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RESEARCH ARTICLE

Electroactivity of polyphenols in sweet sorghum (*Sorghum bicolor* (L.) Moench) cultivars

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Abstract

Polyphenols and other potential health-promoting components of sorghum (*Sorghum bicolor* (L.) Moench) drove its recent growth in the U.S. consumer food industry. Linear sweep (cyclic voltammetry, CV) and differential (cyclic differential pulse) voltammetry methods were developed to detect target polyphenols and amino acids in sweet sorghum juice without interference from the dominant secondary (trans-aconitic acid) and primary (sucrose) metabolites. Of 24 cultivars investigated, No.5 Gambela showed the highest electron-donating capacity, as indicated by the highest peak area, height, and peak anodic potential. Pearson's correlation analysis indicated the contribution of polyphenols (rather than amino acids) on CV voltammograms of juice samples. The E_h -pH values of 173 sweet sorghum juice samples collected in 2017 aligned with quercetin model polyphenol. Accumulation of quercetin-like polyphenols in No.5 Gambela could offer antioxidant-rich juice for conversion to edible syrup as well as an increased tolerance against a recently emerged pest, sugarcane aphid [(*Melanaphis sacchari* (Zehntner)].

Introduction

Over the past several years, sorghum (*Sorghum bicolor* (L.) Moench) production for the consumer food industry has drastically grown within the U.S [1]. This market trend originates from high phenolics in sorghum grain, along with other attributes including gluten-free and high-fiber characteristics. A wide range of higher plants produce polyphenols as the secondary metabolites for the defense against photooxidative damage and pests [2]. Sorghum genotypes could be bred to accumulate desirable phenolics for a wide range of food products including sweeteners (for sweet sorghum varieties), cereals, and antioxidant food additives. However, the current bottleneck in precision breeding is the access to field-deployable omics fingerprinting tools to phenotype crop varieties without the analyte separation required for metabolomics. Such chemistry tool must be simple, rapid, and user-friendly to the breeders.

Redox chemistry underlies the antioxidant properties of polyphenols [2]. Thermodynamic driving force (standard reduction potential, E^0 in volts) and kinetic lability determine the

redox reactivity of polyphenols [3, 4]. Aromatic rings of polyphenols (1) decrease E^0 by both inductive and resonance effects, (2) stabilize the radical intermediates by resonance stabilization, enabling reversible electron transfer, and (3) protect the phenolic redox center from nucleophilic addition and other side reaction-induced decomposition [4]. Polyphenols (those containing a quinone redox center) and flavins are well-described electron shuttles in biological systems, because of their ability to transfer one or two electrons reversibly between the bulk electron donor and the terminal electron acceptor [5]. Sorghum and other polyphenol-rich cereals, fruits, and vegetables contain a complex mixture of phenolic and non-phenolic electroactive structures [2]. In addition, the polyphenol composition is highly cultivar-dependent [6].

Our previous report used cyclic voltammetry (CV) to oxidize the stem juice of sweet sorghum; polyphenolic structure similar to quercetin was determined to be the primary redoxactive moiety [7]. A sweet sorghum genotype called No.5 Gambela showed resistance to a recently emerged pest in the U.S.: sugarcane aphid [(*Melanaphis sacchari* (Zehntner)]. No.5 Gambela accumulated polyphenols as well as a carboxylate secondary metabolite, trans-aconitic acid [8]. Trans-aconitic acid was highly sensitive to spectrophotochemical detection by fluorescence excitation-emission (EEM) and UV/visible absorption, within the wavelength ranges overlapping with polyphenols [8]. Trans-aconitic acid and polyphenols [9] in juice, riboflavin-like fluorophore in bagasse [8], and dhurrin (tyrosine-derived cyanogenic glycoside) in leaves [10] are all candidate allelochemicals against aphids. Electrochemistry-based fingerprinting methods are necessary to understand the importance of redox-active defense phytochemicals.

Square wave (SWV) [11–13] and differential pulse voltammetry (DPV) [14], and CV [15] have been used to detect aliphatic carboxylates including trans-aconitic, fumaric, maleic, malic, oxalic, and citric acids. The SWV of fumaric and maleic acids [11, 12] were irreversible in neutral and acid media, and the voltammograms resembled that of trans-aconitic acid [13], which was most electroactive in the fully protonated form. Differential electrochemical methods (DPV and SWV) are used to improve the detection limit of linear sweep voltammetry (CV) [16] for the target electroactive structures in food and beverage [17]. Differential electrochemical methods improve the signal to noise ratio by minimizing the capacitive current [16]. This is achieved by (1) sampling current after capacitive current has decayed, or (2) taking the differential currents [16]. Faradaic process transfers charge across the interface to give a steady-state current with distinct peaks [18]. In contrast, capacitive current originates from the charging and discharging of the electrode double layer to give a box-shaped voltammogram [19]. Capacitive current is also called the background current (without redox-active species), double layer current, or non-Faradaic current to describe the rearrangement of ions in the double layer when potential is applied to the working electrode [19]. Magnitude of background current is directly proportional to the scan rate, while the current from diffusion-controlled electrochemical reaction is proportional to the square root of the scan rate [20]. As a result, background (capacitive) current is the major source of noise in CV especially at fast scan rates [16]. In addition, double layer capacitance is controlled by the electrode potential, which depends on the electrode material and the type and concertation of the supporting electrolyte [16].

