

## Laurentixanthonones A and B, antimicrobial xanthonones from *Vismia laurentii*

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### Abstract

A phytochemical investigation of the constituents of the roots of *Vismia laurentii* has resulted in the isolation of two xanthone derivatives named laurentixanthone A (**1**) (6-hydroxy-3,3-dimethyl-11-(3-methylbut-2-enyl)pyrano[2,3-*c*]xanthen-7(3*H*)-one) and laurentixanthone B (**2**) (1-hydroxy-5,6,7,8-tetramethoxyxanthone), along with 11 known compounds: 1,7-dihydroxyxanthone, vismiaquinone, vismiaquinone B, bivismiaquinone, 3-geranyloxy-6-methyl-1,8-dihydroxyanthraquinone, *O*<sup>1</sup>-demethyl-3',4'-deoxypsorospermin-3',4'-diol, 6-deoxyisojacareubin, 1,8-dihydroxy-6-methoxy-3-methylanthraquinone, kaempferol, friedelin and stigmasterol. The structures of compounds were established by means of spectroscopic methods. Furthermore, the compounds were screened for antimicrobial activities in vitro.

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**Keywords:** *Vismia laurentii*; Guttiferae; Xanthonones; Anthraquinones; Antimicrobial activities

### 1. Introduction

The tribe Vismieae belongs to the family of Guttiferae and comprises three genera, *Vismia*, *Harungana* and *Psorospermum*. It is distributed in the tropical and subtropical regions of the world (Hutchinson and Dalziel, 1954). *Vismia laurentii* De Wild is a large shrub or tree which is found in the Centre Province of Cameroon where it is locally called “atondo owse”. The bark and roots are employed in decoctions as tonic and febrifugal (Kerharo, 1974; Bamps, 1966, 1970) and is also used in tropical Africa as a remedy for the treatment of skin diseases (such as derma-

titis, leprosy, scabies, eczemas) and wounds using coconut as vehicle. Previous chemical studies have reported the presence of triterpenoids, prenylated anthrones, anthraquinones, bianthraquinones, benzophenones, xanthonones and lignans from *Vismia* species (Nagem and De Olivera, 1997; Dos Santos et al., 2000; Araújo et al., 1990; Nagem and Alves, 1995; Nagem and Faria De Jesus, 1990; Hussein et al., 2003; Bilia et al., 2000; Botta et al., 1986). Cytotoxic and antifeedant activities have also been reported for various *Vismia* constituents (Cassinelli et al., 1986; Seo et al., 2000). As part of our continuous search for bioactive natural products from Guttiferae family (Nkengfack et al., 2002; Azebaze et al., 2004; Yimdjo et al., 2004), we investigated the constituents of the *V. laurentii*, a plant on which no phytochemical studies has been reported to the best of our knowledge. In the present work, we report the isolation

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and structural elucidation of 13 secondary metabolites from the roots of this plant, two of which are new derivatives, and the results obtained from the evaluation of their antimicrobial properties.

## 2. Results and discussion

Air-dried and ground roots of *V. laurentii* were extracted at room temperature with a mixture of methanol–dichloromethane (1:1). Separation and purification of this extract by successive silica gel column chromatography (see Section 3) led to the isolation and characterisation of two new xanthone derivatives, named laurentixanthones A (**1**) and B (**2**) along with 11 known secondary metabolites. The known compounds were identified by comparison of their physico-chemical and spectroscopic data ( $^1\text{H}$ ,  $^{13}\text{C}$  NMR, 2D NMR and MS) with those of authentic samples and reference data as: 1-hydroxy-6,7-dimethoxyxanthone (Gunatilaka et al., 1982), 1,7-dihydroxyxanthone (Fujita et al., 1992), vismiquinone (Goncalves and Mors, 1981), vismiaquinone B (Miraglia et al., 1981), bivismiaquinone (Hussein et al., 2003), 3-geranyloxy-6-methyl-1,8-dihydroxy anthraquinone (Botta et al., 1983), *O*<sup>1</sup>-demethyl-3',4'-deoxysorospermin-3',4'-diol (Abou-Shoer et al., 1988), 6-deoxyisocareubins (Wu et al., 1998), 1,8-dihydroxy-6-methoxy-3-methylanthraquinone (Goncalves and Mors, 1981), kaempferol (Kouam

et al., 1993), friedelin (Gunatilaka et al., 1982) and stigmasterol (Alan et al., 1996).

