

Phytochemical screening, toxicity and antimicrobial activity of different *Mimosa tenuiflora* extracts on *Aeromonas* strains

Triagem fitoquímica, toxicidade e atividade antimicrobiana de diferentes extratos de *Mimosa tenuiflora* sobre cepas de *Aeromonas*

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Highlights

Aeromonas genus cause disease and mortality in fish.

Methanol proved to be the best solvent for extracting secondary metabolites.

Acute toxicity (LC₅₀) of the methanol extract of *M. tenuiflora* was considered low.

M. tenuiflora extracts are promising for the treatment of diseases in fish farms.

Abstract

The indiscriminate use of synthetic veterinary drugs in fish farms for disease control has caused recurring environmental pollution and reduced productivity; however, the search for ecologically viable alternatives is increasing. Thus, this study aimed to evaluate the phytochemical characterization of the hexanic, methanolic, and aqueous extracts of black jurema (*M. tenuiflora*), and their antimicrobial activity against strains of *Aeromonas*, and acute toxicity (LC₅₀) to fingerlings of *O. niloticus*. The isolates were identified,

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and phenotypic virulence and antimicrobial susceptibility tests were performed. *A. caviae*, and *A. veronii* bv. *veronii* showed 75.0 - 87.5% positivity for the virulence factors tested, and resistance to the antimicrobials ampicillin, amoxicillin, erythromycin, and tetracycline of 67.0% and 50.0%, respectively. Phytochemical screening of black jurema extracts detected phenols, flavonoids, tannins, saponins, alkaloids, and steroids/triterpenoids, with methanol proving to be more efficient in the extraction of metabolites. The methanolic and aqueous extracts showed moderate antimicrobial activity, with minimum inhibitory concentration and minimum bactericidal concentration of 250 $\mu\text{g mL}^{-1}$, and the methanolic extract revealed an LC_{50} of 40 $\mu\text{g mL}^{-1}$ for *O. niloticus*. This study demonstrated the efficiency of the in vitro antimicrobial activity of *M. tenuiflora* extracts, and their use in vivo in the treatment or prophylaxis in fish farming can be investigated to replace the use of synthetic antimicrobials.

Key words: Virulence factors. *Oreochromis niloticus*. Pisciculture. Antimicrobial resistance.

Resumo

O uso indiscriminado de drogas veterinárias químicas em pisciculturas para controle de doenças tem sido um problema recorrente, de poluição ambiental e redução de produtividade. Em contrapartida, a busca por alternativas ecologicamente viáveis também tem crescido. Este estudo teve por objetivo avaliar a caracterização fitoquímica dos extratos hexânico, metanólico e aquoso de jurema preta (*Mimosa tenuiflora*), atividade antimicrobiana em cepas de *Aeromonas* e toxicidade aguda do extrato (CL_{50}) em alevinos de *Oreochromis niloticus*. Também foi realizado a identificação dos isolados de *Aeromonas*, perfil fenotípico de virulência e suscetibilidade antimicrobiana. As cepas de *A. caviae* e *A. veronii* bv. *veronii* apresentaram positividade entre 75,0 - 87,5% para os fatores de virulência testados, e resistência aos antimicrobianos, ampicilina, amoxicilina, eritromicina e tetraciclina de 67,0% e 50,0%, respectivamente. A triagem fitoquímica dos extratos de jurema-preta detectou fenóis, flavonoides, taninos, saponinas, alcaloides e esteroides/triterpenoides, com o metanol se mostrando o solvente mais eficiente na extração dos metabólitos secundários. Os extratos metanólico e aquoso apresentaram moderada atividade antimicrobiana, com Concentração Inibitória Mínima (CIM) e Concentração Bactericida Mínima (CBM) de 250 $\mu\text{g mL}^{-1}$, enquanto o extrato metanólico apresentou CL_{50} para *O. niloticus* de 40 $\mu\text{g mL}^{-1}$. Este estudo mostra a eficiência da atividade antimicrobiana in vitro de extratos de *M. tenuiflora*, podendo ser investigada sua utilização in vivo no tratamento ou profilaxia na piscicultura em substituição ao uso de antimicrobianos sintéticos.

Palavras-chave: Fatores de virulência. *Oreochromis niloticus*. Piscicultura. Resistência antimicrobiana.

