

Potentials of *Melastoma malabathricum* Linn. Flower and Fruit Extracts as Antimicrobial Infusions

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ABSTRACT

Melastoma malabathricum Linn. is a shrub that belongs to the family *Melastomataceae* and a common herbal plant used in folk medicines to treat inflamed wounds. This study was carried out with the aim to evaluate the inhibitory activities of different concentrations of the *M. malabathricum* Linn. flower and fruit crude extracts against a variety of microorganisms. The inhibitory effects of both extracts were tested against the microorganisms using the disc diffusion method. The lowest concentrations of the extracts producing inhibition zones against the test microorganisms were used to determine their Minimum Inhibitory Concentrations (MICs) and Minimum Microbicidal Concentrations (MMCs). Both crude extracts showed strong inhibitory activities against Gram-positive bacteria. The range of MIC values for the crude flower and fruit extracts on all the bacteria tested were 12.5 to 100.0 mg/ml. Overall, Gram-positive bacteria were more susceptible to the crude extracts compared to Gram-negative species, potentiating a possible use of the extracts to inhibit or kill potential pathogens.

Keywords: *Melastoma malabathricum* Linn.; Flower; Fruit; Disc Diffusion Method; Minimal Inhibitory Concentration; Minimal Microbicidal Concentration

1. Introduction

In the present era, modern medicine develops very quickly and makes major contributions to worldwide diseases control. Despite the rapid growth in knowledge and techniques in modern medicine, the populations of developed and developing countries also show an increase interest in traditional medicines. According to the World Health Organization [1], the definition of traditional medicine may be summarized as the sum total of all the knowledge and practices, whether explicable or not, used in the diagnosis, prevention and elimination of physical, mental or social imbalances, and relying exclusively on practical experiences and observations handed down from generation to generation, whether verbally or in writing. Today, almost 80% of the Asian population relies on traditional medicine as a part of their health care regime.

Melastoma malabathricum Linn. is a shrub that belongs to the family *Melastomataceae*. It can be found abundantly in Malaysia. It comes with beautiful pink or purple flowers. The flower consists of five narrow reddish sepals and five purplish petals. The fruits are en-

capsulated and contain many non-endospermous seeds [2] with small embryos inside purplish pulps. *M. malabathricum* Linn. is rich with flavonoid compounds [3,4]. In folklore remedies, different parts of the plants are used for the treatment of various human ailments. The leaf, root and flower crude extracts of *M. malabathricum* Linn. had been used to treat toothache, wounds, diarrhea, anti-infection, scar prevention, and post-partum recovery [5]. Numerous pharmacological studies and clinical practices have reported that various parts of *M. malabathricum* Linn. plant possess biological functions such as antioxidant and anti-cancer [3], antiviral [6], anti-inflammatory, antinociceptive and anti-pyretic [7], and anti-ulcerogenic [8].

In this study we employed thirty-two microbial species to test the possible antimicrobial activity of *M. malabathricum* Linn. crude fruit and flower extracts. These species include the Gram-positive and Gram-negative bacteria and fungi, which are food spoilers but normal human body flora. Most of these species cause serious cases of food poisoning and hence, there is a need for the development of preservatives to eliminate their growth. However, research on antibacterial properties of *M. malabathricum* Linn. is still scarce; only a few scientific

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articles reported some of these activities [9]. Looking at the myriad uses of *M. malabathricum* Linn. in folk medicine, it is therefore worthwhile to scientifically evaluate the antimicrobial activity of the crude flower and fruit extracts of *M. malabathricum* Linn. against the groups of microorganisms earlier mentioned.

2. Materials and Methods

2.1. Plant Materials

The flowers and fruits of *Melastoma malabathricum* L. were collected along the road sides in Universiti Putra Malaysia, Serdang, Selangor, Malaysia. Authentication of the plant was done at the Institute of Bioscience, Universiti Putra Malaysia, where the voucher specimen was conserved under the reference number SK1517/07.

2.2. Extraction

About 100 g of each fresh petals and fruits was extracted separately overnight with 1000 ml of methanol (Merck, Darmstadt, Germany), at $25^{\circ}\text{C} \pm 1^{\circ}\text{C}$. The extracts were ducked through Whatman No. 1 filter paper (Whatman, Maidstone, England), and then vacuum dried in a rotary evaporator (Buchi, Flawil, Switzerland) at $37^{\circ}\text{C} \pm 1^{\circ}\text{C}$.

2.3. Microbial Strains and Culture Conditions

The test microorganisms used in this study are as listed in **Table 1**. ATCC strains were bought from the American Type Culture Collection (USA) and other strains of human origin were obtained from the Institute of Medical Research (IMR), Malaysia.

