

THE EVALUATION OF THE EFFECTIVENESS OF SELECTED ANTIMICROBIAL PLANT EXTRACTS COATING ON TEXTILES

by

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ABSTRACT

Antimicrobial textile development has increased due to microbial growth which is a global health threat facing humanity. Microbial growth can be inhibited by employing an antimicrobial agent or coating application to prevent and treat infections from humans to plants. This study aims to provide information on the effectiveness of the selected plant extracts from South African biomass (*Eucalyptus globulus* and *Artemisia afra*) coated on textiles for application as biomedical textiles. The extracts were prepared by pulsed ultrasound-assisted solvent extraction method. A simple immersion method was used to coat the polyester and cotton fabrics. The extracts have also been analysed for their yields, antimicrobial activity, phytochemical analysis (phytochemical screening) minimum inhibitory concentration, zone of inhibition and washing durability. The methanolic extractions show the highest yields compared to the hexanoic *Artemisia afra* and *Eucalyptus globulus* extractions.

Phytochemical screening of *Artemisia afra* and *Eucalyptus globulus* crude leaf extract was determined. The results showed that all extracts contain the bioactive compounds phenolic compounds, quinones, and sterols.

The *Eucalyptus globulus* hexanoic extracts showed the highest antibacterial activity which has high inhibition zones against Gram-positive bacteria *S. aureus* ATCC 33591 followed by the methanolic *Eucalyptus globulus* extract. *Artemisia afra* methanol extracts showed higher antibacterial activity than that of the hexanoic extract against *S. aureus* ATCC 33591. The methanol extracts had higher inhibition zones against Gram-positive and Gram-negative bacteria.

Among the crude extracts, the highest zone of inhibition (258.4 mm²) was recorded against *S. aureus* ATCC 33591 when the *E. globulus* methanolic extract was applied to the cotton fabric. On the other hand, the lowest inhibition was exhibited on the *E. globulus* methanolic extract coated on polyester fabric (65.97 mm²). Bioactivity was only detected against the *S. aureus* strain. MIC values of the different extracts showed varied results. The MIC for the samples tested is therefore all >500 μ g/ml. The MIC values obtained from antimicrobial tests ranged from 5-500 μ g/ml (Table 5). The results showed that the bacterial strain S. *aureus* ATCC33591 was the most sensitive to *A. afra* methanol extract with a MIC value of 5 μ g/ml to 25 μ g/ml.

Furthermore, the inhibition zone was determined after laundering and no activity was found in the samples. This might be due to the preparation of fabric, inactivation of active compounds during the treatment/preparation of the material samples, or improper binding of the active compounds to the textile material.

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Dedication

This thesis is dedicated to my family. My late mom- Mrs Elizabeth Caroline Nortjie My dad Johannes Nortjie, My brother Heino Nortjie My friend Rainer Absenger The Mannel Family My Friends

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Rest in peace!

"Life is a surf. Love and live life to the fullest." Elvino Nortjie

Thank you!

RESEARCH OUTPUTS

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Contents

| DECLARAT | ΓΙΟΝ | ii |
|------------|---|-----|
| ABSTRACT | ٢ | iii |
| ACKNOWL | EDGEMENTS | iv |
| LIST OF AE | BREVIATIONS AND ACRONYMS | x |
| GLOSSAR | Y | xii |
| CHAPTER | 1: INTRODUCTION | 2 |
| 1.1 | Introduction | 2 |
| 1.2 | Problem Statement | 2 |
| 1.3 | Research Questions | 4 |
| 1.4 | Hypotheses | 4 |
| 1.5 | Aim and Objectives | 4 |
| 1.6 | Delineation of the study | 4 |
| 1.7 | Significance and contribution of the study | 5 |
| 1.8 | General outcomes and contribution of the research | 5 |
| CHAPTER | 2: LITERATURE REVIEW | 7 |
| 2.1 | Introduction | 7 |
| 2.2 | Textiles and Microorganisms | 12 |
| 2.3 | Requirements, Modes of Antimicrobial Action of Antimicrobial Agents | 12 |
| 2.4 | Pre-Treatment and Processing of Biomass | 14 |
| 2.5 | Microorganisms | 17 |
| 2.6 | Brief Description of the Biocide Agents on the Market | 19 |
| 2.7 | Extraction Methods for Studying Phytochemicals | 20 |
| CHAPTER | 3: RESULTS AND DISCUSSION | 31 |
| 3.1 | Introduction | 31 |
| 3.2 | Materials and Methods | 34 |
| 3.3 | Results and Discussion | 40 |
| CHAPTER | 4: CONCLUSIONS AND RECOMMENDATIONS | 63 |
| CHAPTER | 5: REFERENCES | 66 |
| CHAPTER | 6: APPENDICES – RAW DATA | 74 |

LIST OF TABLES

CHAPTER 2

| Table 2-1 Representation of medicinal plant extracts and their applications | 8 |
|--|----|
| Table 2-2 Representation of the mechanism of action of antimicrobial agents | 13 |
| Table 2-3 Representation of commercially available antimicrobials agents on the market | 20 |
| Table 2-4 Extraction methods used in biomass extraction | 24 |
| CHAPTER 3 | |
| Table 3-1 The average yield percentage of the plant extracts obtained by PUAE | 35 |
| Table 3-2 Extraction conditions | 35 |
| Table 3-3 Physiochemical properties of the solvents used | 36 |
| Table 3-4 Phytochemical compounds of A. afra and E. globulus extract | 41 |
| | |

vii

| Table 3.5 Phytochemical compounds and their mechanisms of action | 43 |
|--|----|
| Table 3-6 Summary of the antibacterial activity detected against two ATCC test strains when | 48 |
| a concentration range of 5-500 μ g/ml was evaluated in a 96-well bioactivity assay | |
| Table 3-7 Summary of the bioactivity detected for material samples infused with plant extracts | 49 |
| evaluated against E. coli ATCC 25922 and S. aureus subsp. aureus ATCC 33591 | |

LIST OF FIGURES

CHAPTER 2

| Figure 2-1 <i>Eucalyptus globulus</i> | 10 |
|---------------------------------------|----|
| Figure 2-2 Artemisia afra | 11 |

CHAPTER 3

| Figure 3-1 Process flow diagram of the experimental setup of the antimicrobial textiles | 36 |
|---|----|
| Figure 3-2 Antibacterial Activity of plant extracts <i>A. afra and E. globulus</i> against <i>S. aureus</i> and <i>E. coli</i> . | 46 |
| Figure 3-3 Bioactivity of material samples on agar plates containing the ATCC test strains. A: Light blue/grey cotton material control sample on <i>S. aureus</i> | 50 |
| Figure 3-4 FTIR spectra of polyester fabrics uncoated (U) and coated (1-3) with <i>A. afra</i> hexane coating. Samples 1-12 were performed in triplicates. | 52 |
| Figure 3-5 FTIR spectra of polyester fabrics uncoated (U) and coated (4-6) with <i>A. afra</i> methanol coating | 53 |
| Figure 3-6 FTIR spectra of polyester fabrics uncoated (U) and coated (7-9) with <i>E. globulus</i> hexane coating | 54 |
| Figure 3-7 FTIR spectra of polyester fabrics uncoated (U) and coated (10-12) with <i>E. globulus</i> methanol coating | 56 |
| Figure 3-8 FTIR spectra of cotton fabrics uncoated (U) and coated (1-3) with A. afra Hexane coating | 57 |
| Figure 3-9 FTIR spectra of cotton fabrics uncoated (U) and coated (4-6) with <i>A. afra</i> methanol coating | 58 |
| Figure 3-10 FTIR spectra of cotton fabrics uncoated (U) and coated (7-9) with <i>E. globulus</i> hexane coating | 59 |

Figure 3-11 FTIR spectra of cotton fabrics uncoated (U) and coated (10-12) with *E. globulus* 60 methanol coating.

The layout of the thesis

Chapter 1

Chapter one covers the background and motivation of this study, it describes the problem statement, hypothesis, research aim and objectives and further outlines the delineation of the study.

Chapter 2

This chapter covers the literature review for this study, it focuses on the analysis of the various aspects of producing antimicrobial finishings, the microorganisms, their mechanisms of attachment on natural and synthetic fibres, the effect of microbial growth, and the principal and mechanism of microbial activity of the medicinal plants. Furthermore, the extraction methods, qualitative and quantitative phytochemical evaluations and antimicrobial efficacy and developments of antimicrobial-treated textiles using various agents are covered in the literature review. The chapter was published in the MDPI Plant Journal for 2022.

Chapter 3

This chapter presents the methodology and experimental results of the study. The chapter also discusses the – results, problems faced during the laboratory work and any observations made that could have impacted the outcome of the results.

Chapter 4

This chapter gives recommendations and conclusions based on, emphasizing what can be improved for the experimental work to achieve better results.

Chapter 5

References and Additional information

Chapter 6 Appendices

LIST OF ABBREVIATIONS AND ACRONYMS

| PPE | Personal Protective Equipment | |
|-----------------|--|--|
| A. afra | Artemisia afra | |
| E. globulus | Eucalyptus globulus | |
| S. aureus | Staphylococcus aureus | |
| E. coli | Escherichia coli | |
| A. eupatoria | Agrimona eupatoria | |
| H. alpestre | Hypericum alpestre | |
| R. obrusifollus | Rumex obrusifollus | |
| S. officinalis | Sanguisorba officinalis | |
| PPE | Personal Protective Equipment | |
| WHO | World Health Organization | |
| FTIR | Fourier-transform infrared spectroscopy | |
| HCAIs | Healthcare-related Infections | |
| CDC | Centres for Disease Control and Prevention | |
| AATCC | American Association of Textile Chemists and | |
| | Colorists | |
| rpm | Revolutions per minute | |
| ml | Millilitre | |
| mg | Milligram | |
| nm | Nanometer | |
| μg | Microgram | |
| ISO | International Standards Organization | |
| AATCC | American Association of Textile Chemists and | |
| | Colourists | |
| JIS | Japanese Industrial Standards | |
| PC | Paper Chromatography | |
| TLC | Thin Layer Chromatography | |
| GC | Gas Chromatography | |
| HPLC | High-performance liquid chromatography | |
| QAC | Quaternary Ammonium Compounds | |
| mRNA | messenger Ribonucleic acid | |
| HAI | Health Associated Infections | |
| RNA | Ribonucleic acid | |

| DNA | Deoxyribonucleic acid | |
|---------------|--|--|
| UTI | Urinary Tract Infection | |
| ESKAPE | Enterococcus faecium, Staphylococcus aureus, | |
| | Klebsiella pneumoniae, Acinetobacter baumanni, | |
| | Pseudomonas aeruginosa, Acinetobacter | |
| | aerogenes. | |
| PUAE | Pulsed Ultrasound-Assisted Extraction | |
| MIC | Minimum Inhibitory Concentration | |
| ZOI | Zone of Inhibition | |
| UV | Ultraviolet | |
| COVID-19 | Coronavirus Disease of 2019 | |
| W/V | Weight-to-volume ratio | |
| FTIR | Fourier transform infrared spectrometry | |
| N/TT | (3-(4,5-Dimethylthiazol-2-yl)-2,5- | |
| MTT | Diphenyltetrazolium Bromide) | |
| S. pyogenes | Streptococcus pyogenes | |
| H. influenzae | Hemophilus influenzae | |
| PPL | Priority Pathogen List | |
| GRAS | Generally Recognized as Safe | |
| | | |

GLOSSARY

| a) | AATCC147 Method | The AATCC 147 technique evaluates the | |
|----|-------------------------|---|--|
| | | antibacterial action of diffusible antimicrobial | |
| | | agents on treated fabric and assesses the | |
| | | capability of the treated fabric to inhibit the | |
| | | multiplication of microorganisms. | |
| b) | Antibacterial: | Can be defined as an agent that inhibits bacterial | |
| | | growth. | |
| C) | Antimicrobial activity: | Can be defined as a collective term for all active | |
| | | agents that inhibit the growth of bacteria, prevent | |
| | | the formation of microbial colonies | |
| d) | Antimicrobials: | Is a subclass of antibacterial that kills or inhibits | |
| | | microbial growth of microorganisms including | |
| | | bacteria, viruses, fungi, mould, and protozoans. | |
| e) | Pathogens: | Can be defined as an organism causing disease | |
| | | to its host. Every living organism is affected by | |
| | | pathogens, including bacteria. | |
| f) | Phytochemicals: | Can be defined as chemical compounds that are | |
| | | biologically active compounds found in plants. | |
| g) | Sonicator Apparatus: | Is an apparatus used to extract several types of | |
| | | material with volatile solvents. The sonicator | |
| | | applies sound energy to agitate particles in liquid | |
| | | media reaching ultrasonic frequencies of more | |
| | | than 20 kHz. | |
| h) | Zone of Inhibition: | It is the area of media where bacteria do not grow. | |

CHAPTER 1

INTRODUCTION

CHAPTER 1: INTRODUCTION

1.1 Introduction

The development of antimicrobial textiles has become of utmost importance for enhanced durability of the fabrics' intended use. Antimicrobial textiles are defined as functional textiles that may kill or inhibit the growth of pathogens (Gulati et al., 2022). Cotton and polyester are commonly used textiles being used in everyday life due to their properties that include comfort, strength, handling, and durability. They are also prone to microbial attacks. Cotton is a natural fabric that bears an environment for microbial growth depending on the growth of microorganisms e.g., pH, temperature, time, oxygen, and moisture (Gao et al., 2008). Bacterial growth leads to a bad odour and a reduction mechanical strength of fibres. The microbial growth on polyester fabric is slower than that of cotton due to its polymer backbone. Pathogens responsible for cross-contamination and infections include *Escherichia coli, Staphylococcus aureus, Klebsiella pneumoniae, Acinetobacter baumannii, Pseudomonas aeruginosa* and *Enterobacter* spp.

The ideal antimicrobial coating on textiles against pathogens should satisfy the following requirements (Chauhan & Kumar, 2020):

- 1. be effective against a broad spectrum of pathogens but exhibit low toxicity to the user.
- 2. be durable to laundering.
- 3. It should not negatively affect the quality or appearance of the textile.
- 4. be cost-effective

Antimicrobial textiles are required in the apparel, commercial, healthcare and household sector (Gulati et al., 2022), namely:

- Caps, jackets, sanitary pads, sportswear, undergarment, and winter wear in the apparel sector.
- Carpets, covering for seats, windows, vehicles, dusting cloths, military fabrics, tents, and uniforms in the commercial sector.
- Bandages, earbuds, scrubs, masks, lab coats and protective kits in the healthcare sector.
- Bedding, carpets, covers, curtains, mops, pillows, and towels in the household sector.

1.2 Problem Statement

Textiles are regarded as the only barrier between humans and pathogens. Antimicrobialresistant pathogens are a threat to human health, especially healthcare-associated infections. The COVID-19 pandemic crisis proved the importance of personal hygiene and PPE. The World Health Organization suggested and implemented the use of Personal Protective Equipment (triple-layered face masks, bodysuits), social distancing (Chauhan & Kumar, 2020) and disinfectant recipes to sanitize surface areas etc.

Large surface areas and moisture content favour the growth of microbes and lead to odour. In textiles, microbial growth is also led by moisture, pigmentation, and deterioration in quality, which affect the products and the wearer (Reshma et al., 2022). The attachment of the bacteria onto the fabric is dependent on the characteristics of the fabric and the contact time of the microbe onto the fabric, surface roughness and moisture retention. Natural and synthetic fibres react differently to microbial growth. Natural fibres are more prone to microbial attack because they retain water easily. Microbial growth on synthetic fibres like polyester is slower due to their polymer backbone (Gupta & Bhaumik, 2007). Cotton and polyester fabrics were chosen for this study due to their high absorbency and being the most widely used available fabrics in the world.

Antimicrobial textile agents and coatings allow the re-use of face masks, clothing, and textiles, reducing domestic laundering that results in a reduction in water consumption, and curtailing the worldwide pandemic, global warming, and environmental degradation.

This study focused on natural plant-based antimicrobial agents because it is less poisonous, less irritant, environmentally safe, and therefore used as antimicrobial finishes on textiles. Consumers have become aware of the harmful antimicrobial agents and the exposure of harmful microorganisms to humans. There are limitations in extracting from natural plants e.g., the bioactive compounds could be destroyed when subjected to high temperatures and extraction times and might influence the yield. The Pulsed Ultrasound-Assisted Extraction (PUAE) method was used in this study.

This research aimed at developing an eco-friendly plant-based antimicrobial coating from *Artemisia afra* and *Eucalyptus globulus* plant extracts coated on textile surfaces to combat the growth of pathogens. The extracts were applied to the polyester and cotton fabric. The yield, phytochemical screening, antibacterial analysis, minimum inhibition concentration and zone of inhibition of the extracts and washing durability were studied. Due to the nature of the extraction and application process of the plant extracts onto fabric and testing against Gramnegative and Gram-positive bacteria, this study qualifies under bioengineering. Overall, studies show there is a need for more effective ways to produce antimicrobial agents and coatings. This study is aligned with the same vision of providing a greener approach to antimicrobial plant extractions.

1.3 Research Questions

- 1.3.1 Do the drying conditions, extraction method, time and temperature affect the antimicrobial activity of the extracts?
- 1.3.2 Which extraction method that uses polar or non-polar solvents is more effective?
- 1.3.3 Which medicinal plant extract from the *Artemisia afra* and *Eucalyptus globulus* plant shows the best antimicrobial activity?
- 1.3.4 How effective are the antimicrobial coatings against the S. aureus and E. coli bacteria?
- 1.3.5 How effective is the application of the coating on polyester and cotton fabric for laundering?