Introduction

Aquatic animals and plants grow faster than any other animal production sector worldwide; therefore, aquaculture is an important production system (Romero, Feijoo, & Navarrete, 2012). In Brazil, 722,560 thousand tons of fish were produced in 2018,

which is an increase of 4.5% compared to 2017, following the average global annual growth rates (Tacon, 2020). Although fish farming has several advantages, the development of aquaculture activity results in farming systems with high stocking densities and increased animal stress, which lead to changes in water quality (Monteiro

et al., 2015). These conditions contribute to the emergence of microbial and parasitic diseases, requiring control measures such as the use of synthetic or natural antimicrobials (Doan, Soltani, Ingelbrecht, & Soltani, 2020).

Antimicrobials are commonly used in the aquaculture production cycle, both for bacterial infection treatment and prophylactic measures. The excessive use of these drugs causes accumulation of their residues in the environment, which can travel through the food chain and contribute to the emergence of opportunistic and infection-resistant microorganisms that can be toxic to aquatic biota or cause allergic reactions in sensitive individuals (Monteiro et al., 2015).

In Brazil, only the antimicrobials florfenicol and oxytetracycline are licensed for aquaculture use (Sindicato Nacional da Indústria de Produtos para Saúde Animal [SINDAN], 2014). However, bacteria have become increasingly resistant to antibiotics, resulting in more expensive and longer treatments, creating a strong appeal to reduce their use (Huang & Nitin, 2019).

Given this scenario, researchers have been looking for alternative measures to control pathogens, such as biologically active compounds present in medicinal plants with a broad spectrum of antibacterial actions (Bezerra et al., 2011; Kuebutornye & Abarike, 2020). Black jurema (*Mimosa tenuiflora*) is a native plant of the Caatinga, that is widely distributed in Northeastern Brazil and known for its anti-inflammatory, curative, and antibiotic properties (Bezerra et al., 2011; Borges et al., 2017). High levels of tannins and flavonoids are associated with its antimicrobial potential (S. A. N. M. Silva et al., 2020).

The aim of the present study was to evaluate the antimicrobial activity and toxicity of different extracts of *M. tenuiflora* on *Aeromonas* strains isolated from ornamental fish with clinical signs of bacterial infection, to produce a possible alternative to the use of synthetic antimicrobials in the treatment of infections in fish farms.

Material and Methods

The research was conducted with authorization from the Ethics Committee in the Use of Animals (CEUA) of the Federal University of Recôncavo da Bahia (UFRB) under registration no. 23007.027506/2017-58.

Obtaining fish and isolating microorganisms

Dead fish with clinical signs of bacterial infection (eye lesions on the surface of the body and fins, and erratic swimming) were acquired from an ornamental fish farm in the municipality of Dom Macedo Costa, Bahia, Brazil.

Swabs were taken from the lesions and inoculated on glutamate agar *Pseudomonas/Aeromonas* (GSP) media (30 °C/24 h), MacConkey agar (30 °C/24 h) and agar enriched with 5% of sheep blood (35 °C/24 h) (Alexandrino et al., 1999; Salvador et al., 2005; N. Silva et al., 2010b). After incubation, typical *Aeromonas* colonies were seeded on tryptic soy agar (ATS) for phenotypic (oxidase and catalase) and morphotintorial (gram stain) characterizations.

Molecular identification of isolates and phylogenetic reconstruction

DNA was extracted according to the methodology described by Lee, Kim, Liul and Lee (2003) with modifications. The isolates were grown in nutrient broth for 24 h, polyvinylpyrrolidone and 2-mercaptoethanol were removed from the lysis solution, and potassium acetate was added. To precipitate the DNA, isopropanol and ammonium acetate were included. Primers 8F (5'-AGAGTTTGATCCTGGCTCAG-3') and 1492R (5'-ACGGCTACCTTGTTACGACTT-3') (Miyashita et al., 2009) were used to amplify and sequence the 16S rRNA gene (Leite et al., 2013). The PCR products were sequenced at the company ACTGene Analisis Moleculares Ltda, using an AB 3500 sequencer (Applied Biosystems, California, USA). The sequences were edited using Sequencher v.5 program. 4.6, and compared using the BLASTn program with other sequences deposited in public databases.