2.4. Antimicrobial Assays

2.4.1. Preparation of Extract Test Samples

Vacuum dried crude flower and fruit extracts of the following amounts: 600 mg, 500 mg, 400 mg, 300 mg, 200 mg, 100 mg were dissolved respectively in 1ml of water. Then, all the dissolved extracts were filtered through a $0.22\ \mu\text{m}$ membrane. A total of 10 μl of each sample extracts was loaded onto a sterile 5 mm diameter paper discs. Sterilized water was used as a negative control. All impregnated discs were allowed to dry overnight at room temperature in a laminar flow hood. The commercial Tetracycline 30 (Oxoid, Hampshire, England) and Nystatin 100 (Oxoid, Hampshire, England) discs were used as positive controls for bacteria and fungi, respectively.

2.4.2. Disc Diffusion Assay

The antimicrobial activity was measured using the disc diffusion method [10]. All bacterial were grown for 18 - 24 h at 37°C in Nutrient broth (Merck, Darmstadt, Germany) and the cultures were adjusted to match 0.5

McFarland standard. The fungi were cultured for 24 - 48 h in Potato Dextrose broth (BD Difco, Maryland, USA) at $30^{\circ}\text{C} - 37^{\circ}\text{C}$. Petri dishes containing Nutrient agar (Merck, Darmstadt, Germany) and Potato Dextrose agar (BD Difco, Maryland, USA) were each swabbed with 100 μl of the microbial suspensions (contained approximately 10^6 to 10^7 cfu/ml). Sterile 5 mm diameter paper discs impregnated with 10 μl each of 100, 200, 300, 400, 500 and 600 mg/ml extracts were placed on the agar surface. A disc impregnated with water was used as a negative control. All the plates were incubated at 37°C for 24 h for the bacteria and $30^{\circ}\text{C} - 37^{\circ}\text{C}$ for 24 - 48 h for the fungi. All tests were performed in triplicates and repeated three times.

2.4.3. Macrodilution Method for Minimum Inhibitory Concentration (MIC) Determination

The Minimum Inhibitory Concentration (MIC) was determined only for microbial species which showed growth inhibition zones in the disc diffusion method described above. The inoculum was prepared from a 18 - 24 h broth culture using the disc diffusion method described above. A 18 - 24 h broth culture of each respective microbial species, adjusted to 0.5 McFarland standard, was used as the inoculum in this experiment. Approximately 100 mg each of the crude flower and fruit extracts were dissolved in Nutrient broth separately in a test tube to a final concentration of 100 mg/ml. A two folds dilution was prepared to give final concentrations of 0.39, 0.78, 1.56, 3.12, 6.25, 12.5, 25.0, 50.0 and 100.0 mg/ml of the crude flower and fruit extracts. Then, 500 μl (contained approximately 10^6 to 10^7 cfu/ml) of bacterial inoculum was added into each corresponding tube. Crude flower and fruit extracts without bacterial inoculum were used as negative controls. Tetracycline (Oxoid, Hampshire, England) was used as a positive control.

Culture growth was determined macroscopically and recorded after 18 - 24 h incubation at 37°C . The MIC was determined as the lowest concentration corresponding to the test tube presenting no turbidity changed after incubation. All tests were performed in triplicates and repeated three times.

2.4.4. Minimum Microbicidal Concentration (MMC) Determination

For the determination of the Minimum Microbicidal Concentration (MMCs), 100 μl of the content from each tested tube in the MIC assay that showed no turbidity changed was used to spread onto fresh Nutrient agar. The plates were further incubated for 18 - 24 h at 37°C . The lowest concentration that yielded no growth, at which 99.9% of the bacterial was killed, was determined as the MMC [11]. All tests were performed in triplicates and repeated three times.

Table 1. Microbial species, their cultivation and assays media used in this study.

