



Polyphenols from *Allanblackia floribunda* seeds: Identification, quantification and antioxidant activity



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ABSTRACT

Oil rich seeds of *Allanblackia floribunda*, a tree from tropical Africa, have traditionally been used in food preparation. Furthermore, the therapeutic properties of various parts of this tree have long been exploited in traditional medicine. As both food and pharmaceutical industries show growing interest in tropical tree crops, this study aimed to investigate whether *A. floribunda* seeds could also be used as a source of potentially bioactive compounds. The polyphenol profile revealed six predominant compounds which were identified by HPLC-PDA-ESI/MSⁿ as the biflavonoids morelloflavone, Gb-2a and volkensiflavone and their respective glucosides. A range of less abundant flavones, flavonols and flavan-3-ols was also detected. All six major compounds showed antioxidant activity, with the activity of morelloflavone, its glucoside and Gb-2a-glucoside comparable with that of ascorbic acid. The main compounds accounted for approximately 10% of dry weight, making the seeds used for oil production a rich source of biflavonoids as a by-product.

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1. Introduction

Allanblackia floribunda Oliv (Clusiaceae or Guttiferae) is an ever-green tree which grows in the tropical rainforests of Africa to a height of up to 30 m and is traditionally used in a variety of ways (Orwa & Munjuga, 2007; Orwa et al., 2009). Whilst the seeds consist to over 60% of oil and provide edible vegetable fat (Orwa et al., 2009; Wilfred, Adubofuor, & Oldham, 2010), various parts of the tree, including roots and bark, are used in traditional medicine for the treatment of a range of ailments such as toothache, dysentery and coughs (Betti, 2004; Kayode, 2006; Olowokudejo, Kadiri, & Travihi, 2008). Furthermore, *Allanblackia floribunda* extracts have been reported to exhibit antimicrobial, anti-inflammatory and antioxidant activity (Ajibesin, Rene, Bala, & Essiett, 2008; Ayoola

et al., 2008; Ayoola et al., 2009; Kuete et al., 2011) as well as potential antimalarial (Azebaze et al., 2015) and anticancer effects (Fadeti, Fadeti, Adejumo, Okoro, & Myles, 2013). To date, a range of potentially bioactive compounds have been reported in a variety of extracts from *Allanblackia floribunda* and related species, including benzophenones, xanthenes and biflavonoids (Fuller, Blunt, Boswell, Cardellina, & Boyd, 1999; Locksley & Murray, 1971; Nkengfack, Azebaze, Vardamides, Fomum, & van Heerden, 2002). In the seed, the focus has mainly been on oil, with some information available on antioxidative properties of *A. floribunda* seed cake (Boudjeko, Ngomoyogoli, Woguia, & Yanou, 2013). With growing interest in *Allanblackia* species as tree crop in particular for the food industry (IUCN, 2014), the aim of this study was to investigate the polyphenol profile of *A. floribunda* seeds in order to establish whether seed remnants after plant oil extraction could be a source of bioactive compounds with interest to the pharmaceutical sector.

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2. Materials and methods

2.1. Plant material and reagents

Fruits of *Allanblackia floribunda* were collected in three different years from uncultivated farmland in Oke Igbo, Ondo State, South Western Nigeria. The fruits were authenticated at the Forestry Research Institute of Nigeria (FRIN), Ibadan (voucher specimen number FHI107929). Seeds were separated from the mesocarp, and air-dried seeds were crushed, freeze-dried and milled into a fine powder (≤ 1.0 mm) with an automated bespoke ball mill robot (Labman Automation, Middlesbrough, UK). Briefly, three stainless steel balls (3 mm diameter) were added to approximately 200 mg seeds in a 2 mL microtube and shaken in the robot at the predetermined speed and duration to achieve the desired particle size.

HPLC grade solvents for extraction and analysis of phenolics were purchased from VWR (Lutterworth, Leicestershire, UK). Response factors for quantification were obtained using flavonoid standards with a minimum of 98% purity (Carbosynth Ltd, Compton, Berkshire, UK). All other analytical standards and chemicals were purchased from Sigma Aldrich (Gillingham, Dorset, UK).

2.2. Extraction of soluble phenolics

Soluble phenolics were extracted from approximately 20 mg seed powder by shaking with 5 mL 70% methanol for 15 min at room temperature. The sample was then centrifuged for 10 min at 1700g, the supernatant decanted and the pellet extracted twice more. Methanol was removed from the combined supernatants under vacuum at 60 °C before extracts were partially purified by solid phase extraction (SPE) using Sep-Pak C₁₈ cartridges (Waters Ltd, Elstree, UK) as described by Hauck, Gallagher, Morris, Leemans, and Winters (2014) and dried under vacuum at 60 °C. Prior to further analysis, samples were typically dissolved in 0.5 mL 70% methanol and diluted as necessary.