The 16S rRNA type species sequences were retrieved from the *List of Prokaryotic names with Standing in Nomenclature* (LPSN) in August 2018 (Euzéby, 1997). MEGA v. 6.0 was used to align and perform the phylogenetic analysis of the sequences with the maximum likelihood method and with 1000-repeat bootstrap analysis. The Hasegawa-Kishino-Yano model with gamma distribution and invariant sites was selected as the best nucleotide replacement model.

Antimicrobial susceptibility

The susceptibility profile of the isolates was determined using the disk diffusion

technique with a bacterial density of 10^8 CFU mL⁻¹ (Clinical and Laboratory Standards Institute [CLSI], 2018). The antimicrobials amoxicillin (10 µg), tetracycline (30 µg), ampicillin (10 µg), chloramphenicol (30 µg), streptomycin (10 µg) and erythromycin (15 µg) were tested. The inhibition zones were classified as sensitive (≥ 18 mm), intermediate (13-17 mm), and resistant (≤ 13 mm) (Odeyemi & Ahmad, 2015) for the interpretation of the results. The multiple antimicrobial resistance index (MAR) was also used (Osundiya, Oladele, & Oduyebo, 2013).

Phenotypic virulence tests

To detect exoenzymes, the strains were grown on TSA agar containing 1% NaCl and supplemented with 0.5% gelatin (gelatinase), 5% skimmed milk powder (casein), 1% egg yolk emulsion (phospholipase), 1% Tween 80 (lipase), 20% sheep erythrocyte solution (hemolysin), 0.1% starch (amylase), urea broth (urease), and DNase agar supplemented with 0.01% toluidine chloride (DNase). Halo formation indicated test positivity (Wagatsuma, 1968; Hongping, Jilun, Ting, Yixi, & Xiaoming, 2011; I. P. Silva et al., 2018).

Preparation of plant extracts

The plant material was collected in the municipality of Cruz das Almas, Bahia (12°39'13.0"S, 39°05'06.1"W) and its exsiccate is deposited in the Recôncavo da Bahia Herbarium under the number (HURB 18333). The stem bark of *M. tenuiflora* was dried at room temperature (25 ± 2 °C), crushed in a knife mill, and subjected to maceration in hexane and methanol three times for each solvent.

The material was filtered and concentrated using a rotary evaporator at 30 °C. To obtain the aqueous extract, the dried and ground material was macerated in sterile distilled water at room temperature, refrigerated for 24 h, filtered, and lyophilized at -20 °C.

Antimicrobial activity of Mimosa tenuiflora extract

The microdilution plate test was performed using two standard strains (*Pseudomonas aeruginosa* ATCC 27853 and *Staphylococcus aureus* ATCC 25923) and *Aeromonas*, following the methodology of CLSI (2012). For the minimum inhibitory concentration (MIC), the three extracts were diluted in 10% DMSO concentrations from 7.81 to 1000 µg mL⁻¹. The minimum bactericidal concentration (MBC) was defined as the lowest concentration capable of causing cell death (CLSI, 2012).

Phytochemical screening

The extracts were subjected to qualitative phytochemical analysis to detect the main classes of secondary metabolites, following the methodologies of N. L. A. Silva, Miranda and Conceição (2010a) and Joshi, Bhoje and Sattarkar (2013). Preliminary techniques were performed to detect flavonoids (Shinoda test), tannins (iron precipitation test), saponins and alkaloids (Drangendorff), and steroids and triterpenoids (Liebermann-Burchard and Salkowski).

Determination of the content of total phenols, tannins and flavonoids

Total phenol content was determined using the Folin-Ciocalteu method from a solution containing 100 µL of each extract diluted in 25 mL of methanol. A 100 µL aliquot was stirred into 500 µL of Folin-Ciocalteu reagent and 6 mL of distilled water for 1 min. Then, 2 mL of 20% Na₂CO₃ was added and the solution was stirred for 1 min, rested for 2 h and the results read at 750 nm. Gallic acid (0.9 to 6.3 mg L⁻¹) was used for the calibration curve: $y = 0.0983x + 0.0355$ ($R^2 = 0.992$). The results were expressed in mg of gallic acid equivalents (GAE) per gram of extract (Meda, Lamien, Romito, Millogo, & Nacoulma, 2005).