The objective of this study was to understand the redox reactivity of secondary metabolites putatively contributing to the pest resistance of sweet sorghum cultivar, No.5 Gambela [8]. No.5 Gambela accumulated trans-aconitic acid and polyphenols, but not amino acids [8]. Because those classes of chemicals have overlapping peaks in fluorescence and UV/visible detection [8], electrochemistry was investigated in the present study as a tool to differentiate polyphenols from amino acids and trans-aconitic acid. Such redox method would allow breeding of sorghum cultivars utilizing different mechanisms: antioxidants vs. organic acids. First,

DPV was employed to increase the sensitivity of detection for the peak previously attributed to quercetin-like structures in sweet sorghum juice. Peak identification relied on the comparison of juice against electroactive structures in juice or bagasse of sweet sorghum [7, 8]: tryptophan, tyrosine, trans-aconitic acid, quercetin, and riboflavin.

Materials and methods

Sweet sorghum cultivars were planted in April, May, or June 2017, and harvested at the hard dough stage of grain fill, which occurred during July through November, depending on cultivar and planting date. Field experimental design and juice characterization procedures were described in detail previously [7, 21, 22], and are summarized in Section I of <u>S1 File</u>. Group membership and Plant Introduction (PI) number [23] of 24 sweet sorghum genotypes examined in this study are provided in S1 Table in <u>S1 File</u>.

Juice and bagasse characterization

As described in detail previously [7, 21], distilled, deionized water (DDW) with a resistivity of 18 M Ω cm (APS Water Services, Van Nuys, CA) was used in all laboratory procedures. All chemical reagents were obtained from Sigma-Aldrich (Milwaukee, WI) with the highest purity available. Because sweet sorghum juice is unstable at room temperature [24], samples were defrosted in a refrigerator, and were immediately diluted (large-orifice pipette tips, Fisher, Hampton, NH) by DDW, filtered (0.45 µm PVDF, Fisher), and analyzed. Brix (estimate of g sucrose/100g solution) was measured using a portable digital refractometer (300053; Sper Scientific, Scottsdale, AZ). Sucrose, glucose, fructose, and citric, oxalic, trans- and cis-aconitic acid concentrations were quantified using an HPLC system with refractive index and diode array detectors (Agilent Technologies, Santa Clara, CA) and a Hi-Plex H column (Fisher) with 5 mM sulfuric acid mobile phase at 0.6 mL min⁻¹ flow rate and 20 µL injection volume. Total organic carbon (TOC in gC/L) and total nitrogen (TN in gN/L) were analyzed using a Torch combustion TOC/TN analyzer (Teledyne Tekmar, Mason, OH). UV/visible spectra (HP8452A, Hewlett-Packard, Palo Alto, CA) of juice (20-fold diluted and 0.45 µm filtered) were obtained with DDW blank subtraction.

Electric conductivity (EC), pH, and oxidation reduction potential (ORP; E_h in mV) of juice samples were determined without dilution using YSI 3200 conductivity meter (YSI, Yellow Springs, OH) for EC; ORION ROSS sure-flow glass combination pH electrode for pH; and an epoxy sure-flow combination redox/ORP electrode with built-in temperature probe (9678BNWP, Fisher) for E_h . Sartorius Professional PP-15 meter (Sartorius, Bohemia, NY) was used to record the electrode response for both pH and E_h . The ORP electrode is composed of platinum sensing electrode and Ag/AgCl reference electrode. At 25 °C and 4 M KCl filling solution employed, E_h value relative to the standard hydrogen electrode (SHE) is calculated by adding 200 mV (±60 mV, error originating from variations in sample ionic strength and reference junction potentials) to the absolute value measured using ORP electrode.

Fluorescence EEM spectra of juice and bagasse samples were collected using F-7000 spectrofluorometer (Hitachi, San Jose, CA) at 220–500 nm excitation and 280–730 nm emission wavelengths in 3 nm intervals; 2.5 nm excitation and emission slits; auto response time; and 2400 nm min⁻¹ scan rate [8]. After blank subtraction and correction for Rayleigh and Raman peaks, parallel factor (PARAFAC) analysis was performed with non-negativity constraint using MATLAB version 8.6.0.267246 (R2015b; Mathworks, Natick, MA) with PLS toolbox version 8.6.2 (Eigenvector Research, Manson, WA). PARAFAC models contributions of primary fingerprints on each sample by minimizing the sum of squares of the residuals. For 2017 juice samples, three PARAFAC components (based on residual/leverage analysis) were

assigned to tryptophan-like (hereby denoted EEM/PARAFAC factor 1), tyrosine-like (factor 2), and phenolic structure with high aromaticity (aromatic factor 3) [8]. EEM/PARAFAC results were interpreted as follows: a juice sample (representing a cultivar and a planting month) with high absolute contribution from factor 1 is estimated to have high "concentration" of tryptophan-like structure. Standard curve calibration of colorimetric method is not used in EEM/PARAFAC, because of a complex mixture of contributing structures, e.g., different phenolic molecules. The purpose of fluorescence technique is to estimate the relative fraction of target analyte in a large set of samples, without analytical separation required for LC-MS used in metabolomics.