2.3. Liquid chromatography-tandem mass spectrometry

Secondary metabolites were analysed by reverse-phase high performance liquid chromatography with online photodiode array detection and electrospray ionisation-ion trap tandem mass spectrometry (HPLC-PDA-ESI/MSⁿ). Structural elucidation was performed on a Thermo Finnigan LC-MS system (Thermo Electron Corp, Waltham, MA, USA) comprising a Finnigan PDA Plus detector, a Finnigan LTQ linear ion trap with ESI source and a Waters C₁₈ Nova-Pak column (3.9 × 100 mm, particle size 4 μm), with column oven temperature maintained at 30 °C. The PDA scan range was set to 240–400 nm, and injection volume was typically 10 μL. The mobile phase consisted of water with 0.1% formic acid (solvent A) and methanol with 0.1% formic acid (solvent B). The column was equilibrated with 95% solvent A at a flow rate of 1 mL min⁻¹, with 10% going to the mass spectrometer, and the percentage of solvent B increased linearly to 65% over 60 min. MS parameters were as follows: sheath gas 30, auxiliary gas 15 and sweep gas zero (arbitrary units), spray voltage –4.0 kV in negative and 4.8 kV in positive ionisation mode, capillary temperature 320 °C, capillary voltage –1.0 V and 45 V, respectively, tube lens voltage –68 and 110 V, respectively, and normalised collision energy (CE) typically 35%.

Accurate mass measurements only were carried out on an Orbitrap Fusion Tribrid mass spectrometer (Thermo Fisher Scientific, Waltham, MA, USA) with heated ESI source coupled to a Dionex UltiMate 3000 ultra high performance liquid chromatography (UHPLC) system (Thermo Fisher Scientific). Chromatographic sepa-

ration was performed on a reverse-phase C₁₈ Hypersil Gold column (20 × 2.1 mm, particle size 1.9 μm; Thermo Fisher Scientific) which was maintained at a temperature of 60 °C and with solvents A and B as described above. The column was equilibrated with 95% solvent A at a flow rate of 0.4 mL min⁻¹, and the percentage of solvent B increased linearly to 100% over 7 min, followed by isocratic elution at 100% solvent B for 3.5 min. High resolution mass spectra were acquired in both negative and positive ionisation mode at a resolution of 120,000 with an automatic gain control (AGC) target of 2e⁵ and a maximum injection time of 100 ms, with other MS parameters as follows: vaporiser temperature 358 °C, spray voltage –2.5 kV in negative and 3.5 kV in positive ionisation mode, sheath gas 45, auxiliary gas 13 and sweep gas 1 (arbitrary units) and capillary temperature 342 °C.

2.4. Quantification of predominant compounds

For quantification of the predominant compounds, seed samples harvested in three different years were extracted in triplicate as described in Section 2.2, followed by reverse-phase HPLC. The system comprised a Waters 996 PDA detector and a Waters C₁₈ Nova-Pak radial compression column (8 × 100 mm, particle size 4 μm), with chromatographic conditions as described by Hauck et al. (2014) and detection wavelength set to 280 nm. Response factors (RF) for the flavonoid monomers naringenin, eriodictyol, apigenin and luteolin were obtained from linear standard curves over the range of 0.2–10 μg of standard and calculated as

$$RF = \text{amount } [\mu\text{g}] / \text{peak area}_{280\text{nm}}$$

All response curves had linear regression coefficients >0.996.

Response factors of biflavonoid aglycones were calculated as the mean of the relevant flavonoid monomer RFs, and for the quantification of biflavonoid glucosides these values were multiplied by the following correction factor to allow for the difference in molecular weight (M_r):

$$(M_r \text{ aglycone} + 162) / M_r \text{ aglycone}$$

2.5. Acid hydrolysis and analysis of sugar moieties and flavonoid aglycones

To identify the aglycone part of the flavonoid glycosides, acid hydrolysis of total extract was carried out by combining equal volumes of aqueous extract and 2 mol L⁻¹ HCl. After heating the solution to 90 °C for 1 h, the pH was adjusted to 3–4 with NaOH prior to partial purification by SPE as described in Section 2.2. Flavonoid aglycones were then analysed by HPLC-PDA-ESI/MSⁿ as described in Section 2.3 and identified by direct comparison with relevant flavonoid standards.

For identification of the sugar moieties of the main glycosylated metabolites, compounds were purified by collecting the HPLC eluent of individual peaks from the Waters system described in Section 2.4. After reducing the volume under vacuum at 60 °C acid hydrolysis was performed by combining equal volumes of aqueous extract and 2 mol L⁻¹ HCl. The solution was heated to 90 °C for 1 h and subsequently the pH was adjusted to 6–7 with NaOH. The samples were then loaded onto Strata-X-A strong anion exchange columns (Phenomenex, Macclesfield, UK) which had been conditioned with 100% methanol and equilibrated with water. The sugar containing non-binding fraction of the samples was collected for further analysis.

Monosaccharides were analysed by high-performance anion-exchange chromatography with pulsed amperometric detection (HPAEC-PAD). Separation of monosaccharides was achieved on a Dionex ICS-5000 system (Thermo Fisher Scientific, Waltham, MA, USA) fitted with an Eluent Generator and a Dionex CarboPac

SA10 column (250 × 4 mm; Thermo Fisher Scientific) with an SA-10 guard column (50 × 4 mm). The operating temperature of the column was 45 °C with an isocratic mobile phase of 1 mmol L⁻¹ KOH pumped at a rate of 1.5 ml min⁻¹. Monosaccharides were identified by comparison of retention times with glucose and galactose standards.

