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Pheophorbide Derivatives Isolated from Açai Berries (*Euterpe oleracea*) Activate an Antioxidant Response Element *In Vitro*

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Abstract

Activity-guided fractionation was used to isolate and identify two components of the Brazilian açai berry (*Euterpe oleracea* Mart.) with the ability to induce antioxidant response element (ARE)-dependent gene transcription in human hepatoma (HepG2) cells. Using an ARE-Luciferase reporter construct in cultured HepG2 cells, a suite of fractions from dried and powdered açai berries were evaluated for transcriptional up-regulation of the luciferase gene. Active fractions were further refined until several pure compounds were isolated and identified. These compounds belong to the pheophorbide class of molecules, and are composed of the methyl and ethyl esters of the parent pheophorbide A, all of which are classified as photosensitizers. Using standard pheophorbides, dose response studies were carried out, and ARE-activation could be observed at concentrations as low as 8.2 μ M and 16.9 μ M for pheophorbide A methyl ester and pheophorbide A, respectively. These studies not only suggest a possible source of antioxidant properties for the açai berry, but may also explain the recently identified photosensitizing abilities of açai products as well.

Keywords

Euterpe oleracea Mart; açai; Nrf2; anti-oxidant; pheophorbide

The açai [*Euterpe oleracea* Mart. (Arecaceae)] berry has become very popular in recent years as a functional food resulting from its health promoting effects, and in particular, its ability to reduce oxidative stress. Although açai products have been shown to act effectively as an antioxidant *in vitro* and *in vivo*, it has not been well-characterized with regard to the mechanisms associated with these effects in humans [1]. Studies have demonstrated certain types of antioxidant effects in a variety of model systems [2,3] and whole organisms [4,5] that are consistent with the Nrf2-dependent mode of action. In addition, Monge-Fuentes *et al.* [6] demonstrated that formulations containing açai oil were effective photosensitizers when applied to B16F10 melanoma cells and in tumor bearing C57BL/6 mice, indicating certain compounds in the berry may be effective in photodynamic therapy for melanoma.

This is attributed to production of singlet oxygen, which has been shown to stimulate Nrf2-dependent pathways. Other studies in mice have shown that açai products reduce lipid peroxidation [7], increase antioxidant gene expression [8], and protect against carcinogen induced oxidative stress [9].

Açai extract can reduce oxidative stress induced by cigarette smoke in mice, which could also be partially due to activation of anti-oxidant gene expression. In addition, human trials have indicated efficacy in reducing pain and inflammation, along with improvements in range of motion, all of which were highly correlated to antioxidant status [10,11]. Furthermore, chemical analysis of the freeze dried açai [12] has indicated the presence of several types of polyphenolic compounds with known antioxidant properties, including anthocyanins and flavones [3]. Schauss *et al.* [12] reported the pulp of açai juice to contain high levels of anthocyanins and other classes of polyphenolic compounds, and according to Milbury *et al.* [13], the bioavailability of these compounds may be sufficient to effect biological targets.

The luciferase reporter used in the study, pGL3-*wtARE*, was constructed according to methods reported by Kim *et al.* [14] using the pGL3 luciferase vector (Figure 1). This vector incorporates a functional Antioxidant response element (ARE) in front of an active promoter controlling luciferase gene expression. Thus, Nrf2 activators will induce luciferase expression and activity in the cultured cells. A second construct in which the ARE sequence was mutated (pGL3-mARE) was also produced as a negative control. The data shown in Figure 2 demonstrate that luciferase activity was substantially induced at 100 μ M tBHQ using the *wtARE* construct, while pGL3 and pGL3-mARE were unresponsive to the inducer. Furthermore, the açai chloroform extract at 200 μ g/mL produced an inductive effect that was even more pronounced than tBHQ, suggesting the presence of ARE inducers in this extract. Induction by the butanol, aqueous and hexane extracts, at the same concentration, were less than 1.5-fold. It should be noted that the level of induction by the positive control, tBHQ, varied from experiment to experiment, however, it generally fell within the range of 2.5- to 5-fold.