Electrochemistry

The CV and CDPV of diluted (2-fold by DDW) juice samples (buffered at pH 5 with 40 mM phosphate in 0.1 M KCl) were obtained using a WaveNow potentiostat with disposable screen-printed carbon electrodes (SPE with Ag/AgCl reference electrode; Pine Research Instrumentation, Durham, NC) of two different surface areas: 2 mm diameter disk, and 4×5 mm rectangle patterned carbon working electrode. The larger surface area of the 4x5 mm carbon SPE is expected to enhance the current signal relative to the 2 mm SPE used in our previous report [7]; however, the background current could increase as well. Both SPEs are compatible with aqueous solution, and are composed of polyethylene terephthalate plastic screen with Ag/AgCl reference electrode and carbon working/counter electrodes on conductive silver layer. For each juice sample, a new SPE was used to first collect background CV (0.1 M KCl and 40 mM phosphate buffer at pH 5) by applying -0.5 V for 60 s, and subsequently increasing the potential to 1.2 V, and then decreasing to -0.5 V at 100 mV s⁻¹ sweep rate in both directions. Immediately after collecting the background voltammogram, CV of the juice sample was obtained using the same SPE.

To identify electroactive structures in juice samples, 1 mM of following standards were prepared in DDW to perform CV as described above: tryptophan, tyrosine, trans-aconitic acid, sucrose, and citric acid. CV of quercetin (as a model polyphenol) was obtained from the literature employing 0.5 mM quercetin dissolved in 50 vol% ethanol [7]. Quercetin is sparingly soluble in water and reported aqueous solubility varies by orders of magnitude [25]. Carbon SPE employed in this study is designed for aqueous solutions, and compatibility with organic solvents decreases in the following order: ethanol>acetonitrile>methylene chloride. The layer containing working/reference/counter electrodes begins to swell after a few minutes in methylene chloride, but not in acetonitrile. Blank subtraction was utilized to account for co-solvent and other background effects, as described in detail below.

CDPV employed a procedure analogous to CV with the following pulse parameters: 50 mV height, 10 ms width, 100 ms period, 10 mV increment, and 3 ms pre- and post-pulse widths. This parameter setting corresponds to the effective scan rate (increment*(1/period)) of 100 mV/s. Higher mV height (90 instead of 50) or wider mV (upto 1500 mV) did not improve the peak sensitivity or resolution. For CV as well as CDPV, current (in mA) range was first determined using the auto-range option of the electrochemical software (AfterMath version 1.5.9568, Pine Research Instrumentation) to auto-adjust current and voltage. This procedure prevents low sensitivity leading to the lack of peak, or excessive sensitivity resulting in the clipped/truncated peak. Once the amplitude range was determined, the range was manually set to 10 μ A for both CDPV and CV to prevent current drifts caused by switching cell/circuit during auto-ranging. In addition, 60 s induction period (at -500 mV) was used to allow the cell to equilibrate with the initial signal level applied to the working electrode. Above 1 V, solvent (water) oxidation and reactions involving O₂ will influence the anodic current [26].

Each raw anodic CV and anodic/cathodic CDPV (in current (A) vs. potential (V)) was processed using OriginPro 2019 (OriginLab, Northampton, MA) by (1) background subtraction to minimize background signals from conductive carbon ink and other electrochemical cell components, (2) Savitzky-Golay smoothing of the first derivative to enhance peak resolution, and (3) trapezoidal integration at 0.5–1.0 V range. The derivative CV and CDPV of juice sample was additionally processed by the Gaussian integration (OriginPro 2019) to obtain the peak anodic/cathodic potential (E_{pa} and E_{pc} as the center of peak in volts) and corresponding peak area and maximum current height in mA. Gaussian fitting of peaks employed user-selected baseline anchor points connected by interpolation; peak selection by the second derivative; Gaussian fit until convergence; and error analyses by the reduced Chi-square and residual plots.

Statistical analysis

Main effects and interactions were examined by factorial (cultivar×planting month) ANOVA using Statistica version 12 (Statsoft, Tulsa, OK) at a significance level of p<0.05. Type VI sums of squares was used to test the effective hypothesis for unbalanced observations. If significant differences existed, post hoc comparison by Tukey's honestly significant difference (HSD) test was performed with cultivar or planting month as the categorical factor.

To examine linear relationships between all investigated parameters for the 2017 juice and bagasse samples, a Pearson's correlation matrix pseudocolor map was constructed using MATLAB version 8.6.0.267246 (R2015b; Mathworks, Natick, MA) with PLS toolbox version 8.7 (Eigenvector Research, Manson, WA). Variables were reordered by the similarity in Pearson's r values using a modified k-nearest neighbor algorithm.

Eq 1 [27] was used to explore the linear relationships between the anodic current (1st derivative) of juice samples against the following parameters that previously [8, 28] showed correlation with electrochemical parameters, and could directly or indirectly contribute to redox reactivity of juice samples (x_i): absolute contributions of aromatic, tryptophan, and tyrosine EEM/PARAFAC fingerprints; sucrose, trans-aconitic acid, and citrate concentrations; UV/visible absorbance at 270 and 340 nm; and Brix, pH, EC, E_h, TN (in gN/L).

$$r = \frac{n(\sum x_i y_i) - (\sum x_i)(\sum y_i)}{\sqrt{[n(\sum x_i^2) - (\sum x_i)^2][n(\sum y_i^2) - (\sum y_i)^2]}}$$
(1)

where x_i is the juice property, y_i is the current in mA (first derivative) at a given potential (in V), and n is 24 for juice samples representing each cultivar listed in Section II of S1 File.

