Two New Antimicrobial Dimeric Compounds: Febrifuquinone, a Vismione-Anthraquinone Coupled Pigment and Adamabianthrone, from two *Psorospermum* Species

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Febrifuquinone (**1**), a new vismione-anthraquinone coupled pigment and a new bianthrone named adamabianthrone (**2**), were isolated respectively, from the roots of *Psorospermum febrifugum* and from the bark of *Psorospermum adamauiense* along with eight known compounds including: two bianthrone [(bianthrone A₁ (**3**) and bianthrone A_{2b}), one vismione [(vismione D (**4**)), one anthrone (3-geranyloxyemodin anthrone) and four anthraquinones [(1,8-dihydroxy-3-isoprenyloxy-6-methylanthraquinone, emodin (**5**), 3-geranyloxy-1,8-dihydroxy-6-methylanthraquinone and 2-geranyl-1,8-dihydroxy-6-methylanthraquinone]. Their structures were determined using modern spectroscopic methods including one and two dimensional-NMR techniques as well as MS. Compounds **1** and **2** showed significant antimicrobial activities against a wide range of bacteria and fungi.

Key words *Psorospermum febrifugum*; *Psorospermum adamauiense*; quinone; antimicrobial test

The genus *Psorospermum*, belonging to the tribe Vismieae (family Guttiferae, subfamily Hypericoideae), comprises 55 species found in tropical Africa and Madagascar.¹⁾ *Psorospermum* species are shrub or small trees that are largely used in the African folk medicine as febrifugal, antidote against poison and purgative. They are also used as a remedy for the treatment of leprosy, skin diseases (such as dermatitis, scabies and eczemas) and subcutaneous wounds.^{2–4)} Previous chemical studies of different species of this genus have led to the isolation of vismiones,^{5–8)} anthraquinones,^{5,9)} anthrones and bianthrone,^{7,10–12)} xanthonolignoids¹³⁾ and xanthenes.^{14–16)} On the other hand, the ethanolic extract of *Psorospermum febrifugum* was found to exhibit significant *in vivo* cytotoxicity and antitumor activity against the P-388 lymphocytic leukemia (3PS) in mice and *in vitro*, activity against a cell culture derived from human epidermoid carcinoma of the nasopharynx (9KB).¹⁵⁾ Further biological activity guided investigation, was undertaken¹⁴⁾ and led to the isolation of the first dihydrofuranoxanthone named psorospermin, which exhibited significant cytotoxicity and antitumor activity.¹⁶⁾ In addition to the other bioactive classes of secondary metabolites. Consequently, prompted by this result, the investigation on two Cameroonian *Psorospermum* species: *Psorospermum febrifugum* and *Psorospermum adamauiense* was carried out. On *Psorospermum adamauiense*, no phytochemical study has yet been reported to the best of our knowledge. The present work deals with the isolation, structure elucidation of secondary metabolites as well as the evaluation of their antimicrobial activities.

Results and Discussion

Air dried and powdered roots of *P. febrifugum* and bark of *P. adamauiense* were extracted at room temperature with a mixture of methanol/methylenechloride (1 : 1), respectively. The residue, obtained from each plant after evaporation of

the solvents, was respectively subjected to repeated silica gel column chromatography (see Experimental) and eluted with a mixture of hexane/AcOEt of increasing polarity. From the roots extract of *P. febrifugum*, four compounds were obtained among which, one new derivative named febrifuquinone (**1**) and three known compounds: bianthrone A₁ (**3**), vismione D (**4**) and 3-geranyloxyemodin anthrone. While, from the bark of *P. adamauiense*, one new compound named adamabianthrone (**2**), was isolated along with six known others:

3-Geranyloxyemodin anthrone, 1,8-dihydroxy-3-isoprenyloxy-6-methylanthraquinone, bianthrone A_{2b}, emodin (**5**), 3-geranyloxy-1,8-dihydroxy-6-methylanthraquinone and 2-geranyl-1,8-dihydroxy-6-methylanthraquinone. The structures of the known compounds were determined from their ¹H-, ¹³C-NMR and mass spectral data and confirmed by comparison of these data and their physical constants with those published in the literature.^{5–18)}

Compound **1**, named febrifuquinone, mp 132–133 °C, was isolated as red powder. It reacted positively to the FeCl₃ reagent suggesting its phenolic nature. Its “high resolution electrospray ionization-time-of-flight (HR-ESI-TOF)” MS showed a protonated molecule [M+H]⁺ at *m/z* 679.2514 (Calcd for C₄₀H₃₉O₁₀: 679.2532) corresponding to the molecular formula C₄₀H₃₈O₁₀ with 22 degrees of unsaturation.