Compound **1**, laurentixanthone A, m.p. 156–157 °C, was obtained as yellow needles from a mixture of cyclohexane–EtOAc (4:1). It reacted positively to  $\text{FeCl}_3$  reagent, suggesting the presence of at least one phenolic hydroxyl group in the molecule. Its molecular formula was determined to be  $\text{C}_{23}\text{H}_{22}\text{O}_4$  using positive high-resolution (HR) electrospray ionisation time of flight (ESI-TOF) mass spectroscopy which showed a pseudomolecular ion peak  $[\text{M}+\text{H}]^+$  at  $m/z$  363.1589 (calcd. for  $\text{C}_{23}\text{H}_{23}\text{O}_4$ , 363.1596). The broad-band proton decoupled  $^{13}\text{C}$  NMR spectrum of compound **1** (Table 1) showed 22 carbon signals which were attributed by APT and HSQC techniques as four methyls, one methylene, seven methines and 11 quaternary carbons including one carbonyl group ( $\delta_{\text{C}}$  181.8), four oxygenated  $\text{sp}^2$  carbons, five  $\text{sp}^2$ , and one  $\text{sp}^3$  carbons. The IR spectrum of compound **1** displayed vibration bands characteristic of a chelated hydroxyl group ( $\nu_{\text{max}} = 3356\text{ cm}^{-1}$ ), a conjugated and chelated carbonyl group ( $\nu_{\text{max}} = 1664\text{ cm}^{-1}$ ) and a C–C double bonds of an aromatic ring ( $\nu_{\text{max}} = 1650\text{ cm}^{-1}$ ). These data together with those obtained from the UV spectrum of compound **1** [ $\lambda_{\text{max}}$  (log  $\epsilon$ ): 250 (4.08), 322 (3.71) nm], were consistent with the presence of a xanthone skeleton (Cardona et al., 1990). In the  $^1\text{H}$  NMR spectrum (acetone- $d_6$ ) (Table 1) of compound **1**, analysed by a combination of  $^1\text{H}$ – $^1\text{H}$  COSY

Table 1  
 $^1\text{H}$  and  $^{13}\text{C}$  NMR data for compounds **1** and **2**

Position	<b>1</b> (Acetone- $d_6$ )		<b>2</b> ( $\text{CDCl}_3$ )	
	$\delta_{\text{H}}$ (mult. $J$ Hz)	$\delta_{\text{C}}$ (APT)	$\delta_{\text{H}}$ (mult. $J$ Hz)	$\delta_{\text{C}}$ (APT)
1	–	164.1	–	162.0
2	5.67 (1H, s)	99.7	6.75 (1H, dd, 7.9, 1.8)	110.7
3	–	161.7	7.51 (1H, t, 7.9)	136.1
4	–	102.1	6.90 (1H, dd, 7.9, 1.8)	106.4
4a	–	152.1	–	155.3
5	–	107.9	–	137.2
6	7.28 (1H, dd, 7.8, 1.9)	122.2	–	147.7
7	7.22 (1H, t, 7.8)	125.0	–	143.1
8	7.75 (1H, dd, 7.8, 1.9)	116.5	–	149.4
8a	–	121.7	–	117.0
4b	–	147.0	–	153.2
9	–	181.8	–	181.6
8b	–	104.2	–	108.8
2'	–	79.1	–	–
3'	5.60 (1H, d, 10)	128.1	–	–
4'	6.74 (1H, d, 10)	115.7	–	–
5',6'	1.49 (6H, s)	28.4	–	–
1-OH <sup>a</sup>	13.07 (1H, s)	–	13.07 (1H, s)	–
5-OMe	–	–	3.90 (3H, s)	61.6 <sup>b</sup>
6-OMe	–	–	3.95 (3H, s)	61.7 <sup>b</sup>
7-OMe	–	–	3.95 (3H, s)	62.0 <sup>b</sup>
8-OMe	–	–	4.10 (3H, s)	62.8
1''	3.49 (2H, d, 7.9)	23.4	–	–
2''	5.22 (1H, t, 7.9)	125.1	–	–
3''	–	133.6	–	–
4''	1.86 (3H, s)	26.1	–	–
5''	1.71 (3H, s)	19.3	–	–