The bioactivity-guided fractionation approach was then followed using the chloroform extract in an attempt to identify components in this fraction with the ability to activate the ARE-luciferase reporter system in the HepG2 cells. Figure 3 represents the scheme used to track active fractions through each generation of purification. The chloroform extract was fractionated into 10 sub-fractions designated 34-1 through 34-10. Each fraction was assessed for its ability to activate the reporter plasmid in HepG2 cells at a concentration of 50 μ g/mL; this concentration was chosen based on previous experience working on the bioactivity-directed fractionation of herbals [15-17]. Fractions highlighted by the vertical shading represent those fractions that were carried forward in the purification based on activity and yield. Since fraction 34-8 resulted in the most significant activation, it was subsequently fractionated into 8 sub-fractions designated as 38-1 through 38-8. Sub-fractions 38-1 through 38-4 displayed modest induction at 50 μ g/mL, however, potent induction was not observed for any of these sub-fractions. These sub-fractions were therefore fractionated further producing the 61-series, the 62-series, the 63-series, and the 64-series, all indicated by the vertical hashed shading. The 38-8 fraction (checkered shading

in Figure 3) actually produced a significant response in the reporter assay, however, due to paucity of sample further purification was not pursued. Importantly, multiple criteria were used at each stage to select fractions to carry forward in the process. In addition to activity in the luciferase assay, the total mass of material in the fraction was an important consideration in the fractionation approach, along with the chromatographic complexity of the samples at later stages of the process. For example, although the fraction 64–1 (shown as checkered shading) was very active, the amount of material found in this fraction was very low, and subsequent purification was not feasible.

Following this approach to completion, fractions 97-B and 97-D were shown to contain single active compounds (Figure 4). Fraction 97-B (3.3 mg) was identified as pheophorbide A methyl ester using high resolution mass spectrometry with obsd. m/z 607.2917 $[M+H]^+$ (calcd. for $C_{36}H_{39}N_4O_5$, 607.2915) and comparison to literature NMR data [18]. Fraction 97-D (2.7 mg) was identified as pheophorbide A ethyl ester using high resolution mass spectrometry with obsd. m/z 621.3074 $[M+H]^+$ (calcd. for $C_{37}H_{41}N_4O_5$, 621.3071), and comparison to literature NMR data [19].

Finally, pure standards of pheophorbide a (the parent compound) and pheophorbide a methyl ester were obtained commercially, and a dose response was generated for each using the ARE-driven luciferase reporter. As shown in Figure 5, both compounds resulted in increased luciferase expression at concentrations as low as 8.2 μ M, and there was a clear dose-dependent increase observed for each.

Pheophorbides are known plant products related to the chlorophyll molecule and have been shown to have interesting medical properties and applications. For example, a major application of these photoactive compounds is in photodynamic therapy [20]. Indeed, Monge-Fuentes *et al.* [6] demonstrated both in cultured cells and in tumor induced mice that açai berry oil possess compounds with photosensitizing properties, which may be useful for photodynamic therapy. Furthermore, Liu *et al.* [21] have shown the parent compound isolated from the açai berry, pheophorbide A, reduces proliferation and metastasis of human prostate cancer cells in vitro when used as a photodynamic agent. In addition, Hagiya *et al.* [22] demonstrated that pheophorbides could induce Nrf2-dependent gene expression, consistent with our findings in HepG2 cells.

Due to the light sensitive nature of the compounds, all treatments were carried out under black light. That induction of the luciferase reporter was observed in a dose dependent manner in the absence of light suggests that the pheophorbides possess some direct activating effects on the Nrf2-dependent anti-oxidant pathway, however, additional experiments to evaluate the effects of light on this induction would help to clarify the mechanisms involved.

