
Means	12 ^B	6 ^C	41 ^A	3 ^D
Pn3G (mg/100g DW)				
KGW1	3 ^{ghi}	3 ^{ghi}	5 ^{fg}	2 ⁱ
KGW2	15 ^c	7 ^{ef}	35 ^a	3 ^{ghi}
KGW3	9 ^{de}	5 ^{fg}	26 ^b	3 ^{ghi}
KNDM4	26 ^b	11 ^d	35 ^a	5 ^{fgh}
FANCY111	9 ^{de}	3 ^{ghi}	28 ^b	2 ^{hi}
FANCY121	2 ⁱ	1 ⁱ	2 ⁱ	2 ⁱ
Means	11 ^B	5 ^C	22 ^A	3 ^D
DPPH (µmol/gDW)				
KGW1	11 ⁱ	10 ⁱ	18 ^{cde}	8 ^{jk}
KGW2	17 ^{ef}	14 ^h	18 ^{cd}	10 ⁱ
KGW3	17 ^{de}	13 ^h	19 ^{cd}	9 ^j
KNDM4	22 ^a	18 ^{cd}	15 ^g	16 ^{fg}
FANCY111	19 ^c	13 ^h	21 ^b	11 ⁱ
FANCY121	5 ^l	5 ^l	4 ^l	7 ^k
Means	15 ^A	12 ^B	16 ^A	10 ^C
TEAC (µmol/gDW)				
KGW1	0.8 ^{mn}	2.8 ^h	5.3 ^e	1.3 ^{kl}
KGW2	6.4 ^d	3.8 ^g	8.1 ^b	1.3 ^{lk}
KGW3	7.3 ^c	3.7 ^g	6.9 ^c	2.2 ^{ij}
KNDM4	1.1 ^{lm}	5.7 ^e	8.2 ^b	4.2 ^f
FANCY111	1.4 ^k	2.5 ^{hi}	10.2 ^a	2.1 ^j
FANCY121	0.3 ^{op}	0.0 ^p	0.0 ^p	0.5 ^{no}
Means	2.9 ^B	3.1 ^B	6.4 ^A	1.9 ^C

Means of varieties in the same column and means of locations in the same row with the same letter(s) are not significantly different at $P \leq 0.05$ by LSD

As the interactions between location and genotype were significant for all parameters, identification of the best genotypes for each parameter was difficult. The varieties that showed good performance in one location did not perform well in other locations. However, KNDM4 showed consistently the highest values for TAC,

C3G, Pg3G and Pn3G across locations. High values for these parameters in KNDM4 would be due to the fact that it is an open-pollinated variety and other hybrids had one parent with white kernels. For DPPH, KNDM4 was highest in Saraburi, Khon Kaen and Chiang Rai, whereas FANCY111 was highest in Nakhon Ratchasima. For

TEAC, KNDM4 was the winner in two locations in Khon Kaen and Chiang Rai. FANCY111 was still highest in Nakhon Ratchasima, whereas KGW3 was highest in Saraburi. FANCY111, which is the waxy corn, showed consistently the lowest values for all parameters across locations.

Stability analysis for anthocyanins and antioxidant activity

The genotypes with stable yield should have high mean yield across locations, regression coefficient close to one ($b_i=1$) and deviation from regression (Sd^2) close to zero (Eberhart and Russell, 1966). KNDM4 had the

highest TAC (903 mg/100 gDW) and regression coefficient close to one (0.9), but it showed high deviation from regression (254) (Table 7). KNDM4 was the highest variety for TAC across locations. It did not performed well under unfavorable environment in Chiang Rai but it was still better than others (Table 6). However, anthocyanins of KNDM4 was rather fluctuated across environments as it has high Sd^2 . KNDM4 was suitable for favorable environments. KGW2 had the second rank for TAC (536 mg/100g DW), regression coefficient close to one (1.1) and low deviation from regression (67). This genotype was most stable for TAC across environments.

Table 7. Stability parameters for total anthocyanin content (TAC), cyanidin 3-glucoside (C3G), pelargonidin 3-glucoside (Pg3G), and peonidin 3-glucoside (Pn3G) of six waxy corn lines/cultivars.