Microbial species	Strain	Cultivation medium	Assay medium
<i>Alcaligenes faecalis</i>	IMR A111	Nutrient agar	Nutrient agar
<i>Escherichia coli</i>	IMR E113	Nutrient agar	Nutrient agar
<i>Enterobacter aerogenes</i>	IMR E153	Nutrient agar	Nutrient agar
<i>Enterobacter aerogenes</i>	ATCC 13048	Nutrient agar	Nutrient agar
<i>Proteus mirabilis</i>	IMR P184	Nutrient agar	Nutrient agar
<i>Proteus vulgaris</i>	IMR P147	Nutrient agar	Nutrient agar
<i>Proteus vulgaris</i>	ATCC 8427	Nutrient agar	Nutrient agar
<i>Klebsiella pneumonia</i>	IMR K36	Nutrient agar	Nutrient agar
<i>Klebsiella pneumonia</i>	ATCC 33495	Nutrient agar	Nutrient agar
<i>Salmonella typhimurium</i>	IMRS1205	Nutrient agar	Nutrient agar
<i>Salmonella typhi</i>	IMR S346	Nutrient agar	Nutrient agar
<i>Shigella dysenteriae</i>	IMR S340	Nutrient agar	Nutrient agar
<i>Shigella sonnei</i>	ATCC 2593	Nutrient agar	Nutrient agar
<i>Serratia marcescens</i>	IMR S913	Nutrient agar	Nutrient agar
<i>Pseudomonas aeruginosa</i>	IMR P187	Nutrient agar	Nutrient agar
<i>Staphylococcus aureus</i>	IMR S244	Nutrient agar	Nutrient agar
<i>Staphylococcus aureus</i>	IMR S942	Nutrient agar	Nutrient agar
<i>Staphylococcus epidermidis</i>	IMR S168	Nutrient agar	Nutrient agar
<i>Bacillus subtilis</i>	IMR B145	Nutrient agar	Nutrient agar
<i>Bacillus cereus</i>	IMR B43	Nutrient agar	Nutrient agar
<i>Streptococcus pyogenes</i>	IMR S1269	Blood agar	Nutrient agar
<i>Streptococcus pyogenes</i>	ATCC 19615	Blood agar	Nutrient agar
<i>Listeria monocytogenes</i>	IMR L55	Nutrient agar	Nutrient agar
<i>Enterococcus faecalis</i>	IMR E227	Nutrient agar	Nutrient agar
<i>Enterococcus faecalis</i>	ATCC 29212	Nutrient agar	Nutrient agar
<i>Micrococcus luteus</i>	IMR M174	Nutrient agar	Nutrient agar
<i>Micrococcus luteus</i>	ATCC 10240	Nutrient agar	Nutrient agar
<i>Candida albicans</i>	IMR C451	Potato dextrose agar	Potato dextrose agar
<i>Candida albicans</i>	ATCC 10231	Potato dextrose agar	Potato dextrose agar
<i>Saccharomyces cerevisiae</i>	IMR S1224	Potato dextrose agar	Potato dextrose agar

2.5. Data Analysis

All the experiments were repeated three times. The statistical data such as means and standard deviations was calculated using ANOVA (SPSS software for windows, version 17.0). The significance of the differences was determined by Tukey's test at $P < 0.05$ [12].

3. Results and Discussion

The average yield of the crude flower extract obtained from a starting 100 g of fresh petals was 6.0 g. While, the average yield of the crude fruit extract obtained was 9.6 g from 100 g of fresh fruits.

Table 2 shows the inhibition zones of *Melastoma malabathricum* Linn. extracts against Gram-negative bacte-

ria at 100 - 600 mg/ml. The results showed that *S. sonnei* ATCC 2593 was the most sensitive candidate towards the crude flower extract, with a minimal inhibition zone of 13.2 ± 0.2 mm. While *P. mirabilis* IMR P184 and *P. vulgaris* IMR P147 exhibited the highest sensitivity towards the crude fruit extract with a 8.0 ± 0.0 mm inhibition zone size. *E. coli* IMR E113, *E. aerogenes* IMR E153, *E. aerogenes* ATCC 13048, *K. pneumoniae* IMR K36, *K. pneumoniae* ATCC 33495, *S. typhi* IMR S346 and *S. typhimurium* IMR S1205 were not inhibited by both extracts at all the concentrations tested.

Overall, the results showed that the flower and fruit extracts were more effective against the Gram-positive bacteria tested (**Table 3**). The highest inhibition was attributed by the flower extract against *M. luteus* ATCC

Table 2. Inhibition zones (mm) of *Melastoma malabathricum* Linn. extracts against Gram-negative bacteria.

