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Crassiflorone, a new naphthoquinone from Diospyros crassiflora (Hien)

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Abstract—A new naphthoquinone, 11-hydroxy-1,9-dimethyl-6H-naphtho[2',3':4,5]furo[3,2-c]chromene-6,7,12-trione, named crassiflorone, was isolated from the stem bark of *Diospyros crassiflora* together with the known compounds plumbagin, cyclocanaliculatin, gerberinol, lupeol, lupenone and betulinic acid. The structures of the compounds were established on the basis of 1D and 2D NMR spectroscopic data, as well as co-TLC with authentic samples. Some of the above compounds exhibited significant antimicrobial activity against bacteria and yeasts. $© 2006 Elsevier Ltd. All rights reserved.$

The genus Diospyros contains as many as 350 species, some of which are widely used for the treatment of many ailments.^{[1](#page-3-0)} Due to its use in folk medicine, the above taxon has attracted the attention of many scientists who have investigated the chemical constitution of many species and reported the presence of various classes of compounds including hydrocarbons, terpenes, naphthoquinones and coumarins.^{[1–4](#page-3-0)} In our systematic search for antimicrobial agents from the plant kingdom, we examined the chemical constitution of Diospyros crassiflora (a tree up to 25 m high and 80–120 cm width). 5 We report here a phytochemical investigation of the stem bark of D. crassiflora Hien collected from Mount Fébé in the Central Province of Cameroon in May 1998, as well as the antimicrobial activities of some of the above compounds against bacteria and yeasts. A voucher specimen (4924/SRFK) has been deposited at the Cameroon National Herbarium, Yaoundé.

The methylene chloride extract of the stem bark of D. crassiflora afforded, after repeated chromatography,

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seven compounds among which are, a new naphthoquinone derivative 1, together with two naphthoquinone derivatives, plumbagin 3^6 3^6 and cyclocanaliculatin 4^1 4^1 , a coumarin, gerberinol 5, [7](#page-3-0) three triterpenes, lupeol, lupenone and betulinic acid.[1,8](#page-3-0)

Keywords: Diospyros crassiflora; Ebenaceae; Crassiflorone; Bacteria; Yeasts.

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Crassiflorone 1 was obtained as the major component of the dichloromethane extract of the stem bark of D. crassiflora, and isolated by flash chromatography (solvent CH_2Cl_2 –MeOH) using silica gel. It was soluble in dimethyl sulfoxide and crystallized from CH_2Cl_2 – MeOH as orange crystals, mp $230-232$ °C (uncorrected). The EI-MS suggested the molecular formula of crassiflorone to be $C_{21}H_{12}O_6$ (*m/z* 360.1 [M]⁺, 99.6%) with 16° of unsaturation. The UV spectrum showed absorption maxima at λ_{max} (MeOH) 283 and 341 nm (log ε 2.45 and 2.53, respectively) typical of 5-methyl-4-oxy-coumarin/2-oxychromenones,^{[3,9,10](#page-3-0)} and at 414 nm (log ε 2.63) typical for the juglone derivatives.^{[3,10](#page-3-0)} The IR spectrum displayed a broad band at v_{max} (KBr) 3421 cm⁻¹ (chelated OH), a band at $1597-1634$ cm⁻¹ (aromatic ring and quinonoid carbonyl), and a further band at 1678 cm^{-1} (coumarin carbonyl).^{[9,11](#page-3-0)} The ¹H and ¹³C spectral data, by means of COSY, DEPT, HMQC, HMBC and NOESY experiments, allowed a full assignment of the NMR signals and led to structure 1.

