

EXTRACTS FROM *Acmella caulirhiza* POTENTIATE THE ACTIVITY OF CIPROFLOXACIN AGAINST MULTIDRUGRESISTANT BACTERIA EXPRESSING EXTENDED-SPECTRUM BETA-LACTAMASES

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ABSTRACT

Faced with the disturbing phenomenon of bacterial resistance, the search for new efficient and available natural antibacterial agents becomes primordial. In the present study we assessed the antibacterial potential of four extracts (hexane, methylene chloride, ethyl acetate and ethanol) from *Acmella caulirhiza* (a plant that belongs to the *Asteraceae* family) in restoring the activity of three common antibiotics on multidrug-resistant and extended spectrum beta-lactamase-positive bacteria. Extracts were primarily subjected to chemical screening according to standard protocols. The minimal inhibitory concentration (MIC) and minimal bactericidal concentration (MBC) tests were conducted with a modified rapid *p*-iodonitrotetrazolium chloride (INT) colorimetric assay. Subsequently, association tests were performed between these extracts and two beta-

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lactams (Amoxicillin, Ampicillin) on one hand and a fluoroquinolone (Ciprofloxacin) on the other. The phytochemical assays revealed that all crude extracts contained alkaloids, triterpenes and sterols. The ethyl acetate and ethanol extracts further contained all secondary metabolites investigated. Only the hexane extract displayed moderate activity on multidrug-resistant strains (*Klebsiella pneumoniae*, *Klebsiella oxytoca* and *Serratia odorifera*) with the MIC values found between 512 and 1024 $\mu\text{g/mL}$. After combination, improved amoxicillin activity was observed with all extracts though this was observed on a single clinical isolate (*Staphylococcus* spp.). Ciprofloxacin was potentiated on 5 out of 7 (71.43%) multidrug-resistant isolates by all the extracts, including those that did not have any activity when they were tested alone (FIC values: 0.125 through 0.5). These findings suggested that the extracts might contained inhibitors which act on other resistance mechanisms like efflux pumps. In addition, it appeared that constituents that are responsible for the antibacterial activity of the extracts when they were used alone are not the ones that induced the synergistic effects once used in combination. It might be legitimate to pay more attention on these combinations, especially in the caretaking of infections due to extended-spectrum β -lactamases-positive bacteria which also express resistance to Ciprofloxacin.

KEYWORDS: *Acmella caulirhiza*, multidrug-resistance, extended spectrum beta-lactamase, ciprofloxacin.

INTRODUCTION

Beta-lactam antibacterial agents belong to one of the most diverse and widely used families of antibiotics worldwide. The high utilization rates is partly due to their broad spectrum of action, low toxicity, and cost-effectiveness.^[1] With their frequent use in therapeutics, bacterial resistance rapidly developed to become a major health challenge at the global scale, reducing the drug effectiveness of the available therapeutic arsenal.^[2] Though a natural phenomenon^[3] bacterial resistance is known to be exacerbated by inappropriate use of antimicrobial agents in humans, animals and plants.^[4-7] It is estimated that 90% of the overall deaths caused by infectious diseases were recorded in Africa. This rate is partly associated with drug resistance expressed by the microbes incriminated. In fact, multidrug resistance observed in Gram-negative bacteria are responsible for a large proportion of hospital acquired infection (60%) and can be masterminded by sets of bacterial strategies like those involving the production of inactivating enzymes such as beta-lactamases.^[1] which are amongst the most common and most important resistance mechanisms against beta-lactams.^[6,8-10] Broad