Gram-negative bacteria	Inhibition zone (mm)												Antibiotics ^A
	Flower extract (mg/ml) [extract yield (%) in solvent]						Fruit extract (mg/ml) [extract yield (%) in solvent]						
	100	200	300	400	500	600	100	200	300	400	500	600	
<i>A. faecalis</i> IMR A111	6.7 ± 0.2 ^a	10.2 ± 0.2 ^b	11.2 ± 0.7 ^{b,c}	12.0 ± 0.5 ^c	12.5 ± 0.5 ^c	14.3 ± 1.1 ^d	7.5 ± 0.0 ^a	8.7 ± 0.2 ^{a,b}	10.7 ± 0.2 ^b	11.5 ± 0.5 ^{b,c}	14.0 ± 1.0 ^{c,d}	14.5 ± 0.5 ^d	38.6 ± 1.1
<i>E. coli</i> IMR E113	-	-	-	-	-	-	-	-	-	-	-	-	8.6 ± 0.2
<i>E. aerogenes</i> ATCC 13048	-	-	-	-	-	-	-	-	-	-	-	-	20.0 ± 0.0
<i>E. aerogenes</i> IMR E153	-	-	-	-	-	-	-	-	-	-	-	-	18.1 ± 0.2
<i>K. pneumoniae</i> ATCC 33495	-	-	-	-	-	-	-	-	-	-	-	-	25.0 ± 0.0
<i>K. pneumoniae</i> IMR K36	-	-	-	-	-	-	-	-	-	-	-	-	23.6 ± 2.0
<i>P. mirabilis</i> IMR P184	11.8 ± 0.7 ^a	12.8 ± 0.7 ^{a,b}	13.5 ± 0.5 ^{b,c}	14.2 ± 0.7 ^{c,d}	14.8 ± 0.7 ^{d,e}	15.7 ± 0.5 ^e	8.0 ± 0.0 ^a	8.3 ± 0.5 ^a	10.0 ± 1.0 ^{a,b}	11.2 ± 1.0 ^{b,c}	12.3 ± 0.2 ^c	13.2 ± 0.2 ^c	8.0 ± 0.0
<i>P. vulgaris</i> IMR P147	7.8 ± 0.2 ^a	10.7 ± 0.2 ^b	12.3 ± 0.2 ^c	13.3 ± 0.2 ^d	13.8 ± 0.2 ^d	15.3 ± 0.2 ^e	8.0 ± 0.2 ^a	8.3 ± 0.5 ^a	9.7 ± 0.5 ^{a,b}	10.0 ± 0.0 ^{a,b,c}	11.3 ± 0.5 ^{b,c}	12.3 ± 0.5 ^c	23.6 ± 0.5
<i>P. vulgaris</i> ATCC 8427	-	10.8 ± 0.2 ^a	12.2 ± 0.2 ^a	13.2 ± 0.2 ^b	14.8 ± 0.5 ^c	18.2 ± 1.1 ^d	-	9.0 ± 0.5 ^a	10.3 ± 0.2 ^b	11.8 ± 0.2 ^c	13.2 ± 0.2 ^d	14.5 ± 0.5 ^d	27.0 ± 0.0
<i>S. typhimurium</i> IMR S1205	-	-	-	-	-	-	-	-	-	-	-	-	28.0 ± 0.0
<i>S. typhi</i> IMR S346	-	-	-	-	-	-	-	-	-	-	-	-	28.6 ± 1.1
<i>S. dysenteriae</i> IMR S340	12.0 ± 0.0 ^a	13.5 ± 0.5 ^{a,b}	14.2 ± 0.2 ^b	15.7 ± 0.5 ^c	17.3 ± 0.5 ^d	18.0 ± 0.0 ^d	-	9.7 ± 0.5 ^a	11.0 ± 0.0 ^a	11.7 ± 0.5 ^a	12.3 ± 0.5 ^a	13.3 ± 1.1 ^a	32.6 ± 4.6
<i>S. sonnei</i> ATCC 2593	13.2 ± 0.2 ^a	14.2 ± 0.2 ^b	15.0 ± 0.0 ^b	15.8 ± 0.2 ^{b,c}	16.3 ± 0.5 ^{c,d}	17.5 ± 0.5 ^d	-	7.7 ± 0.2 ^a	8.0 ± 0.0 ^a	8.8 ± 0.2 ^b	9.8 ± 0.2 ^c	10.0 ± 0.0 ^c	18.0 ± 0.0
<i>P. aeruginosa</i> IMR P187	9.7 ± 0.2 ^a	12.3 ± 0.2 ^b	13.7 ± 0.5 ^{b,c}	15.3 ± 0.2 ^d	17.0 ± 0.0 ^e	17.7 ± 0.5 ^e	-	8.0 ± 0.0 ^a	9.7 ± 0.5 ^{a,b}	10.2 ± 0.7 ^b	11.0 ± 1.0 ^{b,c}	12.3 ± 0.5 ^c	14.3 ± 1.1
<i>S. marcescens</i> IMR S913	-	8.8 ± 0.2 ^a	9.8 ± 0.2 ^b	10.7 ± 0.2 ^c	11.3 ± 0.2 ^{c,d}	11.8 ± 0.2 ^d	-	-	-	-	-	-	19.6 ± 0.5

-: No inhibition zone; ^AStandard antibiotic (Tetracycline 30 for bacteria species); Negative control (100% water) did not show inhibitory activity; Results are the means of inhibition zone values followed by standard deviations; Different letters within the same row indicate means at different concentrations for the same species are significantly different ($P < 0.05$).