In summary, we have demonstrated the effective use of bioactivity-guided fractionation for the isolation and identification of individual constituents of the Brazilian açai berry with potential antioxidant properties related to their ability to stimulate ARE-dependent gene induction. Identification of these active compounds as photosensitizers suggests alternative therapeutic applications for this versatile natural product.

Experimental

Açaí berry freeze dried extracts

Freeze dried powdered açaí berry was purchased from Optimally Organic Inc, (Westlake Village, CA); all of the material was from a single lot, and a voucher specimen was deposited in the Herbarium of the University of North Carolina at Chapel Hill (NCU601336). Moreover, a sample of the berry powder was submitted for genetic analysis to Authentech Inc, (Richmond, CA), which showed the most likely identity of the sample to be *Euterpe oleracea* (Açaí), based on comparison with authentic samples and divergence from all closely related species analyzed.

Construction of the antioxidant reporter vector

The firefly luciferase reporter plasmid pGL3 was purchased from Promega and was genetically engineered to include a known ARE sequence that is present in the human thioredoxin promoter. Standard cloning techniques were employed to produce the corresponding pGL3-ARE vector described by Kim *et al.* [14] which has been used effectively to monitor ARE induction in a variety of human cell lines, including HepG2.

Induction of ARE-reporter in HepG2 cells

Expression of luciferase was determined using a commercial Dual-Glo® Luciferase Assay System from Promega according to the established protocols and a BMG Labtech PolarStar Optima 96 well plate reader for luminescence measurements. HepG2 cells were obtained from ATCC and have been grown and maintained in Dulbecco's modified Eagle medium, high (4.5 g/L) glucose, 4.0 mM glutamine and 1.0 mM sodium pyruvate [21]. For all subsequent experiments, cells were cultured and grown to 60% confluence and transfected with the pGL3-ARE reporter plasmid using FuGene® reagent for 24 hours. Cells were treated with extract or pure compounds for 24 hours. The activities of luciferase expressed in HepG2 cells were normalized by co-transfecting HepG2 cells with pRL-TK Renilla reporter vector as an internal control. All cell culture experiments were carried out in triplicate, and independently reproduced. Positive control experiments were also carried out in parallel in which *tertiary* butylhydroquinine (tBHQ), at 100 µM, was used to induce luciferase activity in the pGL3-ARE containing cells. Active fractions were identified as those inducing a minimum of 2x luciferase activity relative to the negative control.

Preparation of extracts

For the initial fractionation, 1820 g of freeze dried açaí powder was extracted in 4L MeOH, by stirring at room temperature, to generate a methanol extract (Figure 2). The methanol fraction was filtered and the volume reduced to 2L and subsequently partitioned with 9:1:10 MeOH:H₂O:hexane in a total volume of 4L. Aqueous MeOH is immiscible with hexane, which allows for the “defatting” of the original MeOH extract, thereby generating a hexane fraction (1) containing 62.5 g dried material (Figure 2). The aqueous MeOH fraction was dried and partitioned with 4:1:5 chloroform:MeOH:H₂O in a total volume of 4L. The organic layer produced 18.0 g of dried chloroform extract (2). Finally, the remaining aqueous fraction was partially evaporated to remove methanol and partitioned between equal

volumes of n-BuOH and H₂O (total volume 4L) resulting in 29.6 g and 78.1 g of dried butanol (3) and aqueous (4) fractions, respectively.

Total fractionation protocol through the 97 series

34 series—The chloroform extract (18 g) was adsorbed onto celite and chromatographed on a 330-gram silica column utilizing a Combiflash Rf 200 automated flash chromatography system (Teledyne-Isco, Lincoln, NE). A gradient from 100 % n-hexanes to 100 % chloroform then to 100 % MeOH was used over 28 column volumes to obtain 10 fractions labeled the 34 series.