The analysis of the 13 C NMR spectral data of 1 (Table 1) through DEPT revealed 21 carbon atoms among which are two methyl groups (δ_c 21.5 and 23.2), five methine and 14 quaternary carbon atoms. Of the 14 quaternary carbon atoms, two were those of α , β -unsaturated ketones (δ _C 181.4 and 178.8), one was a chromenone carbonyl $(\alpha, \beta$ -unsaturated lactone) which resonated at δ_c 158.0 and four were linked to oxygen atoms in view of their deshielded chemical shifts at δ_C 153.9, 160.2, 156.2 and 165.5. Among the four oxygenated quaternary carbon atoms, one was linked to a

hydroxyl group and thus accounted for the 12 hydrogen atoms present in compound 1.

High field ¹H NMR (DMSO- d_6 , 500 MHz) signals assigned with the aid of ${}^{1}H-{}^{1}H$ COSY and NOESY (Fig. 1) experiments, showed one methyl group at $\delta_{\rm H}$ 2.46 and two aromatic protons at $\delta_{\rm H}$ 7.23 (br s, H-10) and 7.47 (br s, H-8), which were typical of a 7-methyljuglone unit without protons on $C-2$ and $C-3$, 10,12 10,12 10,12 and justified the presence of unit A [\(Fig. 2\)](#page-2-0) in compound 1. From the HMBC experiment [\(Table 2\)](#page-2-0), the proton at δ_H 7.23 showed correlations with C-11 (δ_C 160.2), C-11a (δ_c 113.4) and C-15 (δ_c 21.5) ([Table 2\)](#page-2-0). Also, the proton at $\delta_{\rm H}$ 7.47 showed correlations with C-7 (δ _C 181.4), C-7a (δ _C 132.8), C-10 (δ _C 122.8), C-11a and C-15. The signal at δ_H 11.91 (s, 1H) which exhibited HMBC correlations with C-10, C-11 and C-11a appeared to be that of a chelated hydroxyl group.

 1 H $-{}^{1}$ H COSY and NOESY (Fig. 1) experiments, led to the assignment of the C-methyl group appearing at δ_H 2.61 to be that of a coumarin/chromenone system and

Figure 1. NOESY correlations of compound 1.

Figure 2. Unit A.

Table 2. ¹*J* (from HMQC), ²*J* and ³*J* correlations of crassiflorone (1)

Proton $\delta_{\rm H}$ (ppm)		carbon	Position ¹ J-Correlated ² J- and ³ J-correlated carbons
11.91	11-OH		C-10, C-11, C-11a
7.47	8	119.5	C-7, C-7a, C-10, C-11a, C-15
7.45	3	132.1	C-1, C-2, C-4, C-4a
7.23	10	122.8	C-8, C-11, C-11a, C-15
7.11	2	127.5	C-1, C-3, C-4, C-13b, C-14
7.09	4	115.2	C-2, C-3, C-4a, C-13b
2.61	14	23.2	C-1, C-2, C-13b
2.46	15	21.5	$C-8$, $C-9$, $C-10$

must therefore, be located on a benzene ring. The signals for the three remaining protons occurred at $\delta_{\rm H}$ 7.09 (br d, $J = 8$ Hz, H-4), 7.11 (br d, $J = 8$ Hz, H-2) and 7.45 (t, $J = 8$ Hz, H-3), the above pattern being consistent only with a 1,2,3-trisubstituted benzene ring of an oxychromenone (5-methyl-4-oxycoumarin) and justified the presence of unit B (Fig. 3) in compound 1. The signal at $\delta_{\rm H}$ 7.45 appeared as a triplet due to coupling with H-2 and H-4, and the link to the carbon at δ_c 132.1 (C-3) was confirmed by 2D experiments. From the HMBC experiment, H-3 showed correlations with C-1 (δ C 137.4), C-2 (δ _C 127.5), C-4 (δ _C 115.2) and C-4a (δ _C 153.9) (Table 2), while, H-2 showed correlations with C-1, C-3 (δ _C 132.1), C-4, C-13b (δ _C 114.5) and C-14 $(\delta_{\rm C} 23.2).$