<sup>a</sup> Exchangeable in  $\text{D}_2\text{O}$ .

<sup>b</sup> Assignments bearing the same superscript in the same column may be reversed.

and HSQC experiments, an ABC spin system, indicated by two double doublets at  $\delta_{\text{H}}$  7.28/ $\delta_{\text{C}}$  122.2 (1H, *dd*,  $J = 7.8$ , 1.9 Hz) and  $\delta_{\text{H}}$  7.75/ $\delta_{\text{C}}$  116.5 (1H, *dd*,  $J = 7.8$ , 1.9 Hz) and a triplet at  $\delta_{\text{H}}$  7.22/ $\delta_{\text{C}}$  125.0 (1H, *t*,  $J = 7.8$  Hz), corresponding to a 1,2,3-trisubstituted benzene ring, was observed in addition to a chelated hydroxyl signal at  $\delta_{\text{H}}$  13.07 and one-proton singlet at  $\delta_{\text{H}}$  5.67/ $\delta_{\text{C}}$  99.7 due to an aromatic proton. Furthermore, the  $^1\text{H}$  and  $^{13}\text{C}$  NMR spectra also displayed the presence of two sets of signals. The first set, comprising a six-proton singlet at  $\delta_{\text{H}}$  1.49/ $\delta_{\text{C}}$  28.4 and two *cis*-olefinic protons ( $\delta_{\text{H}}$  5.60/ $\delta_{\text{C}}$  128.1 and  $\delta_{\text{H}}$  6.74/ $\delta_{\text{C}}$  115.7, each *d*,  $J = 10.0$  Hz) due to a dimethylchromene bicyclic system. The second set of signals, consisting of a doublet of two protons at  $\delta_{\text{H}}$  3.49/ $\delta_{\text{C}}$  23.4 (2H, *d*,  $J = 7.9$  Hz), a triplet of one proton at  $\delta_{\text{H}}$  5.22/ $\delta_{\text{C}}$  125.1 (1H, *t*,  $J = 7.9$  Hz), a carbon signal at  $\delta_{\text{C}}$  133.6, three proton singlet at  $\delta_{\text{H}}$  1.86/ $\delta_{\text{C}}$  26.1 (3H, Me) and another three proton singlet at  $\delta_{\text{H}}$  1.71/ $\delta_{\text{C}}$  19.3 (3H, Me), established the presence of a  $\gamma,\gamma$ -dimethylallyl moiety. A combination of the COSY and HSQC experiments permitted the assignment of all the protonated carbons (Table 1). It remained to establish the positions of the substituents on the xanthone skeleton. This was done by analysis of the HMBC spectrum. In effect, the chelated hydroxyl group at  $\delta_{\text{H}}$  13.07 correlated with carbons at  $\delta_{\text{C}}$  164.1 (C-1),  $\delta_{\text{C}}$  99.7 (C-2) and  $\delta_{\text{C}}$  104.2 (C-8b), while the aromatic proton at  $\delta_{\text{H}}$  5.67 correlated with quaternary carbons at  $\delta_{\text{C}}$  164.1 (C-1),  $\delta_{\text{C}}$  161.7 (C-3) and  $\delta_{\text{C}}$  104.2 (C-8b). These correlations indicated clearly that the aromatic proton at  $\delta_{\text{H}}$  5.67 is located at C-2 position. On the other hand, the *cis*-olefinic proton of the dimethylchromene bicyclic system at  $\delta_{\text{H}}$  6.74, showed correlations with the quaternary carbons at  $\delta_{\text{C}}$  102.1 (C-4) and  $\delta_{\text{C}}$  152.1 (C-4a), while the other *cis*-olefinic proton at  $\delta_{\text{H}}$  5.60 was correlated with the quaternary carbons at  $\delta_{\text{C}}$  102.1 (C-4) and  $\delta_{\text{C}}$  79.1 (C-2'). These results demonstrated clearly that the dimethylchromene moiety was fused in an angular manner to the aromatic ring A of xanthone skeleton bearing the hydroxyl group. The position of the remaining  $\gamma,\gamma$ -dimethylallyl group was established as follows. Once more, the HMBC spectrum of **1** (Fig. 1), showed that, one of the ABC spin proton at  $\delta_{\text{H}}$  7.28 displayed cross-peaks with carbons C-5 ( $\delta_{\text{C}}$  107.9) and C-1' ( $\delta_{\text{C}}$  23.4), whereas the allylic proton at  $\delta_{\text{H}}$  3.49 of the  $\gamma,\gamma$ -dimethylallyl group gave cross-peaks with carbons C-4b ( $\delta_{\text{C}}$  147.0), C-5 ( $\delta_{\text{C}}$  107.9), C-6 ( $\delta_{\text{C}}$  122.2) and C-2'' ( $\delta_{\text{C}}$  125.1), indicating clearly that the  $\gamma,\gamma$ -dimethylallyl group was located at the C-5 position. This was further confirmed by the NOESY spectrum which showed on one hand, correlations peaks between H-2 at  $\delta_{\text{H}}$  5.67 and the chelated hydroxyl group at  $\delta_{\text{H}}$  13.07, on the other hand, between allylic proton H-1'' at  $\delta_{\text{H}}$  3.49 and H-6 at  $\delta_{\text{H}}$  7.28. On the basis of the above results, the structure of laurentixanthone A (**1**), was determined to be 6-hydroxy-3,3-dimethyl-11-(3-methylbut-2-enyl)pyrano[2,3-*c*]xanthen-7(3*H*)-one.