The broad band proton decoupled ¹³C-NMR spectrum (DMSO-*d*₆) of compound **1**, revealed the presence of 40 carbon signals which were sorted by attached proton test (APT) and heteronuclear single quantum coherence (HSQC) techniques as five methyls, four methylenes, nine methines and twenty-two quaternary carbon atoms among which, three carbonyl groups at δ_C 181.2, δ_C 189.6 and δ_C 203.6 (see Table 1).

The UV spectrum of compound **1** showed absorption bands at 235, 278, 320, 390 nm and 408, 435 nm characteristic of vismione and 1,8-dihydroxyanthraquinone chromo-

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phores.^{5,19} These assumptions were supported by the ^1H -NMR spectrum ($\text{DMSO}-d_6$) of compound **1** which showed two singlets, of one proton each, at δ_{H} 9.78 and δ_{H} 15.79 exchangeable with D_2O and assigned to two hydroxyl groups, located, respectively at C-1 and C-9 of a vismione moiety²⁰ and two other singlets of one proton each at δ_{H} 11.99 and δ_{H} 12.06 also exchangeable with D_2O and corresponding to two chelated hydroxyl groups at C-1 and C-8 of an anthraquinone moiety.¹⁹ These data, combined with the protonated molecule $[\text{M}+\text{H}]^+$ at m/z 679.2514, suggested that compound **1** is a dimer containing in its skeleton, a vismione and an anthraquinone moieties.

This assumption was confirmed by the ^1H -NMR spectrum ($\text{DMSO}-d_6$) of **1** which showed two sets of signals. The multiplicities and chemical shifts of the first set of signals were very close to those observed in the ^1H -NMR spectrum of vismione **D**⁵) (**4**) (see Table 1), while those of the second set, attributed to fragment “B,” were quite similar to signals appeared in the ^1H -NMR spectrum of emodin (**5**)¹⁷) (see also Table 1).

The junction between the two moieties was established using heteronuclear multiple bond correlation (HMBC) techniques (Fig. 1) which showed, ^2J , ^3J correlations peaks between the proton singlet H-5 (δ_{H} 2.96) belonging to vismione D moiety and C-5' (δ_{C} 137.3), C-6' (δ_{C} 148.1) and C-10'a (δ_{C} 108.3) belonging to emodin moiety. This result clearly indicated that the junction between the two moieties was through C-5 and C-5' linkage. On the other hand, the geranyloxy moiety was located at C-3 according to the cross peak observed on its nuclear Overhauser enhancement and exchange spectroscopy (NOESY) spectrum (Fig. 1) between the methylene proton at δ_{H} 4.62 and H-2 (δ_{H} 6.38) and H-4 (δ_{H} 6.70).

This was further supported by the HR-ESI-TOF-MS/MS of the protonated molecule which showed the presence of two important fragment at m/z 409 $[\text{M}-\text{C}_{15}\text{H}_9\text{O}_5]^+$ (of vis-

Fig. 1. Structure, Key HMBC and NOESY Correlations of Compound **1**

mione D fragment) and m/z 269 $[\text{M}-\text{C}_{25}\text{H}_{29}\text{O}_5]^+$ (of emodin fragment) resulting from the breaking of the C-5, C-5' bond. On the basis of the above results, compound **1** was identified as 1-(6-((*E*)-3,7-dimethylocta-2,6-dienyloxy)-1,2,3,4-tetrahydro-3,8,9-trihydroxy-3-methyl-1-oxoanthracen-4-yl)-4,5,7-trihydroxy-2-methylantracene-9,10-dione named febrifuquinone (see Fig. 1).