The oxygen atom of unit B can be linked either to the C-2 or C-3 carbon of unit A. On this basis, the structure of crassiflorone could be either 1 or 2; however, the quinonoid carbonyl resonance at C-7 (δ _C 181.4) of compound 1, which is close in value to the coumarin carbonyl group as found for the quinone carbonyl C-12 (δ C 180.6) of cyclocanaliculatin 4 ([Table 1](#page-1-0)) and the HMBC correlation between H-8 and C-7 in structure 1 favour the C-3 linkage (leading to 1). Also, the strong shielding effect on C-12 (δ _C 178.8) in compound 1 which normally appears at δ _C 189.7 (C-4) in a 7-methyljuglone (unit A with protons on C-2 and C-3)^{[13](#page-3-0)} and at δ_C 181.0 (C-7 with an hydroxylated methylene at the beta-position)

in a cyclocanaliculatin (Table 2) can been seen as the result of the β -effect of the furan oxygen.^{[14](#page-3-0)} From the above data, 1 is considered to be the structure of crassiflorone. These data are also evidence for the linkage of unit B on C-2 and C-3 of unit A as shown in structure 1 in which the resulting furan ring constitutes the remaining degree of unsaturation. Crassiflorone gave fragmentations on EI-MS compatible with the suggested structure, notably the ion at m/z 176.1 [M+2H-186]⁺ (50%) attributed to a 5-methyl-4-coumarin fragment; ions at m/z 134.1 $[M-226]^+$ (69%), 332.1 $[M-CO]^+$ (22%) and 106.1 $[M-254]^+$ (17%) were other important ions which are typical internal cleavages of quinonoid and coumarol rings.[7,15](#page-3-0)

The antimicrobial activity of 1 and 5 was studied using the microdilution assay on a total of six microbial cultures belonging to four aerobic bacterial species (Escherichia coli LMP0101U, Shigella dysenteriae LMP0208U, Staphylococcus aureus LMP0206U, Salmonella typhi LMP0209U) and two Candida species (Candida albicans LMP0204U and Candida krusei LMP0311U). These strains were clinically isolated from the urogenital discharges of patients in the Centre Pasteur du Cameroun health institution and monitored in the Laboratory of Applied Microbiology and Molecular Pharmacology (LMP) of the University of Yaoundé I. The strains were activated at 37 \degree C for 24 h on nutrient agar (NA), Sabouraud Glucose Agar (SGA) (fungi).

The antimicrobial activity was evaluated on the basis of the minimal inhibition concentration (MIC). The inocula of microorganisms were prepared from 12 h broth culture and the suspensions were adjusted to 0.5 McFarland turbidity. The tested compounds were first dissolved in dimethyl sulfoxide 10% to the highest dilution $(625 \mu g/mL)$, and serial twofold dilutions were made in a concentration range from 2.44 to $625 \mu g$ / mL in the 96-wells microplate containing nutrient broth. MIC values of the tested compounds against the above pathogens were determined according to the microdilution method.[16](#page-3-0) Gentamycin (bacteria) and nystatin (yeasts) diluted in water were used as reference antibiotics. As shown in Table 3, compound 5 exhibited the strongest activity against Shigella dysenteriae and Salmonella typhi whilst compound 1 showed weak activities against the tested microbials (Table 3). The antimicrobial activity of compound 3 was described in our previous work on *D. canaliculata.*^{[3](#page-3-0)}

Table 3. In vitro antimicrobial activity of compounds 1 and 5

Tested microrganisms	1 ^a	ζ^a	GM/N^b
Bacteria			
Escherichia coli	19.53	19.53	10
Shigella dysenteriae	78.12	4.88	5
Salmonella typhi	19.53	4.88	10
Staphylococcus aureus	78.12	19.53	10
Yeasts			
Candida albicans	78.12	39.06	30
Candida krusei	78.12	39.06	30

^a Results of the MIC recorded as the mean of triplicated experiments. ^b GM: gentamycin; N: nystatin.

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