spectrum beta-lactamases encompass a group of enzymes that confer resistance to penicillins, 1st, 2nd, 3rd and 4th generations of cephalosporin and Aztreonam. In addition, they do not confer resistance to Cephamecime and Carbapenems and are inhibited by beta-lactamase inhibitors like clavulanic acid.^[1] The selection, growth and spread of antibiotic resistance therefore, render healthcare unaffordable to many populations around the world, especially in low-income countries where traditional medicine practices are common.^[11,12] According to the WHO, more than 80% of African populations rely on traditional medicine for their health issues. Otherwise the higher infectious rates in most low-income countries, especially in Africa couples with natural resources that can be valorised and used sustainably for the neediest communities. Valorising this natural inheritance requires researches for standards protocols that could be recommended to traditional healers in their daily activity. In this vein, several initiatives on medicinal plants are on the way in Cameroon.^[12,13] More specifically, some of these investigations focus on the phytochemical,^[14] the ethnobotanical and pharmacological aspects^[15] of the extracts from *Acmella caulirhiza* (*Asteraceae*). From the pharmacological point of view, promising results against infectious agents have been recorded on bacteria. According to Sinei *et al.*, (2013), *Acmella caulirhiza* is widely spread in Cameroon.^[16] The ethnobotanical survey conducted in the Ndé Division (in West Cameroon) reported the use of parts of *Acmella caulirhiza* in the caretaking of several human disorders, including infectious diseases).^[17] Another one undertaken two years later (Tatah *et al.*, 2015) revealed the potential of its extracts on antibiotic-resistant bacteria that are aetiologies of urinary tract infections and on *Salmonella* Typhi.^[18] To further investigate the antibacterial potential of this plant, the present survey was carried out to address the extended spectrum beta-lactamase (ESBL) inhibitor potential of some of *Acmella caulirhiza* extracts. Confirming this inhibitory potential could guide associating the extracts to conventional antibacterial agents in healthcare for infections due or related to ESBL-positive bacteria. In other words, the present work was designed to evaluate the ability of *Acmella caulirhiza* extracts to restore the activity of common antibiotics by inhibiting broad-spectrum beta-lactamases expression in multidrug-resistant bacteria.

MATERIAL AND METHODS

Plant material and extracts

Material used in this work consisted of the whole *Acmella caulirhiza*, collected in Bangangté (West Cameroon) in June 2015. Identification was thereafter conducted at the National Herbarium under reference Voucher number 42 040 HNC. Each specimen type was air dried

and powdered. For 72 hours at room temperature, 300 g of each powder was used separately in the extraction process (1: 10 m/v) with ethyl acetate, methylene chloride, ethanol and hexane. The extraction products were then concentrated under reduced pressure to have the necessary crude extracts. All extracts were subsequently kept at 4°C until use.

Bacteria strains

Eight bacterial types were chosen either for their frequent involvement in human pathologies and/or ESBL positivity. These included 5 clinical isolates (*Staphylococcus* spp., *Klebsiella oxytoca*, *Bacillus* spp., *Serratia odorifera* and *Citrobacter* spp.) and 3 reference strains (*Klebsiella pneumoniae* ATCC 700603, *Escherichia coli* ATCC 35218 and *Staphylococcus aureus* ATCC 25923). The clinical isolates were provided the Laboratory of Microbiology of the “Université des Montagnes” Teaching Hospital and the reference strains by the Unit of Microbiology and Research in Natural Antimicrobial Agents, University of Dschang.

Pure powers of conventional antibacterial agent consisted of Ampicillin (1 g), Amoxicillin (500 mg), Ciprofloxacin (500 mg) and Amoxicillin + clavulanic acid (625 mg). These were used as references for antibiotics during the investigations. Also, 0.2 % *p*-Iodonitrotetrazolium chloride (INT) and Dimethylsulfoxide (DMSO) (Sigma-Aldrich (St. Quentin Fallavier, France) were used respectively as microbial growth indicator and surfactant for compound dissolution, respectively.