Table 3. Inhibition zones (mm) of *Melastoma malabathricum* Linn. extracts against Gram-positive bacteria.

Gram-positive bacteria	Inhibition Zone (mm)												Antibiotics ^A
	Flower extract (mg/ml) [extract yield (%) in solvent]						Fruit extract (mg/ml) [extract yield (%) in solvent]						
	100	200	300	400	500	600	100	200	300	400	500	600	
<i>S. aureus</i> IMR S244	9.7 ± 0.5 ^a	10.5 ± 0.5 ^{a,b}	11.7 ± 0.5 ^{b,c}	12.8 ± 0.2 ^{c,d}	13.2 ± 0.2 ^d	13.8 ± 0.2 ^d	8.0 ± 0.5 ^a	8.2 ± 0.7 ^a	8.8 ± 1.0 ^a	10.3 ± 0.5 ^{a,b}	11.2 ± 0.7 ^b	12.0 ± 1.0 ^b	30.0 ± 0.0
<i>S. aureus</i> IMR S942	7.0 ± 0.0 ^a	8.2 ± 0.2 ^{a,b}	9.0 ± 0.0 ^{b,c}	9.8 ± 0.2 ^{b,c}	10.7 ± 1.1 ^{c,d}	12.0 ± 1.0 ^d	7.0 ± 0.2 ^a	8.2 ± 0.2 ^a	9.0 ± 0.0 ^{a,b}	9.8 ± 0.2 ^b	10.7 ± 0.7 ^{b,c}	12.0 ± 1.0 ^d	26.0 ± 1.0
<i>S. epidermidis</i> IMR S168	10.3 ± 0.2 ^a	11.7 ± 0.2 ^b	12.3 ± 0.2 ^{b,c}	12.8 ± 0.2 ^{c,d}	13.3 ± 0.2 ^{d,e}	13.8 ± 0.2 ^e	7.8 ± 0.2 ^a	7.8 ± 0.2 ^a	7.8 ± 0.2 ^a	9.7 ± 0.2 ^b	10.3 ± 0.5 ^{b,c}	11.3 ± 0.5 ^c	10.6 ± 0.2
<i>B. cereus</i> IMR B43	6.2 ± 0.2 ^a	6.8 ± 0.2 ^b	9.0 ± 0.0 ^c	10.0 ± 0.0 ^d	10.0 ± 0.0 ^d	11.0 ± 0.0 ^e	7.8 ± 0.2 ^a	8.8 ± 0.2 ^b	10.0 ± 0.5 ^c	10.5 ± 0.5 ^c	11.5 ± 0.5 ^d	12.5 ± 0.7 ^d	30.0 ± 0.0
<i>B. subtilis</i> IMR B145	9.2 ± 0.2 ^a	10.2 ± 0.2 ^b	10.8 ± 0.2 ^{b,c}	11.2 ± 0.2 ^{c,d}	11.8 ± 0.2 ^d	12.0 ± 0.5 ^d	6.5 ± 0.0 ^a	6.5 ± 0.0 ^a	8.8 ± 0.2 ^b	10.3 ± 0.5 ^{b,c}	11.7 ± 1.1 ^{c,d}	12.8 ± 0.7 ^d	30.6 ± 0.5
<i>S. pyogenes</i> IMR S1269	8.2 ± 0.2 ^a	9.2 ± 0.2 ^b	10.3 ± 0.2 ^c	11.2 ± 0.2 ^d	11.7 ± 0.2 ^{d,e}	12.3 ± 0.2 ^e	-	7.0 ± 0.0 ^a	7.5 ± 0.5 ^{a,b}	8.7 ± 0.5 ^{b,c}	9.8 ± 0.5 ^{c,d}	10.3 ± 0.5 ^d	26.6 ± 0.5

