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To cite this article: Larissa M. Magnibou, Steven C. N. Wouamba, Abel J. G. Yaya, Judith F. Mbougna, Guy S. S. Njateng, Ghislain W. Fotso, Celine Henoumont, Sophie Laurent & Talla Emmanuel (17 Nov 2023): Chemical profiling by UHPLC-Q-TOF-HRESI-MS/MS and antibacterial properties of *Entada abyssinica* (Fabaceae) constituents, Natural Product Research, DOI: [10.1080/14786419.2023.2280171](https://doi.org/10.1080/14786419.2023.2280171)

To link to this article: <https://doi.org/10.1080/14786419.2023.2280171>

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 Published online: 17 Nov 2023.

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Chemical profiling by UHPLC-Q-TOF-HRESI-MS/MS and antibacterial properties of *Entada abyssinica* (Fabaceae) constituents

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ABSTRACT

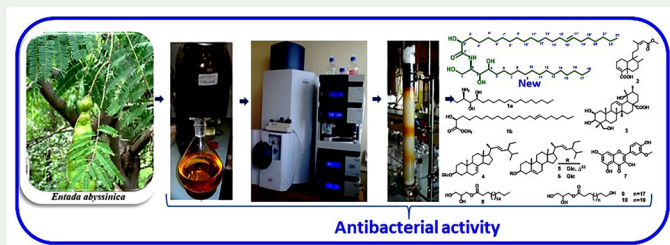
A rapid untargeted UHPLC-Q-TOF-ESI-MS/MS-Based metabolomic profiling of the medicinal plant *Entada abyssinica* was performed. A total of 18 metabolites were detected, of which 10 could not be identified. Based on this result, an extensive chemical investigation of the CH₂Cl₂-MeOH (1:1) extract of this plant was carried out, leading to the isolation of a new ceramide, named entadamide (1), together with nine known compounds: monomethyl kolavate (2), 24-hydroxytormentic acid (3) chondrillasterol (4), 3-O-β-D glucopyranosylstigmasterol (5), 3-O-β-D glucopyranosylsitosterol (6), quercetin 3'-methylether (7), 2,3-dihydroxypropyl icosanoate (8), 2,3-dihydroxypropyl 23-hydroxytricosanoate (9) and 2,3-dihydroxypropyl 24-hydroxytetracosanoate (10). Their structures were elucidated by the analyses of their spectroscopic and spectrometric data (1D and 2D NMR, and HRESI-MS) in comparison with those reported in the literature. Furthermore, the crude extract and some isolated compounds were tested against non-ciprofloxacin resistant strains *viz.* *Pseudomonas aeruginosa* (ATCC 27853), *Escherichia coli* (ATCC 25922), *Samonella thyphi* (ATCC 19430) and *Samonella enterica* (NR4294). The tested samples demonstrated significant activity against all the tested bacteria (MIC values: 3.12–12.5 μg/mL).