Adamabianthrone (**2**), mp 152–154 °C, was obtained from the bark of *P. adamauense* as a yellow powder in a mixture of hexane/ethylacetate (1 : 9). It reacted positively to the FeCl_3 reagent and magnesium acetate, indicating the presence in its structure of phenolic hydroxyl groups and 1,8-dihydroxyanthrone skeleton, respectively. The molecular formula of compound **2** was determined as $\text{C}_{45}\text{H}_{46}\text{O}_8$ from its “high resolution ESI-TOF (HR-ESI-TOF)” MS, which showed a protonated molecule $[\text{M}+\text{H}]^+$ at m/z 715.3231. The UV and ^1H -NMR spectra of **2** were very similar to those recorded for bianthrone **A**₁¹⁰) (**3**).

The major difference between the two compounds appeared in the ^1H -NMR spectrum ($\text{DMSO}-d_6$) of compound **2** which, showed one set of signals [$(\delta_{\text{H}}$ 1.60 (3H, s), δ_{H} 1.62 (3H, s) and δ_{H} 1.81 (3H, s); δ_{H} 2.05 (4H, m); δ_{H} 4.64 (2H, d, $J=6.0\text{ Hz}$)/ δ_{C} 68.2; δ_{H} 5.30 (1H, t, $J=8\text{ Hz}$) and δ_{H} 5.46 (1H, t, $J=6.0\text{ Hz}$)/ δ_{C} 119.3], corresponding to one *O*-geranyloxy group (instead of two, as in bianthrone **A**₁ (**3**)) and an-

Table 1. ^1H -NMR Data of Compounds **1**, **4** and **5** and ^{13}C -NMR Data of Compound **1**

Position	1 DMSO- d_6		4 ¹⁶⁾ CDCl_3	5 ¹⁷⁾ Acetone- d_6
	δ_{H} (mult. J Hz)	δ_{C} (APT)	δ_{H} (mult. J Hz)	δ_{H} (mult. J Hz)
1-OH	9.78 (1H, s)	158.7 (s)	9.79 (1H, s) ^{c)}	—
2	6.38 (1H, d, 2.2 Hz)	107.8 (d)	6.49 (1H, d, 2.3 Hz)	—
3	—	161.6 (s)	—	—
4	6.70 (1H, d, 2.2 Hz)	100.7 (d)	6.44 (1H, d, 2.3 Hz)	—
4a	—	135.5 (s)	—	—
5	2.96 (1H, s)	42.4 (d)	3.03 (2H, br s)	—
6-OH	4.30 (1H, s)	69.4 (s)	—	—
7	2.65 (1H, d, 17 Hz, H_{7a}) ^{a)} 2.88 (1H, d, 17 Hz, H_{7b}) ^{a)}	50.7 (t)	2.82 (2H, br s)	—
8	—	203.6 (s)	—	—
8a	—	108.3 (s)	—	—
9-OH	15.79 (1H, s)	164.3 (s)	16.12 (1H, s) ^{c)}	—
9a	—	108.3 (s)	—	—
10	6.89 (1H, s)	116.7 (d)	6.83 (1H, s)	—
10a	—	108.9 (s)	—	—
11	1.23 (3H, s)	28.8 (q)	1.44 (3H, s)	—
12	4.62 (2H, d, 6.4 Hz)	64.6 (t)	4.62 (2H, d, 6.6 Hz)	—
13	5.45 (1H, t, 6.4 Hz)	119.2 (d)	5.50 (1H, t, 6.6 Hz)	—
14	—	140.7 (s)	—	—
15	2.05 (4H, m)	38.8 (t)	2.11 (4H, m)	—
16	—	25.7 (t)	—	—
17	5.06 (1H, t, 7.5 Hz)	123.6 (d)	5.30 (1H, t, 8 Hz)	—
18	—	130.9 (s)	—	—
19	1.72 (3H, s)	16.3 (q)	1.75 (3H, s)	—
20	1.56 (3H, s) ^{b)}	17.4 (q)	1.69 (3H, s) ^{a)}	—
21	1.60 (3H, s) ^{b)}	25.3 (q)	1.60 (3H, s) ^{a)}	—
1'-OH	12.06 (1H, s) ^{a,c)}	164.6 (s)	—	12.15 (1H, s) ^{a,c)}
2'	6.57 (1H, d, 2.5 Hz)	108.6 (d)	—	6.66 (1H, d, 2.5 Hz)
3'-OH	11.33 (1H, s) ^{c)}	165.4 (s)	—	—
4'	7.11 (1H, d, 2.5 Hz)	120.4 (d)	—	7.25 (1H, d, 2.5 Hz)
4'a	—	135.0 (s)	—	—
5'	—	137.3 (s)	—	7.58 (1H, s)
6'	—	148.1 (s)	—	—
7'	6.77 (1H, s)	124.0 (d)	—	7.15 (1H, s)
8'-OH	11.99 (1H, s) ^{a,c)}	161.3 (s)	—	12.05 (1H, s) ^{a,c)}
8'a	—	113.3 (s)	—	—
9'	—	189.6 (s)	—	—
9'a	—	132.7 (s)	—	—
10'	—	181.2 (s)	—	—
10'a	—	108.3 (s)	—	—
11'	2.40 (3H, s)	21.4 (q)	—	2.47 (3H, s)