The tannin content was determined using the vanillin/HCl method. One milliliter of each extract was diluted in methanol, added to 5 mL of a 1% vanillin and 8% HCl solution (1:1), rested for 20 min at 30 °C, and read at 500 nm. For the calibration curve, the catechin standard (200 - 1000 mg mL⁻¹) and the calibration equation: $y = 0.0004x + 0.0128$ ($R^2 = 0.998$) were used. The results were expressed in grams of catechin equivalent 100 g⁻¹ extract (Missio et al., 2017).

The total flavonoid content was determined according to Meda et al. (2005). Fifty microliters of each extract was homogenized in a methanolic solution of AlCl₃ (2%), kept in the dark for 30 min, and read at 415 nm. For the calibration curve, rutin (10 - 100 µg mL⁻¹) was used based on $y = 0.0146x + 0.2147$ ($R^2 = 0.999$). The results are expressed as percentages.

Acute toxicity test (LC_{50})

The MIC of the *M. tenuiflora* extract responsible for causing 50% mortality of *O. niloticus* fingerlings was determined by first exposing the fish to different concentrations of the extract diluted in 10% DMSO ($\mu\text{g mL}^{-1}$) (T1 = 250, T2 = 75, T3 = 50, T4 = 25, and T5 = 10), a positive control with 10% DMSO (T6), and a negative control using only water (T7). All treatments were performed with three replicates and five fish per repetition in a completely randomized design (CRD). The experiment was conducted in aquariums containing 5 L of water, light, and constant aeration in an air-conditioned environment, with individuals weighing 1 g and a maximum density of 1 g L⁻¹ (Instituto Brasileiro do Meio Ambiente e dos Recursos Naturais Renováveis [IBAMA], 1987). Mortality was assessed daily, and the LC_{50} at 48 h was determined using the trimmed Spearman-Kärber method (Hamilton, Russo, & Thurston, 1977).

Results and Discussion

The main bacteria responsible for losses in fish farms belong to the *Aeromonas* genus, which has complex pathogenicity related to multiple factors, such as the production of enzymes and extracellular toxins (Sreedharan, Philip, & Singh, 2013), indicating the virulent potential of these microorganisms. In this study, genetic analysis

showed that strains A4 and E5 were closely related to *A. caviae* (ATCC15468) and *A. veronii* bv. *veronii* (ATCC 35624), respectively, with 100% homology (Figure 1). The presence of *A. veronii* bv. *veronii* was also identified by 16S rRNA in diseased fish that had internal hemorrhage and necrosis of the tail and fins, which were acquired from an ornamental fish farm in Kerala, India (Sreedharan et al., 2013).

Aeromonas strains showed resistance to ampicillin and intermediate resistance to tetracycline, which can be attributed to the wide use of these antimicrobials due to their broad spectrum of action, low toxicity, and low cost (Scarano et al., 2018). *Aeromonas veronii* showed greater susceptibility (50%) to antimicrobials than *A. caviae* (33.3%). When grouping the resistance and intermediate resistance categories, *A. caviae* presented 67% resistance to antimicrobials, and *A. veronii* showed 50% resistance (Table 1). Similar results were reported by Suhet, Schocken-Iturrino and Amaral (2011), who observed 100% resistance to amoxicillin and 50% to erythromycin for *A. caviae*, and 100% to amoxicillin, and 50% to tetracycline for *A. veronii*. According to Jagoda, Honein, Arulkanthan, Ushio and Asakawa (2017), *A. veronii* has increasingly attracted the attention of researchers due to its virulence potential in a wide range of hosts, both as a primary and an opportunistic pathogen, as well as its capacity to develop multidrug resistance phenotypes.