2.6. Antioxidant activity of selected metabolites

The predominant metabolites were purified by collecting the HPLC eluent of individual peaks from the Waters system described in Section 2.4 and then dried down and resuspended in 50% methanol at a concentration of 100 µg mL⁻¹. The antioxidant activity of individual compounds was measured in triplicate by 1,1-diphenyl-2-picrylhydrazyl (DPPH) radical scavenging assay (Brand-Williams, Cuvelier, & Berset, 1995) and by ferric ion reducing antioxidant power (FRAP) assay (Benzie & Strain, 1996).

For the DPPH assay, 0.5 mL of DPPH solution (140 µmol L⁻¹ in 50% methanol) was combined with an equal volume of sample at a concentration of 5–100 µg mL⁻¹ and left to stand at room temperature for 30 min. After this time absorbance (A) was measured on a Pharmacia Biotech Ultraspec 4000 UV/Visible spectrophotometer (Amersham Pharmacia Biotech, Little Chalfont, UK) at 517 nm, including 50% methanol as control, and % inhibition (I) was determined as

$$I = \frac{(A - A_{\text{blank}})_{\text{control}} - (A - A_{\text{blank}})_{\text{sample}}}{(A - A_{\text{blank}})_{\text{control}}} \times 100$$

with IC₅₀ defined as the concentration which resulted in 50% inhibition.

The FRAP assay measures reduction of the Fe³⁺/tripyrindyl-s-triazine complex (TPTZ) to the blue ferrous form and was performed as follows: acetate buffer (300 mmol L⁻¹, pH 3.6), TPTZ (10 mmol L⁻¹ in 40 mmol L⁻¹ HCl) and ferric chloride (20 mmol L⁻¹) were mixed in the ratio of 10: 1: 1 to obtain the FRAP reagent. Sample volumes of 125 µL at a concentration of 100 µg mL⁻¹ were added to 1 ml FRAP reagent and after 10 min absorbance was measured at 593 nm. A ferric chloride calibration curve ranging from 100 to 1000 µmol L⁻¹ was prepared to estimate Fe²⁺ concentration.

2.7. Statistical analysis

Quantitative analyses were carried out in triplicate and results are presented as mean ± standard error of the mean. Where appropriate results were statistically analysed by one-way ANOVA and significant differences were determined by the Bonferroni post hoc test, GenStat 14th edition.

3. Results and discussion

3.1. Phenolic profile

Fig. 1 shows the profile of soluble polyphenols extracted from *Allanblackia floribunda* seeds which showed little variation between years. Analysis by HPLC-PDA-ESI/MSⁿ revealed six abundant and a range of minor compounds which were tentatively identified based on their UV/vis absorbance and mass spectral characteristics summarised in Tables 1 and 2 and Fig. 2. The compounds which were identified all belonged to the flavonoids, with biflavonoids as the main class. The main compounds were collected individually from peaks 1 to 6 (Fig. 1) and purified as pale yellow to orange crystalline powders. Molecular weights (*M_r*) of the main compounds are based on accurate mass measurements in both negative and positive ionisation mode, whilst *M_r* of less abundant compounds are nominal mass based on general MS scans

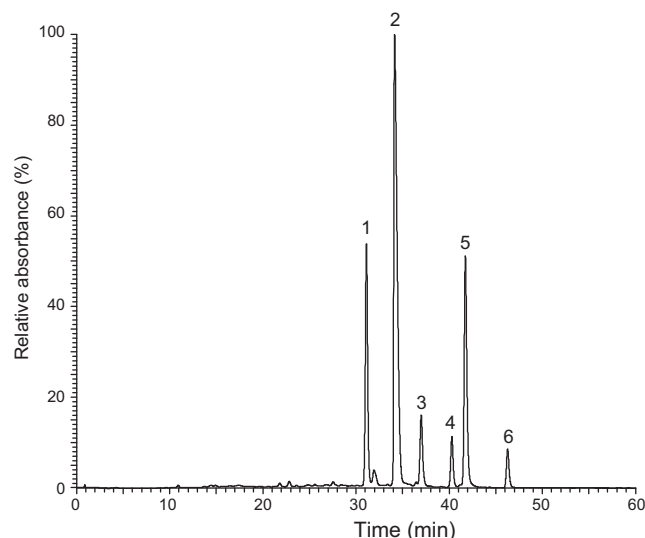


Fig. 1. HPLC chromatogram of soluble secondary metabolites from *Allanblackia floribunda* seeds detected by photodiode array at 280 nm. Peak numbering refers to the six predominant compounds.

in both ionisation modes. MSⁿ analyses were carried out in negative ionisation mode unless stated otherwise.