a, b) Assignments bearing the same superscript in the same column may be reversed. *c*) Exchangeable in D_2O . ^1H -NMR was measured at 400 MHz for **1**, at 200 MHz for **4** and at 300 MHz for **5**. ^{13}C -NMR of **1** was measured at 100 MHz.

other set of signals [$(\delta_{\text{H}} 3.35$ (2H, d, $J=7.9$ Hz)/ $\delta_{\text{C}} 22.7$, $\delta_{\text{H}} 5.10$ (1H, t, $J=7.9$ Hz)/ $\delta_{\text{C}} 123.9$ and $\delta_{\text{H}} 1.86$ (3H, s)], establishing the presence of a γ,γ -dimethylallyl moiety, in addition to a phenolic hydroxyl proton at $\delta_{\text{H}} 9.54$ exchangeable with D_2O (see Table 2). The absence in the ^1H -NMR spectrum of adamabianthrone (**2**) of H-2' signal (which appeared at $\delta_{\text{H}} 6.35$ (1H, d, $J=2.2$ Hz) in bianthrone **A**₁ (**3**)), indicated that the γ,γ -dimethylallyl group was located at C-2'. This structure was further supported by analysis of the HMBC spectrum of compound **2** (Fig. 2) which showed cross peaks between the proton at $\delta_{\text{H}} 3.35$ and carbons C-1' ($\delta_{\text{C}} 161.8$), C-2' ($\delta_{\text{C}} 136.0$) and C-3' ($\delta_{\text{C}} 162.6$) indicating that the γ,γ -dimethylallyl group is located at C-2'. On the other hand, the geranyloxy moiety was located at C-3, according to the cross peak observed on its NOESY spectrum (Fig. 2) between the methylene proton at $\delta_{\text{H}} 4.64$ and H-2 ($\delta_{\text{H}} 6.10$) and H-4 ($\delta_{\text{H}} 6.30$).

The relative stereochemistry of H-10 and H-10' was not

Fig. 2. Structure, Key HMBC and NOESY Correlations of Compound **2**

determined. But, from the chemical shifts of the chelated hydroxyl groups at C-1/C-1' ($\delta_{\text{H}} 12.37/\delta_{\text{H}} 12.05$) and C-8/C-8' ($\delta_{\text{H}} 11.84/\delta_{\text{H}} 11.74$) and also from those of the methyl groups at C-6/C-6' ($\delta_{\text{H}} 2.25$), we can confirmed on the basis of literature data,¹⁰⁾ that compound **2** was a mixture of *meso* and ($\pm$)-racemic diastereoisomers. In fact, when only one of

Table 2. ¹H-NMR Data of Compounds **2** and **3** and ¹³C-NMR Data of Compound **2**