1.4 Hypotheses

The plant extracts (*A. afra* and *E. globulus*) will inhibit microbial growth on textiles (*E. coli* and *S. aureus*).

1.5 Aim and Objectives

1.5.1 Aim

The aim is to evaluate the effectiveness of the selected plant extracts from South African biomass (*Eucalyptus globulus* and *Artemisia afra*) for possible textile coating or antimicrobial agents for surfaces.

1.5.2 Objectives

The objectives of this study are:

- 1.5.2.1 Prepare extracts containing antimicrobial compounds by pulsed ultrasound-assisted extraction using polar and non-polar solvents separately.
- 1.5.2.2 To conduct qualitative phytochemical screening analysis on *Eucalyptus globulus* and *Artemisia afra* to identify the bioactive compounds from the selected plants.
- 1.5.2.3 Apply the extracts onto cotton and polyester fabrics and assess the antimicrobial activity before and after laundering. The coated samples will be analysed using Fourier-transform infrared spectroscopy (FTIR).

1.6 Delineation of the study

This study will not look at:

- An economic evaluation of the process
- Composition of the compounds
- The toxicity of the compounds
- Quantitative analysis of the extracts

1.7 Significance and contribution of the study

The development of antimicrobial agents and coating for textiles is necessary to combat pathogens. Researchers are focusing on the development of natural antimicrobial agents and enhancement of the coating for washing durability of natural antimicrobial coated fabrics after washing (Reshma et al., 2022).

Antimicrobial coatings have become a highly active field of research, strongly stimulated by the increasing urgency of identifying alternatives to the traditional administration of antibiotics. This study plays a pivotal role in advancing green extraction techniques and shedding light on the antimicrobial effects when employing the pulsed ultrasound-assisted extraction approach, in contrast to conventional extraction methods.

1.8 General outcomes and contribution of the research

The purpose of the study was to evaluate the efficacy of the antimicrobial coating using selected medicinal plant extracts coated on the fabric. This study investigated the antimicrobial activity and the laundering effects of the antimicrobial coating in clothing and textiles. The research findings were written up as a master's thesis and published in scientific journals.

CHAPTER 2

LITERATURE REVIEW

Extraction Methods, Quantitative and Qualitative Phytochemical Screening of Medicinal Plants for Antimicrobial Textiles: A Review

Nortjie, E.; Basitere, M.; Moyo, D.; Nyamukamba, P. Extraction Methods, Quantitative and Qualitative Phytochemical Screening of Medicinal Plants for Antimicrobial Textiles: A Review. *Plants* 2022, *11*, 2011. <u>https://doi.org/10.3390/plants11152011</u>

This chapter was published in the MDPI Plants Journal 2022

CHAPTER 2: LITERATURE REVIEW

2.1 Introduction

Antibiotics play a vital role in fighting bacterial infections, but antibacterial resistance has caused havoc in the healthcare and pharmaceutical sector that accelerates socio-economic losses (O'Neill, 2016). Multidrug Resistance is said to increase by 10 million deaths per year by 2050 (O'Neill, 2016; Kraker et al., 2016). Biological screening, separation of the phytochemicals, and clinical trials of the medicinal plants have advanced over the years unfolding the secrets of ancient herbal remedies (Bibi et al., 2017). Traditional medicine is effective in dealing with diseases caused by bacteria or oxidative stress (Baydoun et al., 2015; Singh et al., 2017; Restuati & Diningrat, 2018).

Natural compounds have been extensively explored for new drug discoveries (Chandra et al., 2017). Humanity has always been fascinated by natural compounds from pre-biotic, microbial, plant, and animal sources. Extracts of different parts of plants contain bioactive compounds that fight against diseases such as alkaloids, steroids, tannins, glycosides, volatile oils, fixed oils, resins, phenols, terpenoids, and flavonoids (Kutama, 2018) .The phenolic phytochemicals from plants play a key role as antimicrobial agents (Restuati & Diningrat, 2018; Smeriglio et al., 2017). Antimicrobial agents decay the protein components of the cell wall, disrupting the work of enzymes and DNA and RNA replication (Bakal et al., 2017) . Table 2.1 shows a selection of plants, their phytochemicals responsible for antimicrobial activity, and their applications.

| Plant Name | Phytochemicals | Applications |
|-------------------------|--|---|
| Sutherlandia fructecens | Saponins, pinitols, flavonoids, triterpenoids, canavanine, cycloartane glycosides, flavanols glycosides, and aminobutyric acid (Dube et al., 2017). | Wound treatment, cancer treatment, diabetes, skin diseases, rheumatism, urinary tract infection, fever, gonorrhea, kidney, and liver problems (Dube et al., 2017). |
| Eucomis autumnalis | Homoisoflavanones, terpenoids, and dibenzo-α- pyrones (Alaribe et al., 2018). | Reducing fever, urinary diseases, stomach, lower backaches, and syphilis. <i>Eucomis autumnalis</i> is sometimes used to induce labour (Alaribe et al., 2018). |
| Plumbago auriculata | Tannins, phenols, alkaloids, saponins, flavonoids, plumbagin, α-amyrin, capensisone, and diomuscinone (Lediga et al., 2018) | Treating headaches, warts, skin infections, wounds, and fractures (Lediga et al., 2018) |
| Catharanthus roseus | Vinblastine, deoxyvinblastin, vincoline, cathanranthamine, rosicine, leurosine, vindoline and vincristine (Gajalakshmi et al., 2011). | Treating rheumatism, venereal diseases, skin infections, high blood pressure, and diabetes (Oguntibeju et al., 2019) |
| Aspalathus linearis | Spalathin, orientin, isoquercitrin, and luteolin hyperoside (Oguntibeju et al., 2019). | Treat insomnia, stomach cramps, allergies, and digestive problems as well as improve appetite (Mordeniz, 2019b). |
| Centella asiatica | Triterpenoids, centellose, medacassoside, triaponosides, flavonoid quercetin, rutin, | Treating fever, leprosy, syphilis, tuberculosis, leprosy, asthma, epilepsy, mental |

| | kaempferol, patuletin, apigenin, polyacetylenes, phenolic acids, sterols (Rout et al., 2013). | disorder, and minor wounds. Consumed as a vegetable and used as a spice (Rout et al., 2013). |
|------------------------|---|---|
| Sclerocarya birrea | Steroids, glycosides, flavonoids, fatty oils, alkaloids, phenols, resins, calcium, and phosphorus (Joubert et al., 2011). | Treating dysentery, rheumatism, malaria, and diarrhoea (Joubert et al., 2011). |
| Hypoxis hemerocallidea | Rooperol, β-sitosterol (Owira & Ojewole, 2009) . | Immune booster, purgative, and laxative tonic. Treat tuberculosis, urinary tract infection, infertility, cancer, diabetes, and wounds (Owira & Ojewole, 2009). |
| Galenia africana | Trihydroxyflavanone, trihydroxychalcone, dihydroxychalcone, and trihydroxy-3-methoxychalcone (Elbagory et al., 2017). | Treat venereal sores, eye infections, asthma, tuberculosis, cough, wounds, and skin infections and relieve toothache (Elbagory et al., 2017). |

2.1.1 Eucalyptus globulus

The Eucalyptus species is a well-known plant species for its bioactive and pharmacological properties. It is the most represented species in the international pharmacopoeia. The essential oil compounds of the genus Eucalyptus (Myrtaceae) have been well characterized (Bakkali et al., 2008). *Eucalyptus globulus* is known to be the main furnisher of essential oils and is high in demand in the soap and cosmetic industries (Bajaj, 1995). It is a tree that grows to a height of 60-80 m with smooth bark and v yellow-brownish in colour and sheds in long strips leaving a white-greyish colour on the surface of the bark (Pereira et al., 2014; Araújo et al., 2010). Flowering occurs between May and January and the flower buds appear to be in leaf axils. The fruits are woody circular capsules that close at the edge in Figure 2.1 below (Mason et al., 2000; Mulyaningsih et al., 2010).



Figure 2.1: Eucalyptus globulus

The Eucalyptus oil extracted from the leaf contains volatile terpenes and aromatic compounds and the most abundant monoterpenoid 1,8-cineole (Pino et al., 2006; Brophy et al., 2009; Luís et al., 2016). British and European pharmacopoeias guide to using eucalyptus oil for medicinal purposes is that it must least contain at least 70% 1,8-cineole. The medicinal use of the Eucalyptus globulus is based on its biological activities shown by the oils e.g., antioxidant (González-Burgos et al., 2018), anti-inflammatory, analgesic (Silva et al., 2003), and antimicrobial activities (O. Oyedeji et al., 1999; Takahashi, Kokubo and Sakaino, 2004; Gilles et al., 2010; Sidana et al., 2010; Djenane et al., 2011; González-Burgos et al., 2018). Clinical trials with eucalyptus oil and the major component 1,8-cineole (eucalyptol) have been conducted to evaluate their efficacy in the treatment of a diverse range of conditions and diseases, including respiratory disorders (Worth, Schacher and Dethlefsen, 2009), and head lice infestation (Barker and Altman, 2011) The antimicrobial capabilities of the essential oils have been reported by the different species (Mulyaningsih et al., 2011; Ghaffar et al., 2015). The Eucalyptus globulus leaf extract shows antibacterial activity against bacteria responsible for respiratory tract infections such as Staphylococcus aureus ATCC 33591, Streptococcus pyogenes (S. pyogenes), and Hemophilus influenzae (H. influenzae).

2.1.2 Artemisia afra Jacq.

Artemisia afra is known as the Wild wormwood, African wormwood (Eng.); Wilde-Als (Afr.); umhlonyane (isiXhosa); mhlonyane (isiZulu), a soft aromatic shrub that grows up to 0.6-2 m (Figure 2.2). It is one of the most popular medicinal plants in South Africa. *A. afra* is a common species in South Africa with a wide distribution from the Cederberg Mountains in the Cape, northwards to tropical East Africa and stretching as far north as Ethiopia (van Wyk, 2008).



Figure 2.2: Artemisia afra Jacq.

Artemisia afra covers a wide range of ailments such as coughs, colds, fever, loss of appetite, colic, headache, earache, intestinal worms to malaria (van Wyk, van Oudtshoorn and Gericke, 1997). The *Artemisia afra* species can be administered in a variety of ways for personal medicinal use like enemas, infusions to cosmetics (Patil et al., 2011). It's also used as a preservative in food (Muyima et al., 2002) as well as an insecticide (Patil et al., 2011) The uses of the *Artemisia afra* have been studied for its antimicrobial activities (More et al., 2012; Martini et al., 2020) and its antifungal properties, It was found that it possesses antifungal activity (José Abad et al., 2000) as well as inhibitory activity against a broad spectrum of Gram-positive and Gram-negative bacteria (Jager, 2003; More et al., 2012; Martini et al., 2020; Graven et al., 1992; Muyima et al., 2002).

2.2 Textiles and Microorganisms

Textiles are carriers of microorganisms and are subjected to the growth of microorganisms, such as bacteria and fungi, depending on the food, acidic Ph, temperature, time, oxygen, and moisture (Dube et al., 2017). Bacteria interrelates with fibres in phases, from initial attachment onto fibres to the growth and damage to the fibres (Alaribe et al., 2018). Cotton is one of the ideal natural fibre fabrics for the growth of pathogens than polyester. Neely & Maley (2000) has shown the survival of several Gram-positive bacteria (*Staphylococcus aureus, Enterococcus faecalis*) on standard hospital fabrics made of 100% cotton clothing, 100% cotton terry towels, 60:40 cotton/polyester-scrub suits and lab coats, and 100% polyester drape.

A study by Neely & Maley (2000) showed the growth of bacteria within 48 hours, most bacterial growth survived at least a day, and some survived more than 90 days. Natural fibre textiles are more prone to microbial growth and could lead to the spread of infections (Gajalakshmi et al., 2011). A study by Gupta & Bhaumik (2007) reports that the attachment of the bacteria onto the fabric is dependent on the characteristics of the fabric, the contact time of the microbe onto the fabric, surface roughness, and moisture retention for natural and synthetic fibres reacting differently to microbial growth (Kraker et al., 2016). Natural fibres are more prone to microbial attack because they retain water easily. Microbial growth on synthetic fibres like polyester is slower due to their polymer backbone (Alaribe et al., 2018).

2.3 Requirements, Modes of Antimicrobial Action of Antimicrobial Agents

The ideal antimicrobial treatment for textiles must be effective against a broad spectrum of pathogens but exhibit low toxicity to the user. It must be cost-effective, durable to launder, and not alter the quality or appearance of the textile (Chauhan & Kumar, 2020).

A study by Gao (2008) reported that microbes are microscopic organisms that exist as unicellular, multicellular, or cell clusters. They consist of an outermost cell wall that constitutes polysaccharides. The cell wall maintains the integrity of cellular components and shields the cell from the extracellular environment. Beneath the cell wall is a semi-permeable membrane that encloses intracellular organelles and multiple enzymes and nucleic acids. The enzymes are responsible for the chemical reactions within the cell, followed by the storage of nucleic acid genetic information of the organism. The purpose of antimicrobial agents is to destroy the cell wall or alter cell membrane permeability, denature proteins, inhibit enzyme activity, or inhibit lipid synthesis so that the cell does not survive.

Various classes of phytochemicals and antimicrobial agents possess different mechanisms of action against microbes.

As for phytochemicals tested in this study, their activity and mechanism of action are described below (Tiwari et al., 2011):

- Quinones: possess antimicrobial activity which binds to adhesins and inactivates enzymes.
- Flavonoids: possess antimicrobial activity which is and binds to adhesins
- Phenol and tannins: possess antimicrobial activity which binds to adhesins, enzyme inhibition, substrate deprivation, membrane disruption, and metal ion complexation.
- Terpenoids: disrupts the membrane of microorganisms.
- Saponins: inhibits Gram-positive and Gram-negative bacteria.

Table 2.2 shows the different antimicrobial classes, the mechanisms of action, and the activity spectrum, respectively (Tille, 2014). Adapted with permission from Ref. (Tille, 2014). 2014, Dr Patricia Tille.

Table 2.2:Representation of mechanisms of action of antimicrobial agents (Tille, 2014).Adapted with permission from Ref. (Tille, 2014). 2014, Dr Patricia Tille.

| Antimicrobial Class | Mechanism of Action | Activity Spectrum | |
|---------------------|---------------------------------------|-------------------------------------|--|
| | They inhibit cell wall synthesis by | Gram-negative bacteria and | |
| β-lactams | binding enzymes in peptidoglycan | Gram-positive bacteria could differ | |
| | production | with individual antibiotic | |
| Aminoglycosides | Hinders the protein synthesis by | Gram-negative bacteria and | |
| Ammogrycosides | binding 30S ribosomal subunits | Gram-positive bacteria | |
| Chloramphonical | Inhibits the protein synthesis by | Gram-negative bacteria and | |
| Chioramphenicol | binding 50S ribosomal subunits | Gram-positive bacteria | |
| | | Gram-negative bacteria and | |
| Flueroquinelence | Inhibits DNA synthesis by binding | Gram-positive bacteria, but it | |
| riuoroquinoiones | the DNA gyrase topoisomerase IV | could differ with individual | |
| | | antibiotic | |
| | Inhibite the protein synthesis by | A wide spectrum of Gram-negative | |
| Glycylglycines | hinding EOS ribosomal units | bacteria and Gram-positive | |
| | binding 505 hbosonial units | species | |
| | Inhibita protain aunthonia bu hinding | Gram-positive cocci including | |
| Ketolides | | certain macrolide resistance | |
| | 50S ridosomai sudunits | strains and Gram-negative strains | |
| Linonontidoo | Binding and disruption of cell | Gram-positive bacteria including | |
| Lipopeptides | membrane | β-lactams and glycopeptides | |
| | The mechanism is unknown and | Crom pogative basteria and | |
| Nitrofurantoin | may have bacterial enzyme targets | Gram positive bacteria | |
| | and damaging DNA. | Grani-positive bacteria | |

| | Hinders the initiation of protein | Wide variety of Gram-positive | | |
|----------------|---------------------------------------|------------------------------------|--|--|
| Oxazolidinones | synthesis by binding 50S ribosomal | bacteria including those resistant | | |
| | subunits | antimicrobial classes | | |
| | | Poor activity against most Gram- | | |
| Polymyxins | Disrupts cell membrane | positive bacteria. Gram-negative | | |
| | | bacteria | | |
| Pifamnin | Hinders RNA synthesis by binding | Gram-positive and certain Gram- | | |
| Rilampin | DNA dependent, RNA polymerase. | negative bacteria | | |
| | Hinders the protein synthesis by | Gram-positive bacteria | | |
| Streptogramins | binding two separate sites on the | | | |
| | 50S ribosomal subunit | | | |
| | | Gram-negative bacteria and | | |
| Tetracycline | Inhibits protein synthesis by binding | Gram-positive bacteria and | | |
| retrucyonne | of 30S ribosomal subunit | several intracellular bacterial | | |
| | | pathogens | | |
| | Hinders the folic acid pathway, | Gram-negative bacteria and | | |
| Sulfonamides | binding the enzyme dihydropteroate | Gram-positive bacteria | | |
| | synthase | | | |
| | Hinders with the folic acid pathway | Gram-negative bacteria and | | |
| Trimethoprim | by binding the enzyme dihydrofolate | Gram-positive bacteria | | |
| | reductase | • | | |

Textiles are regarded as a barrier between humans and pathogens. Plant-based extracts and materials provide efficient and natural microbial resistivity. Antimicrobial textiles are essential in the apparel, commercial, and healthcare sector (Gulati et al., 2022). A study by Vastrad et al., (2016) reported on the evaluation of total phenolic content and flavonoid content using leaf extracts (eucalyptus and lemon grass) with methanol, ethanol, chloroform, and distilled water extract indicated the potential of antimicrobial application of textiles. The antimicrobial agents and finishing on textiles may allow the re-use of face masks, and clothing, reducing PPE kits in health care, and reducing domestic laundering that may lead to a reduction in water consumption.