Genotypes	TAC (mg/100 gDW)			C3G (mg /100 gDW)			Pg3G (mg /100 gDW)			Pn3G (mg /100 gDW)		
	Mean	b ^{1/}	Sd ²	Mean	b	Sd ²	Mean	b	Sd ²	Mean	b	Sd ²
Purple waxy corn												
KGW1	263 ^e	0.9	34	12 ^e	0.3**	2	4 ^e	0.2**	1	3.5d	0.1**	0.4
KGW2	536 ^b	1.1	67	71 ^b	1.8	14	18 ^c	1.2	2	15.1b	1.4**	0.3
KGW3	367 ^d	0.5	45	30 ^c	0.7	4	11 ^d	0.6**	1	10.9c	1.0	1.7
KNDM4	903 ^a	0.9	254	93 ^a	1.7	24	35 ^a	1.7**	3	19.2a	1.3	5.4
FANCY111	468 ^c	1.6	120	21 ^d	0.5	4	22 ^b	1.4**	1	10.4c	1.1	1.9
White waxy corn												
FANCY121	4			1			2			1.9		

** Significant different from 1 at $P \leq 0.01$

Means in the same column with the same letter(s) were not significantly different at $P \leq 0.05$ probability level by LSD.

^{1/}Regression coefficients were analyzed without white waxy corn.

KNDM4 had the highest C3G (93 mg /100 gDW) followed by KGW2 (71 mg /100 gDW) and their regression coefficients were not different from one (1.7 and 1.8, respectively). However, their deviations from regression were high, indicating that they performed well under favorable conditions but performed poorly under unfavorable conditions (Chiang Rai). KGW1 had the lowest C3G (12 mg/100 gDW) and regression coefficient (0.3**) was significantly lower than one, indicating that this genotype had low C3G and stability for C3G across environments.

Regression coefficients for Pg3G for most genotypes were significantly different from one except KGW2, indicating that these genotypes were not stable for this trait. KNDM4 had the highest Pg3G of 35 mg/100 g DW followed by FANCY111 (22 mg/100g DW) but regression coefficients (1.7** and 1.4**, respectively) were significantly higher than one, indicating that these genotypes performed well under favorable conditions but they had low Pg3G under unfavorable conditions.

KNDM4 had the highest Pn3G (19.2 mg/100 gDW), regression coefficient close to one (1.3) and high deviation from regression. However, it performed best for all locations but it had low Pn3G under unfavorable conditions (Chiang Rai) (Table 5). KGW2 showed high Pn3G (15.1 mg/100 gDW) but it had regression coefficient higher than one (1.4**) and low deviation from regression, indicating that it performed well for Pn3G under favorable conditions but performed poorly under unfavorable conditions.

KNDM4 had the highest DPPH (18.0 μ mol/gDW), regression coefficient smaller than one (0.3**) and high deviation from regression (13.6) (Table 8). This genotype showed good adaptation to favorable conditions but it had poor adaptation to unfavorable conditions. KGW3 and KGW2 had high TEAC and regression coefficient close to one with high deviation from regression. Therefore, they may be suitable for some specific conditions.

Table 8. Stability parameters for DPPH radical scavenging activity of phenolics (DPPH) and the Trolox equivalent anti-oxidant potential (TEAC) of six waxy corn lines/cultivars.

Genotypes	DPPH ($\mu\text{mol} / \text{gDW}$)			TEAC ($\mu\text{mol} / \text{gDW}$)		
	Mean	b	Sd ²	Mean	b	Sd ²
Purple waxy corn						
KGW1	11.8 ^d	1.0	3.0	2.6 ^d	0.8	0.9
KGW2	14.6 ^c	1.1	0.3	4.9 ^{ab}	1.0	2.0
KGW3	14.3 ^c	1.3**	0.3	5.0 ^a	0.6	2.4
KNDM4	18.0 ^a	0.3**	13.6	4.8 ^b	0.9	2.5
FANCY111	15.9 ^b	1.3	1.3	4.1 ^c	1.6**	1.5
White waxy corn						
FANCY121	5.2			0.2		

** Significant different from 1 at $P \leq 0.01$

Means in the same column with the same letter(s) were not significantly different at $P \leq 0.05$ probability level by LSD.

Khampas et al. (2013) and Harakotr et al. (2014) reported that kernels of purple waxy corn had high TAC and antioxidant activity (DPPH, FRAP and TEAC). In this study, cobs of purple waxy corn also had high TAC and antioxidant activity, and cobs can be used for production of natural colorant with health benefits.

CONCLUSION

Location (L), genotype (G) and GxL interaction were highly significant for all characters. Location contributed to large portion of total variation in anthocyanin content and antioxidant activity in corn cobs, whereas genotype and genotype by location interaction contributed smaller portions of total variations in these traits. Nakhon Ratchasima, which was characterized by low temperature and low phosphorus, was the highest location for TAC and C3G. KNDM4 gave the highest TAC and C3G, and its regression coefficients (b) were close to one. However, KNDM4 also had the highest Sd², indicating that it had specific adaptation to favorable environments. KNDM4 performed well under unfavorable environments for all traits. This information is useful for breeding programs in multi-location trails and production of anthocyanins from purple waxy corn.

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