3.1.1. Identification of biflavonoids

Based on accurate mass data (Table 2) compound 5 was assigned the elemental composition C₃₀H₂₀O₁₁ with *M_r* 556.1005 Da. The corresponding UV/vis spectrum showed absorbance maxima at 275(sh), 290 and 348 nm, a combination of the absorbance spectrum typical of a flavone (with two major absorption bands in the region of 300–380 nm for band I and 240–280 nm for band II; Mabry, Markham, & Thomas, 1970) with that of a flavanone (with predominant absorption band II and a much reduced band I) and in close agreement with the absorbance characteristics reported for the biflavonoid morelloflavone, a naringenin-luteolin conjugate (Herbin, Jackson, Locksley, Scheinmann, & Wolstenholme, 1970). MS² fragmentation of the parent ion at *m/z* 555 in negative mode resulted in loss of 126 Da, yielding one main product ion with *m/z* 429. This can be attributed to cleavage of ring C (Fig. 3a), typical of the fragmentation of flavonoids (Cuyckens & Claeys, 2004; Fabre, Rustan, de Hoffmann, & Quetin-Leclercq, 2001), here breaking bonds 1 and 4 and producing fragment ion [M–H–^{1,4}A]⁻ with the remainder of the upper moiety of the dimer still attached to the lower moiety. A low intensity fragment with *m/z* 403 was also observed. This corresponds to loss of 152 Da and stems from cleavage of bonds 1 and 3 of ring C. Fragmentation of naringenin standard in negative mode showed cleavage of corresponding bonds, yielding ion ^{1,3}A⁻ with *m/z* 151 as base peak and ion ^{1,3}B⁻ with *m/z* 119 among the less intense fragment ions. In the dimer, ring B is still attached to the lower moiety, thus resulting in fragment [M–H–^{1,3}A]⁻. The minor fragments also included ions at *m/z* 449 (loss of 106 Da) and *m/z* 461 (loss of 94 Da, [M–H–B]⁻), and corresponding losses were also seen when fragmenting naringenin standard.

With normalised collision energy (CE) set to default (35%), MS³ fragmentation of the predominant MS² ion at *m/z* 429 occurred only to a small extent, an observation also made with luteolin standard. Raising CE to 70% resulted in the neutral loss of a number of small molecules such as CO, CO₂ and H₂O, also typical of flavonoid fragmentation (Cuyckens & Claeys, 2004; Fabre et al., 2001). A further MS³ product ion with *m/z* 295 (loss of 134 Da) can be

Table 1
Polyphenolic compounds detected in *Allanblackia floribunda* seed extract by HPLC-PDA-ESI/MSⁿ.

<i>t_R</i> (min)	<i>λ</i> _{max} (nm)	<i>M_r</i>	MS ² fragments in –ve mode unless stated otherwise (base peak in bold)	MS ² fragments of MS ² base peak (base peak in bold)	Additional diagnostic MS ⁿ fragments (parent ion in bracket)	Tentative ID (Fig. 1 peak No. in brackets)
10.9	279	578	287, 289, 407, 425 , 451, 559		179, 205, 245 (289)	Epicatechin dimer
13.7	279	290	179, 205, 245			Epicatechin ¹
14.4	279	866	407, 425, 451, 575, 577, 695 , 713, 739		287, 289 (577) 179, 205, 245 (289)	Epicatechin trimer
14.9	279	1154	575, 577, 739, 863, 865 , 983, 1001, 1027, 1135		245, 287, 289 (577)	Epicatechin tetramer
21.8	nd	736	429, 447 , 573	269, 385, 403, 419, 429	419, 421, 429, 447, 467, 479 (573)	Hexoside of biflavonoid at <i>t_R</i> 30.9 min
22.8	nd	442	139, 143, 161, 179, 205, 217, 235, 330 / 331 , 397			Unidentified compound
23.6	nd	578 624	287, 289, 407, 425 , 451, 559 285 , 327, 489		179, 205, 245 (289)	Epicatechin dimer Luteolin-hexosyl-glucuronate
24.9	nd	578	287, 463 (+ve mode) 293, 311, 341, 413 , 457			Apigenin-2''-O-rhamnosyl-C-hexoside
25.6	nd	462	285			Luteolin-glucuronate
26.8	nd	1008 608	431, 557, 719, 837, 845, 855 , 881 269	431, 649, 675, 811, 837	431, 557, 593 (719)	Gb-2a conjugate
27.5	nd	880	271, 447 (+ve mode) 717	403, 429, 493, 537, 555, 565 , 591	295 (429)	Apigenin-hexosyl-glucuronate
28.4	nd	610	301			Morelloflavone-dihexoside
28.4	nd	1006	303, 449, 465 (+ve mode) 717, 809, 835, 853 , 879	809, 835	429, 493, 537, 555, 565, 591, 623 (717)	Quercetin-rhamnosyl-hexoside
29.6	nd	756	285 , 575, 593			Morelloflavone conjugate
30.2	nd	1008	287, 449, 595, 611 (+ve mode) 431, 557, 719, 837, 845, 855 , 881	431, 649, 675, 811, 837	431, 557, 593 (719)	Luteolin-rhamnosyl-dihexoside
30.9	nd	574	447	269, 295, 325, 403, 419 , 429	421, 467, 479 (573)	Gb-2a conjugate
31.1	288	720	431 , 557, 593	269, 295 , 413, 321	405, 431, 451, 463 (557) 625 (719)	Unidentified biflavonoid
31.3	nd	578	269			Gb-2a-glucoside (1)
31.9	nd	594	271, 433 (+ve mode) 285			Apigenin-rhamnosyl-hexoside
31.9	nd	594	287, 449 (+ve mode) 285			Kaempferol-rhamnosyl-hexoside (leading edge)
34.1	274, 290, 351	718	287, 449 (+ve mode) 429, 493, 537, 555, 565 , 591, 623	403 , 445	403, 449, 461 (555) 295 (429) 623 (717)	Luteolin-rhamnosyl-hexoside (trailing edge)
36.4	nd	704	389, 415 , 447, 523, 541, 551, 577, 609	295 , 321, 269	389, 415, 435, 447 (541)	Morelloflavone-glucoside (2)
37.0	293	558	431	269, 295 , 321, 413	405, 451, 463 (557)	Dinaringenin-hexoside
37.2	nd	610	463 , 301			Gb-2a (3)
38.0	nd	286	151, 175, 197, 199, 201, 213, 217, 241 , 243, 257, 267			Quercetin-rhamnosyl-hexoside
40.3	274, 290, 330	702	413, 433, 445, 521, 539 , 575	387, 413 , 433, 445	295 (413)	Luteolin ¹
41.2	nd	594	285 , 447			Volkensiflavone-glucoside (4)
41.6	nd	542	389, 415 , 435, 447	269, 295 , 309, 321		Kaempferol-rhamnosyl-hexoside
41.7	275(sh), 290, 348	556	429	295, 357, 385, 401	403, 449, 461 (555)	Dinaringenin
46.3	274(sh), 290, 328	540	387, 413 , 433, 445	293, 295, 369, 385		Morelloflavone (5)
						Volkensiflavone (6)