2.4 Pre-Treatment and Processing of Biomass

2.4.1 Drying of Biomass

The selection of pre-treatment and processing methods may influence the reduction in extraction time, an increase in extraction yield, the quality of the biological compounds, and a reduction in input energy (Safarzadeh Markhali, 2021). The drying of any biomass inhibits microbial growth (Joubert et al., 2011), and it aids in the longer shelf life and transportation costs due to the weight and space of dry products (Pham et al., 2015; Saifullah et al., 2019;

Shrestha et al., 2007). Drying can affect the phytochemical components of the thermally sensitive components (Pham et al., 2015; Nóbrega et al., 2015; Nadi, 2017; Nguyen et al., 2015), and the process can also contribute to improved conservation of the bioactive compounds against oxidative (Ahmad-Qasem et al., 2013) and enzymatic activities (Babu et al., 2018) and spoilage bacteria (Saifullah et al., 2019; Afaneh et al., 2015; Orphanides et al., 2013), enabling cellular destruction (Saifullah et al., 2019; Ahmad-Qasem et al., 2013; Safarzadeh Markhali, 2021). There are many different drying methods, e.g., thermal through natural convection (shade and open sun drying), forced convection (oven drying, solar drying, and heat pump drying), freeze-drying, greenhouse drying, microwave drying, and infrared drying (Safarzadeh Markhali, 2021).

The freeze-drying method retains the bioactive compounds of the dried product due to minimal thermal damage to the cell tissue, thermolabile compounds, and its porous surface, enabling increased penetration of solvents (Saifullah et al., 2019; Harnkarnsujarit et al., 2016; Ahmad-Qasem et al., 2013). Olive leaf extracts pre-treated with a hot air drier at 120 °C showed higher phenolic recovery compared to freeze-drying (loss of polyphenols reached up to 39% in dry weight). Freeze drying shows great potential in the extraction of the total phenolic content (Ahmad-Qasem et al., 2013; Ghelichkhani et al., 2019).

Ahmad-Qasem (Ahmad-Qasem et al., 2013) reported that temperature plays a key role in the drying process as it may be beneficial or unfavourable to the microstructure of the biomass and the use of hot air drying at a high temperature. The study by Ahmad-Qasem (Ahmad-Qasem et al., 2013) also reported better extraction efficacy of some phenolic compounds in olive leaves when compared to samples dried at lower temperatures and by drying at a moderate to low temperature may need a longer drying time to reach the desired moisture content of the biomass.

2.4.2 Choice of Solvents

The solvent selection is crucial in determining the bioactive compounds of plants used for extractions. Ideal extraction solvent properties include low toxicity, evaporating easily at low temperatures, having good solubility of the target compound, and being sufficiently volatile. The factors affecting the selection of solvents are the rate of extraction, diversity of compounds extracted, ease of handling of extracts, and the cost-effectiveness of the extraction solvents and targeted compounds. Plants contain diverse bioactive compounds with different polarities. A variety of techniques have been devised and employed to isolate pure compounds, thereby identifying their structures and biological activities (Sasidharan et al., 2011).

Many solvent extractions have been done to obtain phytochemical compounds for their activity against pathogens. Various phytochemicals possess distinct structural characteristics and comprise diverse compounds, each exhibiting unique action mechanisms, as elaborated below:

- Phenols and polyphenols are obtained from acetone and ethanol solvent extractions which consist of a C₃ sidechain, hydroxyl groups and a phenol ring e.g., catechol, epicatechin, cinnamic acid that has antimicrobial, anthelmintic, and antidiarrheal activity (Cowan, 1999; Tiwari et al., 2011).
- Chloroform, methanol, and ethanol solvents extract mainly quinones. They consist of aromatic rings, and two ketone substitutions e.g., hypericin which has antimicrobial activity (Cowan, 1999; Tiwari et al., 2011).
- Ethanol and water mainly extract tannins which consist of polyphenols e.g., ellagitannin which has antimicrobial anthelmintic and antidiarrheal activities (Cowan, 1999; Tiwari et al., 2011).
- Chloroform solvents extract mainly flavonoids which consist of a phenolic structure, a carbonyl group, hydroxylated phenols C₃-C₅ unit linked to an aromatic ring, flavones and a +3-hydroxyl group that has antimicrobial, anthelmintic and antidiarrheal activity (Cowan, 1999; Tiwari et al., 2011).
- Ether solvent extracts mainly coumarins and it consists of phenols made up of fused benzenes e.g., warfarin with antimicrobial activity (Cowan, 1999; Tiwari et al., 2011).
- Water, ethanol, chloroform, and ether solvents extract mainly terpenoids which consist of fatty acids and acetate units with antimicrobial activity (Cowan, 1999; Tiwari et al., 2011).
- Lectins and polypeptides can be extracted by water which consists of mainly extracts proteins e.g., mannose-specific agglutinin, and fabatin that has antimicrobial activity (Cowan, 1999; Tiwari et al., 2011).
- Alkaloids can be extracted by ethanol and ether solvents which consist of heterocyclic nitrogen compounds e.g., berberine, piperine, palmatine and tetrahydropalmatine which has antimicrobial, anthelmintic and antidiarrheal activity (Cowan, 1999; Tiwari et al., 2011).
- Saponins can be extracted by methanol, water, and hydro-alcoholic 70 % methanol which consists of amphipathic glycosides e.g., vina-ginsenosides R5-R6 with antidiarrheal activity. (Cowan, 1999; Tiwari et al., 2011).

During extraction, solvents diffuse in the plant material and dissolve compounds with similar polarity. The different extraction methods will also affect the composition of the secondary metabolites of the extracts namely, type of extraction, time of extraction, temperature and nature of the solvent, solvent concentration, and polarity.

The determination of biologically active compounds from plant materials is crucial and dependent on the type of solvent used (Velavan, 2015). Solvents are selected based on their availability, low toxicity, boiling point and solvent polarity (Das et al., 2010; Kumar, 2015).

The FAO/WHO Expert Committee reported seventeen solvents that are generally recognized as safe (GRAS) to use for food and personal-care products.

2.5 Microorganisms

Resistance to antibiotics has become a serious problem globally. ESKAPE are multidrugresistant pathogens such as *Enterococcus faecium, Staphylococcus aureus, Klebsiella pneumoniae, Acinetobacter baumannii, Pseudomonas aeruginosa*, and *Enterobacter* species that are responsible for Hospital-Acquired Infections (HAI). New antibiotics have been produced over the years. The resistance by the ESKAPE pathogens to the drugs has accelerated tremendously (FAO/WHO, 2018). A Priority Pathogen List (PPL) was released by the WHO in 2016 as a guide to research, discovery, and development of new antibiotics globally (Bhatia et al., 2021). Pathogens occupy the surfaces of fabrics depending on the contact time, moisture retention, and surface roughness. *Staphylococcus aureus and Escherichia coli* pathogens cause hospital infections leading to pneumonia and sepsis. It is, therefore, important to keep track of the availability of alternative medicinal plants and herbs to conquer this challenge (World Health Organisation., 2017). The discovery of new drugs that can be mastered with the use of plant extracts is a hoard of a spectrum of secondary metabolites (Ghareeb et al., 2015; Ugwoke et al., 2020; Cragg & Newman, 2013; Madubuonu et al., 2019; Aisida et al., 2020; Kandemir et al., 2020; Ginting et al., 2020).

2.5.1 Enterococcus faecium

Enterococcus faecium Enterococcus faecium is a Gram-positive bacterium that causes infections; it is increasingly resistant compared to *Vancomycin-resistant Enterococci faecium* (Aisida et al., 2019; Lebreton et al., 2013). *E. faecium* lives in the gut microbiome of animals (Pendleton et al., 2013; Marques et al., 2018). Food is an excellent hideout for the strains to remain dormant (Fiore et al., 2019). Treatment is dependent on second-line antibiotics (Ubeda et al., 2010). Urinary tract infections, bacteraemia, and endocarditis are caused by this bacterium (Marques et al., 2018).

2.5.2 Staphylococcus aureus

Staphylococcus aureus is a Gram-positive bacterium prevalent on the human skin, particularly in immune-compromised individuals. This bacterium causes infections on medical implants and forms biofilms that make it extremely difficult to treat with antibiotics. The Methicillin-resistant *Staphylococcus aureus* developed resistance against β -lactam antibiotics (Lebreton et al., 2013). Community-associated MRSA lineages are associated with skin and soft tissue infections. The strains are associated with pneumonia and bloodstream infections (Prasad et al., 2012).

2.5.3 Klebsiella pneumonaie

Klebsiella pneumonaie is a Gram-negative bacterium that causes urinary tract infections septicaemia, surgical wound infections, pneumonia, endocarditis, pyogenic liver abscess cystitis, and endogenous endophthalmitis (DeLeo et al., 2010). Cephalosporin- and carbapenem-class antibiotics have been the base treatment for *Enterobacterales* infections, such as *Klebsiella pneumoniae*. The efficacy of the antibiotics is compromised by the widespread acquisition of genes and encoding enzymes that aid in the respective resistance to these critical drugs (Navon-Venezia et al., 2017).

2.5.4 Acinetobacter baumannii

Acinetobacter baumannii is a Gram-negative bacterium that is more common in hospital settings (Paterson & Bonomo, 2005). It is aerobic and non-fermenting pleomorphic. This bacterium can resist dehydration. It forms biofilms, surface adhesins, that help this bacterium thrive in its environment (Munoz-Price & Weinstein, 2008). The infection rates of *A. baumannii* are low compared to the ESKAPE pathogens (Magill et al., 2014).

2.5.5 Pseudomonas aeruginosa

Pseudomonas aeruginosa is a Gram-negative bacterium associated with respiratory infections and displays resistance to multiple classes of antibiotics (Magill et al., 2014). *P. aeruginosa* grows and colonizes in moist environments, especially in healthcare settings in the context of chronic wounds, respiratory support, or urinary tract devices, immune evasion, and antimicrobial resistance (Yayan et al., 2015).

2.5.6 Enterobacter spp.

It's a Gram-negative bacterium, anaerobic in nature. *Enterobacter aerogenes, known as Klebsiella aerogenes* are responsible for the increasing number of hospital-acquired infections [79]. Immunocompromised individuals are more susceptible to urinary and respiratory tract infections of this bacterium (Malek et al., 2019).

2.5.7 Escherichia coli

Escherichia coli, which is not part of the ESKAPE pathogens is the major cause of bloodstream and urinary tract infection (UTI) in both community and healthcare settings globally. Sepsis is one of the most common manifestations of *E. coli* urinary tract infection. *E. coli* is the most common Gram-negative bacterial species isolated from blood and urine cultures (Reza et al., 2019).

2.6 Brief Description of the Biocide Agents on the Market

Many biocide agents already exist on the market. They are classified into the following compounds (Morais et al., 2016):

Quaternary Ammonium Compounds: These compounds represent a group of compounds. They consist of a subgroup of alkyl linear ammonium compounds, composed of hydrophobic alkyl chain and hydrophilic-counterpart. Quaternary Ammonium Compounds damage cell membranes, modify proteins and inhibit DNA production. They are applied in cotton, polyester, nylon, and wool fibres. These compounds are active against a wide range of pathogens but lack physical bonding in textiles.

Triclosan: These compounds are odourless chlorinated bisphenol and improve the durability of laundering. They are active against a wide range of pathogens. They block lipid biosynthesis and hinder the integrity of the cell membranes. They are applied to polyester, nylon, cellulose acetate, and polypropylene.

Metals and metallic salts: At low concentrations, they are exceptionally active against pathogens. They generate reactive oxygen species, damaging cellular proteins lipids and DNA. Silver, copper, zinc, and cobalt are used has been widely used as antimicrobial agents and applied to cotton, wool, nylon, and polyester.

Chitosan: is a natural hydrophilic copolymer. It's a linear polysaccharide that is biocompatible, non-toxic, non-carcinogenic, and antimicrobial. They are applied to cotton, wool, polyester, and nylon fibres. The low molecular weight results in inhibiting the synthesis of mRNA, preventing protein synthesis, and the high molecular weight cause leakage of intracellular substances or blocks the transport of essential solutes.

Poly (Hexamethylene Biguanide): These agents are polycationic amines biguanide repeat units separated by aliphatic chains. They interact with membrane phospholipids, resulting in disturbance and the fatal leakage of cytoplasmic materials. They are applied to cotton, nylon, and polyester fibres.

N-halamines: They are heterocyclic organic compounds. N-halamines prevent the cell enzymatic and metabolic processes, causing the consequent microorganism destruction. They are applied in cotton, nylon, polyester, and wool fibres and are active against a wide range of pathogens.

Many plant-based compounds with a wide range of antimicrobial activity spectrum have been identified and are commercially available. Table 2.3 shows the wide range of commercially available antimicrobial agents on the market (Morais et al., 2016).

| Product Name | Company | Description | | |
|---------------------------|--|--|--|--|
| Agion® | Sciessent, Massachusetts , United States Silver and zeolite-based | | | |
| AlphaSan® | Milliken Chemical, Spartanburg, South Carolina Silver-based additive | | | |
| BioGaurd® | AEGI Microbe Shield, Huntersville, North Carolina | Finishing agent based on 3- trimethoxysilylpropyldimethyloctadec ylammonium chloride | | |
| Biozac ZS | Zschimmer & Schwarz Mohsdorf GmbH, Burgstadt, Germany | PHMB-based finishing agent | | |
| Cosmocil CQ™ | Lonza, Basel, Germany | Polyaminopropyl biguanide- based additive | | |
| Eosy® | Unitika, Osaka, Japan | Finishing agent based on chitosan | | |
| Irgaurd [®] 1000 | BASF, Ludwigshafen, Germany | Finishing agent based on triclosan | | |
| Irgasan | Sigma Aldrich, Missouri, United States | Finishing agent based on triclosan | | |
| Microban® | Microban International, North Carolina, United States | Triclosan-based agent | | |
| Reputex™ | Lonza, Basel, Germany | PHMB-based finishing agent | | |
| Sanigard KC | L. N. Chemical Industries, Maharashtra, India | Finishing agent belonging to the QAC group | | |
| Saniguard Nano-ZN | L. N. Chemical Industries, Maharashtra, India | Finishing solution based on aqueous nano-dispersion of zinc oxide | | |
| Sanitised® | SANITIZED AG, Burgdorf, Germany | Finishing agent based on 3- trimethoxysilylpropyldimethyloctadec ylammonium chloride | | |
| Silpure® | Thomson Research Associates, Toronto, Canada | Silver particles-based finishing agent | | |
| Silvadur™ | The Dow Chemical Company, Michigan, United States | Interpenetrating polymer network with silver ions | | |
| SmartSilver/;® | Nanohorizon Inc. Philadelphia, United States | Silver nanoparticles-based agent | | |
| Silverion 2400 | Pure Bioscience, Inc.California, United States | Stabilised silver complex-based agent | | |

Table 2.3: Representation of commercially available antimicrobial agents on the market

2.7 Extraction Methods for Studying Phytochemicals

2.7.1 Introduction

There are various extraction methods, e.g., solvent extraction, distillation method, pressing, and sublimation. Solvent extraction is the most widely used method where the natural products undergo a process where the solvent penetrates through the plant cell wall and the solute

dissolves in the solvents the solute followed by collecting the extract. It has been reported that the plant material, properties of the solid-to-solvent ratio, extraction temperature, and extraction time will affect the extraction efficiency (Li et al., 2008; Mordeniz, 2019a). The selectivity of the solvents, solubility, cost and safety play a crucial role in solvent extraction. Solvents with similar polarity of the solute will result in a greater yield. High temperature affects dispersion and solubility. High temperatures may result in solvents lost and the degradation of thermolabile compounds. The extraction yield increases with extraction time. The greater the solvent-to-solid ratio, the greater the extraction yield (Zhang et al., 2018). Solvent extraction is the most widely used extraction method from plant material.

2.7.1.1. Cold Extraction

In this extraction process, the plant parts are dried in a controlled environment at low temperatures and milled into a powder and weighed. The powder is added to a beaker with solvents and kept at room temperature for thirty minutes. The contents are shaken every twenty-four hours for seven days. The extract is filtered using Whatman filter paper under vacuum and dried at room temperature in a watch glass dish. The weight of the powder is recorded before and after drying (Harborne JB, 1998).

2.7.1.2 Plant Tissue Homogenization

Fresh plant parts are grounded in a blender. The solvent is added and shaken vigorously for 5–10 min or left for 24 h followed by filtration of the extract. The filtrate can be dried under reduced pressure and redissolved in the solvent to determine the concentration, or it can be centrifuged for clarification for further studies (Velavan, 2015).

2.7.1.3 Serial Exhaustive Extraction

In this extraction method, the solvent of increasing polarity from a non-polar solvent (hexane) to a polar solvent (methanol) is used to ensure a broad polarity range of compounds being extracted and to prepare crude extracts (Velavan, 2015).

2.7.1.4 Soxhlet Extraction

In this extraction method, solid material is placed in a thimble in the extractor. The solvent is heated until reflux. The vapour rises, and the solvent is condensed and fills up the thimble. The extraction is repeated (Handa et al., 2008; Patel et al., 2019).

2.7.1.5. Maceration

A whole or coarsely powdered plant is soaked in the solvent in a container for a period of continuous mixing until agitation until the biomass matter is dissolved (Velavan, 2015).