Results and discussion

Electroactive components: Amino acids vs. polyphenols

Fig 1g shows the anodic peak response recorded by CV for sweet sorghum juice sample. For comparison, analogous voltammograms for primary chemical structures in sweet sorghum juice are provided in Fig 1a–1f, where y-axis scale is fixed (except sparingly water-soluble quercetin in Fig 1a) to allow a visual comparison of peak heights. Trans-aconitic acid, sucrose, and citric acid did not show anodic peaks (Fig 1d–1f) within the potential range where sweet sorghum juice had a peak (Fig 1g). In contrast, both tryptophan and tyrosine showed a single peak within the voltage range overlapping with juice sample. In addition, a peak attributable to–OH group on the A-ring of quercetin occurred within the potential range of juice sample (Fig 1a and 1g). Quercetin contains five oxidizable–OH substituents. Experimentally observable peaks will depend on pH and other electrochemical parameters as well as data processing



Fig 1. Raw CV (circles; after background subtraction and first derivative with smoothing) and Gaussian fits (colored lines) of 1 mM standards in DDW (a-f, quercetin (literature value) is 0.5 M in ethanol) [7] and representative juice sample (g) using 2 mm carbon SPE. Horizontal line represents E_{pa} of juice (0.81V in Table 2).

methods. Hendrickson et al. (1994) attributed three observable anodic peaks of quercetin to hydroxyl groups on the catechol B-ring, C-ring, and A-ring, towards more positive potentials [29]. Brett and Ghica (2003) observed a fourth peak originating from the second–OH on the catechol B-ring [30]. In the literature, total number of CV peaks for a given compound

(quercetin in this case) is determined by visual inspection of a voltammetric curve lacking clear baseline. In Fig 1a, overlapping peaks at the least positive potential are attributable to two hydroxyl substituents on catechol B-ring of quercetin. Additional peaks towards more positive potentials are attributable to hydroxyls on C-ring and A-ring of quercetin. There are several complexities in assigning a polyphenol structure to anodic peak of juice in Fig 1g. First, sweet sorghum juice is expected to contain a complex mixture of polyphenols, rather than a single chemical structure such as quercetin. Second, hydroxyl substituents on a given phenolic structure give rise to multiple peaks at different potentials, as observed for quercetin in Fig 1a. Because of those complexities, bulk electron donating capacity of each juice sample will be quantified as the peak area at a given potential to compare different sorghum cultivars.

Fig 1 indicates that E_{pa} (potential at the center of anodic peak) could be used to differentiate polyphenols from amino acids in juice samples. S3 Table in S1 File provides E_{pa} and Gaussianintegrated peak area for tryptophan, tyrosine (1 mM in DDW), quercetin, and catechin (in 50 vol% ethanol) authentic standards [7]. The CV-derived E_{pa} values were lower for the amino acids (0.77–0.83 V) than juice (0.88±0.03 V as mean±s.d. for n = 170), and tryptophan had higher peak area than tyrosine. Analogous trends were observed for CDPV-derived E_{pa} values that are consistently lower than the CV. Model polyphenols catechin and quercetin had E_{pa} above and below that of juice.

To understand the relative contribution of polyphenols and amino acids in juice, S1 Fig in S1 File shows the anodic CV (a) and CDPV (c) of a cultivar containing high tryptophan- and tyrosine-like structures (N109AxChinese/April), based on the absolute contributions determined by fluorescence EEM/PARAFAC [8] described in Materials and Methods. This proteinrich cultivar was compared against another cultivar having high polyphenol-like structures (No.5 Gambela/June) in S1b and S1d Fig in S1 File. For CV (left panels in S1 Fig in S1 File), higher E_{pa} and peak area were observed for No.5 Gambela/June (0.93 V E_{pa} and 4.2e⁻⁶ peak area) than N109AxChinese/April (0.85 V E_{pa} and 1.4e⁻⁶ peak area). Analogous trends were observed for CDPV (right panels in S1 Fig in S1 File) comparing the same two cultivars. In conclusion, S1 Fig in S1 File indicates higher E_{pa} and peak area of a juice sample accumulating polyphenols (No.5 Gambela/June) than amino acids (N109AxChinese/April).

In addition to comparing individual juice samples in S1 Fig in S1 File, replicate samples of a given cultivar (across planting months) were used to observe the contribution of polyphenols, as opposed to amino acids. Cultivars having highest polyphenol-like structures (No.5 Gambela) had the highest E_{pa} by CV (anodic 0.5–1 V) of 0.93±0.01 V (mean±standard error, n = 7), as opposed to 0.87 ± 0.01 V (n = 8) for N109AxChinese containing the highest amino acids [8]. For CDPV anodic current having wider E_{pa} range originating from Faradaic current, No.5 Gambela (0.84 ± 0.01 V, n = 7) and Isidomba (0.82 ± 0.01 V, n = 9) were significantly higher than N109AxN98 (0.76 ± 0.01 V, n = 9). In conclusion, a cultivar enriched with polyphenols (No.5 Gambela) [8] had CV peak with higher E_{pa} attributable to polyphenol oxidation, while an amino acid-enriched cultivar (N109A×Chinese) had significantly lower E_{pa} that could originate from tryptophan and tyrosine. Subsequent section will further examine the genotype effects on relative importance of amino acid and polyphenol electroactive structures.