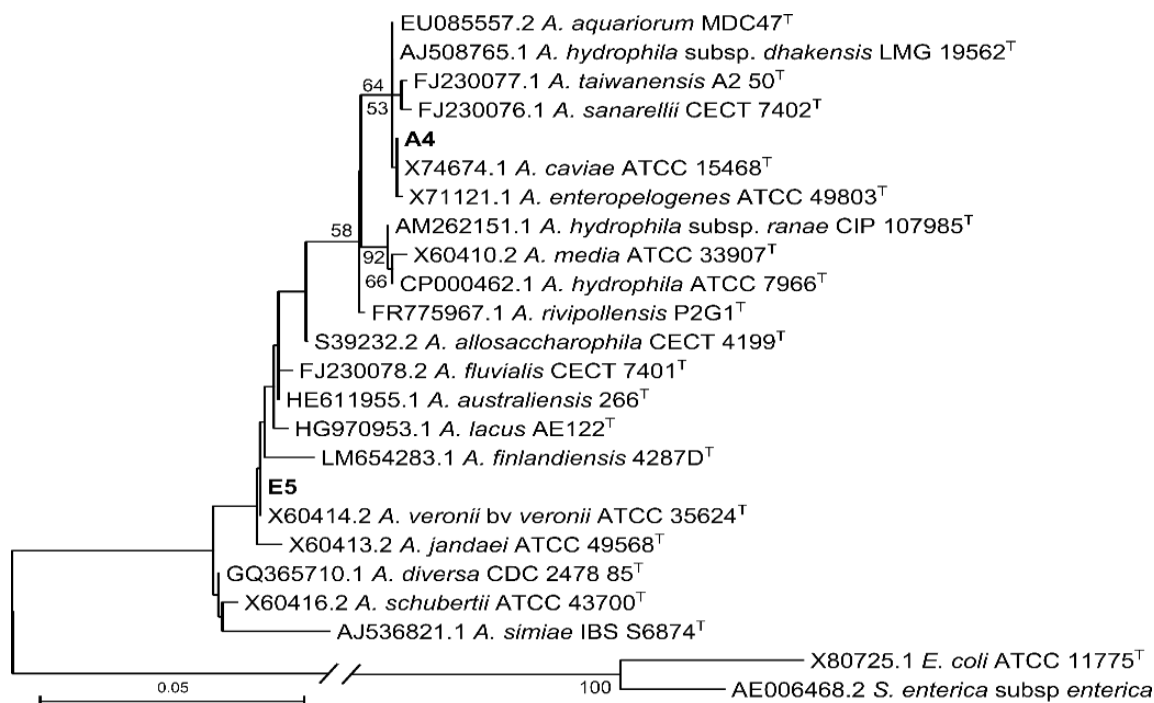


Figure 1. Phylogenetic tree showing the relationship of bacterial isolates obtained (in bold) with the species of the genus *Aeromonas*. Only bootstrap values above 50 are shown. *E. coli* (X80725) and *Salmonella enterica* subsp. *enterica* (AE006468) were used as outgroups. The scale indicates the number of replacements per site.

Table 1

Antimicrobial resistance profile of *Aeromonas caviae* and *A. veronii* bv. *veronii* against synthetic antimicrobials

Antimicrobials	Bacteria	
	<i>A. caviae</i>	<i>A. veronii</i> bv. <i>veronii</i>
Amoxicillin	15.83 (I)	11.44 (R)
Tetracycline	15.12 (I)	15.76 (I)
Ampicillin	5.91 (R)	6.93 (R)
Chloramphenicol	29.4 (S)	45.68 (S)
Streptomycin	23.88 (S)	22.09 (S)
Erythromycin	12.03 (R)	18.82 (S)

S = sensitive. I = intermediate resistance. R = resistant.

The antimicrobial multidrug resistance (AMR) index for *A. caviae* was 0.67, indicating resistance to four antimicrobials and 0.50

for *A. veronii*, denoting resistance to three antimicrobials. Multiple resistance to at least four antimicrobials of more than three

structural classes was also reported by Hossain et al. (2019) with *Aeromonas* strains isolated from healthy ornamental fish in Korea, demonstrating the excessive use of antimicrobials in that industry. According to Shuang et al. (2020), the increase in antimicrobial resistance in cultured *Aeromonas* strains has increased due to the abusive use of these drugs in rearing systems, causing the bacteria to carry the resistance genes through horizontal transfer.

For phenotypic virulence tests, *A. caviae* was negative for urease, and *A. veronii* for DNase and urease of the eight factors tested. Despite the multiple virulence factors found in the strains, J. L. S. Silva et al. (2016) reported that determining the relevance of the various factors is infeasible, as they all are relevant for establishing infection. The production of microbial exoenzymes promotes tissue destruction, immune system cell lysis, and host cell invasion (Otto, 2014). This justifies the hypothesis that the bacteria *A. caviae* and *A. veronii* were responsible for the infection that caused fish mortality in the studied fish farm.