38 series—Fraction eight of the 34 series (34–8, 6.14 g) was adsorbed onto celite and chromatographed on a 120-gram “gold” (high performance) silica column utilizing a Combiflash Rf 200. A gradient from 100 % n-hexanes to 100 % acetone then to 100 % MeOH was used over 18 column volumes to obtain 8 fractions hereby labeled the 38 series.

61 series—Fraction one of the 38 series (38–1, 2.00 g) was adsorbed onto celite and chromatographed on a 30-gram diol column utilizing a Combiflash Rf 200. A gradient from 100 % n-hexanes to 100 % ethyl acetate then to 20 % methanol was used over 22 column volumes to obtain 7 fractions hereby labeled the 61 series.

62 series—Fraction two of the 38 series (38–2, 1.18 g) was adsorbed onto celite and chromatographed on a 30-gram diol column utilizing a Combiflash Rf 200. A gradient from 100 % n-hexanes to 100 % ethyl acetate then to 20 % methanol was used over 22 column volumes to obtain 5 fractions hereby labeled the 62 series.

63 series—Fraction three of the 38 series (38–3, 0.96 g) was adsorbed onto celite and chromatographed on a 30-gram diol column utilizing a Combiflash Rf 200. A gradient from 100 % n-hexanes to 100 % ethyl acetate then to 20 % methanol was used over 24 column volumes to obtain 6 fractions labeled the 63 series.

64 series—Fraction four of the 38 series (38–4, 0.73 g) was adsorbed onto celite and chromatographed on a 30-gram diol column utilizing a Combiflash Rf 200. A gradient from 100 % n-hexanes to 100 % ethyl acetate then to 100 % methanol was used over 26 column volumes to obtain 5 fractions labeled the 64 series.

78 series—Fraction two of the 62 series (62–2, 739 mg) was chromatographed on a Gemini-NX C18 reverse phase HPLC column (250 × 21.2 mm) utilizing a ProStar preparative HPLC system (Varian Inc., Palo Alto, CA) running at 21.2 mL/min. A gradient from 50:50 to 100:0 acetonitrile:water was used over 50 min to obtain 9 fractions labeled the 78 series.

97 series—Fraction nine of the 78 series (78–9, 64.7 mg) was chromatographed on a Gemini-NX C18 reverse phase HPLC column (250 × 21.2 mm) utilizing a ProStar preparative HPLC system running at 21.2 mL/min. A gradient from 70:30 to 100:0 acetonitrile:water over 15 min followed by isocratic elution at 100:0 for 15 min to obtain 9 fractions labeled the 97 series.

MS and NMR characterization—High resolution electrospray ionization mass spectrometry (HRESIMS) data were collected using an electrospray ionization source coupled to a Q Exactive Plus system (Thermo Fisher Scientific, San Jose, CA) in positive ionization mode via a liquid chromatography/autosampler system comprised of an Acquity UPLC system (Waters Corp., Milford, MA).

The NMR data were collected using a JEOL ECS-400 spectrometer (JEOL USA, Inc., Peabody, MA) operating at 400 MHz for ^1H and 100 MHz for ^{13}C , and equipped with JEOL normal geometry broadband Royal probe.

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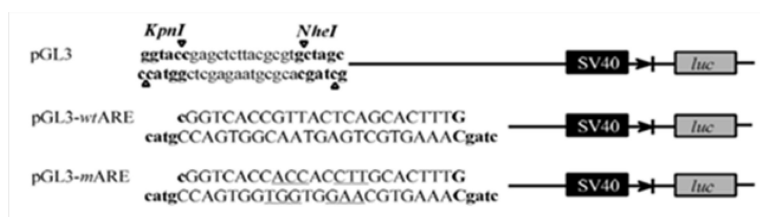


Figure 1:
Schematic illustration of promoter region of pGL3 plasmid vector, wild-type and mutant ARE constructs.