Fig 2 shows Pearson's correlation (Eq 1) between anodic current of juice and the following parameters: EEM/PARAFAC absolute contributions for aromatic, tryptophan, and tyrosine fingerprints; concentrations for sucrose, trans-aconitic acid, and citrate; UV/visible absorbance at 270 and 340 nm; and Brix, pH, EC, E_h, and TN (in gN/L). Fig 2 was constructed using anodic voltammograms of CV for juice samples representing each of 24 cultivars (Section II of S1 File). Fig 2 only considers the potential range (0.7–1 V) of the primary CV peak in juice (Fig 1g). Fig 2 indicates strongly linear (red areas indicating positive correlation) relationships between anodic peak of juice samples at 0.9–1 V and following parameters: aromatic



Fig 2. Pearson's correlation coefficient between CV peak current and juice properties. Only juice peak voltage range was used.

fluorescence fingerprint [28], sucrose and trans-aconitic acid concentrations, Brix, and UV/ visible absorbance at 270 and 340 nm. Of those parameters, aromatic fluorescence fingerprint and UV/visible absorbance are directly related to polyphenols. Other parameters (sucrose, trans-aconitic acid, and Brix) are indirectly related to polyphenols, because concentrations of organic carbon products in juice are linearly correlated [28], as described in the subsequent paragraph dedicated to Fig 3. All other potential ranges and parameters evaluated in Fig 2 had negative r values or no significant linear correlations.

Fig 3 presents a Pearson's correlation map of r values between all parameters investigated on juice and bagasse samples from the planting year 2017 in this and previous report [8]. Similarly to the planting year 2015 [28], parameters originating from the organic carbon products were positively correlated (red square labeled +r in Fig 3): concentrations of fructose, glucose, sucrose, and trans-aconitic acid; Brix, total organic carbon; UV/visible absorbance at 270 and 340 nm; contribution to aromatic fluorescence fingerprint; and E_{pa} by CV and CDPV. Those organic carbon parameters were negatively correlated (blue region marked-r in Fig 3) with tryptophan-like fluorescence fingerprint, electric conductivity, and area and height of small anodic (0.3–0.6 V) peak by CV and CDPV. In conclusion, E_{pa} is positively correlated with



Fig 3. Pearson's r values between all chemical analyses conducted on sweet sorghum juice and bagasse samples collected in 2017. Red regions indicate positive correlation (+r), and darker red indicates higher linearity; blue regions indicate negative correlation (-r). Parameters in +r region: fructose, glucose, sucrose, trans-aconitic acid, Brix, TOC, UV/visible absorbance, aromatic fluorophore, and E_{pa} . Parameters in-r region: pH, TN, electric conductivity, and protein-like fluorophores.

polyphenol, and negatively correlated with amino acid. This finding is in agreement with the contribution of polyphenols, rather than amino acid, on anodic peak of juice samples in Fig 2.

Cultivar and planting month effects on the redox reactivity of polyphenols

Table 1 presents significant (p<0.05) cultivar (24 total in S1 Table in S1 File) and planting month effects and cultivar x planting month interactions on the electrochemical properties (CV and CDPV parameters and bulk E_h). Both CV and CDPV employed 2 mm carbon SPE at 100 mV/s scan rate (an estimate for CDPV, as described in Materials and Methods). Planting month effects by post-hoc Tukey HSD (p<0.05) are presented as the increase (\uparrow) or decrease (\downarrow) from April to May (first arrow), and from May to June (second arrow) planting dates; one arrow indicate a significant difference only between April and June. Each variable is presented as the mean, standard deviation (s.d.), minimum, maximum, and number of non-zero values for n number of samples across three (April, May, and June) planting months for August-September harvest in 2017.

Trapezoidal integration was performed on CV and CDPV of each juice sample for both large (0.5–1 V) and small (0.3–0.6 V) anodic current peaks (S1 Fig in S1 File). Significant planting and cultivar (p<0.0001 for both) effects were observed for the large anodic peak (0.5–1 V) of CV. No.5 Gambela was significantly more electron-rich (larger integrated area of the primary anodic peak; marked * in Table 1) than all other cultivars. Later planting increased the electron donating capacity of juice (quantified as the anodic peak area), in agreement with the decreasing bulk E_h trend from April to May (Table 1). Opposite planting month effect (decrease from May to June) was observed for the small peak (0.3–0.6 V) area that was the

Table 1. Trapezoidal integration of major (0.5–1 V) and minor (0.3–0.6 V) anodic CV, CDPV peaks; E_h in mV; and UV/visible absorbance at wavelengths attributable to trans-aconitic acid and polyphenol [8]. Only significantly different p-value (<0.05) are given for the cultivar and planting month effects with post-hoc Tukey's HSD test. Total number of samples (n for 24 cultivars × planting months (April, May, and June of 2017) × triplicate field plots), mean, standard deviation (s.d.), minimum, maximum, and number of non-zero values for each variable. Cultivar×planting interaction was observed only for the UV/visible absorbance at 270 nm (p = 0.02).