nd: not detected, either due to low absorbance or coeluting compounds.

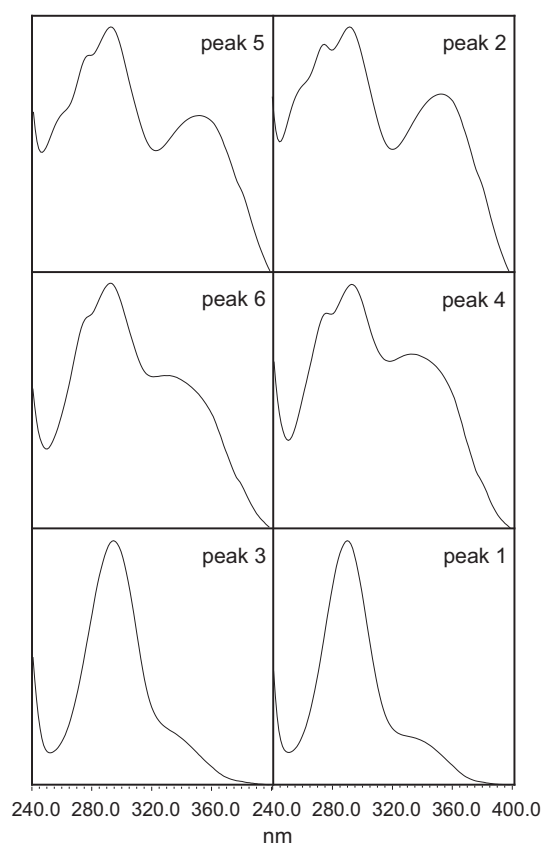
¹ Identified by direct comparison with reference compounds.

accounted for by cleavage of bonds 1 and 3 of the heterocycle of the luteolin moiety, producing $[M-H-^{1,4}A-^{1,3}E]^-$. Cleavage of corresponding bonds was also observed in luteolin standard, resulting in a fragment with *m/z* 151. Overall, fragmentation of the flavonoid dimer mirrored a combination of the fragmentation patterns