Susceptibility tests

All MIC values throughout the study were assessed with the rapid INT colorimetric assay methods as described by Eloff (1998) with some modifications.^[19,20] All extracts primarily underwent dissolution in DMSO/Mueller Hinton Broth (MHB) to the final concentration of DMSO lower than 2.5%.^[21] The resulting mixture was thereafter, added to Mueller Hinton Broth, and serially diluted two folds (in a 96-wells microplate). One hundred microliters (100 µL) of the bacterial inoculum ($\approx 1.5 \times 10^6$ CFU/mL) prepared in appropriate broth was subsequently added to the preparation.^[20,21] The plates containing these preparations were covered with a sterile sealer, thoroughly mixed by gentle agitation with a plate shaker and incubated for 18 h at 37°C. The assay was repeated three times in each case. Wells containing adequate broth, 100 µL of inoculum and DMSO to a final concentration of 2.5% served as negative control. The MIC were recorded upon completion of incubation with addition of 40 µL of the INT that was followed by additional incubation at 37°C for 30 min. Viability was indicated by the dye change from yellow into pink. MIC was defined as the concentration at

which no color change developed, testifying complete inhibition of microbial growth.^[19] For the MBC, 50 μ L from the well where no growth was reported in the MIC assay was added to 150 μ L of MHB and re-incubated for 48 h in convenient bacterial growth environment. Upon completion, the MBC was regarded as the lowest concentration of extract at which no color change appeared upon addition of INT and incubation, as done for the MIC.

To evaluate the potentiating effect of extracts, their sub-inhibitory concentrations (MIC/2 and MIC/4) were combined with antibiotics and used on the subjected bacterial isolates. The fractional inhibitory concentration (FIC) of each combination was thereafter calculated as the ratio of MIC of antibiotic in combination *versus* the MIC of the antibiotic used alone (MIC_{Antibiotic in combination}/MIC_{Antibiotic alone}) and the association were regarded as synergistic when FIC values were ≤ 0.5 . It was said to be indifferent for FIC values found between 1 and 4 and antagonistic when they were larger than 4.^[22,23] All assays were performed in triplicate.

RESULTS

The yield and phytochemical screening of the four extracts tested provided ranges of indications that could help in attesting their activity. Summary of these results was presented as shown in table I.

Table I. Yield and phytochemical screening of each extract (qualitative estimate).

Yield/ Phytochemical categories	Extracts			
	Ethyl acetate	Methylene chloride	Ethanol	Hexane
Yield	1.46	2.06	2.97	0.65
Alkaloids	+	+	+	+
Anthraquinones	+	+	+	-
Flavonoids	+	-	+	-
Polyphenols	+	-	+	-
Saponins	+	+	+	-
Sterols	+	+	+	+
Tannins	+	+	+	-
Triterpenes	+	+	+	+

(-): Absent; (+): Present;

The overall picture highlighted the presence of all secondary metabolites investigated in the ethyl acetate and ethanol extracts. In addition, alkaloids, sterols, and triterpenes were detected in all extraction products.

***In vitro* antibacterial potential**

When tested, the antibacterial potential varied from one extract to the other on the isolates subjected to the experiment. The MIC, MBC and the MBC/MIC ratio (R) recorded from various essays were tabled as displayed below (Table II).

Table II. Antibacterial activity of the extracts from *Acmella caulirhiza* ($\mu\text{g/mL}$).

Bacterial isolates	Hexane Extract			Methylene chloride extract			Ethyl acetate extract			Ethanol extract		
	MIC	MBC	R	MIC	MBC	R	MIC	MBC	R	MIC	MBC	R
<i>Staphylococcus</i> spp.	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>Citrobacter</i> spp.	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>K. oxytoca</i>	512	1024	2	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>Serratia odorifera</i>	1024	>1024	-	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>Bacillus</i> spp.	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>K. pneumoniae</i> ATCC 700603	512	1024	2	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-
<i>S. aureus</i> ATCC 25923	512	1024	2	1024	-	-	>1024	>1024	-	>1024	>1024	-
<i>E. coli</i> ATCC 35218	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-	>1024	>1024	-

R: MIC/MBC; -: not determined.