Compound **2**, m.p. 112–114 °C, was obtained as brown needles from cyclohexane. It reacted positively to  $\text{FeCl}_3$  reagent, suggesting the presence of at least one phenolic

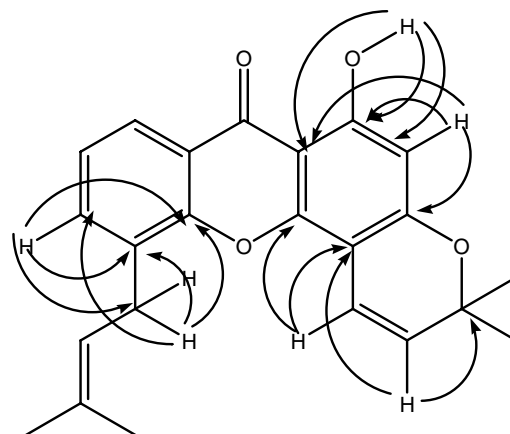


Fig. 1. HMBC correlations of compound **1**.

hydroxyl group in the molecule. Its molecular formula,  $\text{C}_{17}\text{H}_{16}\text{O}_7$ , was determined by high-resolution (HR) electrospray ionisation time of flight (ESI-TOF) mass spectroscopy which showed in positive mode, a pseudomolecular ion peak  $[\text{M}+\text{H}]^+$  at  $m/z$  333.0972 (calcd. for  $\text{C}_{17}\text{H}_{17}\text{O}_7$ , 333.0974) implying 10° of unsaturation. The broad-band proton-decoupled  $^{13}\text{C}$  NMR spectrum of compound **2** (Table 1), sorted by APT and HSQC spectra revealed the presence of four methoxyl groups at  $\delta_{\text{H}}$  3.90/ $\delta_{\text{C}}$  61.6,  $\delta_{\text{H}}$  3.95/ $\delta_{\text{C}}$  61.7,  $\delta_{\text{H}}$  3.95/ $\delta_{\text{C}}$  62.0 and  $\delta_{\text{H}}$  4.10/ $\delta_{\text{C}}$  62.8, three  $\text{sp}^2$  methines and 10  $\text{sp}^2$  quaternary carbons, one of which corresponded to a carbonyl group at  $\delta_{\text{C}}$  181.6. These data together with those obtained from IR spectrum [ $\nu_{\text{max}}$  (C=O) 1661  $\text{cm}^{-1}$ ; ( $\nu_{\text{max}}$  (OH) 3212  $\text{cm}^{-1}$ ] and UV spectrum [ $\lambda_{\text{max}}$  (log  $\epsilon$ ): 221 (4.10), 245sh (4.32), 254 (4.41), 316 (3.85) and 375 (3.70) nm], suggesting that compound **2** possesses a xanthone skeleton in its structure (Cardona et al., 1990). Its  $^1\text{H}$  NMR spectrum ( $\text{CDCl}_3$ ) showed the presence of one proton signal at  $\delta_{\text{H}}$  13.07 characteristic of a chelated hydroxyl group located at *peri* position (C-1) to carbonyl group. This spectrum further exhibited typical signals of three vicinal aromatic protons of an ABC spin system characteristic of 1,2,3-trisubstituted benzene ring at  $\delta_{\text{H}}$  6.75/ $\delta_{\text{C}}$  110.7 (1H, *dd*,  $J = 7.9$ , 1.8 Hz),  $\delta_{\text{H}}$  6.90/ $\delta_{\text{C}}$  106.4 (1H, *dd*,  $J = 7.9$ , 1.8 Hz) and  $\delta_{\text{H}}$  7.51/ $\delta_{\text{C}}$  136.1 (1H, *t*,  $J = 7.9$  Hz) and four three proton singlet confirming the presence of four methoxyl groups in compound **2**. From the above spectroscopic data, it appears that all the four methoxyl

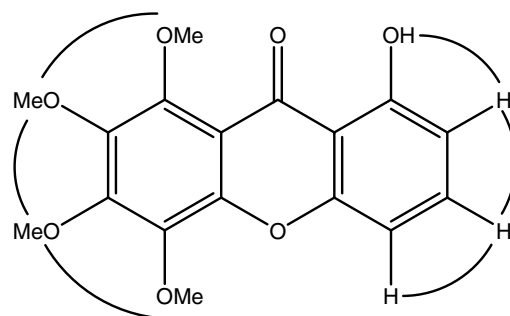
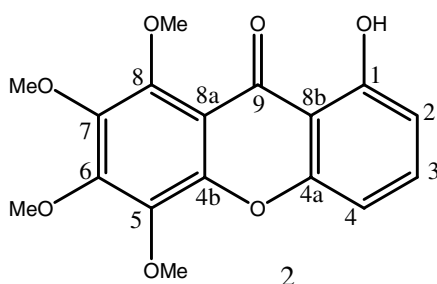
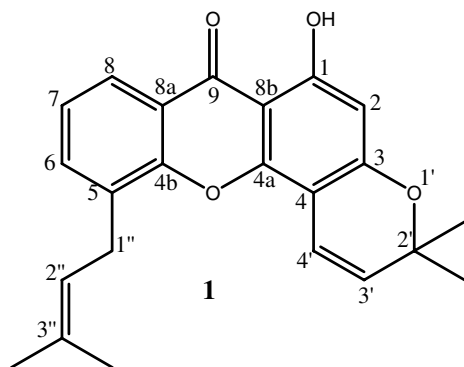


Fig. 2. NOESY correlations of compound **2**.

groups are located on ring B. This was further confirmed by correlations observed on NOESY spectrum between the different methoxyl groups as shown in Fig. 2. Thus, the structure of compound **2** was assigned to be 1-hydroxy-5,6,7,8-tetramethoxyxanthone.