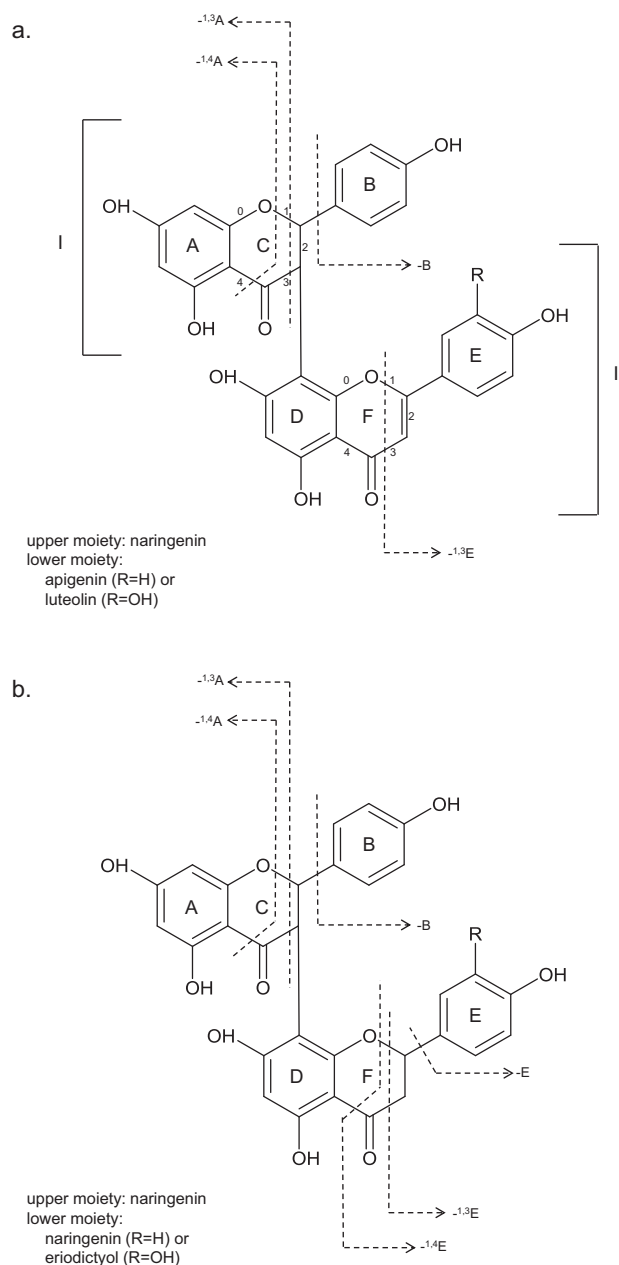
observed for naringenin and luteolin standards. The data presented here is in close agreement with mass spectral data reported by Carrillo-Hormaza et al. (2016) and supports the tentative identification of compound 5 as morelloflavone (naringenin-I,3-II,8-luteolin).

Table 2Accurate mass and content of predominant biflavonoids in seeds of *Allanblackia floribunda*.

Fig. 1 peak no. (compound ID)	t_R (min)	Measured mass $[M-H]^-$	Calculated mass $[M-H]^-$	Mass difference (ppm)	Measured mass $[M-H]^+$	Calculated mass $[M-H]^+$	Mass difference (ppm)	Content ($\mu\text{g mg}^{-1}$ seed powder)
1 (Gb-2a-glucoside)	30.1	719.1615	719.1618	0.42	721.1774	721.1763	1.53	16.73 ± 0.906
2 (morelloflavone-glucoside)	33.7	717.1459	717.1461	0.28	719.1616	719.1607	1.25	58.26 ± 3.872
3 (Gb-2a)	35.4	557.1087	557.1089	0.36	559.1238	559.1235	0.54	2.75 ± 0.146
4 (volkensiflavone-glucoside)	39.2	701.1514	701.1512	0.29	703.1666	703.1657	1.28	4.03 ± 0.297
5 (morelloflavone)	40.3	555.0929	555.0933	0.72	557.1079	557.1078	0.18	16.54 ± 0.935
6 (volkensiflavone)	44.5	539.0984	539.0984	0.00	541.1134	541.1129	0.92	2.02 ± 0.136

The content of individual compounds is expressed as mean $\pm$ standard error of the mean ($n = 9$).**Fig. 2.** UV/vis spectra of biflavonoids from *Allanblackia floribunda* seeds.

The UV/vis spectrum of peak 6 showed absorbance maxima at 274, 290 and 328 nm (Fig. 2), similar to those of peak 5 but with absorbance band I at a lower wavelength, consistent with the absorbance characteristics reported for volkensiflavone, a naringenin-apigenin conjugate (Herbin et al., 1970). Accurate mass measurements (Table 2) suggest the elemental composition $C_{30}H_{20}O_{10}$ with M_r 540.1056 Da for this compound. The fragmentation pattern in negative mode was also similar to that of morelloflavone (Table 1, Fig. 3a) and yielded product ions consistent with a naringenin moiety, namely $[M-H-^{1,4}A]^-$ with m/z 413, $[M-H-B]^-$ with m/z 445, $[M-H-106]^-$ with m/z 433 and $[M-H-^{1,3}A]^-$ with m/z 387, whereas fragmentation of the MS^2 base peak at m/z 413 with raised CE resulted in losses similar to those observed with apigenin standard, yielding product ions at m/z 295 (loss of

**Fig. 3.** Structures and fragmentation patterns of (a) flavanone-flavone dimers (volkensiflavone, morelloflavone) and (b) flavanone-flavanone dimers (Gb-2a, dinaringenin) extracted from *Allanblackia floribunda* seeds.

118 Da, $[M-H-^{1,4}A-^{1,3}E]^-$ and 293 (loss of 120 Da). The findings for compound 6 are in close agreement with the fragmentation pattern reported for volkensiflavone (naringenin-1,3-II,8-apigenin) by Carrillo-Hormaza et al. (2016).