It appeared that the hexane extract was most active, as shown in 50% of bacteria under study. The methylene chloride extract exhibited a slightly weak potential on a single strain (*S. aureus* ATCC 25923). Further insights indicated that the MIC values for other extracts were greater than 1024 $\mu\text{g/mL}$, while the bactericidal effects were recorded when the hexane extract was used on *K. oxytoca*, *K. pneumoniae* and *S. aureus* ATCC 25923.

***In vitro* antibacterial potential of common antibiotics on the studied bacterial isolates**

The test was carried out to determine on one hand, the inherent MIC values for the antibiotics in order to attest the resistant phenotype expressed by the tested bacterium, and evaluate the effect of their association with the plant extracts on the same isolate on the other. For this test, antibacterial agents from two families were used: beta-lactams and fluoroquinolones. The findings recorded were displayed as shown in table III.

Table III. Antibacterial activity of some antibiotics ($\mu\text{g/ml}$).

Bacterial isolates	Amoxicillin			Ampicillin			Amoxicillin/ clavulanic acid			Ciprofloxacin		
	MIC	MBC	R	MIC	MBC	R	MIC	MBC	R	MIC	MBC	R
<i>Staphylococcus</i> spp.	4	8	2	>256	>256	-	4	32	8	4	16	4
<i>Citrobacter</i> spp.	>256	>256	-	256	>256	-	32	256	8	32	128	4
<i>K. oxytoca</i>	>256	>256	-	>256	>256	-	32	64	2	32	32	1
<i>S. odorifera</i>	>256	>256	2	>256	>256	-	2	8	4	4	16	4
<i>Bacillus</i> spp.	>256	>256	-	>256	>256	-	32	32	1	4	128	32
<i>K. pneumoniae</i> ATCC 700603	>256	>256	-	32	128	4	16	64	4	8	128	16
<i>S. aureus</i> ATCC 25923	<0.5	>256	-	32	>256	-	<0.5	>256	-	4	4	1
<i>E. coli</i> ATCC 35218	>256	>256	-	128	256	2	32	32	1	16	64	4

R: MIC/MBC; -: not determined.

Tests on *S. aureus* ATCC 25923 (the susceptible reference) yielded the expected results. This strain is known to be devoid of resistance mechanisms. At the same level of susceptibility (MIC and MBC) for *Staphylococcus* spp (a multidrug-resistant-positive isolates), the values recorded were approximately 32-fold higher than those obtained with *S. aureus* ATCC 25923 when Amoxicillin, Amoxicillin/clavulanic acid were used, and 8-fold higher when Ampicillin and Ciprofloxacin were used.

For *K. pneumoniae* ATCC 700603 (an ESBL-positive strain), the MIC values with amoxicillin were similar to those observed for 3 out of the 4 isolates that belonged to the *Enterobacteriaceae* family. With Ampicillin, the MIC for the other isolates were very large compared to the value obtained with the reference ESBL-positive strain (32 for ≥ 256 $\mu\text{g/mL}$). With Ciprofloxacin, all MIC values were equal or larger than four (≥ 4). A significant change was recorded when the MIC's documented with amoxicillin and amoxicillin/clavulanic acid were compared. This significant decrease (by a factor of at least 8) was likely related to the action of ESBL inhibitor (clavulanic acid).

Antibacterial activities of crude extracts in combination with common antibiotics.

When conventional antibiotics were associated with the plant extracts, sets of findings were gathered. Subtle details on these results with respect to bacterial isolates and extracts/antibiotics combinations used were summarized and displayed as shown in table IV.