Position	2 DMSO- <i>d</i> ₆		3 ²⁷⁾ CDCl ₃
	δ _H (mult. <i>J</i> Hz)	δ _C (APT)	δ _H (mult. <i>J</i> Hz)
1-OH	12.37 (1H, s) ^{a,c)}	4.9 (s)	12.10 (1H, s) ^{a,c)}
2	6.10 (1H, d, 2.5 Hz)	102.5 (d)	6.35 (1H, d, 2.2 Hz)
3	—	162.9 (s)	—
4	6.30 (1H, d, 2.5 Hz)	109.8 (d)	6.00 (1H, d, 2.2 Hz)
4a	—	137.5 (s)	—
5	6.60 (1H, d, 1.8 Hz)	123.2 (d)	6.12 (1H, br s)
6	—	147.3 (s)	—
7	6.20 (1H, d, 1.8 Hz)	118.0 (d)	6.69 (1H, br s)
8-OH	11.84 (1H, s) ^{a,c)}	162.8 (s)	11.90 (1H, s) ^{a,c)}
8a	—	112.9 (s)	—
9	—	190.7 (s)	—
9a	—	111.9 (s)	—
10	4.50 (1H, s)	59.0 (d)	4.33 (1H, s)
10a	—	139.9 (s)	—
11	2.25 (3H, s)	24.0 (q)	2.31 (3H, s)
12	4.64 (2H, d, 6.0 Hz)	68.2 (t)	4.54 (2H, d, 6.6 Hz)
13	5.46 (1H, t, 6.0 Hz)	119.3 (d)	5.47 (1H, t, 6.6 Hz)
14	—	138.9 (s)	—
15	2.05 (4H, m)	39.6 (t)	2.12 (2H, m)
16	—	28.4 (t)	2.14 (2H, m)
17	5.30 (1H, t, 8 Hz)	123.3 (d)	5.12 (1H, m)
18	—	131.9 (s)	—
19	1.81 (3H, s)	18.2 (q)	1.79 (3H, br s)
20	1.60 (3H, s) ^{b)}	19.4 (q)	1.70 (3H, br s) ^{b)}
21	1.62 (3H, s) ^{b)}	27.1 (q)	1.63 (3H, br s) ^{b)}
1'-OH	12.05 (1H, s) ^{c,d)}	161.8 (s)	12.10 (1H, s) ^{c,d)}
2'	—	136.0 (s)	6.35 (1H, d, 2.2 Hz)
3'-OH	9.54 (1H, s)	162.6 (s)	—
4'	6.27 (1H, s)	108.6 (d)	6.00 (1H, d, 2.2 Hz)
4'a	—	135.1 (s)	—
5'	6.66 (1H, d, 2.5 Hz)	122.4 (d)	6.12 (1H, br s)
6'	—	148.0 (s)	—
7'	6.37 (1H, d, 2.5 Hz)	117.6 (d)	6.69 (1H, br s)
8'-OH	11.74 (1H, s) ^{c,d)}	162.6 (s)	11.90 (1H, s) ^{c,d)}
8'a	—	113.0 (s)	—
9'	—	190.3 (s)	—
9'a	—	111.6 (s)	—
10'	4.50 (1H, s)	59.1 (d)	4.33 (1H, s)
10'a	—	143.5 (s)	—
11'	2.25 (3H, s)	24.2 (q)	2.31 (3H, s)
12'	3.35 (2H, d, 7.9 Hz)	22.7 (t)	—
13'	5.10 (1H, t, 7.9 Hz)	123.9 (d)	—
14'	—	133.5 (s)	—
15'	1.86 (3H, s) ^{e)}	20.4 (q)	—
16'	1.86 (3H, s) ^{e)}	27.1 (q)	—

a, b, c, d, e) Assignments bearing the same superscript in the same column may be reversed, c) Exchangeable in D₂O. ¹H-NMR was measured at 400 MHz for **2** and **3**. ¹³C-NMR of **2** was measured at 100 MHz.

those two forms is obtained, the chelated hydroxyl groups at C-1/C-1' and C-8/C-8' appears as a singlet of one proton. In the *meso* forms, chelated hydroxyl groups at C-1/C-1' and C-8/C-8' resonated at δ_H 11.86 and δ_H 11.78, respectively,¹⁰⁾ and in the (±)-racemic forms, they appears at δ_H 11.92 and δ_H 11.68, respectively.¹⁰⁾

From the above results, the structure of compound **2** was determined as 9-(3-((*E*)-3,7-dimethylocta-2,6-dienyloxy)-9,10-dihydro-1,8-dihydroxy-6-methyl-9-oxoanthracen-10-yl)-2,4,5-trihydroxy-7-methyl-3-(3-methylbut-2-enyl)anthracen-10(9*H*)-one, named adamabianthrone (see Fig. 2).