ARTICLE HISTORY

Received 6 April 2023
Accepted 30 October 2023

KEYWORDS

Entada abyssinica;
metabolomics profiling;
ceramide; antibacterial
activity



1. Introduction

In Africa, alongside malaria, infections induced by bacterial pathogens are one of the major causes of mortality. Water-borne and enteric diseases come to the fore in Cameroon (Wouamba et al. 2020b; Nguiam et al. 2021; Tali et al. 2022). The access to conventional medicines being expensive for the Cameroonian indigenous population, they rely on medicinal plants for their primary health care (Kuate and Efferth 2010; Njanpa et al. 2021; Kemgni et al. 2021; Magnibou et al. 2022). *Entada abyssinica* Steud. ex A. Rich (Fabaceae), the subject of this study, is listed among these plants. It is a small to medium-sized, deciduous tree, 3–15 m high, with a flat, spreading crown; bark grey to reddish, slightly fissured, flaking off in irregular patches; slash pink with streaks of red; branchlets pendulous, glabrous or sometimes pubescent (Orwa et al. 2009; Magnibou et al. 2022). It is found throughout tropical Africa where it is used for the treatment of miscarriage and against fever (Dzoyem et al. 2017). A decoction of the bark is taken against cough, chronic bronchial engorgement, rheumatic and abdominal pains (Haile and Delenasaw 2007). In general people with low income such as farmers, people of small isolated villages and native communities use folk medicine for the treatment of common enteric infections (Fabricant and Farnsworth 2001). Bacterial infectious diseases caused by these multidrug resistant strains remain the leading cause of death. Thus people are turning their attention to alternative novel antimicrobial agents to combat such pathogens (Jain et al. 2011). Some pharmacological properties of *E. abyssinica* have been previously reported, including anti-inflammatory, antileishmanial, antimicrobial and antioxidant (Olajide and Alada 2001; Nyasse et al. 2004; Teke et al. 2011; Melong et al. 2014; Dzoyem et al. 2017; Magnibou et al. 2022; Mbougna et al. 2022). Previous chemical investigations of *E. abyssinica* have reported the presence of secondary metabolites such as flavonoids, diterpenoids, triterpenoids, steroids, steroidal saponins, phenantrenes, cerebrosides, kolavivic acid derivatives and fatty acids (Freiburghaus et al. 1998; Tchinda et al. 2007; Melong et al. 2014; Magnibou et al. 2022).

In our continuous search for secondary metabolites with biological importance from Cameroonian medicinal plants, we undertook the investigation of the CH₂Cl₂-MeOH (1:1) extract of the stem bark of *E. abyssinica*. Considering the vast potential of plants as sources of antimicrobial drugs, the objective of this research was to perform rapid, untargeted LC-MS/MS-based metabolomic profiling of this plant by LC-MS/MS, followed by isolation of the compounds and finally evaluation of their antibacterial activity. The final goal being the discovery of new molecules of interest for future generations.

2. Results and discussion

2.1. Qualitative determination of compounds contents of *E. abyssinica* using UHPLC(DAD)-Q-TOF-ESI-MS/MS

The present study, which contributes to phytochemical knowledge of the *Entada* genus, evaluated the chemical composition of *Entada abyssinica* stem bark and screened it for antibacterial activity.

Stem bark extract of *E. abyssinica* was extracted using the mixture of CH₂Cl₂/MeOH (1:1, v/v). It is worth noting that this solvent system was chosen because the CH₂Cl₂/MeOH (1:1, v/v) extract displayed the best antibacterial activity after micro-extraction compared to the hydroethanolic extract. The resulting crude extract was firstly subjected to liquid chromatography coupled to mass spectrometry (LC–MS/MS).

Figure S1 and Table 1 present the chromatographic profiles of the extract and the main peaks corresponding to compound detected by UHPLC-Q-TOF-(HR)ESI-MS/MS (positive ionisation mode). The blue curve represents the profile of the crude extract studied and the black curve represents the profile of the blank (see Figure S1). The superposition of these two chromatographic traces revealed the presence of several compounds. Eighteen of them showed good ionisation and fragmentation and therefore were detected. The putative identification of these compounds present in crude extract was performed according to the fragmentation profiles of the spectra generated in comparison with the data from the literature of compounds previously isolated from the *Entada* genus, and available in the databases (SciFinder, NIST/EPA/NIH Mass Spectral Library (NIST 14) and Mass Bank of North America (MoNA)) (Table 1., supplementary data). This strategy led to the identification of 8 compounds (2, 11–17) among the 18 detected by LC-MS (see supplementary data, Figure S2). However, compound 17, diheptylphthalate, although isolated from this plant by Melong et al. (2014), could be a contaminant. In fact, phthalates are increasingly being considered as contaminants in plant extracts (Bianco et al. 2014; Venditti 2020; Thiemann 2021). The other compounds we have not identified were simply not found in the databases. They could be new classes of natural compounds present in the *Entada* genus in very low proportions.

Table 1. Metabolites assigned in *E. abyssinica* stem bark extract via UHPLC-Q-TOF-ESI-MS/MS in positive ion mode.