Table IV. Potential of the extracts/antibiotics combinations

Bacteria isolates	Antibiotics	MIC value	Extracts (µg/mL)							
			Ethanol		Ethyl acetate		Methylene chloride		Hexane	
			MIC/2	MIC/4	MIC/2	MIC/4	MIC/2	MIC/4	MIC/2	MIC/4
<i>K. pneumoniae</i> ATCC 700603		8	1(0.125) ^S	2(0.25) ^S	1(0.125) ^S	1(0.125) ^S	1(0.125) ^S	1(0.125) ^S	2(0.25) ^S	2(0.25) ^S
	Amoxicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I
	Ampicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I
<i>E. coli</i> ATCC 5218	Ciprofloxacin	16	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A	>256(≥16) ^A
	Amoxicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I
	Ampicillin	128	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I	>256(≥2) ^I
<i>S. aureus</i> ATCC 25923	Ciprofloxacin	4	1(0.25) ^S	2(0.5) ^S	1(0.25) ^S	1(0.25) ^S	1(0.25) ^S	1(0.25) ^S	1(0.25) ^S	1(0.25) ^S
	Amoxicillin	<0.5	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A	>256(≥512) ^A
	Ampicillin	32	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A
<i>S. phylococcus</i> <i>us</i> spp.	Ciprofloxacin	4	1(0.25) ^S	2(0.5) ^S	0.5(0.125) ^S	1(0.25) ^S	0.5(0.125) ^S	0.5(0.125) ^S	1(0.25) ^S	1(0.25) ^S
	Amoxicillin	4	0.5(0.125) ^S	0.5(0.125) ^S	0.5(0.125) ^S	0.5(0.125) ^S	0.5(0.125) ^S	1(0.25) ^S	1(0.25) ^S	1(0.25) ^S
	Ampicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I
<i>Serratia</i> <i>odorifera</i>	Ciprofloxacin	4	4(1) ^I	4(1) ^I	2(0.5) ^S	2(0.5) ^S	1(0.25) ^S	2(0.5) ^S	2(0.5) ^S	4(1) ^I
	Amoxicillin	>256	128(≥0.5) ^S	>256(≥1) ^I	128(≥0.5) ^S	128(≥0.5) ^S	128(≥0.5) ^S	128(≥0.5) ^S	128(≥0.5) ^S	128(≥0.5) ^S
	Ampicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I
<i>Citrobacter</i> <i>spp.</i>	Ciprofloxacin	32	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A	>256(≥8) ^A
	Amoxicillin	>256	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I	>256(≥1) ^I

	Ampicillin	256	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I
<i>K. oxytoca</i>	Ciprofloxacin	32	8(0.25) ^S	8(0.25) ^S	4(0.125) ^S	4(0.125) ^S	4(0.125) ^S	4(0.125) ^S	8(0.25) ^S	8(0.25) ^S
	Amoxicillin	>256	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I
	Ampicillin	>256	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I
<i>Bacillus</i> spp.	Ciprofloxacin	4	8(2) ^I	8(2) ^I	2(0.5) ^S	4(1) ^I	8(2) ^I	4(1) ^I	4(1) ^I	4(1) ^I
	Amoxicillin	>256	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I
	Ampicillin	>256	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I	>256(≥ 1) ^I

(I): FIC (Fractional Inhibitory Concentration) of the antibiotics after association with plants extract; S: Synergy, I: Indifference; The values in bold represent the cases of synergy between extract and antibiotic; A: antagonist

Amongst beta-lactams, amoxicillin was the only one that was potentiated by all extracts on *Staphylococcus* spp. All others isolates remained “indifferent” to the drug antibiotic/extract combination.

Ciprofloxacin was also potentiated by the extracts on 5 strains which previously expressed resistance to Ciprofloxacin when it was used alone. Also fascinating, the ethyl acetate and methylene chloride extracts displayed relatively stronger synergistic effects when they were used in combination with Ciprofloxacin compared to ethanol and hexane extracts.

Comparing the MIC values of Amoxicillin/clavulanic acid (MIC = 4 μ g/mL) and Amoxicillin/plant extract (MIC < 0.25 μ g/mL) for the *Staphylococcus* spp., it was obvious that the MIC’s of the antibiotic /extract combination was relatively lower, indicating a higher inhibitory potential.