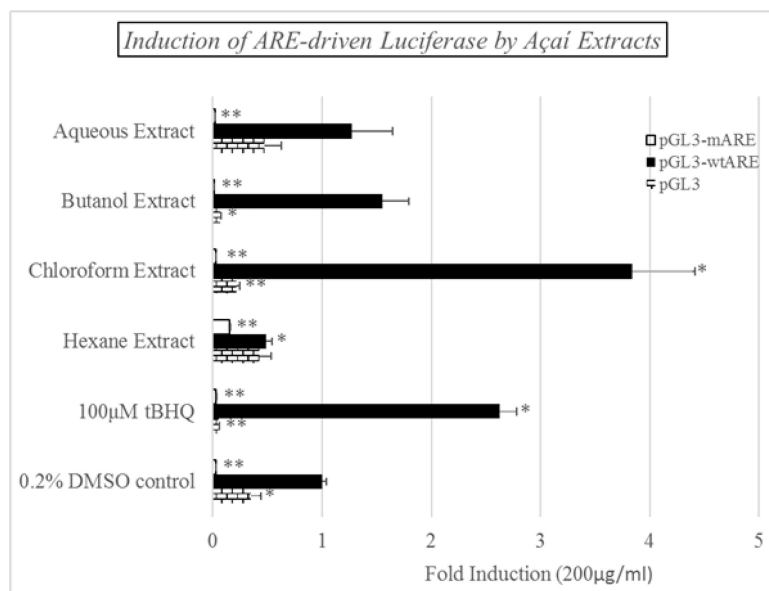


Figure 2: Induction of ARE-driven luciferase expression in cultured HepG2 cells by the açai-97-Series. The HepG2 cells were treated with each fraction at a dose of 50 µg/mL for 24 hrs.

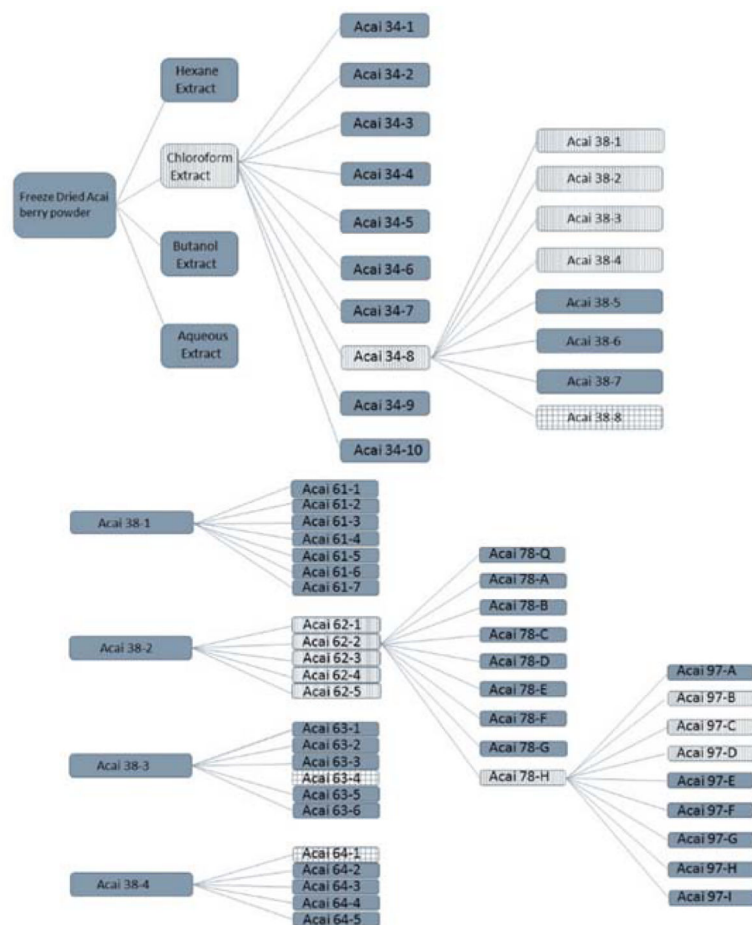


Figure 3: Fractionation scheme for the active chloroform extract from açai berry powder. The fractions highlighted with vertical lines were those selected for further fractionation in this scheme. The checked fractions indicate those that were active but could not be further fractionated..