N°	Tr (min)	[M + Na] ⁺ or [M + H] ⁺		MS/MS	Pseudo-molecular ion formula	Name of structure
		Exp.	Calcd.			
1	0.31	198.0535	198.0525	60	C ₁₀ H ₉ NO ₂ Na ⁺	Acetic1H-Indole-3-acid (11)
2	1.58	226.1812	223.1802	125; 110	C ₁₃ H ₂₄ NO ₂ ⁺	2-methyl-N-(4-methylpentan-2-yl)hexa-2,4-dienamide (12)
3	1.68	246.1500	246.1489	60; 195	C ₁₅ H ₂₀ NO ₂ ⁺	2-(5-(tert-Butyl)-2-methyl-1H-indol-3-yl) acetic acid (13)
4	2.80	381.0590	381.0581	–	C ₁₈ H ₁₄ O ₈ Na ⁺	Not Identified
5	2.92	263.1651	263.1642	–	C ₁₆ H ₂₃ O ₃ ⁺	Not Identified
6	3.14	345.2407	345.2400	–	C ₂₀ H ₃₄ O ₃ Na ⁺	Not Identified
7	3.28	316.2849	316.2846	288	C ₁₈ H ₃₈ NO ₃ ⁺	Not Identified
8	3.41	318.3008	318.3003	300	C ₁₈ H ₄₀ NO ₃ ⁺	Not Identified
9	3.80	303.1525	303.1567	246	C ₁₆ H ₂₄ O ₄ Na ⁺	parathyrinsoidin G (14)
10	3.90	373.2355	373.2349	–	C ₂₁ H ₃₄ O ₄ Na ⁺	Not Identified
11	4.04	499.3286	499.3265	–	C ₂₇ H ₄₇ O ₈ ⁺	Not Identified
12	4.12	329.2457	329.2451	242; 236	C ₂₀ H ₃₄ O ₂ Na ⁺	Ethyl linolenate (15)
13	4.55	371.2155	371.2193	326; 350	C ₂₁ H ₃₂ O ₄ Na ⁺	Monomethyl kolavate (2)
14	4.87	353.2407	353.2087	224	C ₂₁ H ₃₀ O ₃ Na ⁺	Not Identified
15	5.06	325.2152	325.2138	258; 244	C ₂₀ H ₃₀ O ₂ Na ⁺	4-Epicommunic acid (16)
16	5.19	377.2674	377.2662	258	C ₂₁ H ₃₈ O ₄ Na ⁺	Not Identified
17	5.34	367.0433	367.0424	–	C ₁₇ H ₁₂ O ₈ Na ⁺	Not Identified
18	5.45	385.2363	385.2349	385	C ₂₂ H ₃₄ O ₄ Na ⁺	Diheptylphthalate (17)

While comparing the retarding factor, the experimental and calculated ion peak masses and the molecular formula, the result revealed that the CH₂Cl₂/MeOH (1:1, v/v) extract of *E. abyssinica* contained several secondary metabolites. This result paved the way for the isolation of secondary metabolites of this extract. This approach constitutes on the one hand an interesting lead towards the discovery of new compounds and on the other hand the evaluation of their antimicrobial activity could shed light on the ethno-pharmacological use of this plant.

2.2. Isolation and structure elucidation

After analysis of the LC/MS data, the crude extract (~30.5g) was subjected to repeated silica gel and Sephadex LH-20 column chromatography (CC) and recrystallization to afford a previously undescribed ceramide named Entadamide [(**1**), 8.40 mg], together with nine known compounds viz, (8S, 13E)-kolavivic acid 15-methyl ester [(**2**), 6 mg] (Tchinda et al. 2007), 24-hydroxytormentic acid [(**3**), 10.0 mg] (Houghton and Lian 1986), Chondrillasterol [(**4**), 7 mg] (Bergmann and McTigue 1948), mixture of 3-O-β-D glucopyranosylstigmasterol/3-O-β-D glucopyranosylsitosterol [(**5/6**), 15.0 mg] (Wouamba et al. 2020a), quercetin 3'-methylether [(**7**), 9.00 mg] (Fu et al. 2005), 2,3-dihydroxypropyl icosanoate [(**8**), 7.00 mg] (Rocchetti and O'Callaghan 2021), 2,3-dihydroxypropyl 23-hydroxytricosanoate [(**9**), 12.0 mg] and 2,3-dihydroxypropyl 24-hydroxytetracosanoate [(**10**), 8.00 mg] (Sharma et al. 2016) (Figure 1).