DISCUSSION

This investigation on the phytochemical screening and anti- β -lactamase potential of *Acmella caulirhiza* extracts on ESBL-positive isolates generated questionable results in several ways. The phytochemical screening revealed that the chemical composition of extracts were solvent-dependent. Thus, with ethyl acetate and ethanol, all the target metabolites were detected, consistent with the polarity and indicating their suitability as the most effective extraction solvents. However, the extraction yield with ethanol was twice higher than the one obtained with ethyl acetate. Otherwise, ethanol would extract more polar compounds than ethyl acetate. The yield with methylene chloride that was also better than the one obtained with ethyl acetate could have in addition to its polar characteristic, a greater corrosive effect that allows better fractionation, which in turn, facilitates release of the plant cell's contents.^[24] However, the fact that, unlike ethyl acetate, hexane did not extract polyphenols that are polar compounds is yet to address comprehensively. Overall on this issue, availability of ethanol is a great asset in developing cost-effective techniques that could be used in traditional medicine.

With a glance on the antibacterial potential of extracts on target bacteria isolates, it was observed that only the hexane and methylene chloride extracts expressed a certain degree of activity. Further insight indicated that the hexane extract was more potent than the methylene chloride's. It could therefore, be anticipated that the chemicals that are present in the extract have preferentially been extracted with hexane. The fact that the poorest extract in terms of metabolites (the hexane extract) has the highest antibacterial potential could be consistent with anticipation that its overall antibacterial potential strongly correlated its contents in secondary metabolites, more specifically, the sterols and the triterpenes. This extract was in fact, the poorest in target secondary metabolites, but exhibited the highest antibacterial activity compared to the others on 3 out of 7 multidrug-resistant isolates (MIC values ranging from 512 to 1024), with moderate activity from one isolate to the other. According to former authors, a plant extract is regarded as very active when the MIC < 100 μ g/mL, moderate when $100 \leq$ MIC < 625 μ g/mL and low when the MIC > 625 μ g/mL. Accordingly, the methylene chloride and hexane extracts exhibited moderate activity on *S. aureus* ATCC 25923, a susceptible reference strain. This agrees with the findings by previous investigations (Ramsewak *et al.*, 1999; Tatah *et al.*, 2015) which reported the antibacterial potential of the hexane and methylene chloride extracts of *Acmella caulirhiza*.^[18,25] According to Tatah *et al.* (2015), however, this activity observed with *Acmella caulirhiza* is likely associated with the

presence of soluble lipid and more specifically β -stigmaterol and hexadecanoic acid (palmitic acid).^[18] Hexane is a nonpolar solvent that can easily extract the soluble lipids such as essential oils, sterols and triterpenes in the plant.^[18] This is once again, consistent with findings from the present investigation and the above discussion on the detection of sterols and triterpenes implying that these chemicals (sterols and triterpenes), β -stigmaterol and hexadecanoic acid might possess the antibacterial potential. Extracts from ethyl acetate and ethanol were richer in secondary metabolites but did not express any antibacterial activity, also in line with earlier investigation.^[26] Unlike that study however, MIC values were obtained by agar dilution technique for aerial parts of *Acmella caulirhiza*. Inactivity of these extract could be attributed to several factors including negative interactions that might develop amongst the chemicals in the crude product. Flavonoids detected in higher concentration in the ethyl acetate and ethanol extract could also play critical roles in the overall activity observed and beyond.