2.7.1.6 Decoction

In this extraction method, the plant parts are brought to a boil in water followed by cooling, straining, and passing sufficient cold water through the drug to produce the required volume (Handa et al., 2008).

2.7.1.7 Infusion

In this extraction method, the plant parts are macerated with either cold or boiling water (Handa et al., 2008).

2.7.1.8 Digestion

In this extraction method, the plant parts are macerated under gentle heating (Handa et al., 2008).

2.7.1.9 Percolation

In this extraction method, the raw material is placed in an appropriate amount of solvent for approximately 4 hours in a closed container. Additional solvent is added to the top of the raw material and macerated in a closed container for 24 hours. The percolator is opened, and the extract is poured out drip-wise. Additional solvent is added until the percolate measures about three-quarters of the required volume of the finished product. The marc is pressed, and the pressed liquid is added to the percolate. Additional solvent is added to produce the required volume, and the mixed liquid is clarified by filtration or by decanting (Handa et al., 2008).

2.7.1.10 Sonication

This method uses ultrasound technology to assist in the extraction of bioactive compounds under frequencies ranging from 20 kHz to 2000 kHz. The ultrasound increases the permeability of cell walls and produces cavitation and ruptures the plant cell wall (Handa et al., 2008).

2.7.1.11 Enzymatic Extraction

In this extraction method, enzymes are used to increase the yields during the extraction. Enzymes are used to soften the tissues of biomass and facilitate the degradation of the cells (Lu et al., 2017).

2.7.1.12 Microwave-Assisted Extraction

This extraction method uses microwave radiation and solvents to extract bioactive compounds. Microwave energy is generated through microwave radiation that heats the solvents whilst enhancing the kinetics of the extraction. Moisture occurs in the plant cells when heat is applied and evaporates. The pressure on the cell wall results in cell rupture. Exudation occurs and leads to an increase in extraction yield (Lu et al., 2017).

2.7.1.13 Pulsed Ultrasonic-Assisted Extraction

This is an extraction method using ultrasonic sound waves that pass through the solvent, producing energy by enhancing the diffusion of the solvent into the sample array. Pulsed Ultrasonic-Assisted Extraction is cost-effective in terms of the quantity of solvent used, temperature, and time (Lu et al., 2017). Increasing the intensity of ultrasound in a liquid lead to the breaking down of molecular structures and the formation of bubbles through a process called cavitation (Baig et al., 2010). The collapse of these bubbles produces physical, chemical, and mechanical effects, facilitating the disruption of biological membranes. This disruption aids in the release of extractable compounds, improves solvent penetration into cellular materials, and enhances mass transfer (Cares et al., 2010; Metherel et al., 2009) thus, this extraction method was used for this study.

2.7.1.14 Supercritical Fluid Extraction

In this extraction method, supercritical fluids at temperatures and pressures above the critical values are applied to the extraction and the supercritical fluids return to their gas phase and evaporate without leaving solvent residues (Lu et al., 2017).

2.7.1.15 Pressurised Liquid Extraction

This extraction method is conducted under high pressures and temperatures that aid in the high solubility of the compounds in the solvent and result in high diffusion of the solvent into the sample array (Lu et al., 2017).

Table 2.4 shows the various extraction methods used when extracting (Zhang et al., 2018).

| Method | Solvent | Temperature | Pressure | Time | Volume Consumed | The Polarity of Natural Products |
|---------------------------------------|---|---|-------------|----------|--------------------|--|
| Maceration | Water, Aqueous and non- aqueous solvents | Room temperature | Atmospheric | Long | Large | Dependent on extracting solvent |
| Percolation | Water, Aqueous and non- aqueous solvents | Room temperature, occasional heat | Atmospheric | Long | Large | Dependent on extracting solvent |
| Decoction | Water | Under heat | Atmospheric | Moderate | None | Polar compounds |
| Reflux extraction | Aqueous and non-aqueous solvents | Under heat | Atmospheric | Moderate | Moderate | Dependent on the extracting solvents |
| Soxhlet extraction | Organic solvents | Under heat | Atmospheric | Long | Moderate | Dependent on extracting solvent |
| Pressurised liquid extraction | Water, aqueous and non- aqueous solvents | Under heat | High | Short | Small | Dependent on extracting solvent |
| Supercritical fluid extraction | CO ₂ | Near room temperatures | High | Short | None or small | Non-polar to moderate compounds |
| Ultrasound- assisted extraction | Water, aqueous and non- aqueous solvents | Room temperature or under heat | Atmospheric | Short | Moderate | Dependent on extracting solvent |
| Microwave- assisted extraction | Water, aqueous and non- aqueous solvents | Room temperature | Atmospheric | Short | Moderate | Dependent on extracting solvent |
| Pulsed electric field extraction | Water, aqueous and non- aqueous solvents | Room temperature or under heat | Atmospheric | Short | Moderate | Dependent on extracting solvent |
| Enzyme assisted extraction | Water, aqueous and non- aqueous solvents | Room temperature or heated after enzyme treatment | Atmospheric | Moderate | Moderate | Dependent on extracting solvent |

| Table 2.4: Extraction methods used in plan | nt extractions (Zhang et al., 2018) |
|--|-------------------------------------|
|--|-------------------------------------|
2.7.2. Chromatographic Techniques

2.7.2.1. Introduction

Chromatography is a technique used to separate molecules based on their size, shape, and charge. The analyte in the solvent passes through a molecular sieve which leads to its separation. Paper and thin-layer chromatography readily provides qualitative information.

Paper Chromatography (PC)

In this technique, a sheet of paper is used to carry out separations which act as both support as well a medium for separation. The sample is placed near the bottom of the filter paper and the filter paper is placed in the chromatographic chamber with solvent. The solvent moves forward by capillary action carrying soluble molecules along with it. Low-porosity paper will produce a slow rate of movement of the solvent and thick papers have increased sample capacity (Coskun, 2016).

2.7.2.1.1. Thin Layer Chromatography (TLC)

This technique is used to separate the samples based on the interaction between a thin layer of adsorbent attached to the plate with low molecular weight compounds. Different adsorbents are used to separate various compounds (Coskun, 2016).

2.7.2.1.2. Gas Chromatography (GC)

This technique is used to separate volatile compounds. The rate of kinetics for the chemical species is determined through its distribution in the gas phase. Gas chromatography involves a sample being vaporized and injected into the head of the chromatographic column. The sample is transported through the column by the flow of the inert, gaseous mobile phase. The column itself contains a liquid stationary phase which is adsorbed onto the surface of an inert solid (Coskun, 2016).

2.7.2.1.3. High-Performance Liquid Chromatography (HPLC)

This technique separates compounds based on their interactions with solid particles of a tightly packed column and the solvent of the mobile phase. The Diode Array Detector measures the absorption spectra of the analytes to aid in their identification of the compounds (Coskun, 2016).

2.7.3 Qualitative and Quantitative Phytochemical Screening

2.7.3.1 Introduction

The study of bioactive compounds encompasses phytochemical and pharmacological approaches (Foye et al., 2008). Many plant parts contain bioactive components, e.g., bark, leaves, stems, fruits, and seeds (Cragg & Newman, 2001). Phytochemicals are chemicals produced by the various parts of the plants namely, alkaloids, flavonoids, terpenoids, steroids, tannins, glycosides, etc. The bioactive compounds have various antimicrobial and antibacterial properties (Santhi & Sengottuvel, 2016). Qualitative phytochemical screening plays a crucial role in identifying various biochemical compounds produced by plants. The quantification of those metabolites may assist in the extraction, purification, and identification of the bioactive compounds for human use (Santhi & Sengottuvel, 2016).

Qualitative Phytochemical Analysis

The preliminary qualitative phytochemical screening is carried out as per standard methods described by Trease & Evans 1989:

Detection of Alkaloids

The extracts are dissolved in dilute hydrochloric acid and filtered individually and tested for the presence of alkaloids.

Mayers test: The extraction is added to the Mayers reagent. A yellow-cream precipitate formation indicates the presence of alkaloids.

Wagner's test: Wagner's reagent is added to the extraction of a brown-reddish brown formation is observed, and it indicates the presence of alkaloids.

Detection of Flavonoids

Lead acetate test: A few drops of lead acetate solution are added to the extracts. A yellowcolour precipitate indicates the presence of flavonoids.

Sulfuric acid test: A few drops of sulfuric acid are added to the extracts, and the formation of orange colour indicates the presence of flavonoids.

Detection of Steroids

A few drops of acetic anhydride are added to the extracts and the formation of violet to blue to green in some samples indicates the presence of steroids.

Detection of Terpenoids

Salkowski's Test: Extract 5 mg of the selected plant part is mixed it with 2 ml chloroform and 3 ml concentrated sulfuric acid added carefully to form a layer. A reddish-brown colour indicates the presence of terpenoids.

Detection of Anthraquinones

Bontrager's Test: About 5 mg of the extract is boiled with 10% HCl for a few minutes in a water bath. It's filtered and allowed to cool. An equal volume of $CHCl_3$ is added to the filtrate. A few drops of 10% NH₃ are added to the mixture and heated. The formation of pink colour indicates the presence of anthraquinones.

Detection of Phenols

Ferric chloride test: A few drops of ferric chloride are added to the 10 ml extract. A bluishblack colour indicates the presence of phenol.

Lead acetate test: A few drops of lead acetate solution are mixed with 10 mg extract. A yellow colour indicates the presence of phenol.

Detection of Saponins

0.5 mg of the extract is mixed vigorously with 5 ml of distilled water. The formation of frothing indicates the presence of saponins.

Detection of Tannins

A few millilitres of the extract are mixed with a few millilitres of water and heated in a water bath. The mixture is filtered. Ferric chloride is added to the filtrate. The dark green colour indicates the presence of tannins.

Detection of Carbohydrates

A 0.5 mg of the extract is dissolved individually in five ml of distilled water and filtered. The filtrate is used to test the presence of carbohydrates (Trease & Evans, 1989).

2.7.3.2 Quantitative Phytochemical Analysis

2.7.3.2.1 Estimation of Total Alkaloids

One gram of extract sample is added to a 250 ml beaker, and 200 ml of 10% acetic acid in ethanol is added, covered, and left for settling for 4 hours. The extract is filtered and concentrated in a water bath to one-quarter of the original volume. Concentrated ammonium hydroxide is added dropwise to the extract until the precipitation is complete. The solution is

allowed to settle, and the precipitate is collected and washed with dilute ammonium hydroxide, followed by filtration. The residue is dried and weighed (Harborne, 1984).

2.7.3.2.2 Estimation of Total Flavonoids

A gram of sample is extracted repeatedly with 100 ml of 80% aqueous methanol. The mixture is filtered through Whatman no.1 filter paper into a pre-weighed 250 ml beaker. The filtrate is transferred to a water bath and allowed for evaporation to dryness and followed by weighing off the sample (Santhi & Sengottuvel, 2016).

2.7.3.2.3 Estimation of Total Phenols

The sample is placed in a beaker and boiled for 15 minutes with 50 ml of ether for the extraction of phenolic compounds. Five ml of the extract is pipetted out into a 50 ml flask followed by the addition of 10 ml of distilled water, 2 ml of ammonium hydroxide solution, and 5 ml of concentrated amyl alcohol. The samples are left to react for 30 minutes for colour development and measured by spectrophotometry, reading the absorbance at 505 nm (Santhi & Sengottuvel, 2016).

2.7.4 Textiles Analysis

2.7.4.1 Biocidal Analysis

The biocidal analysis evaluates the effectiveness of antimicrobial textiles. Several test methods have been established through quantitative antimicrobial tests. The number of microbes present on the finished fabrics can be counted and expressed as a percentage or as a log reduction. The test methods for quantitative determination are ATCC TM100, JIS L1902, AATCC90 percentage reduction, and ISO 20743 shake flask reduction methods (Behary et al., 2020).

The Parallel Streak Method (AATCC TM147) is a qualitative method used to determine the antibacterial activity of diffusible antimicrobial agents on treated textile materials. The Parallel Streak Method has proven to be effective. This method shows antibacterial activity against both Gram-positive and Gram-negative bacteria. The sterilized agar is dispensed (cooled to 47 °C (117 °F) by pouring 15 ml into each standard (15 × 100 mm) flat-bottomed petri dish. Allow agar to gel firmly before inoculating. The inoculum is prepared by transferring 1.0 ml of a 24-hour broth culture into 9.0 ml of sterile distilled water and containing it in a test tube or small flask. A 4 mm inoculating loop is used, loaded with one loopful of the diluted inoculum and transferred to the surface of the sterile agar plate by making five streaks approximately 60 mm in length, spaced 10 mm apart by covering the central area of a standard petri-dish without refilling the loop. The specimen is pressed onto the agar surface with a sterile spatula. After 18 to 24 hours of incubation at 37 °C, the plates are examined for bacterial growth directly

underneath the textiles and around the edges of the textiles. If the antimicrobial substance diffuses into the agar, an inhibition area is formed, and its size indicates the effectiveness of the antimicrobial effect or the rate at which the active agent is released (Neely & Maley, 2000; Gao et al., 2008). AATCC 100 (Suspension Test) is a quantitative antimicrobial test method used to determine the antibacterial activity of textiles and fabrics against bacteria. The bacterial counts are recorded, and a percent reduction is measured using the initial count and remaining count data (Gao et al., 2008).

2.7.3.2 Durability Analysis

Durability by washing method (ASTM E3162-18 or AATCC61-2A) is used to determine the durability of laundering. This test method is an accelerated laundering test method to measure the durability of antibacterial agents applied to textiles under simulated home laundering conditions. Ten grams of the coated fabric for laundering is prepared, followed by adding a 500 ml defined detergent solution. Set the washing machine at a temperature of 50 °C under abrasive action using stainless steel balls to simulate five home launderings for a 45-minute laundering cycle at 40 revolutions per minute. After each cycle, remove the fabric and rinse it with water thoroughly by hand. Repeat, depending on the total number of washes required.

Plants offer a unique source of bioactive compounds with medicinal properties and biological activities. The selection of solvents during extraction plays a crucial role in obtaining these bioactive chemicals. While synthetic antimicrobial agents have proven effective against pathogens, they also pose risks to the environment and human health. Therefore, there is a growing interest in exploring plant-based antimicrobial agents and textile finishes. By researching and developing these natural alternatives, we can enhance their longevity and durability on textile substrates, providing sustainable solutions for combating harmful microbes.

The emergence of antibiotic-resistant "super germs" presents a significant global health concern. To address this challenge, there is a need for extensive research on medicinal plants as potential sources of alternative medicines. By harnessing the bioactive properties of unexplored medicinal plants and using generally regarded as safe (GRAS) solvents during extraction, we can develop safer and eco-friendly therapeutic options. Moreover, it is essential to conduct in-depth studies to identify the most cost-effective pre-treatment, drying, and extraction methods for future therapeutics. By optimizing these processes, we can make bioactive compounds more accessible and contribute to the development of effective treatments.

In conclusion, delving deeper into the potential of medicinal plants, exploring eco-friendly extraction methods, and improving antimicrobial agents and textile finishes will have a positive impact on global health and the environment.

Chapter 3

Assessing the Efficiency of Antimicrobial Plant Extracts from *Artemisia afra* and *Eucalyptus globulus* as Coatings for Textiles.

Manuscript to be submitted for publication related to this publication.

Nortjie, E.; Basitere, M.; Moyo, D.; Nyamukamba, P. Assessing the Efficiency of Antimicrobial Plant Extracts from *Artemisia afra* and *Eucalyptus globulus* as Coatings for Textiles

CHAPTER 3: RESULTS AND DISCUSSION

3.1 Introduction

Textiles serve as a barrier between the human skin and pathogens. The COVID-19 pandemic crisis made us aware of the importance of personal hygiene and personal protective equipment (PPE). The use of PPE, disinfectants and social distancing was important for our safety as championed by the World Health Organization (Chauhan & Kumar, 2020). Textile coatings improve the functional properties of textiles, creating a value-added product that offers comfort, strength, durability, and better handling. Other functional finishes on textiles may include flame retardancy, water repellency, anti-dirt, antibacterial, and antiviral properties (Ibrahim et al., 2017).

Many synthetic fibers are more resistant to microbial attack than natural fibers. Natural fiberbased fabrics, such as cotton, contain keratinous fibers and carbohydrates that serve as nutrients under optimal conditions, making them prone to microbial attack (Hipler & Elsner, 2006). According to a report by Fortune Business Insights (2020), the global antimicrobial textile market size was valued at USD 9.04 billion in 2020 and is expected to reach USD 13.63 billion by 2028, with a compound annual growth rate of 5.3% from 2021 to 2028. The report attributes the growth of this market to the increasing demand for antimicrobial textiles in medical and apparel applications (Fortune Business Insights, 2021).

There is a wide range of natural to synthetic antimicrobial products on the market. Synthetic antimicrobial agents pose a great concern due to side effects and water pollution hence the great demand for antimicrobial agents that fulfil the requirements of the regulating authorities (Ramachandra & Rao, 2008). While herbal extracts have demonstrated effective antibacterial properties, the longevity of the coating when subjected to laundering is inadequate (Bouchekrit et al., 2016).

Natural compounds have been extensively explored for new drug discoveries (Chandra et al., 2017). Extracts of different parts of plants contain bioactive compounds, such as alkaloids, steroids, tannins, glycosides, volatile oils, fixed oils, resins, phenols, terpenoids, and flavonoids that fight against diseases (Kutama, 2018). The phenolic phytochemicals from plants play a key role as antimicrobial agents (Restuati & Diningrat, 2018; Smeriglio et al., 2017).