variable		mean	s.d.	min	max	non- zero	significant (<0.05) p value		
							cultivar	planting ^a	
Trapezoidal integrated areas of anodic derivative current in juice									
CV (0.5-1V)	172	2.1E- 06	7.5E- 07	7.8E- 07	5.5E- 06	172	<0.00001 (No.5 Gambela>all others)*	<0.00001 (-↑)	
CV (0.3–0.6V) ^b	163	6.5E- 08	3.4E- 08	1.4E- 08	2.0E- 07	163	0.00001 (Chinese>No.5 Gambela, Atlas, N109AxAtals, Isidomba, N111AxAtlas, N110AxAtlas;	<0.00001 (-↓)	
							N110AxN98>No.5 Gambela, Atlas, Isidomba, N110AxAtlas; N98>Atlas; N111B>Atlas)		
CDPV (0.5-1V)	172	2.7E- 06	8.4E- 07	7.3E- 07	7.2E- 06	172	0.00002 (No.5 Gambela>N98, N111AxDale, N111AxIsidomba, Isidomba, N110AxDale, N111AxN98, Dale, N109AxIsidomba, N109AxChinese, N110AxAtlas;* Atlas>Isidomba, N111AxN98, Dale; Isidomba <n109axatlas, chinese,="" dale<n109axatlas,="" n109b)<="" n109b;="" td=""><td><0.00001 (↓↑)</td></n109axatlas,>	<0.00001 (↓↑)	
CDPV (0.3-0.6V)	171	2.1E- 07	8.7E- 08	3.5E- 08	6.1E- 07	171	0.00002 (N98>N110AxChinese, N111AxDale, Atlas, N109AxAtlas, Isidomba, N111AxN98, N109AxIsidomba, N110AxAtlas, N110AxIsidomba; N110AxN98>Isidomba, N111AxN98, N109AxIsidomba, N110AxAtlas; N111B>Isidomba)	<0.00001 (-↓)	
bulk reduction potential and UV/visible absorbance of juice									
E _h (mV)	173	281	32	98	373	173		0.02 (↓-)	
UV/vis 270nm (trans-aconitic acid)	174	0.95	0.48	0.20	2.44	174	<0.00001 (No.5 Gambela>all others;* Isidomba, Dale>Chinese, N111AxChinese, N111B; N109AxDale>Chinese, N111AxChinese, N111AxN98, N110B, N111B; Atlas>N111AxChinese)	<0.0001 (↑↑)	
UV/vis 340nm (polyphenol)	174	0.18	0.09	0.05	0.67	174	<0.00001 (No.5 Gambela>all others)*	<0.00001 (^-)	

^aArrows indicate time trend (p<0.05 by post-hoc Tukey): increase (\uparrow), decrease (\downarrow), or no change (-) from April to May (first arrow), and from May to June (second arrow) planting dates; one arrow indicates significant difference only between April and June.

*Maximum in No.5 Gambela (p<0.05 by Tukey).

^bIf peak exists.

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highest for Chinese, compared to 6 other cultivars including No.5 Gambela. Trapezoidal integration of CDPV gave slightly different results from CV. For the large peak (0.5–1 V), No.5 Gambela (p<0.00001) was significantly more electron-rich than 9 other cultivars (as opposed to 23 others by CV). For the small peak (0.3–0.6 V), N98 had significantly (p = 0.0002) higher peak area than 9 other cultivars. Based on Fig.1, primary CV peak of juice could originate from tryptophan, tyrosine, or polyphenols. This peak was highest in No.5 Gambela, which contains the highest concentrations of trans-aconitic acid and aromatic structures, and has among the lowest concentrations of amino acids (as % contribution of tryptophan and tyrosine by EEM/PARAFAC) [8]. Because trans-aconitic acid is not electroactive with the method employed in the present study (Fig 1), the measured anodic current was determined to primarily originate from polyphenols. Using 2 mm SPE, CDPV enhanced the small peak (0.3–0.6 V) relative to CV (Section V of S1 File). The larger surface area (4x5 mm) of SPE further enhanced the minor peak (Section V of S1 File). Because only a small return (cathodic) peak was obtained for juice samples investigated in this study, CDPV was used for the Gaussian fitting of the cathodic peak (Section VI of S1 File).