Continued

<i>S. pyogenes</i>	8.3 ±	9.8 ±	12.5 ±	15.5 ±	17.2 ±	18.3 ±	7.8 ±	9.0 ±	10.2 ±	11.8 ±	12.1 ±	12.4 ±	31.3 ± 0.2
ATCC 19615	0.2 ^a	0.2 ^b	0.5 ^c	0.5 ^d	0.2 ^e	0.5 ^f	0.5 ^a	0.0 ^b	0.2 ^{b,c}	0.2 ^{c,d}	0.5 ^{d,e}	0.5 ^e	
<i>L. monocytogenes</i>	13.0 ±	14.0 ±	15.8 ±	18.2 ±	21.7 ±	24.3 ±	8.8 ±	13.7 ±	15.3 ±	15.7 ±	16.7 ±	17.8 ±	29.3 ± 0.5
IMR L55	0.5 ^a	0.0 ^a	0.2 ^b	0.2 ^c	0.5 ^d	0.5 ^e	1.0 ^a	0.5 ^b	0.2 ^{b,c}	0.5 ^{b,c,d}	0.5 ^{c,d}	1.1 ^d	
<i>E. faecalis</i> IMR	-	-	9.7 ±	10.2 ±	10.7 ±	11.2 ±	-	6.8 ±	7.7 ±	9.2 ±	10.7 ±	11.7 ±	12.8 ± 0.7
E227	-	-	0.5 ^a	0.7 ^a	0.7 ^a	0.7 ^a	-	0.2 ^a	0.2 ^{a,b}	1.0 ^{a,b,c}	0.5 ^{b,c}	0.5 ^c	
<i>E. faecalis</i>	-	8.0 ±	9.2 ±	10.3 ±	11.3 ±	11.0 ±	-	7.8 ±	8.2 ±	9.3 ±	10.2 ±	10.8 ±	23.0 ± 0.5
ATCC 19212	-	0.5 ^a	0.2 ^b	0.5 ^{b,c}	0.2 ^{c,d}	0.2 ^{c,d}	-	0.2 ^a	0.2 ^{a,b}	0.5 ^b	0.2 ^c	0.2 ^{c,d}	
<i>M. luteus</i>	26.7 ±	30.3 ±	31.7 ±	35.0 ±	35.7 ±	36.8 ±	23.0 ±	24.7 ±	25.7 ±	28.0 ±	29.7 ±	31.0 ±	66.330.5
ATCC 10240	0.5 ^a	0.5 ^b	0.5 ^{b,c}	0.0 ^d	0.2 ^d	0.2 ^e	0.5 ^a	0.5 ^{a,b}	0.5 ^b	0.5 ^c	0.2 ^{c,d}	0.2 ^d	
<i>M. luteus</i> IMR	20.7 ±	24.7 ±	26.7 ±	28.3 ±	29.3 ±	31.3 ±	31.0 ±	36.0 ±	37.0 ±	42.7 ±	43.0 ±	46.7 ±	49.6 ± 0.5
M174	0.5 ^a	0.5 ^b	1.1 ^{b,c}	0.5 ^{c,d}	0.5 ^{c,d}	2.0 ^d	0.5 ^a	0.5 ^{a,b}	1.1 ^b	0.2 ^c	0.5 ^{c,d}	2.0 ^d	

-: No inhibition zone; ^AStandard antibiotic (Tetracycline 30 for bacteria species); Negative control (100% water) did not show inhibitory activity; Results are the means of inhibition zone values followed by standard deviations; Different letters within the same row indicate means at different concentrations for the same species are significantly different ($P < 0.05$).

10240 with an inhibition zone of 26.7 ± 0.5 mm. Likewise, *M. luteus* IMR M174 was reported to be the most sensitive candidate towards the crude fruit extract with an inhibition zone of 31.0 ± 0.5 mm. These results validate the traditional medicine claim on *M. malabathricum* Linn. as having therapeutic potency to relief toothache and to treat wounds [5]. *M. luteus* is an opportunistic pathogen, and commonly found in human nasal cavity and on mammalian skin.

Both extracts did not exhibit any effects on the fungal species tested in this study (Table 4). This result is similar to a previous study by [13], who reported that grape seed extract containing pro-anthocyanidin compounds showed no antifungal activity. Similarly, Duraipandiyan and Ignacimutu [14] revealed that the methanol extracts of *Cassia fistula* flower showed no antifungal activity against *C. albicans*. All these findings could be attributed to the nature of the fungal's cell wall structure. *Candida* and *Saccharomyces* are types of yeast which possess polysaccharides, rigid and chemically unique cell walls containing chitin and glucan [15].