Artemisia afra and *Eucalyptus globulus* plant species have been selected for this study as they may possess antimicrobial properties. Studies have shown that the mode of action of the antimicrobial agents applied to textiles can result in various effects, including the agglomeration of cell proteins, harm to the cell membrane, interference with its function, inhibition of protein synthesis, and suppression of nucleic acid synthesis (Kamel & Hassabo, 2021).

Antibiotic resistance to pathogenic bacteria has become a global problem, leading to naturalbased antimicrobial compounds being studied extensively as an alternative therapeutic strategy to fight against microbial growth (Savoia, 2012; Upadhyay et al., 2014).

Antibiotics play a vital role in fighting bacterial infections. They also cause havoc in the healthcare and pharmaceutical sector, accelerating socio-economic losses (O'Neill, 2016). Multidrug resistance is expected to increase by ten million deaths annually by 2050 (O'Neill, 2016; Kraker et al., 2016). Biological screening, separation of the phytochemicals, and clinical trials of medicinal plants have advanced over the years, unfolding the secrets of ancient herbal remedies (Bibi et al., 2017). Traditional medicine is effective in dealing with diseases caused by bacteria or oxidative stress (Baydoun et al., 2015; Singh et al., 2017; Restuati & Diningrat, 2018). Despite the announcement by the World Health Organization in the seventies of *Artemisia afra* being unfit for human consumption. Till today, *Artemisia afra* is used for ethnomedicinal purposes in South Africa (Oyedeji et al., 2009). Eucalyptus species are the most represented species in the international pharmacopoeia in which *E. globulus* are the main supply of essential oils due to their medicinal properties (Bachir & Benali, 2012).

Therefore, the purpose of this study was to investigate whether plant extracts derived from South African biomass, namely *Eucalyptus globulus* and *Artemisia afra*, could be effective as antimicrobial agents or textile coatings. The work involved the preparation of extracts containing antimicrobial compounds by pulsed ultrasound-assisted extraction using polar and non-polar solvents separately, qualitative phytochemical screening analysis on *Eucalyptus globulus* and *Artemisia afra* to identify the bioactive compounds from the selected plants, and the application the extracts onto cotton and polyester fabrics and assess the antimicrobial activity before and after laundering and the characterisation of the coated samples using Fourier-transform infrared spectroscopy (FTIR).

3.2 Materials and Methods

3.2.1 Sources of plant material and fabric

The leaves of the medicinal plant *Artemisia afra*, except for *Eucalyptus globulus*, were acquired from the South African National Biodiversity Institute located at Kirstenbosch Gardens. The biomass of *Eucalyptus globulus* was collected from Deer Park in Vredehoek, Cape Town. Both *Artemisia afra* and *Eucalyptus globulus* plant specimens were identified by Mr. Pieter Winter at the South African National Botanical Institute (SANBI), with the assistance of Mr. Tielmann Haumann. The commercially available woven cotton fabric was obtained from SA Interlining Company, while the knitted polyester fabric was procured from Priontex Company with the assistance of Mr Rainer Absenger.

3.2.2 Plant sample preparation

The leaves of *Artemisia afra* and *Eucalyptus globulus* plants were harvested and thoroughly washed with distilled water to remove any dust particles. The leaves were then air-dried in the shade at room temperature for 30 days. Phytochemicals were extracted from the dried leaves using methanol and n-hexane solvents. The resulting extract was evaporated to determine the yield of the extractions.

3.2.3 Plant extraction

The solvent extracts were obtained using the pulsed ultrasound-assisted extraction (PUAE) method by applying a 1:10 (w/v) solid-solvent ratio (Green, 2004). Methanol and hexane were selected for their ability to extract polar and non-polar compounds, respectively, and their similarity with previous and future studies.

About 10 g of powdered leaves were extracted by sonicating in 100 ml of methanol and hexane. The PUAE was conducted at a solid-to-liquid ratio of 1:10 for 20 minutes at 40°C, at 50% amplitudes with a pulse of 0.5 s on and 5 s off at a 20 kHz frequency with a probe diameter of 25 mm placed at a depth of 10 mm into the medium. Post-extraction, the mixtures were filtered through a Whatman LSC601 70 mm filter paper disc. The solvents were removed using MiVac Sample Evaporation at 40°C.

The extraction yield of the plant extracts obtained through PUAE was determined, as depicted in Table 3.1. Subsequently, these extracts were reconstituted using their respective solvents and then stored at a temperature of -20° C.

| Average Yield % | = Weight of the extract af Dry weight of p | ter solvent rem lant sample | 100% * 100% |
|-----------------|---|--------------------------------|-------------|
| Plants (Leaves) | | A. afra | E. globulus |
| Solvents | Methanol | 9.22 | 22.76 |
| Solvents | Hexane | 2.16 | 3.98 |

| Table 3.1: The average yield percentage of the plant extracts | s obtained by the PUAE method |
|---|-------------------------------|
|---|-------------------------------|

Table 3.2 shows the extraction conditions used in this study.

| No. | Parameter | PUAE extraction |
|-----|---------------------|-----------------------|
| 1 | Sample size (g) | 10 g |
| 2 | Extraction solvent | n-hexane and methanol |
| 3 | Solvent volume (ml) | 100 ml |
| 4 | Temperature (°C) | 40 °C |
| 5 | Time (min) | 20 minutes |

Table 3.2: Extraction conditions

Table 3.3 shows the physiochemical properties of the solvents used.

| No. | Solvent | Polarity index (pi) | Boiling point ℃ | Viscosity @ 20°C (cP) | Density @ 20°C (g/ml) |
|-----|----------|------------------------|-----------------------|-----------------------------|--------------------------|
| 1 | n-Hexane | 0.1 | 67-69.5 | 0.31 | 0.672 |
| 2 | Methanol | 5.1 | 64.7 | 0.55 | 0.791 |

Table 3.3: Physiochemical properties of solvents used (n-Hexane and Methanol) [50]

3.2.4 Preparation of antimicrobial cotton and polyester material

The fabric was cut into squares measuring 3 cm x 3 cm. Afterwards, it was washed with a mixture of ethanol and water using ultrasonication for 30 minutes to eliminate any wax and impurities on its surface. Next, the fabric was dried for an hour at 80 °C in a warm air oven. Plastic containers were then used to immerse the fabric cuttings in the chosen extracts. The cuttings were left to soak for 24 hours at 25 °C in an FHM incubator shaker and then dried in a warm air oven at 80 °C for one hour to eliminate the solvent. Finally, the fabric cuttings were utilized to determine the zone of inhibition against *S. aureus* ATCC 33591 and *E. coli* ATCC 25922.

3.2.5 Experimental method and procedure

The process flow diagram of the study is shown in Figure 3.1 below.



Figure 3.1: Process flow diagram of the experimental setup of the antimicrobial textiles

Experiments were carried out to determine various factors related to the effectiveness of coated fabrics in preventing microbial growth. Specifically, the Minimum Inhibitory Concentration (MIC), the Zone of Inhibition (ZOI), and the durability of the coated fabric after undergoing a laundering process according to ISO 6330:2012E were analyzed. The antimicrobial effectiveness of the coated fabric was then evaluated by measuring the ZOI against the selected strains *S. aureus* ATCC 33591 and *E. coli* ATCC 25922, as well as appropriate controls.

3.2.6 Characterizations

The functional groups present in the woven cotton and knitted polyester fabric were analyzed before and after the application of an antimicrobial finish. Fourier-transform infrared spectroscopy (FTIR) was used to perform the analysis, using a Cary 660 spectrometer from Agilent Technologies (USA). The wavenumber range of 400–4000 cm⁻¹

3.2.7 Determination of the inhibition zone

The nutrient and tryptic soy broth growth media for the selected test strains, *Staphylococcus* aureus ATCC 33591 and Escherichia coli ATCC 25922 respectively, were autoclaved for 25 minutes at 120 °C and under pressure. Once cooled, the liquid media were inoculated with agar plate cultures. The liquid cultures were incubated at 37 °C at 160 rpm for 24 hours. The next day, the cultures were analyzed by Gram staining to ensure no contamination. The optical density (OD) of the cultures was determined at a wavelength of 600 nm using a Perkin-Elmer Lambda-25 spectrophotometer and diluted to an OD600 = 0.2. Extracts were serially diluted in the range of 5 µg - 500µg, and 50µl of each sample was pipetted into a sterile 96-well plate in triplicate, and the plate was sealed with a Breath-Easy seal to allow for respiration. The plates were incubated at 37°C for 24 hours. Following that, 20 µl of 0.25 % (w/v) MTT solution (in phosphate-buffered saline) was added to each experimental well and incubated for 4 hours at 37 °C. The reaction was stopped by adding 100 µl of 100 % DMSO, and the OD was determined using a microtiter plate reader. To minimize errors, each sample was evaluated in its 96-well plate: for twelve samples, twenty-four plates of 96-well plates (two test strains). The minimum inhibitory concentration (MIC) is determined as the lowest concentration at which total inhibition of the test strain occurs. Controls included wells with media and test strains without any additions.

3.2.8 Determination of the zone of inhibition

Step 1: Preparation of liquid cultures

Specific liquid growth media (5 ml per 50 ml glass flask; five flasks per strain) were prepared for each test strain:

| Staphylococcus aureus ATCC 33591 | Nutrient broth |
|----------------------------------|-------------------|
| Escherichia coli ATCC 25922 | Tryptic soy broth |

The cultures were analysed by Gram staining to ensure that the cultures were not contaminated. The optical density of the cultures was determined at 600 nm using a Perkin-Elmer Lambda-25 spectrophotometer and diluted to an OD600 = 0.2. These diluted cultures were used in the agar diffusion test (Marković et al., 2018).

Step 2: Preparation of agar plates

A total of twelve samples, twelve agar plates per test strain were prepared. This required the preparation of the following (also including plates for controls):

25 tryptic soy agar plates = 500 ml of media 25 nutrient agar plates = 500 ml of media

Each sample was assessed in triplicate on each plate and was prepared as follows:

For each agar plate, 100 μ l of the OD600 = 0.2 cultures was spread-plated onto the agar and allowed to diffuse into the agar. Each textile specimen was pre-treated with UV (30 minutes on each side) to kill off any contaminants and then placed onto the agar plate containing the test strain (to ensure that the UV treatment was effective, the material was also placed on agar containing no test strain). For each test strain, a commercial antibiotic was used as a positive control. Gentamicin and ampicillin were used for the bacterial test strains. Samples were plated on agar plates and incubated at 37 °C overnight. The zones of inhibition were measured to determine whether growth inhibition occurred as reported by (Marković et al., 2018).

3.2.9 Durability Tests

The durability of the extract coating applied to textiles under simulated home laundering conditions was assessed using the washing method ISO 6330:2012. The coated fabric samples were cut into 3 x 3 cm squares and weighed to determine the amount of washing powder and water required for laundering. The fabric samples were washed at 60 $^{\circ}$ C using the Pyrotec MB2 washing machine, rinsed in cold water, and dried in a hot air oven at 80 $^{\circ}$ C for 1

hour. Each textile sample was pre-treated with UV for 30 minutes on each side to kill off contaminants and then placed onto the agar plate containing the test strain. The material was also placed on agar containing no test strain to ensure that the UV treatment was effective. The agar plates were incubated at 37 °C overnight, and the zones of inhibition were measured.

3.2.10 Phytochemical screening of Artemisia afra and Eucalyptus globulus

The extracts were subjected to phytochemical screening of bioactive compounds. The methanolic and hexanoic extracts of the leaves were screened for the presence of phenols, flavonoids, quinones, tannins, saponins, terpenoids, and steroids. The quantitative results are expressed as (+) for positive and (-) for negative results.

3.2.10.1 Detection of Phenols

The extract of 1.0 ml was diluted in distilled water to 3.0 ml followed by filtration. Four drops of 5% iron (III) chloride solution were added to the solution. The dark green colour indicated the presence of phenol (Yimam & Desalew, 2022).

3.2.10.2 Detection of Flavonoids

The extract of 1.0 ml was transferred into a test tube and treated with four drops of sodium hydroxide solution. The yellow colour indicated the presence of flavonoids (Yimam & Desalew, 2022).

3.2.10.3 Detection of quinones

The extract of 1.0 ml was placed in a test tube and mixed with 1.0 ml of pure 98% sulphuric acid. The red colour showed the presence of quinones (Yimam & Desalew, 2022).

3.2.10.4 Detection of Tannins

About 1 ml aliquot of the extract was transferred into a 5 ml test tube followed by the addition of three drops of 5% Iron (III)chloride, and a greenish-black precipitate showed the presence of tannins (Yimam & Desalew, 2022).

3.2.10.5 Test for saponins

About 0.5 mg aliquot of the extract was mixed vigorously with 5 ml of distilled water. The formation of stable foam indicated the presence of saponins (Yimam & Desalew, 2022).

3.2.10.6 Test for steroids

About 1.0 ml of extract was shaken with 99% chloroform, and to the chloroform layer, 98% sulphuric acid was added slowly by the sides of the test tube. The red colour showed the presence of steroids (Yimam & Desalew, 2022).

3.2.10.7 Test for terpenoids

About 0.5 ml aliquot of the extract was mixed with 2.0 ml of 99% chloroform followed by 3 ml concentrated of 98% sulphuric acid added to form a layer. The formation of the reddish-brown colour showed the presence of terpenoids (Odutayo et al., 2017).

3.3 Results and Discussion

3.3.1 Extraction Yield

The average yields of methanolic and hexanoic extracts from *Artemisia afra* and *Eucalyptus globulus* are summarised in Table 3.1. All experiments were conducted in triplicates. The extraction yields of *A. afra* and *E. globulus* were investigated by studying the effects of various solvents, including methanol and hexane. The findings demonstrated that the extraction yields differed depending on the solvent used, attributed to variations in solvent polarity. The yield of extraction is a measure of solvent effectiveness in extracting phytochemicals from the material under selected conditions. The results in Table 3.1 show that the maximum yield was obtained with *E. globulus* methanol extraction at 22.76%, followed by hexane extraction at 3.98%. *A. afra* methanol extraction yielded 9.22%, followed by hexane extraction of 2.16%. Higher yields were observed in the methanolic extracts of *A. afra* and *E. globulus* compared to the hexanoic extracts differed significantly. The improved extraction yields could be attributed to cavitation effects caused by high-intensity ultrasound and solvent polarity.

3.3.2 Phytochemical Screening Analysis

Plant parts contain bioactive compounds with various phytochemical compounds, including alkaloids, flavonoids, steroids, terpenoids, tannins, saponins, phenols, and quinones, all of which possess antimicrobial properties. In this study, the crude extracts were screened for the presence of phenols, flavonoids, quinones, tannins, saponins, terpenoids, and steroids. The phytochemical screening of *Artemisia afra* and *Eucalyptus globulus* crude leaf extracts is shown in Table 3.4, and it revealed the presence of different secondary metabolites, such as phenols, quinones, and steroids. Tannins, saponins, and terpenoids were not found in the hexanoic extracts of *Artemisia afra*. Flavonoids were absent in the methanolic extraction of *Eucalyptus globulus*. Tannins and saponins were also not found in the hexanoic extraction of *Eucalyptus globulus*.

| Plant | Artemisia afra | | Eucalyptus globulus | |
|----------------|----------------|----------|---------------------|----------|
| Solvent | Mothanal | n Havana | Mathanal | n Havana |
| Phytochemicals | Methanol | п-пехапе | Wethanor | п-пехапе |
| Phenol | + | + | + | + |
| Flavonoids | + | + | - | + |
| Quinones | + | + | + | + |
| Tannins | + | - | + | - |
| Saponins | + | - | + | - |
| Terpenoids | + | - | + | + |
| Steroids | + | + | + | + |

Table 3.4: Phytochemical compounds of A. afra and E. globulus extract

-: absence, +: presence

Phytochemical screening revealed the presence and absence of different secondary metabolic compounds such as phenols, quinones, terpenoids, flavonoids, saponins, steroids and tannins. This could be attributed to the different plants, plant parts, types of solvents used for extraction, and geographical areas exhibiting climatic conditions that determine their ability to support rain-fed cultivation (Kane et al., 2019).