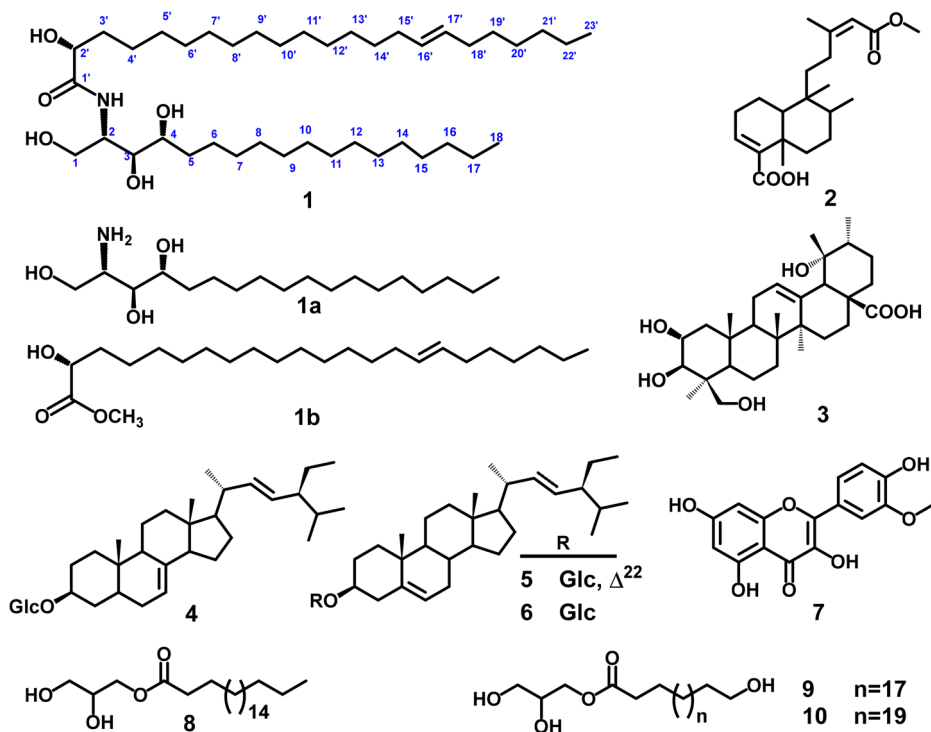


Figure 1. Structures of compounds **1–10** isolated from *E. abyssinica*.

The structures of the known compounds were identified by comparison of their spectroscopic and spectrometric data with those reported in the literature.