An investigation undertaken by Ngoupayo *et al.* in 2015 disclosed that Flavonoids possess anticancer, antioxidant, antiviral, and antimicrobial potentials.^[27] In addition, metabolic contents in plants is dependent upon both biotic and abiotic influences that interact with that plant. Otherwise, plant from the same species that are grown in dissimilar environments would likely have slightly different chemical compositions. Concentration of secondary metabolites may also vary from one part of the plant to the other and, from one plant to the other, the same metabolite may have slightly different characteristics. Frequently shown to display good antibacterial activity on both major bacteria Gram categories ^[27-29], observing that flavonoids were inactive in the present study also still to be addressed, acknowledging that their potential can also be altered by the presence of other polyphenols, 2010).^[24] In their roles in general, polyphenols are known for their antibacterial potential through increased permeabilization of the cell envelope that eventually results in cell lysis like polypeptides, disrupts the super-coiling of bacterial genome like quinolones or acts as anti-metabolite like sulfonamides.^[29] In all cases, the activity was actually observed only with the hexane extract. The present investigation further focused on combinations of antibacterial agents that would inhibit ESBL expression. In this regards, the extract/antibiotic combination appeared to have a positive effect in ESBL expression. It is, therefore, likely that the compounds responsible for the antibacterial activity of the extracts tested alone were not the same ones that induced the synergistic effect that resulted in inhibition of ESBL expression. One might anticipate that these compounds are polar. Further steps into the synergistic studies disclosed that

Ciprofloxacin was potentiated (5/7 strains) by all extracts, including those which did not have any antimicrobial action when they were used alone. The fractional inhibitory concentrations related ranged between 0.125 and 0.5.

This potential might be related to the higher cell membrane permeability facilitated by flavonoids that allowed easy access to the antibiotic target, the topoisomerase. It is also clear that many other mechanisms might be involved, since β -lactam-resistant isolates that are positive for ESBL became susceptible to various drug combinations. Some resistance mechanisms might also make bacteria more susceptible to other antibacterial agents. In this and with regards to *Staphylococcus aureus* ATCC 25923, it would not be exaggerated to allege that the Amoxicillin/extract combination reduced amoxicillin concentration which became less available at the Penicillin Binding Protein (PBP) target in the periplasmic space then, responsible for the inactivity recorded.

With regards to β -lactams, only Amoxicillin was potentiated by all extracts in a single ESBL-positive clinical isolate (*Staphylococcus* spp.), with a FIC lower than 0.25. One could also anticipate that *Staphylococcus* spp., was the only isolate equipped with receptors for inhibitors that were present in the plant extract, or that the ESBL present in this *Staphylococcus* spp., differed from those expressed by the others isolates. The differences might also be related to the localization of β -lactamases in Gram-positive (extracellular) and Gram-negative (periplasmic), or their biogenesis (inducible in *Staphylococcus aureus* and constitutive in Gram-negative).^[30] Comparing the effect of the Amoxicillin/extract combination with that of the Amoxicillin/clavulanic acid association on this clinical isolate it was observed that the MIC of the combination with the extract was lower (MIC < 0.25) than the one recorded with the association with clavulanic acid (MIC = 4). This finding was key evidence that some of the extracts' secondary metabolites had improved inhibitory potential than clavulanic acid.

The fact that the extract did not potentiate the activity of Amoxicillin on ESBL-expressing rods from the *Enterobacteriaceae* family could further imply that the mechanisms enacted by the extracts are different than the one used by clavulanic acid, but these allegations are yet to be fully demonstrated. Subtle focus on the MIC values revealed that the isolates under study expressed resistance to β -lactams.^[31] Furthermore, it was shown that resistance to β -lactams through expression of ESBL is generally associated with resistance to fluoroquinolones.^[32] This latest clue would justify, at least partly, the resistance expressed by isolates to

Ciprofloxacin. However, a synergistic effect was recorded between the extracts and this Ciprofloxacin for certain isolates which were resistant to it, when it was used alone. This might suggest that Ciprofloxacin could be associated with extracts of *Acmella caulirhiza* for the caretaking in some instances where resistance to Ciprofloxacin is observed. This point however, requires further researches to determine the necessary accurate concentrations and resulting contraindications.

CONCLUSION

Acmella caulirhiza is rich in secondary metabolites for which extraction is solvent dependent. The hexane extract was the poorest in secondary metabolites but paradoxically expressed the highest potential on bacteria subjected to susceptibility tests. It might be legitimate to pay more attention on these combinations, especially in the caretaking of infections that involve extended-spectrum β -lactamases-positive bacteria which also express resistance to Ciprofloxacin.

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