The antimicrobial activities of the crude flower and fruit extracts were further confirmed using the microdilution broth assay. The MIC and MMC values were summarized in Table 5. Both extracts exhibited antibacterial activity against the Gram-positive and -negative bacteria, with the MIC values ranging from 12.5 to 100.0 mg/ml for the flower extract and 6.3 to 100.0 mg/ml for the fruit extract. The lowest MIC value of 12.5 mg/ml was obtained using the crude flower extract on *B. cereus* IMR B43, *S. pyogenes* IMR S1269 and *S. pyogenes* ATCC 19615. For the crude fruit extract, the lowest MIC value of 12.5 mg/ml was obtained against *L. monocytogenes* IMR L55, *S. aureus* IMR S942, *S. epidermis* IMR S168, *B. subtilis* IMR B145 and *S. pyogenes* ATCC 19615. The average MMC values obtained varied from 50.0 to 100.0 mg/ml for both extracts.

Overall, the results indicated that Gram-positive bacte-

ria were more sensitive towards the crude flower and fruit extracts. Such observation is consistent with a previous study by Cushnie and Lamb [16], who reported that certain groups of flavonoid compounds exhibited greater inhibition effect on Gram-positive bacteria compared to Gram-negative bacteria. This was also consistent with previous studies on other plant species of *Melastomataceae* [17-19]. The high sensitivity of Gram-positive bacteria may be due to their cell wall and outer membrane structures [20]. Gram-negative bacteria possess an outer membrane and a unique periplasmic space not found in Gram-positive bacteria [21,22]. The resistance of Gram-negative bacteria towards antibacterial substances might also be related to the hydrophilic surface of their outer membrane, composing of lipopolysaccharide molecules, that present a barrier to the penetration of numerous antibiotic molecules and is also associated with the enzymes in the periplasmic space, which are capable of breaking down the molecules introduced from outside [23]. Gram-positive bacteria do not have such outer membrane and cell wall structures. Antibacterial substances can easily destroy the bacteria cell wall and cytoplasmic membrane resulting in a leakage of cytoplasmic materials extracellularly [24].

All concentrations of the extracts tested showed significant differences in the size of the inhibition zone produced (data not shown). This observation revealed that greater inhibition was observed with flower extract compared to fruit extract against all the bacteria tested. This might be due to the presence of different active compounds in the flower and fruit extracts. Susanti [3] reported that the ethyl acetate extract of *M. malabathricum* Linn. flower contained three different compounds such as kaempferol-3-O- β -D-glucoside, kaempferol and naringenin, while the ethyl acetate extract of the fruit contained betulinic acid. Other active compounds such as malvidin-3,5-diglucoside was found in the flowers while cyanidin-3-glucoside and cyanidin-3,5-diglucoside were

Table 4. Inhibition zones (mm) of *Melastoma malabathricum* Linn. extracts against fungi.

Fungi	Inhibition zone (mm)												Antibiotics ^A	
	Flower extract (mg/ml) [extract yield (%) in solvent]						Fruit extract (mg/ml) [extract yield (%) in solvent]							
	100	200	300	400	500	600	100	200	300	400	500	600		
<i>C. albicans</i> ATCC 10231	-	-	-	-	-	-	-	-	-	-	-	-	-	30.0 ± 0.0
<i>C. albicans</i> IMR C451	-	-	-	-	-	-	-	-	-	-	-	-	-	27.0 ± 0.0
<i>S. cerevisiae</i> IMR S1224	-	-	-	-	-	-	-	-	-	-	-	-	-	30.0 ± 0.0

-: No inhibition zone; ^AStandard antibiotic (Nystatin 100 for fungi species); Negative control (100% water) did not show inhibitory activity; Results are the means of inhibition zone values followed by standard deviations; Different letters within the same row indicate means at different concentrations for the same species are significantly different ($P < 0.05$).

Table 5. MIC and MMC (mg/ml) values of *Melastoma malabathricum* Linn. extracts against selected microorganisms.