variable	n	mean	s.d.	min	max	non- zero	significant (<0.05) p value			
							cultivar	planting ^a	interaction	
Gaussian integration of anodic and cathodic juice peaks ^c										
CV anodic area (0.5-1V)	170	2.2E- 06	7.6E- 07	8.6E- 07	4.9E- 06	170	<0.00001 (No.5 Gambela>all others)*	<0.00001 (-↑)		
CV anodic max height (0.5-1V)	170	1.4E- 05	3.9E- 06	7.4E- 06	2.6E- 05	170	0.0001 (No.5 Gambela>all except 8 cultivars ^{d,*} ; Chinese>Isidomba, N111AxN98, Dale)	<0.00001 (↓↑)		
CV anodic E _{pa} (0.5-1V)	170	0.88	0.03	0.82	0.95	170	0.00001 (No.5 Gambela>all except Isidomba, Dale;* Dale>all except No.5 Gambela,	<0.0001 (↑↑) N109AxDale, Atlas, N111AxIsidomba, Isidomba, N110AxAtlas; Isidomba>all except 9 cultivars; ^e N109AxDale>N98)	0.04 ^f	
CDPV anodic area (0.5-1V)	171	2.7E- 06	8.3E- 07	3.3E- 07	6.9E- 06	171	0.0002 (No.5 Gambela>N98, N111AxIsidomba, N109AxN98, Isidomba, N110AxDale, N111AxN98, Dale;* Isidomba <atlas, 109axatlas,="" n109b)<="" td=""><td>0.00006 (\1)</td><td></td></atlas,>	0.00006 (\1)		
CDPV anodic height (0.5-1V)	171	1.7E- 05	5.1E- 06	2.4E- 06	4.4E- 05	171	0.003 (Isidomba <no.5 atlas,<br="" gambela,="">109AxAtlas, Chinese, 109B)</no.5>	0.00001 (↓↑)		
CDPV anodic E _{pa} (0.5-1V)	171	0.80	0.04	0.43	0.86	171	0.04 (N109AxN98 <no.5 gambela,="" isidomba)<="" td=""><td>0.00002 (↑-)</td><td></td></no.5>	0.00002 (↑-)		
CDPV anodic area (0.2–0.6V) ^b	160	2.2E- 07	1.2E- 07	0	8.1E- 07	160	0.00004 (N98>all except N110AxN98, Chinese, N110AxDale, N111B, N109B)	0.01 (↓)		
CDPV anodic height (0.2–0.6V) ^b	160	1.7E- 06	5.8E- 07	2.4E- 07	4.0E- 06	160	0.00007 (N110AxN98>N111AxDale, Isidomba, N111AxN98, N109AxIsidomba, N110AxAtlas; N98>Isidomba, N111AxN98, N110AxAtlas)	<0.00001 (-↓)	0.04	
CDPV anodic E _{pa} (0.2–0.6V) ^b	160	4.5E- 01	3.5E- 02	4.1E- 01	7.9E- 01	160		0.0007 (-↑)		
CDPV cathodic area ^{b,g}	96	1.0E- 07	1.4E- 07	0	8.2E- 07	96		0.009 (↑↓)		
CDPV cathodic height ^{b,g}	96	9.2E- 07	3.5E- 07	9.2E- 08	2.5E- 06	96	0.00002 (No.5 Gambela>all others)*			
CDPV cathodic E _{pa} ^{b,g}	96	0.75	0.03	0.57	0.79	96		0.0004 (-↓)		

Table 2. Gaussian integration of CV anodic and CDPV anodic/cathodic peaks. Significant cultivar, planting month, and interaction p-values (<0.05 with post-hoc Tukey's HSD test) for the fitted area, maximum height, and E_{pa} are provided.

^aArrows indicate time trend (p<0.05 by post-hoc Tukey): increase (\uparrow), decrease (\downarrow), or no change (-) from April to May (first arrow), and from May to June (second arrow) planting dates; one arrow indicates significant difference only between April and June.

^bIf peak exists.

^cE_{pa} is the center of peak in volts; area is Gaussian-integrated current peak area at E_{pa}; max height is the peak current height in amps at E_{pa}.

^dN110AxChinese, Atlas, N109AxAtlas, N111AxAtlas, Chinese, N111AxChinese, N109AxIsidomba, and N109B.

^eNo.5 Gambela, N111AxDale, N109AxDale, Atlas, N111AxIsidomba, N111AxAtlas, N110AxDale, Dale, N110AxAtlas, and N110AxIsidomba. ^fNo separation by Tukey.

^gOne-way ANOVA was employed (instead of factorial), because of low n.

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In Table 2, Gaussian integration was performed on CV and CDPV to obtain electrochemical parameters: (1) peak area to quantify the amount of electron donated or accepted by each juice sample, (2) E_{pa} and E_{pc} to differentiate chemical structures and ring substituents responsible for oxidation/reduction (Fig 1 and S3 Table in S1 File), and (3) maximum peak current height in A (first derivative) at E_{pa} or E_{pc} . For CV, Gaussian integration of anodic peak area (0.5–1 V) gave the same cultivar and planting month effects as trapezoidal integration (Tables 1 and 2): highest area of No.5 Gambela and an increase from May to June planting. Maximum peak height showed similar cultivar and planting month effects as the peak area, and No.5 Gambela was significantly higher than 8 other cultivars. Similarly to the anodic current area and height, No.5 Gambela had significantly higher (p<0.00001) E_{pa} (0.93 ± 0.01 V as mean±standard error across planting months) than all other cultivars, except Dale (second highest E_{pa} , 0.91 ± 0.01 V) and Isidomba (0.90 ± 0.01 V). Higher E_{pa} indicates higher aromaticity of polyphenolic structures, requiring more positive voltage for oxidation [31]. Although the cultivar x planting month interaction was significant (p = 0.04) for E_{pa} , Tukey's HSD did not differentiate the effects. In conclusion, No.5 Gambela had uniquely the highest anodic peak area and height, as well as E_{pa} (Table 2). Collectively, No.5 Gambela was the most electron-rich (peak area and height) and least easily oxidized (E_{pa}) of all 24 cultivars investigated in S1 Table in S1 File. In addition, CV was more sensitive to differentiating cultivar and planting month effects than CDPV by both trapezoidal and Gaussian fits.

For the large peak (0.5–1 V) of CDPV, No.5 Gambela had significantly (p = 0.0002) higher Gaussian-integrated area than 7 other cultivars including Dale and Isidomba. The E_{pa} values obtained by CDPV were lower than CV, and only cultivar effects were observed between the highest (0.84±0.01 V for No.5 Gambela and 0.82±0.01 V for Isidomba) and lowest (0.76±0.01 V for N109AxN98) cultivars. Gaussian integration of the small (0.2–0.6 V) peak showed similar cultivar effects as Trapezoidal integration (Table 1), where N98 had the highest peak area. For the peak height, significant interaction (p = 0.04) was observed, where May planting of N98 significantly differed from most of cultivars in June planting. No significant cultivar effects were observed for the E_{pa} of small (0.2–0.6 V) anodic peak by CDPV.