Phytochemical screening of crude leaf extracts of *A. afra* and *E. globulus* revealed that both plants had phenols, quinones, and steroids. The phytochemical screening indicates that the methanol extract produced positive results for the analysed phytochemicals. Many solvent extractions have been performed to obtain phytochemical compounds for use against pathogens. From the phytochemical assessment, phenols were found in all the extractions. The mechanism of action of polyphenols is to bind to adhesins, inhibit enzyme-substrate deprivation, complex with the cell wall, and disrupt the membrane and metal ion complexation (Cowan, 1999; Tiwari et al., 2011). Quinones were also found in all the extractions. The mechanisms of action of quinones include inactivating enzymes, complexing with the cell wall, and binding to adhesins (Tiwari et al., 2011; Cowan, 1999). Tannins were present in the methanolic extracts, as shown in Table 3.4. Tannins allow the binding of adhesins, inhibit enzyme-substrate deprivation, complex with the cell wall, and form metal ion complexes (Ameer et al., 2017). Flavonoids were present in most of the extracts except for the *Eucalyptus*

globulus methanolic extracts. Flavonoids form a complex with the cell wall and bind to adhesins (Cowan, 1999; Tiwari et al., 2011). The extracts were also screened for terpenoids which disrupt the membrane wall (Tiwari et al., 2011; Cowan, 1999) The extracts were screened for saponins, and they were only present in the methanolic extracts. The mechanism of action of saponins inhibits the growth of Gram-positive bacteria and Gram-negative bacteria by disrupting cell membranes (Tiwari et al., 2011; Cowan, 1999). The extracts were screened for steroids, and they were found in all the extracts. Steroids derived from medicinal plants are known to possess antibacterial and insecticidal properties (Ndezo Bisso et al., 2022). Table 3.5 displays the mechanisms of action of the phytochemical compounds found in the studied plant extracts.

| l able 3.5 Phytochemical compounds and their | mechanisms of action | |
|--|--|---|
| Phytochemical compounds | Chemical profile | Mechanism of action |
| Flavonoids | Ho de la construcción de la cons | The mechanism of action of flavonoids is complex with the cell wall, binds to proteins (adhesins), inhibits the secretion of autocoids and prostaglandins and inhibits contractions caused by spasms |
| | × × | (Cowan, 1999; Tiwari et al., 2011). |
| Tannins | HO | The mechanism of action of tannins allows the binding of proteins (adhesins), inhibits enzyme-substrate |
| | HO + | deprivation, complexes with the cell wall, makes intestinal mucosa more resistant and reduces secretion, increases the supply of digestible proteins by animals by forming protein |

| Guinones | al., 2011). The mechanisms of action of quinones inactivate enzymes, maybe adhere to the cell wall, and bind to proteins (Cowan 1999 Tiwari et al. 2011) |
|---|--|
| Saponins $\begin{pmatrix} \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ $ | The mechanism of action of saponins inhibits histamine release in-vitro (Cowan, 1999; Tiwari et al., 2011). |
| | The mechanism of action of polyphenols binds to proteins (adhesins), inhibit enzyme-substrate deprivation, complex with the cell wall, make intestinal mucosa more resistant and reduce secretion, increase the |

| Bisso et al., 2022). Disrupt the membrane wall (Tiwari et al., 2011; Cowan, 1999). | H H H H H H H H H H H H H H H H H H H | benoids |
|--|---|---------|
| Disrupt the membrane wall (Tiwari et al., 2011; Cowan, 1999). | | oids |
| Bisso et al., 2022). | ROXR | |
| and insecticidal properties (Ndezo | | |
| are known to possess antibacterial | | |
| Steroids derived from medicinal plants | | |
| (Cowan, 1999; Tiwari et al., 2011). | | |
| in gastrointestinal-tract metabolism | | |
| in the rumen, and causes a decrease | | |
| animals by forming protein complexes | | |
| supply of digestible proteins by | | |

3.3.3 Antibacterial Activity of plant extracts

The antibacterial activity of plant extracts *A. afra* and *E. globulus* was investigated against Gram-positive bacteria, *Staphylococcus aureus* ATCC 33591, and Gram-negative bacteria, *Escherichia coli* ATCC 25922, and the results are shown in Figure 3.2.



Figure 3.2: Antibacterial Activity of plant extracts *A. afra and E. globulus* against *S. aureus* and *E. coli*. A.a: *Artemisia afra*; E.g: *Eucalyptus globulus*; H: Hexane; M: Methanol.

The hexanoic *E. globulus* extracts showed good results in suppressing microbial growth of *S. aureus* ATCC 33591 at an average percentage of 18.70% growth due to a higher number of phytochemicals like polyphenolic compounds, present which are known to have antimicrobial activity, followed by the methanolic *E. globulus* extraction at an average of 29.03% growth. The *A. afra* methanolic extractions showed antibacterial activity against *S. aureus* at an average of 54.20%, and the hexanoic *A. afra* extracts at an average percentage growth of 80.90% exhibited the least activity against *S. aureus* ATCC 33591 strains at 250 µg/ml. The *S. aureus* microbial growth was observed to be more susceptible to the *E. globulus* hexanoic extracts, followed by the *E. globulus* methanolic extracts compared to the *A. afra* methanolic and hexanoic extracts. As expected, the control showed no antimicrobial activity (Figure 3). Ampicillin shows antimicrobial activity against *S. aureus* ATCC 33591 at an average growth of 15.27% as expected.

The evaluation of the *A. afra* and *E. globulus* extracts against *E. coli* ATCC 25922 was the least susceptible compared to *S. aureus*. The methanolic *A. afra* shows slight activity against *E. coli* ATCC 25922 at an average percentage growth of 91%, followed by the hexanoic *A. afra* extracts at 99.52% (no effect). The methanolic and hexane *E. globulus* extracts show no effect

against Gram-negative *E. coli* ATCC 25922. Instead, they showed enhanced growth of an average of 159.25% and 155.70% at 250 μ g/ml, respectively. This could be due to compounds from plants that enhance bacterial growth. As expected, the control sample did not show any antimicrobial activity. Ampicillin shows antimicrobial activity against *E. coli* ATCC 25922 at an average growth of 29.34%. The results show that *E. coli* ATCC 25922 was the most resistant strain against the plant extracts. *S. aureus* ATCC 33591 was the most susceptible strain to the plant extracts.

3.3.4 Minimum Inhibition Concentration Analysis

The MIC, which stands for Minimum Inhibitory Concentration, is defined as the lowest concentration of the extract that can prevent the growth of microorganisms. To test the extract's effectiveness against different strains of bacteria, specific liquid growth media were prepared for each test strain in 50 ml glass flasks at a volume of 5 ml per flask, with three flasks per strain. The results showed that only some of the samples displayed activity in the liquid media against the tested ATCC bacterial strains, as detailed in Table 3.6.

Table 3.6: Summary of the antibacterial activity detected against two ATCC test strains when a concentration range of 5-500 μ g/ml was evaluated in a 96-well bioactivity assay. *S. aureus*: *Staphylococcus aureus*; *E. coli*: *Escherichia coli*; A.a: *Artemisia afra*; E.g: *Eucalyptus globulus*; H: Hexane; M: Methanol

| | MIC Values (µg/ml) | | |
|---------------|--|-----------------------------|--|
| | Microorganisms | | |
| Sample No. | Gram-negative bacteria | Gram-positive bacteria | |
| Plant extract | <i>E. coli</i> ATCC 25922 (Tryptic soy broth) | ATCC 33591 (Nutrient broth) | |
| 1 A. aH | 50-500 μg/ml | 5-25 μg/ml; 250-500 μg/ml | |
| 2 A. aH | 10-500 μg/ml | 125-500 μg/ml | |
| 3 A. aH | None | 50-500 μg/ml | |
| 4 A. aM | 5-500 μg/ml | 5-500 μg/ml | |
| 5 A. aM | 5-500 μg/ml | 5 μg/ml; 125-500 μg/ml | |
| 6 A. aM | 10-500 μg/ml | 100-500 μg/ml | |
| 7 E. gH | 5-50 μg/ml* | None | |
| 8 E. gH | 250-500 μg/ml | None | |
| 9 E. gH | 100-500 μg/ml | None | |
| 10 E. gM | None | None | |
| 11 E. gM | None | 250-500 μg/ml | |
| 12 E. gM | None | None | |

*If activity is only observed at lower concentrations, it usually means that a compound inhibitor is being diluted out, resulting in activity only being observed at the lower (more dilute) sample range. Samples 1-12 were performed in triplicates. The study evaluated the MIC values of methanolic and n-hexane extracts of *A. afra* and *E. globulus* leaf extracts against *S. aureus* and *E. coli*. Although some bioactivity was observed, complete inhibition of the cultures did not occur, and all MIC values were above 500 μ g/ml. The antimicrobial tests yielded MIC values ranging from 5 μ g/ml to above 500 μ g/ml, as shown in Table 4. The methanol extract of *A. afra* exhibited sensitivity against the *S. aureus* ATCC33591 strain, with a MIC value ranging from 5 μ g/ml to 25 μ g/ml. However, some extracts, such as *E. globulus* hexanoic extract against *S. aureus* and *E. globulus* methanolic extract against *E. coli* ATCC 25922, did not show any MIC value. Bacterial culture growth exceeded 100% during sample analysis, possibly due to plant compounds that promote bacterial growth. Samples that exhibited antimicrobial activity in liquid-based assays did not show bioactivity in agar-based assays.

3.3.5 Zone of Inhibition Analysis

The zone of inhibition analysis (ZOI) was conducted against both bacterial types *S. aureus* and *E. coli*. The antimicrobial fabric was produced by using the *A. afra* and *E. globulus* methanolic and hexanoic extracts by immersion process for 24 hours. The green polyester and light blue cotton fabric per sample were used for the zone of inhibition analysis. The fabrics were cut into 2 cm diameter circles and subjected to the protocol described above. Only some samples exhibited bioactivity as shown in Table 3.7.

| Sample No. Plant extract | Material type | Disk diameter | Zone diameter | Zone of inhibition |
|-----------------------------|------------------------|---------------|---------------|------------------------|
| 10 E. gMC | Light blue/grey cotton | 20 mm | 23 mm | 101.32 mm ² |
| 10 E. gMP | Turquoise polyester | 20 mm | 22 mm | 65.97 mm ² |
| 11 E. gMC | Light blue/grey cotton | 20 mm | 27 mm | 258.40 mm ² |
| 11 E. gMP | Turquoise polyester | 19 mm | 22 mm | 96.60 mm ² |
| 12 E. gMC | Light blue/grey cotton | 20 mm | 25 mm | 176.71 mm ² |

Table 3.7: Summary of the bioactivity detected for material samples infused with plant extracts evaluated against *E. coli* ATCC 25922 and *S. aureus* subsp. *aureus* ATCC 33591.

* Sample 10 E.gMC: Light blue grey *E. globulus* methanolic coated cotton fabric; 10 E.gMP: Turquoise *E. globulus* methanolic coated polyester fabric; 11E.gMC: Light blue grey methanolic cotton coated fabric; 11E.gMP: Turquoise *E. globulus* methanolic coated polyester fabric; 12 E.gMC: Light grey/ blue methanolic coated cotton fabric. Samples 1-12 were performed in triplicates on the light blue/ grey cotton and turquoise polyester fabrics.

Moist environments promote bacterial growth, and textiles often serve as a breeding ground for bacteria. To assess the antibacterial properties of fabrics, we examined coated and uncoated polyester and cotton samples against Gram-positive *S. aureus* and Gram-negative *E. coli* bacteria. As indicated in Table 6, only samples 10 to 12 exhibited zones of inhibition. Cotton fabrics, which contain cellulose polymer, facilitate bacterial adherence, whereas polyester fabrics, with their polymer backbone, are less susceptible to bacterial attachment. However, the application of *Eucalyptus globulus* methanolic extracts onto fabrics altered the fabric's surface properties. Our study revealed that samples 10 to 12, which were coated with the extract, displayed antibacterial activity, as illustrated in Figure 3.3 by the presence of zones of inhibition around the material.



Figure 3.3 Bioactivity of material samples on agar plates containing the ATCC test strains. A: Light blue/grey cotton material control sample on *S. aureus*; B: Light blue/grey cotton material control sample on *E. coli*; C: Turquoise polyester material control sample on *S. aureus*; D: Turquoise polyester material control sample on *E. coli*; E: Sample 10 light blue/grey cotton material on *S. aureus*; F: Sample 10 turquoise polyester material on *S. aureus*; G: Sample 11 light blue/grey cotton material on *S. aureus*; H: Sample 11 turquoise polyester material on *S. aureus*; I: Sample 12 light blue/grey cotton material on *S. aureus*.

Figure 3.3 shows that the coated fabric demonstrated a larger zone of inhibition compared to the untreated samples. Among the coated fabrics, the highest zone of inhibition was observed

in cotton coated with *E. globulus* methanolic extract (258.4 mm²) against *S. aureus*, while the smallest zone of inhibition was observed in polyester coated with *E. globulus* methanolic extract (65.97 mm²) as depicted in Figure F, sample 10 (turquoise material) on *S. aureus*. The bioactivity was only evident against the *S. aureus* strain. Although some samples displayed activity in liquid media assays, they failed to exhibit bioactivity in the agar-based assay. The inactivation of active compounds during the treatment or preparation of the material samples or the enhanced absorption of other active compounds, when applied to the material, could be the potential reasons for this. The bioactivity seen in such assays also relies on the diffusibility of compounds from the material into the agar medium, leading to the inhibition of growth. It is essential to note that bacterial growth is rampant in humid conditions (Marković et al., 2018). Natural fibers like cotton are more susceptible to microbial attack because they easily retain water. In contrast, the microbial attack is slower in synthetic fibers such as polyester fabric due to their polymer backbone (Alaribe et al., 2018).

3.3.6 Durability to Washing

Frequent washing of coated fabrics may affect the durability of the coating on the fabric as it plays a key role in coated textiles. This study also focuses on the effect of laundering on the coated textiles against the selected bacterial strains. The ZOI of the microorganisms were measured to determine whether growth inhibition occurred.

After laundering, no activity was observed against *S. aureus* and *E. coli*. This could be due to the inactivation of active compounds during sample treatment or preparation, or the improved bioavailability of other active compounds when applied to the material. On agar plates containing ATCC test strains, no bioactivity was observed on the material samples after laundering. The antimicrobial activity of the samples that had previously shown activity against the selected bacterial strains decreased significantly after washing. The inactivity of the coated fabrics may be due to factors such as reduced extract concentration, variation in phytochemicals per extract, solvent polarity, and the coating technique used to apply the extracts onto the fabrics.

To further validate that the fabric was modified by the extracts of *A. afra* and *E. globulus*, an FTIR study was conducted to understand the chemical attachment. The FTIR spectra of uncoated and coated polyester and cotton fabric are presented in Figures 5-12.



Figure 3.4: FTIR spectra of polyester fabrics uncoated (U) and coated (1-3) with *A. afra* hexane coating. Samples 1-12 were performed in triplicates.

FTIR spectra analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and Nandiyanto et al., (2019), for the uncoated polyester fabric, the peak range of 1715 cm⁻¹ indicates the C=O group. The peaks in the region 1502 cm⁻¹ indicate vibrations of an aromatic ring and peaks at 1473 cm⁻¹ indicate the CH₂ bending group, and at 1411-1409 cm⁻¹ indicate vibrations of an aromatic ring. The peaks at 1342-1340 cm⁻¹ represent a CH₂ wagging vibration of trans-co-formation, and the peak 1097 cm⁻¹ indicates CH in-plane bending modes of an aromatic ring. Strong peaks at 1243-1241 cm⁻¹ indicate C–O stretching bonds. Peak 967 cm⁻¹ indicates trans-C-O stretching+ vibrations of the ester group and at peak 872 cm⁻¹ presents an aromatic ring. A peak range of 724 cm⁻¹ indicates ring CH out-of-plane bending + C=O out-of-plane bending group.

For the coated *A. afra* hexane fabric, small peaks around 3427 cm⁻¹ indicate the -OH group bond for samples 1-3. Peaks around 2958 cm⁻¹, 2918 cm⁻¹ and 2852 cm⁻¹ indicate C-H stretching bond from samples 1-3. Sample 3 shows strong peaks around this region whereas sample 1-2 peaks are weak but still indicate of C-H stretching bond for samples 1-3. Peaks from 1715 -1713 cm⁻¹ indicate the C=O stretch group for samples 1-3. The peaks around 1502 cm⁻¹ indicate vibrations of an aromatic ring for samples 1-3 and the peaks at 1473-1471 cm⁻¹ indicate a CH₂ bending bond for samples 1-3. The peaks at 1410 cm⁻¹ indicate an aromatic ring and the peak at 1340 cm⁻¹ indicates a CH₂ bending functional group for samples 1-3. Peaks between 1242- 1232 cm⁻¹ represent the C–O stretch (ester) for samples 1-3. The peaks in the range of 1096, cm⁻¹ 1019-1017 cm⁻¹ indicates C=C stretching trans -C-H stretch outof-plane bend for samples 1-3. Peaks ranging between 725-720 cm⁻¹ indicate ring -C-H out of plane bending + C=O out of plane bending for samples 1-3.

FTIR analysis reported by Arockia et al., (2015), for the coated polyester fabrics, from samples 1-3, the prominent peaks in the region of 2918 cm⁻¹ indicate C-H symmetric stretching vibrations of CH₂ which is an indication of lipids and protein present and the vibrations of 1502 cm⁻¹ indicate vibrations of an aromatic ring stretch indicating an aromatic compound and peaks at 1473 cm⁻¹ indicate the C=C-C aromatic compound present, and at 1411-1409 cm⁻¹ indicate vibrations of O-H bend indicating a phenol compound as well as the peaks at 1342-1340 cm⁻¹. The peak 1097 cm⁻¹ indicates the C-O stretch indicating the presence of cyclic ethers. Strong peaks at 1243-1241 cm⁻¹ indicate C–O stretching bonds. Peak 967 cm⁻¹ and at peak 872 cm⁻¹ presents a P-O-C stretch, indicating aromatic phosphates. A peak range of 724 cm⁻¹ indicates the C-CI stretch aliphatic chloro compound.



Figure 3.5: FTIR spectra of polyester fabrics uncoated (U) and coated (4-6) with *A. afra* methanol coating

In the FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020) and Nandiyanto et al., (2019), the coated *A. afra* methanol fabrics from samples 4-6, the peak ranging from 2965-2851 cm⁻¹ indicates the presence of indicates C-H stretching bond (aliphatic compounds C-H stretching in methyl and methylene groups). They are also identified between this range followed by peaks between 1470-1414cm⁻¹. The peaks at 1714 cm⁻¹ indicate the presence of indicates C=O stretching bond. The small peaks in the range of 1507 cm⁻¹ indicate vibrations from an aromatic ring and the peaks at 1469 cm⁻¹ CH₂ bending functional group. The peaks in 53

the range of 1414 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 1410 cm⁻¹ indicate an aromatic ring. The peak at 1341 cm⁻¹ indicates a CH₂ bending functional group. The wide peaks at the range of 1240 cm⁻¹ indicate the C–O stretch group. The peaks at 1097 cm⁻¹ and 1018 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 972 cm⁼¹ and 872 cm⁻¹ also indicate an aromatic ring. The peaks in the range of 850 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 972 cm⁼¹ and 872 cm⁻¹ also indicate an aromatic ring. The peaks in the range of 970 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 850 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 972 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 850 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 972 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 850 cm⁻¹ indicate vibrations of an aromatic ring. The peaks in the range of 972 cm⁻¹ indicates -C-H out of plane bending + C=O out of plane bending.