Both species were characterized with positive β -hemolysis. Sreedharan et al. (2013) stated that β -hemolysis is one of the most important virulence factors in the pathogenicity process in fish, since hemolysins are extracellular cytolytic proteins capable of destroying membrane permeability barriers by inserting into the lipid bilayer and destroying the red blood cells.

In fish farms, another constraint encountered by producers is an increase in bacterial resistance in environmental isolates. Therefore, the antibacterial activity of *M. tenuiflora* extracts was analyzed and the same MIC and MBC values were obtained for the bacteria, with the methanolic and aqueous extracts showing greater efficiency for *Aeromonas* than for the reference strains (Table 2). According to Simonetti et al. (2016), the methanolic and aqueous extracts showed moderate antimicrobial activity (MIC between 100 and 500 $\mu\text{g mL}^{-1}$) for *Aeromonas* and *S. aureus* and weak antimicrobial activity against *P. aeruginosa* (MIC between 500 and 1000 $\mu\text{g mL}^{-1}$). The hexane extract was inactive against all tested microorganisms (MIC > 1000 $\mu\text{g mL}^{-1}$).

Table 2
Minimum Inhibitory Concentration (MIC) and Minimum Bactericidal Concentration (MBC) of *M. tenuiflora* extracts against *Aeromonas* and reference strains

Pathogens	Extracts ($\mu\text{g mL}^{-1}$)					
	HEMT		MEMT		AEMT	
	CIM	CBM	CIM	CBM	CIM	CBM
<i>A. caviae</i>	>1000	>1000	250	250	250	250
<i>A. veronii</i> bv. <i>veronii</i>	>1000	>1000	250	250	250	250
<i>P. aeruginosa</i> ATCC 27853	>1000	>1000	1000	1000	1000	1000
<i>S. aureus</i> ATCC 25923	>1000	>1000	500	500	500	>1000

HEMT: hexane extract of *M. tenuiflora*; MEMT: methanol extract of *M. tenuiflora*; AEMT: aqueous extract of *M. tenuiflora*. MIC: Minimum Inhibitory Concentration. MBC: Minimum Bactericidal Concentration.

The phytochemical analysis of the extracts (Table 3) showed that the hexane extract did not contain flavonoid compounds or tannins, as observed in the methanolic and aqueous extracts, while steroids/triterpenoids were present in the hexane extract and absence in the other extracts. Phytochemical compounds play a fundamental role in investigating the active principles of medicinal plants, as their presence contributes to the development of new drugs (Cordell, 2014).

The antimicrobial activity of *M. tenuiflora* extract has been associated with the presence of tannins (S. A. N. M. Silva et al., 2020). As noted, these compounds were not found in the hexane extract, which had low antimicrobial activity (MIC > 1000 µg mL⁻¹)

(Table 2). Tannins are mainly responsible for the antimicrobial activity of *M. tenuiflora*, and among the mechanisms of action are enzymes inhibition, cell membrane rupture, and metal ion complexation, which reduce important elements of microbial metabolism and affect its physiology (Ferreira & Evangelista, 2021). The presence of tannins in extracts can be attributed to their polarity and ability to bind to polar solvents such as methanol and water. Hexane is a low-polarity solvent and shows less efficiency in extracting compounds (Meira et al., 2020). Similar results were reported by S. A. N. M. Silva et al. (2020) who observed that ethanol solvent polarity promoted better retrieval of metabolic compounds such as tannins and flavonoids in *M. tenuiflora* extracts.

Table 3
Phytochemical screening of the three extracts obtained from the stem bark of *M. tenuiflora*

Phytochemical tests	HEMT	MEMT	AEMT
Flavonoids	-	+	+
Tannins	-	+	+
Saponins	+	+	-
Alkaloids	+	+	+
Steroids/triterpenoids	+	-	-

HEMT: hexane extract of *M. tenuiflora*; MEMT: methanol extract of *M. tenuiflora*; AEMT: aqueous extract of *M. tenuiflora*; (+): presence; (-): absence.

As flavonoids were not found in the hexane extract screening, the quantification of total phenols and tannins was performed only in the methanolic and aqueous extracts. The methanolic extract had a higher content of secondary metabolites compared to the

aqueous extract, while both contained more than 50% of tannins (Table 4). The methanol extract has a higher content of total phenols and tannins; therefore, it was chosen to test the acute toxicity in Nile tilapia fingerlings.