The two plants were tested for their antimicrobial activities against a wide range of microorganisms, including five Gram-negative bacteria, three Gram-positive bacteria and two yeasts. From the results obtained (Table 3), compound **1** showed significant activity on all the tested microorganisms, *Streptococcus faecalis* and *Bacillus cereus* being the most sensitive pathogens. The minimal inhibition concentration (MIC) values obtained with compound **1** varied from 14.37 μM (on *S. faecalis* and *B. cereus*) to 230.03 μM (on *Microsporum audouinii* and *Citrobacter freundii*). Compound **2** also exhibited a very good activity on 9 of the 10 (90%) tested microorganisms. The lowest MIC values (27.34 μM) observed with this compound was recorded against *S. faecalis*. Despite the fact that the MIC values obtained with the two tested compounds are greater than that of the reference antibiotics (0.45–3.60 μM), their inhibitory potential could be considered as important, taking into account that some of the tested microorganisms were resistant to the commonly used antibiotics. Besides, the MIC values obtained, the tested compounds are generally four fold lesser than their corresponding minimal microbicidal concentration (MMC) values (Table 3). This indicates that the microbicidal effects of the tested compounds could be expected on the tested microorganisms.^{21–25)}

Experimental

General All melting points were measured on a Büchi apparatus and are uncorrected. The UV spectra were obtained on a Kontron Uvikon 932 spectrophotometer. The IR spectra were recorded on a Nicolet Impact 400D FT-IR spectrometer. The ¹H- and ¹³C-NMR data [including HSQC, HMBC, NOESY and ¹H–¹H COSY (correlation spectroscopy experiment)] were measured on a Bruker DRX-400 instrument operating at 400 MHz for ¹H and 100 MHz for ¹³C. The chemical shifts (δ) are reported in ppm downfield from tetramethylsilane (TMS) using TMS or the solvent signal as standard. The HR-ESI-TOF-MS and HR-ESI-TOF-MS/MS were obtained in the posi-

Table 3. Antimicrobial Activity of Compounds **1**, **2** and Reference Antibiotics (RA)

Microorganisms	Minimum inhibition concentration in μg/ml and μM (in parenthesis) of the tested samples			Minimum microbicidal concentration in μg/ml and μM (in parenthesis) of the tested samples		
	1	2	RA ^{a)}	1	2	RA ^{a)}
Gram-negative bacteria						
<i>C. freundii</i>	156.25 (230.03)	156.25 (218.74)	9.76 (1.80)	>156.25 (>230.03)	>156.25 (>218.74)	19.53 (3.60)
<i>E. coli</i>	19.53 (28.75)	156.25 (218.74)	9.76 (1.80)	78.12 (115.02)	>156.25 (>218.74)	19.53 (3.60)
<i>P. aeruginosa</i>	39.06 (57.51)	39.06 (54.68)	19.53 (3.60)	78.12 (115.02)	156.25 (218.74)	39.06 (7.20)
<i>K. pneumoniae</i>	19.53 (28.75)	78.12 (109.37)	19.53 (3.60)	39.06 (57.51)	156.25 (218.74)	39.06 (7.20)
<i>S. typhi</i>	19.53 (28.75)	39.06 (54.68)	19.53 (3.60)	39.06 (57.51)	78.12 (109.37)	39.06 (7.20)
Gram-positive bacteria						
<i>B. cereus</i>	9.76 (14.37)	39.06 (54.68)	2.44 (0.45)	39.06 (57.51)	78.12 (109.37)	4.88 (0.90)
<i>S. aureus</i>	39.06 (57.51)	156.25 (218.74)	9.76 (1.80)	78.12 (115.02)	>156.25 (>218.74)	19.53 (3.60)
<i>S. faecalis</i>	9.76 (14.37)	19.53 (27.34)	4.88 (0.90)	39.06 (57.51)	78.12 (109.37)	9.76 (1.80)
Fungi						
<i>C. albicans</i>	39.06 (57.51)	>156.25 (>218.74)	9.76 (1.04)	78.12 (115.02)	>156.25 (>218.74)	19.53 (2.08)
<i>M. audouinii</i>	156.25 (230.03)	39.06 (54.68)	9.76 (1.04)	>156.25 (>230.03)	156.25 (218.74)	19.53 (2.08)

a) RA: reference antibiotics (gentamicin for bacteria and nystatin for fungi).

tive ion mode on an API QSTAR pulsar mass spectrometer.