Because only 96 of 171 CDPV voltammograms had cathodic return peaks (0.5–1 V), oneway ANOVA (with cultivar and planting month as the categorical factors, instead of factorial ANOVA) was carried out for the cathodic parameters of CDPV in <u>Table 2</u>. Cultivar effects were only observed for the peak height of cathodic return CDPV. No.5 Gambela had significantly higher cathodic current height (p = 0.00002) than all other cultivars, similarly to the Gaussian and trapezoidal integration of corresponding anodic current area by CV at 0.5–1 V. The return peak is attributable to the reduction of oxidized polyphenols in No.5 Gambela.

In Table 1, UV/visible absorbance of juice was analyzed at selected wavelengths to determine the cultivar effects of polyphenols at 340 nm, where trans-aconitic acid does not absorb. At 340 nm where only polyphenols absorb significantly, No.5 Gambela had significantly higher absorbance $(0.37\pm0.02, p<0.00001)$ than all other cultivars. This trend follows that of primary anodic CV areas used to measure polyphenol electrochemistry. At lower wavelength (270 nm) where trans-aconitic acid absorbs significantly, cultivar and planting month effects were similar to 340 nm. Therefore, trans-aconitic acid and polyphenols likely have similar cultivar and planting month effects. At 270 nm, significant cultivar x planting month interaction was observed, especially between April and May plantings (p = 0.02).

Fig 4 presents an E_h -pH diagram for one-electron oxidation of tryptophan (to form indole radical) [31], tyrosine (phenoxyl radical) [32], quercetin (radical of catechol in B ring) [33], and riboflavin (semiquinone) [34], all reported in the literature against normal hydrogen electrode (NHE). Reduction potentials of quercetin and riboflavin in Fig 4 was obtained kinetically (radical generation and spectrophotometric quantification) by pulse radiolysis at fixed pH [33], while CV was employed for tryptophan and tyrosine. Standard hydrogen electrode (SHE) is an ideal (theoretical) electrode with H⁺ at unity activity saturated with H₂ gas at unity fugacity [35, 36]. NHE is a different reference electrode containing 1 atm partial pressure H₂ and 1M H₂SO₄ [37]. Degree of deviation from the standard state is unknown in experiments employing NHE, where activity of different ionic solution components cannot be controlled [35–37].



Fig 4. E_h-pH diagram for one-electron oxidation/reduction of tryptophan (pink line), tyrosine (blue line) [32], quercetin (red line) [33], riboflavin (green line) [34], and sweet sorghum juice samples from 2017 planting year (black symbol) against the standard hydrogen electrode. Error bars for the 2017 juice represents mean±s.d. for 173 samples in Table 2. Literature values were used for amino acids (in 0.2 M KCl, pH 13) [32], quercetin (in 0.5 M KCl; ethylene glycol as co-solute) [31, 33, 39], and riboflavin (in 0.1 M NaHCO₂) [34]. Literature E_h-pH values were reported against NHE. Experimental E_h value of juice samples could be converted to SHE scale by adding 200 ±60mV.

Those literature values are compared against E_h and pH values of sweet sorghum juice samples from 2017 planting year (black symbol as mean±s.d. for 173 samples in Table 1). Experimental E_h value of juice samples could be converted to SHE scale by adding 200±60mV, as described in Materials and Methods. At pH 5.7±0.3, the E_h value of sweet sorghum (0.281±0.032 V) is lower than amino acids, higher than riboflavin, and closest to quercetin. The, E_h , pH, and EC of 2017 juice samples all decreased with the planting month (Table 1) [8]. As shown in Table 1, there was no statistically significant (p<0.05) difference in E_h (and pH provided in our previous report [8]) values among different cultivars.

Quercetin has the lowest one-electron reduction potential among flavonoids (e.g., catechin, galangin), because the electron-donating hydroxyl group in the O-3 position of the C ring is conjugated with the catechol B radical through the 2,3 double bond [33]. Observation of E_{h^-} pH behavior in Fig 4 suggests that semiquinone radical [3, 4] of quercetin-like polyphenolic structure is responsible for the redox reactivity of sweet sorghum. The E_{h^-} pH trend of plant phenolics is traditionally investigated by manually changing the pH of a given sample by adding acid or base [38]. Instead, Fig 4 (filled circle labeled "2017 juice") presents intrinsic E_{h} and pH values of 173 sweet sorghum juice samples collected from the field experiments, without

modifying pH. Although thermodynamic driving force does not predict the kinetic lability, polyphenols in sweet sorghum (Fig 4) are capable of reducing the primary amino acids in juice, and other chemical structures having higher reduction potentials.

Supporting information

S1 File. Sections I-VI: Methods; representative juice samples (24 total matching S1 Table) used to calculate Pearson's correlation coefficient across 0.7–1 V range of CV anodic voltammogram; Gaussian fitting parameters for the authentic standards; representative cultivars enriched with amino acids and polyphenols; influence of SPE area and methods (CV, CDPV) on peak properties; representative anodic and cathodic CDPV peaks subjected to integration.

(DOC)

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