FTIR analysis reported by Arockia et al., (2015), the coated *A. afra* methanol fabrics from samples 4-6, the prominent peaks ranging from 2965-2851 cm⁻¹ indicate the presence of C-H stretching bond (aliphatic compounds). The peaks at 1714 cm⁻¹ indicate the presence of C=O, indicating a carbonyl compound. The small peaks in the range of 1507 cm⁻¹ and 1469 cm⁻¹ indicate C=C-C vibrations, indicating an aromatic compound. The peaks in the range of 1414-1410 cm⁻¹ indicate vibrations of the O-H bend, indicating phenol compounds. The peak at 1341 cm⁻¹ indicates O-H vibrations, indicating the presence of phenol compounds. The wide peaks at the range of 1240 cm⁻¹ indicate the C-O stretching bond, indicating. The peaks at 1097 cm⁻¹ indicate the presence of the C-O stretch group, indicating the presence of cyclic ethers and 1018 cm⁻¹ indicates the presence of phosphate ions. The peaks in the range of 972 cm⁻¹ and 850 cm⁻¹ also indicate the P-O-C stretch group, indicating aromatic phosphates. An aromatic ring. The peak in the range of 725 cm⁻¹ indicates the C-CI stretch group indicating an aliphatic chloro compound.



Figure 3.6: FTIR spectra of polyester fabrics uncoated (U) and coated (7-9) with E. *globulus* hexane coating

The FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and Nandiyanto et al., (2019), the coated *E. globulus* hexane-coated fabric from samples 7-9, peaks at 2952 cm⁻¹ indicate C-H stretch in methyl and methylene groups (aliphatic compounds), followed by peaks at 1471 and 725 cm⁻¹. The peaks at 1714 cm⁻¹ indicate C=O stretching bond. There are peaks at the band range 1506 cm⁻¹, indicating that there are vibrations of an aromatic ring. The peaks around 1479 cm⁻¹ indicate CH₂ bending. The peaks at 1412-1402 cm⁻¹ indicate an aromatic ring and the peaks at 1342 cm⁻¹ indicate a CH₂ bending functional group. Peaks between 1244-1237 cm⁻¹ represent the C–O stretch group. The peaks at 1097 cm⁻¹ indicate vibrations of an aromatic ring along with peaks in the range of 1017- 972 cm⁻¹ and peaks at 871 cm⁻¹. Peaks ranging between 725-720 cm⁻¹ indicate ring -C-H out of plane bending + C=O out of plane bending.

FTIR analysis reported by Arockia et al., (2015), the coated *E. globulus* hexane-coated fabric from samples 7-9, peaks at 2952 cm⁻¹ indicating a C-H stretch group, indicating an alkane compound, followed by a peak at 1471 cm⁻¹ indicating aromatic ring stretch indicating an aromatic compound. The peak at 725 cm⁻¹ indicates the C-Cl stretch group indicating an aliphatic chloro compound. The peaks at 1714 cm⁻¹ indicate the C=O group, indicating the carbonyl group. There are peaks at the band range 1506 cm⁻¹ and indicating the C=C-C group, indicating that there are vibrations of an aromatic ring. The peaks in the range of 1479 cm⁻¹ indicate an aromatic ring stretch C=C-C group, indicating an aromatic ring stretch C=C-C group, indicating phenol compound. The peaks range 1412-1342 cm⁻¹ indicates the O-H bend group. The peaks at 1097 cm⁻¹ indicate the presence of the C-O stretch group, indicating the presence of cyclic ethers and 1018 cm⁻¹ indicates the presence of phosphate ions. The peaks in the range of 972 cm⁻¹ and 850 cm⁻¹ also indicate the P-O-C stretch group, indicating aromatic phosphates. an aromatic ring. The peak in the range of 725 cm⁻¹ indicates the C-Cl stretch group indicating an aliphatic chloro compound.



Figure 3.7: FTIR spectra of polyester fabrics uncoated (U) and coated (10-12) with *E. globulus* methanol coating.

In the FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and Nandiyanto et al., (2019) the *E. globulus* methanol-coated fabric from samples 10-12, the peaks around 2935 cm⁻¹ In indicate in C-H stretching in methyl and methylene groups. The peaks at a range of 1714 -1710 cm⁻¹ C=O stretching bond. The peak 1411 cm⁻¹ indicates an aromatic ring and the peak 1342cm ⁻¹ indicates a CH₂ bending functional group. The peak range 1247-1237 cm⁻¹ indicates the C-O stretch group. Peaks at 1092 cm⁻¹ indicate vibrations of an aromatic ring. The peaks at 1018 cm⁻¹ and 873 cm⁻¹ indicate vibrations of an aromatic ring. A peak range of 847-735 cm⁻¹ indicates vibrations of aromatic rings. Peaks at 725-722 cm⁻¹ indicating CH out of plane bending + C=O out of plane bending.

FTIR analysis reported by Arockia et al., (2015), the *E. globulus* methanol-coated fabric from samples 10-12, the peaks around 2935 cm⁻¹ indicate asymmetric stretching of -CH(CH₂) vibration indicating the presence of lipids and protein compounds. The peaks at a range of 1714 -1710 cm⁻¹ C=O stretching bond, indicating carbonyl compounds. The peaks range 1412-1342 cm⁻¹ indicates the O-H bend group, indicating phenol compounds. Peaks between 1247-1237 cm⁻¹ represent the C–O stretch group. The peaks in the range 1097 cm⁻¹ indicate the presence of the C-O stretch group, indicating the presence of cyclic ethers and 1018 cm⁻¹ indicate the presence of phosphate ions. The peaks in the range of 972 cm⁻¹ and 850 cm⁻¹ also indicate the P-O-C stretch group, indicating aromatic phosphates. An aromatic ring. The peak in the range of 725-722 cm⁻¹ indicates the C-CI stretch group indicating an aliphatic chloro compound.



Figure 3.8: FTIR spectra of cotton fabrics uncoated (U) and coated (1-3) with *A. afra* Hexane coating

The FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020) and Nandiyanto et al., (2019), the uncoated cotton, peaks for cotton fabrics are found at around 3326 cm⁻¹ corresponding to O–H stretching, 2897 cm⁻¹ region for C-H stretching in methyl and methylene groups. The peak at 1611 cm⁻¹ indicates the C=O carbonyl group. Peaks at 1417 cm⁻¹ indicate vibrations of an aromatic ring. A peak at 1311 cm⁻¹ is associated with the O-H group and peaks at 1120 cm⁻¹, 1066 cm⁻¹ and 1120-1029 cm⁻¹ indicate a C-O stretch. Peaks at 895 cm⁻¹ indicate the C=C group and 680-629 cm⁻¹ indicate the alkyne C-H bend.

For the cotton, *A. afra* hexane coated fabric from samples 1-3, with peaks of 3323 cm⁻¹ indicates O-H stretch. A small wide absorption peak in 2889 cm⁻¹ indicates C-H stretching in methyl and methylene groups. A peak in the range of 1429 cm⁻¹ indicates the bending of the CH group and the peak range around 1316 cm⁻¹ is associated with CH₂ groups. Peaks between the range of 1129 -1119 cm⁻¹ indicate C-O stretching of trans conformer. The peak range at 1109 cm⁻¹ indicates a C-O stretch is associated with CH₂ groups -cyclohexane vibrations at 1057 cm⁻¹ and 1000 cm⁻¹. Peaks from 667-620 cm⁻¹ indicate an alkyne C-H bend. For the cotton, *A. afra* hexane coated fabric from samples 1-3, with peaks of 3323 cm⁻¹ indicates O-H stretch. A small wide absorption peak in 2889 cm⁻¹ indicates C-H stretching in methyl and methylene groups. A peak in the range of 1429 cm⁻¹ indicates the bending of the CH group and the peak range around 1316 cm⁻¹ is associated with CH₂ groups. Peaks between the range of 1129 -1119 cm⁻¹ indicate C-O stretching of trans conformer. The peak range at 12057 cm⁻¹ indicates C-H stretching in methyl and methylene groups. A peak in the range of 1429 cm⁻¹ indicates the bending of the CH group and the peak range around 1316 cm⁻¹ is associated with CH₂ groups. Peaks between the range of 1129 -1119 cm⁻¹ indicate C-O stretching of trans conformer. The peak range at 1109 cm⁻¹ indicates a C-O stretch is associated with CH₂ groups -cyclohexane vibrations at 1057 cm⁻¹ and 1000 cm⁻¹. Peaks from 667-620 cm⁻¹ indicate an alkyne C-H bend.

FTIR analysis reported by Arockia et al., (2015), the peaks for cotton fabrics are found at around 3326 cm⁻¹ corresponding to O–H stretching group indicating polyhydroxy compounds. The peak at 1611 cm⁻¹ indicates the C=O stretching vibrations, indicating a ketone group. A peak at 1311 cm⁻¹ is associated with the O-H group indicating a phenol compound and peaks in the range of 1120 cm⁻¹ indicate a C-O stretch indicating a cyclic ether compound. Peaks at 895 cm⁻¹ indicate aromatic phosphates (P-O-C group) and 680-629 cm⁻¹ indicate the C-Br group, indicating aliphatic bromo compounds.



Figure 3.9: FTIR spectra of cotton fabrics uncoated (U) and coated (4-6) with *A. afra* methanol coating

The FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and (Nandiyanto et al., (2019), the cotton *A. afra* methanol coated fabric from samples 4-6, peaks between 3340-3268 cm⁻¹ indicate the -OH group. Peaks at 2897-2823 cm⁻¹ C-H stretching in methyl and methylene groups. Peaks at 1425 cm⁻¹ indicate vibrations of an aromatic ring. Peaks at 1315 cm⁻¹ are associated with the –CH₂ group. Peaks at 1163 cm⁻¹-1123 cm⁻¹ are associated with C-O stretch as well as peaks at 1105 cm⁻¹-1068 cm⁻¹. Peaks at 1000 cm⁻¹ indicate vibrations of an aromatic ring. Peaks at 681- 631 cm⁻¹ indicate an alkyne C-H bend group.

FTIR analysis reported by Arockia et al., (2015), the cotton, *A. afra* methanol coated fabric from samples 4-6, peaks between 3340-3268 cm⁻¹ indicating the O-H stretch group indicating a polyhydroxy compound. Peaks at 2897-2823 cm⁻¹ C-H stretching in methyl and methylene groups. Peaks at 1425 cm⁻¹ indicate vibrations of an aromatic ring. Peaks at 1315 cm⁻¹ are associated with the O-H bend group indicating the presence of a phenol group. Peaks in the range 1105 cm⁻¹-1000 cm⁻¹ indicate phosphate ion group indicating phosphate compounds. of an aromatic ring. Peaks at 681- 631 cm⁻¹ indicate a C-Br stretch (aliphatic bromo compounds).



Figure 3.10: FTIR spectra of cotton fabrics uncoated (U) and coated (7-9) with *E. globulus* hexane coating

The FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and Nandiyanto et al., (2019), the cotton *E. globulus* hexane-coated fabric from samples 7-9, peaks ranging between 3333-3254 cm⁻¹ indicate O-H stretching. Peaks at 2879-2823 cm⁻¹ C-H stretching in methyl and methylene groups. Small peaks ranging from 1636-1626 cm⁻¹ indicate asymmetrical stretching vibration of the COO- bond. Peaks at 1428 cm⁻¹ indicate bending of the CH group and at 1372 cm⁻¹ indicate CH₂ wagging vibration of gauche conformation. Peaks at 1317 -1307 cm⁻¹ asymmetric stretching of C–O–C group. Peaks 1124-1068 cm⁻¹ C-O stretching of trans conformer. Peaks at 1000 cm⁻¹ indicate vibrations of an aromatic ring. Peaks at 622- 612 cm⁻¹ indicate an alkyne C-H bend.

For FTIR analysis reported by Arockia et al., (2015), the cotton *E. globulus* hexane-coated fabric from samples 7-9, peaks ranging between 3333-3254 cm⁻¹ indicates the O-H stretch group indicating polyhydroxy compound. Small peaks ranging from 1636-1626 cm⁻¹ indicate the C-O stretch, indicating the ketone compound. Peaks at 1372- 1307cm⁻¹ indicate O-H bending group indicating a phenol compound. Peaks at 1000 cm⁻¹ indicate vibrations of the phosphate ion group indicating a phosphate compound. Peaks in the range of 622- 612 cm⁻¹ indicate an alkyne C-Br vibration indicating an aliphatic bromo compound.



Figure 3.11: FTIR spectra of cotton fabrics uncoated (U) and coated (10-12) with *E. globulus* methanol coating.

The FTIR analysis reported by Sumantri et al., (2020), Hayat et al., (2020), and Nandiyanto et al., (2019), the cotton *E. globulus* methanol-coated fabric from samples 10-12, peaks around 3332-3270 cm⁻¹ indicate -OH cm⁻¹ group and peaks around 2935 cm⁻¹ were observed and indicate C-H stretching in methyl and methylene group. The peaks at a range of 1714-1710 cm⁻¹ indicate the C=O stretching bond. The peaks at 1411 cm⁻¹ indicate an aromatic ring. The peak at 1342cm⁻¹ indicates a CH₂ bending functional group. The peak range 1247-1237 cm⁻¹ indicates the C-O stretch group. Peaks at 1092 cm⁻¹ indicate vibrations of an aromatic ring. Peaks at 1018 cm⁻¹ and 873 cm⁻¹ indicate vibrations of the aromatic ring. A peak range of 847-735 cm⁻¹ indicates vibrations of aromatic rings. Peaks at 725-722 cm⁻¹ indicating CH out of plane bending + C=O out of plane bending.

The FTIR spectrum results demonstrated absorption For FTIR analysis reported by Arockia et al., (2015), the cotton *E. globulus* methanol coated fabric from samples 10-12, peaks around 3332-3270 cm⁻¹ indicates O-H stretch group indicating polyhydroxy compound and peaks around 2935 cm⁻¹ were observed and indicate asymmetric stretching of -CH(CH₂) vibration indicating saturated aliphatic compound. The peaks at a range of 1714-1710 cm⁻¹ indicate the C=O stretch indicating the carbonyl group. The peaks range 1412-1342 cm⁻¹ indicates the O-H bend group, indicating phenol compounds. Peaks between 1247-1237 cm⁻¹ represent the C-O stretch group. The peaks in the range 1097 cm⁻¹ indicate the presence of the C-O stretch group.
phosphate ions. The peaks in the range of 972 cm⁻¹ and 850 cm⁻¹ also indicate the P-O-C stretch group, indicating aromatic phosphates. an aromatic ring. The peak in the range of 725-722 cm⁻¹ indicates the C-CI stretch group indicating an aliphatic chloro compound.

The FTIR spectrum results reveal that the methanol and n-hexane extract functional groups exhibit absorption signals across multiple wavenumber ranges. These functional groups contain chemical bonds that may contribute to their antimicrobial properties. For instance, the presence of alcohols and phenols (O–H), polyhydroxy compounds (O-H stretch in the extracts), carboxylic acids (C-O stretching), and methyl and aldehyde groups (stretching of C-H bonds) may disrupt cell membranes or inhibit bacterial enzymes, thus potentially enhancing the extracts' antimicrobial activity. Moreover, alkenes (C=C stretching) and aromatics (C-C stretching group) may induce oxidative stress in bacterial cells, contributing to the extracts' anti-inflammatory and antimicrobial properties. Additionally, CH₂ methylene groups and aliphatic compounds (C-H stretching), aliphatic bromo compounds (C-Br), and P-O-C stretch (aromatic phosphates) may also be involved in the biological activity of the extracts. Polyhydroxy compounds (O-H stretch) may contribute to the extracts' anti-inflammatory and immunomodulatory properties, in addition to their antimicrobial activity. It's worth noting that the chemical composition of the extracts may vary depending on factors such as geographical location, climate conditions, drying methods, population variability, and extraction solvents.

The FTIR spectra were found to be similar to those reported by Sumantri et al., (2020), Hayat et al., (2020), and (Nandiyanto et al., (2019) but different from the FTIR spectra reported by Arockia et al., (2015). Overall, based on the FTIR spectrum results, it appears that the methanol and n-hexane extracts of *Artemisia afra* and *Eucalyptus globulus* contain a complex mixture of bioactive compounds with potential antimicrobial properties. (Liu et al., 2009).

Chapter 4

Conclusions and Recommendations

CHAPTER 4: CONCLUSIONS AND RECOMMENDATIONS

4. Conclusion

Eucalyptus globulus and Artemisia afra displayed antimicrobial activity against the Grampositive S. aureus strain but poor activity against the Gram-negative bacteria E. coli strain. The methanolic extractions showed the highest yield compared to hexanoic extractions. The obtained results revealed that the different solvents resulted in different extraction yields due to the differences in solvent polarity. The Eucalyptus globulus hexanoic and methanolic extracts showed good antibacterial activity against the S. aureus strain and poor activity against the E. coli strain. The Artemisia afra hexanoic and methanolic extracts revealed poor antimicrobial activity against the *E. coli* strain. Microbial growths were rather observed when *Eucalyptus globulus* hexane and methanol extracts were tested against *E. coli*. The MIC results revealed that only some samples showed activity in the liquid media against the selected ATCC bacterial strains. The FTIR analysis confirmed the presence of a composition of alcohol and phenols, carboxylic acids, methyl and aldehyde group, alkenes, and aromatics functional groups. The results of this study show that hexane and methanol extract of E. globulus extractions could be a good candidate as an antibacterial agent to fight against Gram-positive bacteria. Only a few coated samples showed a zone of inhibition. No zone of inhibition was detected after one cycle of washing. This could be due coating technique used to apply the extracts onto the fabrics.