Compound **1** was obtained as a white powder. Its molecular formula, $C_{41}H_{81}NO_5$ was established from its HRESI-MS spectrum (Figure S3), showing the pseudo-molecular sodium adduct peak $[M+Na]^+$ at m/z 690.6024 ($C_{41}H_{81}NO_5Na^+$, calcd. 690.6007), indicating two degrees of unsaturation. Its IR spectrum (Figure S4) showed characteristic absorption bands for free OH groups ($3329\text{--}3215\text{cm}^{-1}$) and an amide group (1623cm^{-1}) (Tsamo et al. 2021; Wonkam et al. 2020; Kamdoun et al. 2022). The structure of compound **1** was fully assigned after careful analyses of its ^1H and ^{13}C NMR, $^1\text{H}\text{--}^1\text{H}$ COSY, HMQC, HMBC, tandem MS spectra and methanolysis reaction. Indeed ^1H -NMR spectrum (Figure S5, Table S1) of compound **1** showed the presence of an NH group at δ_{H} 8.60 (1H, d, $J=9.0$), two signals of one proton each in the form of two multiplets between δ_{H} 5.18–5.48 and 5.48–5.55 (2H, m, H-16' and H-17') suggesting the presence of olefinic protons in the structure of compound **1** (Wouamba et al. 2020a; Kamdoun et al. 2022). A broad signal centred at δ_{H} 1.52–1.20 (methylene protons), a distorted triplet at δ_{H} 0.88 (6H, t, $J=7.0$, H-18 and H-23') (two terminal methyl groups) and two oxymethylene protons at δ_{H} 4.48 (dd, $J=10.7$, 5.0, H-1a) and 4.35 (dd, $J=10.7$, 4.8, H-1b), were indicative of a ceramide moiety (Simo et al. 2008; Fouatio Feudjou et al. 2022). This was supported by the presence of signals of a nitrogen-attached sp^3 carbon at δ_{C} 51.5 (C-2) and characteristic olefinic and oxymethine carbons at δ_{C} 130.6 (C-16'), 130.5 (C-17'), 76.5 (C-3), 72.2 (C-4), and 72.6 (C-2'). Further analysis of the ^{13}C NMR spectrum (Figure S6, Table S1) showed signals for oxymethylene carbon at δ_{C} 61.3 (C-1), and an amide carbonyl carbon at δ_{C} 175.0 (C-1'). In addition, conjunction of ^{13}C -NMR and HSQC spectra (Figure S6–S7) confirmed the signal of a nitrogen-attached sp^3 carbon at $\delta_{\text{H/C}}$ 5.14/51.5 (C-2), two diastereotopic protons of an oxymethylene at $\delta_{\text{H/C}}$ 4.35/61.3 (C-1) and 4.35/61.3 (C-1), as well as three oxymethine protons at $\delta_{\text{H/C}}$ 4.38/76.5 (C-3), 4.65/72.2 (C-4) and 4.31/72.6 (C-2') respectively.

The HMBC correlations (Figure S8) of protons at δ_{H} 4.48/4.35 (H-1a/H-1b) to carbons δ_{C} 52.5 (C-2), and 76.5 (C-3); at δ_{H} 5.14 (H-2) to δ_{C} 61.3 (C-1), 76.3 (C-3), and 175.0 (C-1'); and at δ_{H} 4.31 (H-2') to δ_{C} 175.0 (C-1'), allowing us to locate the hydroxy groups at positions C-1, C-3, C-4, and C-2', respectively and to conclude that, the two chains (acid and basic) are linked. Further important HMBC (Figure S8 and S10a) correlations were observed between δ_{H} 5.54 (H-16' and H-17') and δ_{C} 33.5 (C-18'), δ_{H} 1.97 (H-18') and δ_{C} 31.8 (C-19'), and δ_{H} 1.97 (H-19') with δ_{C} 31.8 (C-21') and 14.0 (C-23') suggesting that the double bond was located on the long fatty acid chain. Moreover, $^1\text{H}\text{--}^1\text{H}$ COSY correlations (Figures S9 and S10a) observed between H-1a/H-1b and H-2, H-2 and H-3, H-3 and H-4 allowed us to confirm the position of four hydroxy groups. All these evidences above, and by comparison with data of published related compounds, confirmed that **1** is a ceramide (Wouamba et al. 2020a; Kamdoun et al. 2022). The lengths of the fatty acid and sphingosine moiety as well as the position of double bond were deduced by the methanolysis using 0.9N, HCl/MeOH, at 70°C for 20H (Simo et al. 2008) to yield sphingosine (**1a**) and fatty acid methyl ester (**1b**) (Figure S11). Specifically, the peak at m/z 218.2 $[M_{\text{LCB}}+H]^+ = 318$ corresponding to molecular formula $C_{18}H_{40}NO_3$ was attributable to the long chain base (**1a**) and do not contain degree of unsaturation (Figure S12).