Pure isolated metabolites **1** and **2** were evaluated for their antibacterial and antifungal potential against a wide range of microorganisms: Gram-positive, Gram-negative bacteria and yeasts. As shown in Table 2, compounds **1** and **2** exhibited both antibacterial and antifungal effects. Nevertheless, pronounced activity (MIC lower than 78.12 µg/ml) was observed against a limited number of microorganisms (30%) for compound **2**, whereas compound **1** appeared to be more active. A pronounced activity was observed against 60% of the tested microbial strains. However, the inhibitory effect of the two compounds against some of the tested pathogens was equal or more important than that of the reference drugs (Table 2). *Streptococcus faecalis* and *Candida glabrata* were, respectively, the most sensitive pathogens to the effects of compounds **1** and **2**.

### 3. Experimental

#### 3.1. General experimental procedures

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 <sup>1</sup>H and <sup>13</sup>C NMR data (including HSQC, HMBC, NOESY and <sup>1</sup>H–<sup>1</sup>H COSY) were measured on a Bruker

Table 2  
Minimum inhibition concentration (µg/ml and µM) of **1**, **2** and reference antibiotics

Microorganisms	Tested samples				RA <sup>a</sup>	
	1		2			
Gram-negative bacteria						
<i>Citrobacter freundii</i>	—	—	—	—	4.88	9.0
<i>Enterobacter aerogens</i>	—	—	—	—	9.76	18
<i>Enterobacter cloacae</i>	39.06	107.9	—	—	4.88	9.0
<i>Escherichia coli</i>	—	—	—	—	1.22	2.25
<i>Klebsiella pneumoniae</i>	—	—	—	—	2.44	4.5
<i>Morganella morganii</i>	9.76	26.99	—	—	2.44	4.5
<i>Proteus mirabilis</i>	39.06	107.9	—	—	2.44	4.5
<i>Proteus vulgaris</i>	—	—	—	—	1.22	2.25
<i>Pseudomonas aeruginosa</i>	—	—	4.88	14.70	4.88	9.0
<i>Shigella dysenteriae</i>	4.88	13.49	—	—	2.44	4.5
<i>Shigella flexneri</i>	4.88	13.49	19.53	58.83	2.44	4.5
<i>Salmonella typhi</i>	—	—	—	—	2.44	4.5
Gram-positive bacteria						
<i>Streptococcus faecalis</i>	1.22	3.37	—	—	4.88	9.0
<i>Staphylococcus aureus</i>	—	—	—	—	4.88	9.0
<i>Bacillus cereus</i>	78.12	215.8	78.12	235.3	2.44	4.5
<i>Bacillus megaterium</i>	2.44	6.74	—	—	4.88	9.0
<i>Bacillus stearothermophilus</i>	4.88	13.49	—	—	4.88	9.0
<i>Bacillus subtilis</i>	4.88	13.49	2.44	7.35	2.44	4.5
Yeasts						
<i>Candida albicans</i>	78.12	215.8	19.53	58.83	4.88	5.21
<i>Candida glabrata</i>	2.44	6.74	0.61	1.84	4.88	5.21

No effect of the DMSO used as dilution solvent was observed on the tested microbial strains.

–, Not determined because MIC was greater than 78.12 µg/ml.

<sup>a</sup> RA: reference antibiotics (Gentamycin for bacteria, Nystatin for yeast).