Table 4
Quantification of total phenols, tannins and flavonoids in aqueous and methanolic extracts of *M. tenuiflora*

Extracts	Total phenols (g de GAE/100g de extract)	Tannins (g de CE/100g de extract)	Flavonoids (% de ER)
AEMT	1.42 ± 0.01	59.10 ± 7.58	0.47
MEMT	2.20 ± 0.14	69.25 ± 11.68	0.72

MEMT: methanol extract of *M. tenuiflora*; AEMT: aqueous extract of *M. tenuiflora*; GAE: gallic acid equivalent; CE: catechin equivalent; ER: equivalent of routine.

Regarding acute toxicity (LC_{50}), concentrations of 250 $\mu\text{g mL}^{-1}$ (T1) and 75 $\mu\text{g mL}^{-1}$ (T2) caused 100% mortality of *O. niloticus* fingerlings, while 50 $\mu\text{g mL}^{-1}$ (T3) caused 60% mortality and the control treatments (T6 and T7) showed no mortality. The LC_{50} of the methanol extract of *M. tenuiflora* for fingerlings was considered low (40 $\mu\text{g mL}^{-1}$). According to the classification of acute toxicity for aquatic organisms (Zucker, 1985), the methanol extract of *M. tenuiflora* is considered slightly toxic (values >10 to <100), which helps to determine safe doses for use in fish farming. Thus, the antimicrobial use of plant extracts is a promising method of controlling diseases in fish, as it is easily biodegradable, low-cost, and accumulates less waste in fish (Awad & Awaad, 2017; Doan et al., 2020). Among the secondary metabolites identified in black jurema are the flavonoids apigenin and flavonol (Borges et al., 2017), sakuranetine (5,4'-dihydroxy-7-methoxyflavanone), and sorbifolium (5,6,4'-trihydroxy-7-methoxyflavone) (Hernandez et al., 2021), 5,7,4'-trihydroxy-3-methoxyflavone, 5,4'-dihydroxy-7,8-dimethoxyflavone, 5,7,4'-trihydroxy-6-methoxyflavonol, and 5-hydroxy7,8,4'-trimethoxyflavonol (Cruz et al., 2016), saponins, alkaloids,

polysaccharides, and anthraquinones (Bezerra et al., 2011). However, it is believed that the use of bioactive compounds promotes the slower development of microbial resistance, and the application of inadequate doses of herbal medicines can cause toxicity (Reverter, Bontemps, Lecchini, Banaigs, & Sasal, 2014).

Similar fish reactions to initial exposure of higher extract concentrations (T1, T2, and T3) were observed, such as increased opercular beat, agitation, and erratic swimming. These reactions may be associated with a rapid decrease in dissolved oxygen in water, as evidenced by increased opercular movement, as well as the toxic effect caused by the compounds present in the extract (Kuebutornye & Abarike, 2020). Considering that the extract of *M. tenuiflora* showed acute toxicity for *O. niloticus* at a concentration six times lower than the MIC of the extract for *A. caviae* and *A. veronii*, it is suggested that future studies should test the exposure period of fish to the extract lesser than 48 h, as well as its antibacterial activity *in vivo*. Another alternative would be the use of the aqueous extract, which *in vitro* showed the same MIC for the bacteria, and 15% less tannins when compared to the methanolic extract.

Despite the bioactivity of some plants as an antioxidant, antimicrobial, or immunostimulant agents in the control of fish diseases or weight gain (Durmic & Blache, 2012), further studies on the use of black jurema extract are required in terms of absorption, metabolism, and biological action in animals to assess its potential effect on aquaculture production systems.

Conclusion

Methanolic and aqueous extracts of *M. tenuiflora* showed strong antibacterial activity *in vitro* against pathogenic *A. caviae* and *A. veronii* bv. *veronii*, which is promising for its application in the prophylaxis and treatment of microbial diseases associated with fish production. Another important discovery was the low toxicity of the extracts for Nile tilapia, which could enable their use in sublethal doses. Thus, these extracts can be applied in fish farms as an efficient alternative for the treatment of quarantined fish affected by microbial diseases with the aim of replacing conventional antimicrobials.

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