Furthermore, this molecular formula of sphingosine suggested that the olefinic moiety is located in the long chain acid (LCA). The molecular formula of the methyl ester fatty acid ($C_{24}H_{46}O_3$) was deduced from the molecular formula of compound **1** by subtracting the molecular formula of the sphingosine moiety. The length of the long chain acid (LCA) was confirmed by the presence of some characteristic fragment ions are observed at m/z 389 $[CH_3(CH_2)_5CH=CH(CH_2)_{13}CH(OH)CONHNa]^+$, m/z 369 $[CH_3(CH_2)_5CH=CH(CH_2)_{13}CH(OH)CONH_4]^+$ and m/z 346 $[CH_3(CH_2)_5CH=CH(CH_2)_{13}CH(OH)Na]^+$. Furthermore the MS analysis showed the ions peak of the fragments from the allylic cleavages $[M+NH_4-C_6H_{13}]^+$ at m/z 600 and $[M+NH_4-C_8H_{14}]^+$ at m/z 571 confirming the $\Delta^{16'}$ location of the double bond (Figure S10b). The (*E*)-configuration was deduced from the chemical shift values at δ_c 33.8 and 33.5 of its allylic carbons C-15' and C-18' (Table S1). (Simo et al. 2008; Wouamba et al. 2020a; Tsamo et al. 2021). Generally, the stereochemistry of the olefinic functional group is assigned from the coupling constant values of $J=16$ Hz, and 8 Hz for *trans*- and *cis*-configurations, respectively. Nevertheless, if these coupling constant values are observed as multiplet, this could also be assigned by ^{13}C chemical shift values of allylic carbons, $\delta_c > 30$ for *trans*-configuration and $\delta_c \leq 27$ ppm for *cis*-configuration (Simo et al. 2008; Wouamba et al. 2020a; Tsamo et al. 2021; Kamdoum et al. 2022). Based on their biosynthetic evidence and as observed in all the naturally occurring sphingolipids, the absolute configurations of C-2 and C-3 were assigned theoretically as 2*S*, 3*S*, respectively (Wonkam et al. 2020; Kamdoum et al. 2022). The comparison of the data of compound **1** with those of previously reported compounds (Wouamba et al. 2020a; Wonkam et al. 2020; Kamdoum et al. 2022) further supported the theoretical assignment of absolute configurations *R* and *S* to C-4 and C-2', respectively. From the spectroscopic analysis above, the structure of **1** was unambiguously determined as (2*S*,2'*R*,3*S*,4*R*,16*E*)-*N*-[2'-hydroxytricos-16-enoyl]-2-amino-octadecane-1,3,4-triol and trivially named entadamide.

Compound **2** was obtained as a white powder. Its molecular formula, $C_{41}H_{81}NO_5$ was established from its HRESI-MS spectrum (Figure S3), showing the pseudo-molecular sodium adduct peak $[M+Na]^+$ at m/z 690.6024 ($C_{41}H_{81}NO_5Na^+$, calcd. 690.6007), indicating two degrees of unsaturation. Its IR spectrum (Figure S4) showed characteristic absorption bands for free OH groups ($3329-3215\text{cm}^{-1}$) and an amide group (1623cm^{-1}) (Tsamo et al. 2021; Wonkam et al. 2020; Kamdoum et al. 2022). The structure of compound **1** was fully assigned after careful analyses of its 1H and ^{13}C NMR, $^1H-^1H$ COSY, HMQC, HMBC, tandem MS spectra

Compound **2** is a methyl ester isolated from the methanolic extract. Although its acidic (unmethylated) structural analogue was not detected by UHPLC-(DAD)-Q-TOF-HRESI-MS/MS analysis, this compound **2** could be an artefact. Indeed, compound **2** could be formed during the extraction process by derivatisation of its unmethylated analogue, as the methanol/dichloromethane mixture was used for extraction. However, we do not have sufficient evidence at this stage to conclude with certainty whether this is an artefact or a natural compound from *Entada abyssinica*. But it is interesting to note that this compound **2** was formerly isolated by Nyasse et al. (2004) from the pure methylene chloride extract of this plant. In addition, Tchinda et al. (2007) reported the isolation of compound (**2**) and its unmethylated analogue from the acetone extract of *E. abyssinica*.

varied between 1 and 4. According to Marmonier (1990), pure compounds exert two types of activities: a bacteriostatic ($MBC/MIC \geq 4$) and bactericidal activity ($MBC/MIC \leq 4$). So, globally, all the compounds exerted bactericidal activities against the tested bacteria. These results are in line with those obtained in previous studies (Magnibou et al. 2022).