DRX-400 instrument operating at 400 MHz for  $^1\text{H}$  and 100 MHz for  $^{13}\text{C}$ . The chemical shifts ( $\delta$ ) are reported in ppm downfield from tetramethylsilane (TMS) using TMS or the solvent signal as standard. The HRESI-TOF-MS and ESI-TOF-MS/MS were obtained in the positive ion mode on a API QSTAR pulsar mass spectrometer. Analytical thin layer chromatography (TLC) was performed on Kieselgel 60F<sub>254</sub> 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%  $\text{H}_2\text{SO}_4$  reagent.

### 3.2. Plant material

The roots of *V. laurentii* De Wild were collected in March 2004 at Mbal Mayo, in the Centre Province of the Republic of Cameroon and identified by Mr. Nana (plant taxonomist) of *National Herbarium*, Yaounde. A voucher specimen (No. 1882/SRFK) documenting the collection has been deposited at the *National Herbarium*, Yaounde, Cameroon.

### 3.3. Extraction and isolation

Air-dried powder of the roots of *V. laurentii* (2 kg) was exhaustively extracted at room temperature for 48 h using a mixture of methanol/dichloromethane (1:1) by maceration. The suspension was filtered and the filtrates concentrated under reduced pressure to give 100 g of brown crude residue. This residue was subjected to flash column chromatography on silica gel (Merck, 230–400 mesh) and eluted with cyclohexane, cyclohexane/EtOAc (4:1), cyclohexane/EtOAc (1:1) and EtOAc to give four fractions labelled A (20 g), B (35 g), C (18 g) and D (10 g), respectively. Fraction B (35 g) was further subjected to column chromatography on silica gel (Merck, 70–230 mesh) and eluted with a cyclohexane/AcOEt mixture of increasing polarity. One hundred fractions of 100 ml each were collected and analysed by TLC using cyclohexane– $\text{CH}_2\text{Cl}_2$  (7:3) as mobile phase. Fractions 1–25, eluted with cyclohexane, afforded three compounds which were identified as: 1,8-dihydroxy-3-geranyloxy-6-methylanthraquinone (300 mg), friedelin (25 mg) and stigmasterol (35 mg). Fractions 27–47, eluted with a mixture of cyclohexane/EtOAc (9:1) gave a brown residue (1.3 g) which was subjected to further column chromatography to afford laurentixanthone B (**2**) (25 mg) from sub-fractions 10–25. From sub-fractions 26–29, we obtained 1,8-dihydroxy-3-methoxy-6-methylanthraquinone (25 mg) and sub-fractions 30–40 afforded bivismiaquinone (40 mg). Finally, fractions A (20 g) eluted with cyclohexane/EtOAc (4:1) also afforded brown residue (3.5 g) which was also subjected to column chromatography to give 1,7-dihydroxyxanthone (18 mg), from sub-fractions 10–25, laurentixanthone A (**1**) (50 mg) from sub-fractions 27–37. While, sub-fractions 38–47, 50–60, 65–70, 80–100 and 108–120 were left to crystallize at room temperature to afford kaempferol (16 mg), vismiaquinone (70 mg), vismiaquinone B (50 mg), *O*<sup>1</sup>-demethyl-3',4'-deox-

yporospermin-3',4'-diol (12 mg) and 1-hydroxy-6,7-dimethoxyxanthone (10 mg), respectively.