The UV/vis spectrum of peak 3 (Fig. 2) showed a band II absorbance maximum at 293 nm with band I reduced to a small shoulder, a characteristic of flavanones (Mabry et al., 1970). Based on accurate mass measurements (Table 2) the molecular formula $C_{30}H_{22}O_{11}$ with M_r 558.1162 Da was assigned to this compound. Fragmentation in negative ionisation mode followed a pattern very similar to that described above (Table 1, Fig. 3b). MS^2 analysis of the parent ion at m/z 557 yielded a predominant product with m/z 431 (loss of 126 Da, $[M-H-^{1,4}A]^-$) as well as low abundance ions consistent with the loss of 152 Da (at m/z 405), 106 Da (at m/z 451) and 94 Da (at m/z 463) from a naringenin moiety. MS^3 analysis of the fragment at m/z 431 produced a base peak with m/z 295 (loss of 136 Da, $[M-H-^{1,4}A-^{1,3}E]^-$) and less intense ions at m/z 269 (loss of 162 Da, $[M-H-^{1,4}A-^{1,4}E]^-$; Fabre et al., 2001) and 321 (loss of 110 Da, $[M-H-^{1,4}A-E]^-$), in agreement with the fragmentation pattern observed for eriodictyol standard. Based on the data presented here compound 3 was tentatively identified as Gb-2a (naringenin-1,3-II,8-eriodictyol), a conclusion supported by the mass spectral data of Carrillo-Hormaza et al. (2016).

Peaks 1, 2 and 4 had UV/vis spectra which were almost identical to those of peaks 3, 5 and 6, respectively (Fig. 2) but with M_r 162 Da bigger than the corresponding biflavonoids (Tables 1 and 2), and they were also more polar as shown by the earlier retention times. The MS^2 spectra of the compound 1, 2 and 4 parent ions included products $[M-H-162]^-$, indicating the loss of an *O*-linked hexose (Vukics & Guttman, 2010), while other product ions resulted from fragmentation of the biflavonoid core with or without simultaneous loss of the sugar moiety. In addition, MS^n fragmentation of the presumed aglycone fragments at m/z 557, 555 and 539, respectively, followed the common pattern seen for compounds 3, 5 and 6, thus confirming biflavonoids as core molecules of the glycosides, with neutral loss of 126 Da producing the diagnostic fragments $[M-H-162-^{1,4}A]^-$, while subsequent MS^n analysis included product ions $[M-H-162-^{1,4}A-^{1,3}E]^-$ with m/z 295.

HPAEC analysis of the purified, hydrolysed compounds revealed glucose as the sugar moiety of all three glycosides. Although it is not possible to determine the precise position of the glucose part with the methods employed here, fragmentation data for compounds 1 and 2, which have different substitution patterns on rings B and E, suggests that the glucose may be attached to one of the hydroxyl groups of the lower flavonoid unit. In the MS^2 spectra of both compounds there were product ions characteristic for the fragmentation of the naringenin moiety with the sugar unit still attached, namely $[M-H-^{1,4}A]^-$ (loss of 126 Da), $[M-H-^{1,3}A]^-$ (loss of 152 Da) and low intensity ions indicating the loss of ring B ($[M-H-B]^-$, loss of 94 Da).

Analysis of the smaller peaks revealed further compounds of this type. In particular, a compound with M_r 542 Da eluting at 41.6 min followed a similar fragmentation pattern as described above (Fig. 3b) and was tentatively identified as dinaringenin, and a hexoside of this with M_r 704 Da was found at t_R 36.4 min. A further biflavonoid with M_r 574 Da eluted at t_R 30.9 min and had a fragmentation pattern consistent with naringenin as upper and an unidentified flavonoid with M_r 304 Da as lower unit, and a hexoside of this with M_r 736 was seen at 21.8 min. In addition to morelloflavone and morelloflavone-glucoside, a compound with M_r 880 Da eluting at 27.5 min showed a fragmentation pattern consistent with morelloflavone-dihexoside. Some larger related molecules were also detected amongst the low abundance compounds. For example, fragmentation of a compound with M_r 1006 Da at t_R 28.4 min produced a fragment ion with m/z 717. Further MS^3 analysis of this fragment was consistent with

morelloflavone-hexoside, and the compound was tentatively identified as morelloflavone conjugate. Similarly, MS^2 analysis of two compounds with M_r 1008, eluting at 25.6 and 30.2 min, produced fragments with m/z 719, and MS^3 analysis of this ion product was consistent with the fragmentation pattern of Gb-2a-hexoside.

Both morelloflavone and volkensiflavone have previously been reported in *A. floribunda* heartwood (Locksley & Murray, 1971) and stem bark (Brusotti et al., 2016) as well as in various tissues of related genera such as fruits, leaves and heartwood of *Garcinia* (Carrillo-Hormaza et al., 2016; Herbin et al., 1970; Stark et al., 2015; Yang et al., 2010). Whilst Gb-2a is also well documented in *Clusiaceae* (Carrillo-Hormaza et al., 2016; Herbin et al., 1970; Stark et al., 2015), this is to our knowledge the first report of Gb-2a in *Allanblackia*.