The obtained results support the use of *E. abyssinica* in alternative medicine to treat diseases of bacterial origin with compound **3** being a possible promising drug candidate against bacterial infections.

3. Experimental (supplementary data)

3.6. Entadamide (1)

White powder; IR (KBr): 3329 cm^{-1} , 3215 cm^{-1} and 1623 cm^{-1} ; HRESI-MS: $[M+Na]^+$ at m/z 371.2155 ($C_{41}H_{81}NO_5Na^+$, calcd. 371.2193); 1H -NMR (600 MHz, C_5D_5N) & ^{13}C -NMR (150 MHz, C_5D_5N) see Table S1.

3.7. Monomethyl kolavate (2)

White powder; HRESI-MS: $[M+Na]^+$ at m/z 690.6024 ($C_{21}H_{32}O_4Na^+$, calcd. 690.6007); 1H -NMR (600 MHz, C_3D_6O) δ_H 1.60–1.33 (m, 2H, H-1), 2.11–2.01 (m, 1H, H-2), 6.36 (1H; s, H-3), 1.67–1.42 (m, 2H, H-6), 1.63–1.38 (m, 2H, H-7), 3.45 (s, 1H, H-8), 1.14 (s, 1H, H-10), 1.19–1.21 (m, 2H, H-11), 1.97–2.10 (m, 2H, H-12), 5.54 (1H; d; $J=1.3\text{ Hz}$, H-14), 1.14 (3H; s, H-16), 0.79 (3H; d; $J=0.8\text{ Hz}$, H-17), 0.82 (3H; s, H-19), 1.58 (3H; s, H-20), 3.45 (3H; s, OCH₃); ^{13}C -NMR (150 MHz, C_3D_6O) δ_C 17.9 (C-1), 26.8 (C-2), 136.5 (C-3), 142.4 (C-4), 37.5 (C-5), 37.6 (C-6), 25.2 (C-7), 34.9 (C-8), 37.5 (C-9), 45.0 (C-10), 29.5 (C-11), 34.0 (C-12), 161.2 (C-13), 115.0 (C-14), 166.9 (C-15), 16.9 (C-16), 14.3 (C-17), 166.9 (C-18), 19.8 (C-19), 20.9 (C-20), 15-OMe(50,3)

4. Conclusion

From the total CH_2Cl_2 -MeOH (1:1, v/v) stem bark extract of *Entada abyssinica*, this research allowed the rapid detection of 18 compounds by a metabolomic based on UHPLC-Q-TOF-ESI-MS/MS profiling. Using chromatographic, spectroscopic and spectrometric methods, we isolated and characterised 10 compounds (**1–10**) including entadamide (**1**), a new ceramide. MS fragmentation studies and the use of databases allowed us to identify 8 compounds (**2**, **11–17**) among those detected by LC-MS. Except compound **1**, all the tested samples demonstrated significant or moderate activities against the tested bacteria (MIC values: 3.12–12.5 $\mu\text{g/mL}$). Compound **3** showed significant activity against all the four bacteria with MIC ranging from 1.56 to 6.25 $\mu\text{g/mL}$ and was identified as the most active among all followed by compound **2** which exhibited significant activities against *P. aeruginosa* and *S. enterica*. Compound **3** appears to be a potential candidate for drug discovery against bacterial infections. These results are in accordance with the use of *E. abyssinica* in folk medicine to treat diseases of bacterial origin. *E. abyssinica* may be useful for further investigation in view to develop Improved Traditional Medicines (ITM) to combat bacterial diseases.

Acknowledgement

The authors are thankful to the Department of Organic chemistry of the University of Ngaoundéré for providing some consumables. They also thanks the YaBiNaPA project (Yaounde-Bielefeld Graduate School of Natural Products with Antiparasite and Antibacterial activities) for some MS analysis and the databases made available to them and the NMR and Molecular Imaging Laboratory at University of Mons, Belgium for the NMR analysis of different samples. Finally, the University of Yaoundé I for the antibacterial tests.

Disclosure statement

No potential conflict of interest was reported by the authors.

Funding

This work was supported by the Ngaoundéré University

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