Analytical thin layer chromatography (TLC) was performed on Kieselgel 60F₂₅₄ precoated Al sheet (0.2 mm layer thickness, Merck) and the spots were detected by ultraviolet irradiation (254, 366 nm) and by spraying with 10% H₂SO₄ reagent.

Plant Material The roots of *Psorospermum febrifugum* were collected at Bantoum, West Province (Cameroon), in September 2004, while the barks of *Psorospermum adamaense* were harvested at Batsenga, Centre Province (Cameroon), in March 2007. These two plants were identified by Mr. Nana (plant taxonomist) of National Herbarium, Yaounde where voucher specimens of *P. febrifugum* (N° 9826 SRF Cam) and *P. adamaense* (N° 41149/HNC) documenting the collection, have been deposited.

Extraction and Isolation Air-dried powders roots of *P. febrifugum* (4 kg) and bark of *P. adamaense* (7 kg) were separately extracted by maceration at room temperature for 48 h using a methanol/dichloromethane (1 : 1) mixture. The suspensions were filtered and each filtrate was concentrated under reduced pressure to give 300 g and 580 g of crude residue, respectively. The crude extract from the roots of *P. febrifugum* (300 g) were subjected to flash column chromatography on silica gel (Merck, 230–400 mesh) and eluted with hexane, hexane/AcOEt (4 : 1), hexane/AcOEt (1 : 1), hexane/AcOEt (1 : 4), AcOEt to give five fractions labelled F₁ (53 g), F₂ (40 g), F₃ (25 g), F₄ (12 g), F₅ (10 g). Fractions F₁ and F₂ were mixed, on the basis of their thin layer chromatography (TLC). Fraction F (93 g) obtained was further subjected to column chromatography (CC) on silica gel (Merck, 70–230 mesh) and eluted with hexane/AcOEt mixture of increasing polarity. One hundred and twenty fractions of 150 ml each were collected and analysed by TLC using hexane/AcOEt (7 : 3) as mobile phase. Sub-fractions 2–10, 11–20, 99–101 and 104–120 were left to crystallize at room temperature to afford, respectively: 3-geranyloxyemodin anthrone (30 mg), bianthrone A₁ (3) (40 mg), febrifuquinone (1) (8 mg) and vismione D (4) (20 mg). Similarly, the crude extract of the barks of *P. adamaense* (300 g) were also subjected to flash column chromatography on silica gel (Merck, 230–400 mesh) and eluted with hexane, hexane/AcOEt (4 : 1), hexane/AcOEt (1 : 1), hexane/AcOEt (1 : 4), AcOEt to give five fractions labelled A₁ (52 g), A₂ (80 g), A₃ (7.1 g), A₄ (8 g), A₅ (10 g). Fractions A₁ and A₂ were also mixed, according to the spots observed on their TLC. Fraction F (93 g) A₁ and A₂ were also mixed on the basis of TLC. We obtained fraction A (132 g) which was treated as fraction F. Two hundred and thirty-one fractions of 150 ml each were collected and analysed by TLC using hexane/AcOEt (19 : 1) as mobile phase. Sub-fractions 39–56, 97–109, 132–144 and 161–170, also left at room temperature, were crystallized and gave, respectively: 3-geranyloxyemodin anthrone (80 mg); 1,8-dihydroxy-3-isoprenyloxy-6-methylanthraquinone (20 mg); bianthrone A_{2b} (25 mg); 2-geranylemodin (20 mg) and emodin (5) (14 mg). The residues obtained from sub-fractions 39–59 and 153–170 and labelled A' and A'', were eluted, respectively with the mixture of hexane/AcOEt (9 : 1) and of hexane/AcOEt (19 : 1) and afforded 3-geranyloxy-1,8-dihydroxy-6-methylanthraquinone (20 mg) and adamabianthrone (2) (6 mg), respectively.