#### 3.3.1. Laurentixanthone A (**1**)

Yellow needles, m.p. 156–157 °C. UV  $\lambda_{\text{max}}^{\text{MeOH}}$  nm (log  $\epsilon$ ): 250 (4.08), 260 (3.97), 322 (3.71) and 379 (3.42). IR  $\nu_{\text{max}}^{\text{MeOH}}$ : 3356, 1664, 1650, 1215, 1113  $\text{cm}^{-1}$ .  $^1\text{H}$  and  $^{13}\text{C}$  NMR (see Table 1). ESI-TOF-MS  $m/z$  363  $[\text{M}+\text{H}]^+$ . HRESI-MS  $m/z$  363.1589 (calcd. for  $\text{C}_{23}\text{H}_{22}\text{O}_4+\text{H}$ , 363.1596). MS/MS  $m/z$  (rel. int.): 347  $[\text{M}-\text{CH}_3]^+$  (53), 299  $[\text{M}-\text{C}_5\text{H}_9]^+$  (60), 211  $[\text{M}-\text{CH}_3-\text{C}_7\text{H}_4\text{O}_3]^+$  (47), 188  $[\text{M}-\text{CH}_3-\text{C}_{10}\text{H}_7\text{O}_2]^+$  (35).

#### 3.3.2. Laurentixanthone B (**2**)

Brown needles, m.p. 112–114 °C. UV  $\lambda_{\text{max}}^{\text{MeOH}}$  nm (log  $\epsilon$ ): 221 (4.10), 245 sh (4.32), 254 (4.41), 316 (3.85) and 375 (3.70). IR  $\nu_{\text{max}}^{\text{MeOH}}$ : 3212, 2930, 2845, 1661, 1647, 1245, 1145  $\text{cm}^{-1}$ .  $^1\text{H}$  and  $^{13}\text{C}$  NMR (see Table 1). ESI-TOF-MS  $m/z$  333  $[\text{M}+\text{H}]^+$ . HRESI-MS  $m/z$  333.0972 (calcd. for  $\text{C}_{17}\text{H}_{16}\text{O}_7+\text{H}$ , 333.0974).

### 3.4. Antimicrobial assays

#### 3.4.1. Microbial strains

A total of 20 microbial cultures belonging to six Gram-positive bacterial species (*Bacillus cereus*, *Bacillus megaterium*, *Bacillus subtilis*, *Bacillus stearothermophilus*, *Staphylococcus aureus*, *Streptococcus faecalis*), 12 Gram-negative bacteria (*Escherichia coli*, *Shigella dysenteriae*, *Proteus vulgaris*, *Proteus mirabilis*, *Shigella flexneri*, *Klebsiella pneumoniae*, *Pseudomonas aeruginosa*, *Salmonella typhi*, *Morganella morganii*, *Enterobacter aerogenes*, *Citrobacter freundii*, *Enterobacter cloacae*) and two yeasts from *Candida* species (*Candida albicans* and *Candida glabrata*) were used in this study. Three *Bacillus* species were provided by “l'institut Appert de Paris”, *Bacillus cereus* by the A.F.R.C Reading Laboratory of Great Britain. The other strains were clinically isolated from patients in the Centre Pasteur de Yaoundé-Cameroun (health institution). They were then maintained on agar slant at 4 °C in the LMP where the antimicrobial tests were performed. The strains were activated at 37 °C for 24 h on nutrient agar (NA) (bacteria) or NA supplemented with Bromocresol purple (*Bacillus* strains), Sabouraud glucose agar (fungi). The nutrient broth (NB) was used to determine, respectively, the sensitivity and the minimal inhibition concentration (MIC) of all samples against the tested pathogens.

#### 3.4.2. MIC determination: microdilution method

The MICs of compounds **1**, **2** and reference antibiotics (RA) were evaluated against the pathogens. The inocula of microorganisms were prepared from 12 h both culture and the suspensions were adjusted to 0.5 Mc Farland turbidity. The tested samples were first dissolved in dimethylsulfoxide (DMSO), then in NB to the highest dilution of 78.12  $\mu\text{g/ml}$ . Then, serial twofold dilutions were made in a concentration range from 0.075 to 78.12  $\mu\text{g/ml}$  in the

96 wells microplate. The final concentration of DMSO was less than 1% v/v. MIC values of the tested samples against pathogens were determined based on the microdilution method (Zgoda and Porter, 2001). Gentamycin (Bacteria) and Nystatin (Yeasts) diluted prior in water were used as reference antibiotics. Negative control was made with DMSO.

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