Microbial strains	Flower extract (mg/ml)		Fruit extract (mg/ml)	
	MIC	MMC	MIC	MMC
Gram-negative bacteria				
<i>Alcaligenes faecalis</i> IMR A111	100.0	100.0	50.0	100.0
<i>Escherichia coli</i> IMR E113	N.D.	N.D.	N.D.	N.D.
<i>Enterobacter aerogenes</i> ATCC 13048	N.D.	N.D.	N.D.	N.D.
<i>Enterobacter aerogenes</i> IMR E153	N.D.	N.D.	N.D.	N.D.
<i>Klebsiella pneumoniae</i> ATCC 33495	N.D.	N.D.	N.D.	N.D.
<i>Klebsiella pneumoniae</i> IMR K36	N.D.	N.D.	N.D.	N.D.
<i>Proteus mirabilis</i> IMR P184	50.0	>100.0	25.0	100.0
<i>Proteus vulgaris</i> IMR P147	100.0	100.0	50.0	100.0
<i>Proteus vulgaris</i> ATCC 8427	N.D.	N.D.	100.0	100.0
<i>Salmonella typhimurium</i> IMR S1205	N.D.	N.D.	N.D.	N.D.
<i>Salmonella typhi</i> IMR S346	N.D.	N.D.	N.D.	N.D.
<i>Shigella dysenteriae</i> IMR S340	12.5	100.0	N.D.	N.D.
<i>Shigella sonnei</i> ATCC 2593	12.5	100.0	N.D.	N.D.
<i>Pseudomonas aeruginosa</i> IMR P187	100.0	100.0	N.D.	N.D.
<i>Serratia marcescens</i> IMR S913	N.D.	N.D.	N.D.	N.D.
Gram -positive bacteria				
<i>Staphylococcus aureus</i> IMR S244	100.0	100.0	100.0	100.0
<i>Staphylococcus aureus</i> IMR S942	100.0	>100.0	12.5	100.0
<i>Staphylococcus epidermidis</i> IMR S168	50.0	50.0	12.5	100.0
<i>Bacillus cereus</i> IMR B43	12.5	50.0	25.0	100.0
<i>Bacillus subtilis</i> IMR B145	100.0	100.0	12.5	100.0
<i>Streptococcus pyogenes</i> IMR S1269	12.5	50.0	N.D.	N.D.
<i>Streptococcus pyogenes</i> ATCC 19615	12.5	50.0	12.5	50.0
<i>Listeria monocytogenes</i> IMR L55	12.5	100.0	12.5	100.0
<i>Enterococcus faecalis</i> IMR E227	N.D.	N.D.	N.D.	N.D.
<i>Enterococcus faecalis</i> ATCC 19212	N.D.	N.D.	N.D.	N.D.
<i>Micrococcus luteus</i> ATCC 10240	100.0	100.0	100.0	100.0
<i>Micrococcus luteus</i> IMR M174	50.0	100.0	100.0	100.0
Fungi species				
<i>Candida albicans</i> ATCC 10231	N.D.	N.D.	N.D.	N.D.
<i>Candida albicans</i> IMR C451	N.D.	N.D.	N.D.	N.D.
<i>Saccharomyces cerevisiae</i> IMR S1224	N.D.	N.D.	N.D.	N.D.

Concentrations of extracts range from 0.39 to 100 mg/ml; N.D.: No activity detected; Results are done in triplicates.

found in the fruits [4]. These previous studies implied that the flower and fruit extracts of *M. malabathricum* Linn. contain different flavonoid compounds. According to Cushnie and Lamb [16], the different structural features of flavonoids may target different cellular components thus affecting different metabolic functions in a bacterial cell.

It is critical to note that an important factor that affects the size of the inhibition zone of antimicrobial activities using the disc diffusion method is the molecular weight of the bioactive compounds in the extract or antibiotics tested. Higher molecular weight antibiotics tend to have difficulty in diffusing into the agar medium, resulting in none or small inhibition zones. As was observed in a previous report that cranberry extract, containing low-molecular-weight phenolic acids and flavanoids such as anthocyanins and flavonols, showed the greatest antibacterial properties [25]. But this does not mean the target organism is sensitive to the agents. According to Cushnie and Lamb [16], incubation at any temperatures other than the optimum often leads to slower growth of the organisms with consequent larger zone. It is clear that whatever temperature chosen for incubation, uniformity of the heating rate of the plates is more important than the absolute value of the incubation temperature. The depth of the medium in the Petri dish should also be of consistent thickness (about 3 - 4 mm) in order to obtain a standard result because the inhibition zone will increase as the thickness of the agar decreases, and the effect is more marked with very thin plate. Among the main sources of error often occur in susceptibility test is the concentration of target organism inoculum [26]. Heavy inoculum tends to give false small inhibition zone, while light inoculum results in false large zone [27]. Therefore, broth dilution method was also carried out in the current study to confirm the results of the disc diffusion assay whereby the MIC and MMC values were determined.

4. Conclusion

This study shows that the crude flower and fruit extracts of *M. malabathricum* Linn. exhibit interesting antimicrobial properties. This may partly explain and support the use of various parts of this plant in traditional medicine for the treatment of infectious diseases caused by microorganisms. However, further studies in determining the bioactive compounds composition, pharmacological action and toxicity of these extracts should be addressed to confirm the therapeutic benefits of the plant prior future applications.

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