Febrifuquinone (1): Red powder. mp 132–133 °C. UV $\lambda_{\max}$ (EtOH) nm: 235, 320, 390 and 435. IR (KBr) cm⁻¹: 3630, 3360, 3300, 3050, 3020, 2860, 1668, 1640, 1630, 1560, 1110. ¹H- and ¹³C-NMR (see Table 1). HR-ESI-TOF-MS *m/z*: 679 [M+H]⁺. HR-ESI-TOF-MS *m/z*: 679.2514 (Calcd for C₄₀H₃₉O₁₀: 679.2532). HR-ESI-TOF-MS/MS *m/z*: 409 [M-C₁₅H₉O₅]⁺, 269 [M-C₂₅H₂₉O₅]⁺, 525 [M-C₁₀H₁₇O]⁺.

Adamabianthrone (2): Yellow powder. mp 152–154 °C. UV $\lambda_{\max}$ (EtOH) nm: 320, 270, 363. IR (KBr) cm⁻¹: 3620, 3366, 3040, 2870, 2870, 1630, 1565, 1100. ¹H- and ¹³C-NMR (see Table 2). HR-ESI-TOF-MS *m/z*: 715 [M+H]⁺. HR-ESI-TOF-MS *m/z*: 715.3231 (Calcd for C₄₅H₄₇O₈: 715.3258). HR-ESI-TOF-MS/MS *m/z*: 391 [M-C₂₀H₁₉O₄]⁺, 323 [M-C₂₅H₂₇O₄]⁺, 561 [M-C₁₀H₁₇O]⁺.

Antimicrobial Assays. Microbial Strains The microorganisms tested included *Bacillus cereus* LMP0404G, *Streptococcus faecalis* LMP0207U, methicillin-resistant *Staphylococcus aureus* LMP805, as three Gram-positive bacteria, five Gram-negative bacteria namely β -lactamase positive (β L+) *Escherichia coli* LMP701, ampicillin-resistant *Klebsiella pneumoniae* LMP803, carbenicillin-resistant *Pseudomonas aeruginosa* LMP804, chloramphenicol-resistant *Salmonella typhi* LMP706, chloramphenicol-resistant *Citrobacter freundii* LMP802 and two fungi namely *Candida albicans* LMP709U, *Microsporium audouinii* LMP725D. These microbial species are clinical isolated from patients in the Yaounde General Hospital (Cameroon).

Their identification were confirmed before usage in the Laboratory of Applied Microbiology and Molecular Pharmacology (LMP) (Faculty of Science, University of Yaoundé I) where the antimicrobial tests were per-

formed, using API system as previously reported.²⁶⁾ They were maintained on agar slant at 4 °C at the LMP and sub-cultured on a fresh appropriate agar plate for 24–48 h prior to any antimicrobial test.

MIC and MMC Determinations: Microdilution Method The MICs of compounds 1, 2 and reference antibiotics (RA) (gentamicin for bacteria and nystatin for fungi) were evaluated as follows: the tested samples were first of all dissolved in dimethylsulfoxide (DMSO). The solution obtained was added to muller hinton agar (MHA) to give a final concentration of 156.25 μ g/ml. This was serially diluted two fold to obtain concentration ranged from 1.22 to 156.25 μ g/ml. One hundred microliters of each concentration was added in a well (96-wells microplate) containing 95 μ l of nutrient broth containing 0.05% phenol red and supplemented with 10% glucose (NBGP) and 5 μ l of inoculum [(at 1.5×10⁶ colony forming unit (CFU)/ml by adjusting the optical density to 0.1 at 600 nm (SHIMADZU UV-120-01 spectrophotometer)].^{21–25)} The final concentration of DMSO in the well was less than 1% (preliminary analyses with 1% (v/v) DMSO/NBGP do not show the growth of the tested microorganisms). The negative control well consisted of 195 μ l of NBGP and 5 μ l of the standard inoculum.^{21–25)} The plates were covered with a sterile plate sealer, then agitated to mix the content of the wells using a plate shaker and incubated at 37 °C for 24 h. The assay was repeated twice. The MIC of samples was detected following addition (40 μ l) of 0.2 mg/ml *p*-iodonitrotetrazolium chloride and incubated at 37 °C for 30 min.²⁵⁾ Viable bacteria reduced the yellow dye to a pink colour. The lowest concentration showing no colour change was considered as the MIC. For the determination of MMC, a portion of liquid (5 μ l) from each well that showed no change in colour was placed on MHA and incubated at 37 °C for 24 h. The lowest concentration that yielded no growth after this sub-culturing was taken as the MMC.^{21–25)}

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