3.1.2. Identification of other flavonoids

In addition to the biflavonoids discussed in Section 3.1.1, a range of low abundance compounds from different flavonoid classes were also detected (Table 1). Epicatechin and luteolin were identified by direct comparison with analytical standards. Due to the absence of catechin in the extract, the flavan-3-ol oligomers found at several retention times were assumed to consist of epicatechin units. There was also a range of flavone and flavonol glycosides which were tentatively identified by their fragmentation patterns following the principles outlined by Vukics and Guttman (2010) and Ferreres, Gil-Izquierdo, Andrade, Valentão, and Tomás-Barberán (2007), whilst their flavonoid cores were confirmed by MS^n analyses and acid hydrolysis. Interestingly, *A. floribunda* seed extract did not contain detectable amounts of xanthenes or benzophenones which were previously reported in other tissues of *A. floribunda* and other species of the *Clusiaceae* family (Azebaze et al., 2009; Fuller et al., 1999; Locksley & Murray, 1971; Nkengfack et al., 2002; Yang et al., 2010).

3.2. Characterisation of biflavonoids

3.2.1. Quantification

The six main biflavonoids were quantitated in $\mu\text{g mg}^{-1}$ seed powder using response factors based on HPLC standard curves obtained with analytical standards as described in Section 2.4. The total biflavonoid content was high, constituting approximately 10% of dried seed powder in comparison to seeds of the related species *Garcinia madruno* where Carrillo-Hormaza et al. (2016) reported a total biflavonoid content of less than 2%. Morelloflavone and its glucoside were the main biflavonoids present in *A. floribunda* seeds (Table 2). Similarly, Locksley and Murray (1971) reported morelloflavone as the predominant metabolite in *A. floribunda* heartwood.

3.2.2. Antioxidant activity

The antioxidant activities of the six main biflavonoids were analysed by the DPPH-radical scavenging and the ferric reducing antioxidant power (FRAP) assays. All six compounds demonstrated antioxidant activity which differed significantly ($P \leq 0.05$) and showed a similar ranking of activity with both assays apart from Gb-2a aglycone and volkensiflavone-glucoside whose order was reversed with the DPPH compared with the FRAP assay (Table 3). The highest activities were observed with Gb-2a and morelloflavone-glucosides and the lowest activities with volkensiflavone aglycone, volkensiflavone-glucoside and Gb-2a aglycone. Morelloflavone-glucoside and aglycone and Gb-2a-glucoside showed radical scavenging activities comparable with ascorbic acid (IC_{50} values 16.87, 21.26, 18.38 and 19.36 $\mu\text{g ml}^{-1}$, respectively). Radical scavenging and FRAP activities observed with morelloflavone aglycone and glucoside also reflect results reported by Kuete et al. (2011) and Carrillo-Hormaza et al. (2016) where the

Table 3Antioxidant capacity of biflavonoids from *A. floribunda* seeds.

Fig. 1 peak no. (compound ID)	FRAP ($\mu\text{mol Fe}^{2+} \text{mg}^{-1}$)	DPPH IC ₅₀ ($\mu\text{g mL}^{-1}$)
1 (Gb-2a-glucoside)	13.67 $\pm$ 0.281	18.38 $\pm$ 0.112
2 (morelloflavone-glucoside)	15.35 $\pm$ 0.082	16.87 $\pm$ 0.337
3 (Gb-2a)	3.82 $\pm$ 0.350	27.20 $\pm$ 0.292
4 (volkensiflavone-glucoside)	2.61 $\pm$ 0.012	26.42 $\pm$ 0.006
5 (morelloflavone)	6.33 $\pm$ 0.016	21.26 $\pm$ 0.059
6 (volkensiflavone)	2.12 $\pm$ 0.272	33.92 $\pm$ 0.382
Ascorbic acid	17.91 $\pm$ 0.243	19.36 $\pm$ 0.036

Results are expressed as mean $\pm$ standard error of the mean ($n = 3$).

aglycone showed lower activity than the glucoside. This pattern was observed with all biflavonoids tested here, indicating that glycosylation enhances antioxidant activity. The higher IC₅₀ values reported by Kuete and co-workers may be due to the higher DPPH concentration used in their assays.

Burda and Oleszek (2001) observed a relationship between certain structural features of flavonoids and antioxidant behaviour. Their studies demonstrated that a hydroxyl group in the *para* position on the B-ring is essential for activity and that this activity is enhanced by a second hydroxyl on the B-ring in the *ortho* position and a double bond between C2 and C3 on the C ring. These findings are consistent with aglycone activities observed in the current study. Overall, the data presented here demonstrates that *Allanblackia* seeds are an abundant source of highly active antioxidant phenolic components in common with seeds of the fruits of other tropical species including jabuticaba (*Myrciaria cauliflora*; Hacke et al., 2016) and guaraná (*Paullinia cupana*; Majhenič, Škerget, & Knez, 2007; Marques et al., 2016).

4. Conclusion

The comprehensive profile of soluble phenolics presented here confirms biflavonoids as the main phenolic compound class present and as major constituents of *A. floribunda* seeds. Due to their antioxidant activity and reported therapeutic properties, these compounds are of increasing interest to the pharmaceutical industry. With oil from *A. floribunda* seeds attracting attention from the food sector as a potential ingredient of margarine and other products (Cernansky, 2015), they could provide an excellent source of biflavonoids as a by-product.

Conflict of interest

The authors declare that they have no conflict of interest.

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