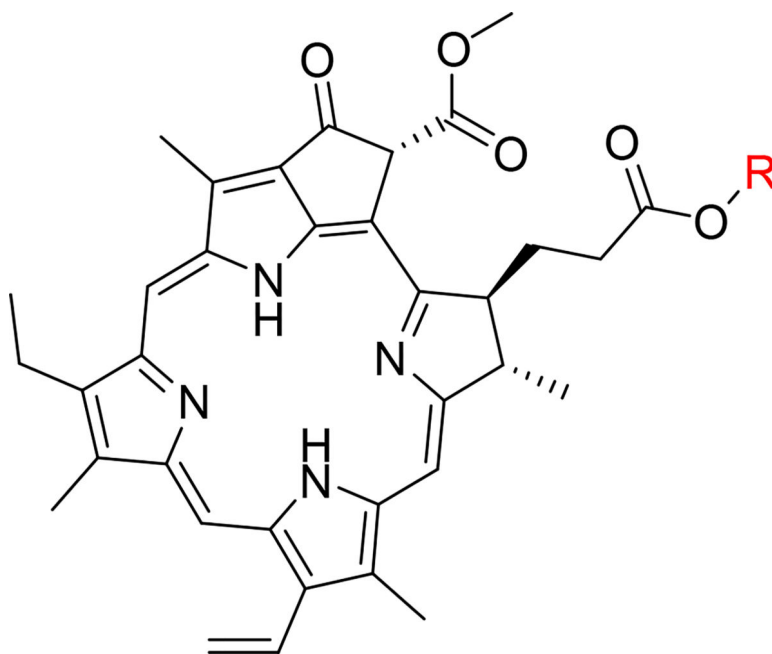


Figure 4:
General structure of the pheophorbide compounds isolated by activity guided fractionation. Açai 97-B was identified as the methyl ester ($R=CH_3$) whereas Açai 97-D was identified as the ethyl ester ($R=CH_2CH_3$).

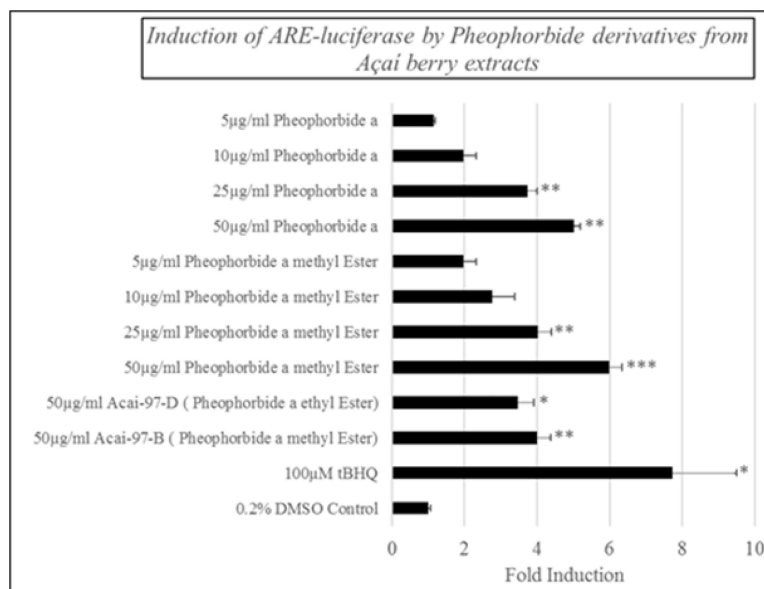


Figure 5. Induction of ARE-driven luciferase expression in cultured HepG2 cells by pheophorbide a methyl ester and pheophorbide a. Cells were treated for 24 hrs at the indicated dose prior to measuring ARE-driven luciferase activity.