Plants play an essential role in the pharmaceutical and healthcare sectors due to their ability to fight pathogens. Unexplored medicinal plants with potent bioactive compounds should be studied, and various plant parts should be investigated to determine the plant part containing the highest bioactive concentration. More research on plant-based antimicrobial agents and finishing should be conducted to extend the antimicrobial activity and durability to laundering on textile substrates. The study's outcome suggests that freeze-drying should be used as a standard drying method at the pre-treatment stage, as it preserves bioactive compounds in the plant material, given that most phytochemicals and bioactive compounds are thermolabile. Freeze-drying is a very energy-consuming process but avoid the thermal decomposition of the valuable bioactive compounds during the drying process.

Finding a single method for a quick, efficient, and eco-friendly extraction has been a struggle due to the limitations of traditional extraction methods. The exploration of green extraction techniques has gained traction due to reduced time, energy, solvent consumption, and low operating costs and its positive impact on global warming and climate change, reducing toxic solvent usage and carbon emissions. Multiple extractions and green extraction methods should be used to increase the extraction yield. Only solvents regarded as safe with strong

polarity should be used for extraction to achieve maximum yield. Additionally, extracts should be further purified by eliminating compounds that might enhance bacterial growth.

Chapter 5

References

CHAPTER 5: REFERENCES

5. References

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Chapter 6

Appendices

CHAPTER 6: APPENDICES – RAW DATA

Addendum A: Antibacterial Activity analysis

| Sample 1 | 1,331 | 1,393 1,393 | Sample 3 1,329 | Sample 4 2,144 | Sample 5 2,203 | Sample 6 2,128 | Sample 7 S | ample 8 Sa | ample 9 1,064 | Sample 10 5 0.222 | Sample 11 S 0.224 | ample12 0.246 |
|----------|----------|-------------|-------------------|-------------------|-------------------|-------------------|------------|------------|------------------|----------------------|----------------------|------------------|
| 1,34/ | - | 1,349 | 1,354 | 2,168 | 2,281 | 2,157 | 0,893 | 0,85 | 1,294 | 0,365 | 0,386 | 0,406 |
| 1,309 | 6 | 1,305 | 1,354 | 1,917 | 2,028 | 2,19 | 1,054 | 0,861 | 1,099 | 0,317 | 0,363 | 0,348 |
| 1,075 | 10 | 1,119 | 1,224 | 2,014 | 2,152 | 1,973 | 0,659 | 0,695 | 0,73 | 0,35 | 0,345 | 0,359 |
| 1,22 | 0 | 1,305 | 1,357 | 1,338 | 1,381 | 1,323 | 0,667 | 0,663 | 0,684 | 1,188 | 1,168 | 1,363 |
| 1,30 | 6 | 1,211 | 1,215 | 0,374 | 0,409 | 0,403 | 0,612 | 0,675 | 0,665 | 0,198 | 0,185 | 0,185 |
| 1,93 | 4 | 2,208 | 2,132 | | | | 0,226 | 0,238 | 0,227 | | | |
| 2,01 | <u> </u> | 2,188 | 2,357 | | | | 0,212 | 0,229 | 0,267 | | | |
| | Á | verage OD | %growth | | NOTE: | | | | | | | |
| | | 1,351 | 100,272143 | | No effect | 0 | | | | 1 8 × 8 | -(+-(| in- and |
| | | 1,349 | 100,123701 | | No effect | | | | | | NO NON | 6 |
| | • | 1,322666667 | 98,1692232 | | | | | | | 入入 | | 6 |
| | • | 1,139333333 | 84,562098 | | | | | | | | かし、この | (05) |
| | • | 1,294 | 96,0415636 | | | 1 | | | | | | |
| | • | 1,245 | 92,4047501 | | | | | | | | | |
| | | 2,091333333 | 155,220188 | | Enhanced gro | wth | | | | | | 6 |
| | | 2,187333333 | 162,345374 | | Enhanced gro | wth | | | | | | 1 () |
| | • | 2,158333333 | 160,192974 | | Enhanced gro | wth | | | | C. C. C | KOMON | |
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| | <u>ا</u> | 2,045 | 151,781296 | | Enhanced gro | wth | | | | | 「「」 | and a second |
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| | . | 0,990666667 | 79,9139554 | | | | | | | | 「シー | |
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| | • | 1,004666667 | 81,0432912 | | | / | A | 1 | | | 1 | |
| | ` | 0,694666667 | 56,036569 | | | | | | | | | |
| | ` | 0,671333333 | 54,1543426 | | | | | | | | | |
| | • | 0,650666667 | 52,4872277 | | | | | | | | | |
| | | 0,230333333 | 18,5802635 | | | | | | | | | |
| | | 0,236 | 19,0373756 | | | | | | | | | |
| | • | 0,230666667 | 18,6071525 | | | | | | | | | |
| | • | 0,385666667 | 31,1105136 | | | | | | | | | |
| | | 0,342666667 | 27,6418392 | | | | | | | | | |
| | | 0,351333333 | 28,3409519 | | | | | | | | | |
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| Analysis | |
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| Sample 2 88,49901 66,0547 100,0595 113,0862 123,5753 112,5753 110,5298 | sample 5 81,30128 83,89806 95,02848 116,1116 123,3227 110,8408 112,2792 93,35843 | Sample 8 129,4705 67,63548 136,08855 196,08855 179,1002 160,3415 175,9396 153,7508 | Sample 11 55,59568 75,59568 201,09568 201,0956 172,0961 172,0046 172,0046 150,1117 | Sample 2 92,11198 90,45172 90,45172 93,57818 94,53768 94,53768 94,78564 94,78564 | Sample 5 92,16836 91,01016 90,209 90,209 102,7925 73,41074 95,74746 101,5559 | Sample 8 75,961975 75,96083 1175,6563 1102,2371 119,906 1102,2531 102,2531 | Sample 11 102,2528 106,5212 160,3984 169,3984 159,3502 171,8233 171,8233 171,8233 171,8233 136,173 |
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| % growth Average 86,51274 85,1969 108,4182 106,5416 100,777 88,41756 60,55641 82,65297 | % growth Average 99,2599 93,17823 84,61563 97,96312 101,4898 98,06931 59,30109 91,11562 | % growth Average 205,0321 233,8208 147,312 147,312 152,0776 147,2252 152,0776 132,2499 | % growth Average 255,9874 255,9874 199,5999 199,5999 158,6315 143,2489 153,6963 153,6963 146,1679 | % growth Average 82,3312 83,68132 93,42039 95,33547 88,04039 111,5284 111,5284 85,01096 104,0249 | % growth Average 78,56023 77,58491 70,58396 80,08999 81,38752 80,07837 73,49202 93,44557 | % growth Average 129,343 134,6558 152,165 152,165 152,165 93,18182 93,18182 99,50592 99,50592 | % growth Average 156,5766 161,5366 151,5366 151,5366 151,5366 151,5366 151,5366 119,8719 1119,8719 94,89764 1113,8171 |
| 80,46305 86,67565 111,7422 112,132 110,6798 100,6798 87,61553 | 90,8968 94,93194 86,5431 94,9431 94,9097 95,88763 60,82633 84,80548 | 215,0838 246,8286 167,2482 157,3761 151,1294 145,5241 138,5431 144,2134 | 205,0777 268,9532 221,9532 194,8238 157,1991 144,0228 147,255 139,1555 | 82,30136 73,47181 93,77824 96,47824 94,94735 117,3932 96,41356 104,6502 | 79,22787 71,05951 87,43106 69,45718 71,94775 71,94775 92,33382 | 122,8938 120,3762 144,6948 114,6948 88,34248 90,16458 96,04232 100,1763 | 149,6799 168,6744 144,8897 144,8897 122,1484 92,47095 84,90795 115,4849 |
| 90,60434 86,60936 110,2572 110,4112 96,76995 75,3971 61,29891 | 106,3713 85,79013 85,79013 85,89013 100,1834 99,72005 95,72353 59,09837 88,87924 | 205,4535 205,4535 228,7671 148,0061 148,0061 148,0247 156,7533 147,2624 121,4297 | 215,464 232,0515 195,3695 197,3695 197,3695 191,2728 141,3041 141,3041 142,8009 | 84,31739 93,63208 97,51221 103,4211 103,4211 103,4217 109,1135 81,18015 108,4774 | 77,62554 75,14369 74,15965 70,47605 93,24819 93,24819 84,53991 78,86212 96,95791 | 128,4483 137,4706 1537,4706 153,4706 153,4706 153,4706 153,43393 95,48393 106,1422 95,43127 | 161,3469 169,3384 169,3384 112,3384 112,3368 112,3548 108,5677 118,5677 |
| Sample 1 88,47082 82,30521 103,2555 103,0765 104,9281 89,17573 89,17573 80,58523 | Sample 4 100,5116 98,81263 98,7436 104,7398 102,5968 57,97857 99,66213 | Sample 7 194,5589 225,8668 149,5580 142,78162 142,7819 153,9553 155,8702 155,8702 131,1065 | Sample 10 210,2607 216,59574 185,5169 206,6051 167,4224 144,4199 154,4091 156,5474 | Sample 1 80,55486 83,94748 88,97474 86,10702 86,76465 108,0785 77,43918 98,94707 | Sample 4 78,82729 78,65152 78,65152 78,65925 71,65925 71,29608 83,74746 69,85776 91,04499 | Sample 7 136,6869 158,1207 158,1207 158,1207 125,5150 99,30447 91,75157 91,75157 101,7535 | Sample 10 158,7029 146,5971 147,7828 136,6493 136,6493 135,1126 102,90202 99,50202 107,3986 |
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| FAXE 1 3 aureus 6 7 8 9 10 11 1200% (exture control) 0.0893 0.9893 0.9883 0.9633 0.9431 100% (exture control) 0.0892 0.9333 0.9431 11 12.00% (exture control) 0.0892 0.9333 0.9431 0.9431 10.9431 10.95866667 10.056 0.0199 0.9333 0.9431 0.9431 0.9431 10.9411 10.95866667 10.056 10.1111 | ALXE 1 2 5 4 0 11 5 4 0 13 5 4 0 13 10 13 10 13 10 13 10 10 13 10 13 10 13 10 13 10 10 13 10 10 13 10 | FASE 1 State 1 State 6 7 7 8 9 10 11 20% (extrue control) 0.0993 1.759 1.4028 1.507 1.4028 1.507 1.4058 1.405 | AA3E 1 3 auroni 5 5 auroni 6 5 auroni 6 6 6 7 7 0 2 2 0 2 1 2 0 2 2 0 2 2 1 2 0 2 2 0 2 <td>RASE 1 9 10 11 2.00% (editure control) 0.445 0.377 0.4415 0.9445 0.412 0.4256 0.425 0.4256 0.4267 0.425 0.4266</td> <td>AA3E 1 9 10 11 22 100% (edd) <th< td=""><td>ALSE 1 0 10 11 E. coll 0.900 6 7 7 9 10 11 12 100% (address control 0.900 6 0.915 0.854 0.885 1.088 1.088 1.02789 0.912 1.07789 0.8454 1.088 1.02789 3.041 1.02789 3.041 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0218 3.12 <t< td=""><td>ALMSE 1 8 9 10 11 12 100% (outluter control control 110% (outluter control 110% (outluter control 110% (outluter control 111 112 100% (outluter control 111 111 110% (outluter control 110% (outluter control 110% (outluter control 111 1</td></t<></td></th<></td> | RASE 1 9 10 11 2.00% (editure control) 0.445 0.377 0.4415 0.9445 0.412 0.4256 0.425 0.4256 0.4267 0.425 0.4266 | AA3E 1 9 10 11 22 100% (edd) 100% (edd) <th< td=""><td>ALSE 1 0 10 11 E. coll 0.900 6 7 7 9 10 11 12 100% (address control 0.900 6 0.915 0.854 0.885 1.088 1.088 1.02789 0.912 1.07789 0.8454 1.088 1.02789 3.041 1.02789 3.041 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0218 3.12 <t< td=""><td>ALMSE 1 8 9 10 11 12 100% (outluter control control 110% (outluter control 110% (outluter control 110% (outluter control 111 112 100% (outluter control 111 111 110% (outluter control 110% (outluter control 110% (outluter control 111 1</td></t<></td></th<> | ALSE 1 0 10 11 E. coll 0.900 6 7 7 9 10 11 12 100% (address control 0.900 6 0.915 0.854 0.885 1.088 1.088 1.02789 0.912 1.07789 0.8454 1.088 1.02789 3.041 1.02789 3.041 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.02789 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0278 3.12 1.0218 3.12 <t< td=""><td>ALMSE 1 8 9 10 11 12 100% (outluter control control 110% (outluter control 110% (outluter control 110% (outluter control 111 112 100% (outluter control 111 111 110% (outluter control 110% (outluter control 110% (outluter control 111 1</td></t<> | ALMSE 1 8 9 10 11 12 100% (outluter control control 110% (outluter control 110% (outluter control 110% (outluter control 111 112 100% (outluter control 111 111 110% (outluter control 110% (outluter control 110% (outluter control 111 1 |
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| Indepint Austriction FALSE 1 Stativest 3 3 5 6 7 8 9 10 11 100% (extrate control) 3 3 5 6 7 8 9 10 11 12.00% (extrate control) 00221 07316 0.8945 1091 0.9848 10093 10 12 100% (extrate control) 03221 0.7320 0.8940 0.9948 1.0993 0.9441 1 10.09566647 10 12 10.05566647 10 12 10.05566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 12 10.95566647 10 <td< td=""><td>Indport Absorbanc FALSE 1 2 5 4 5 4 9 10 11 2 00% 10% 10% 10% 10% 5 5 4 00% 10% 10% 11 10% 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 10 11 10% 10 11 11 10% 10 11 10% 10 11 10% 10 11 10% 10 11 10% 10 10%<</td><td>Implement Abstronk Raw PASS PASS 1 2 5 surrent 2 3 3 1 1 1 100% (4/10 met control) 2 3 3 1 1 1 100% (4/10 met control) 2 3 1 1 1 1 100% (4/10 met control) 2 3 1 1 1 1 1 100% (4/10 met control) 2 3 1</td><td>Old point Absorbanc flaw FALSE 1 9 10 11 2,1,0004 5, aurora control 20101 0.04423 0.9417 1.4366 1.4474 9 10 11 12,00% (outror control) 20101 0.04423 0.9417 1.4366 1.4474 9 10 11 12,00% (outror control) 2,1448 0.7430 0.7730 0.9317 1.2434 1.4474 0.98206667 3 2,1448 1.4474 0.9812 1.4714 1.0461 1.233 1.1046 1.233 1.0416 3 3 0.98206667 3</td><td>Objectivit Absordance (kaw. FAXE 1 8 9 101 11 2100% (editure control) 0.7533 0.7445 0.844 0.345 0.377 0.844 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756 0.9756667 0.975 0.975766667 0.976 0.97566 0.975 0.9756</td><td>Objectin Abbrochanc/feav FALSE 1<!--</td--><td>Jabpent Abcorbanc faw FA15E 1 9 100 11 12 100% (address second seco</td><td>Oppont Ansterbanchaw AAXE 1 2 2 1 2 2 1</td></td></td<> | Indport Absorbanc FALSE 1 2 5 4 5 4 9 10 11 2 00% 10% 10% 10% 10% 5 5 4 00% 10% 10% 11 10% 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 4 5 5 4 10 11 10% 10 11 11 10% 10 11 10% 10 11 10% 10 11 10% 10 11 10% 10 10%< | Implement Abstronk Raw PASS PASS 1 2 5 surrent 2 3 3 1 1 1 100% (4/10 met control) 2 3 3 1 1 1 100% (4/10 met control) 2 3 1 1 1 1 100% (4/10 met control) 2 3 1 1 1 1 1 100% (4/10 met control) 2 3 1 | Old point Absorbanc flaw FALSE 1 9 10 11 2,1,0004 5, aurora control 20101 0.04423 0.9417 1.4366 1.4474 9 10 11 12,00% (outror control) 20101 0.04423 0.9417 1.4366 1.4474 9 10 11 12,00% (outror control) 2,1448 0.7430 0.7730 0.9317 1.2434 1.4474 0.98206667 3 2,1448 1.4474 0.9812 1.4714 1.0461 1.233 1.1046 1.233 1.0416 3 3 0.98206667 3 | Objectivit Absordance (kaw. FAXE 1 8 9 101 11 2100% (editure control) 0.7533 0.7445 0.844 0.345 0.377 0.844 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756667 0.9756 0.9756667 0.975 0.975766667 0.976 0.97566 0.975 0.9756 | Objectin Abbrochanc/feav FALSE 1 </td <td>Jabpent Abcorbanc faw FA15E 1 9 100 11 12 100% (address second seco</td> <td>Oppont Ansterbanchaw AAXE 1 2 2 1 2 2 1</td> | Jabpent Abcorbanc faw FA15E 1 9 100 11 12 100% (address second seco | Oppont Ansterbanchaw AAXE 1 2 2 1 2 2 1 |
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| 16 17 18 18 19 19 19 19 19 19 19 19 19 19 | Survers J.3 Plateform fielpoint Absorbanc flaw FAMSE 1 3 Survers 5 Survers 22.5 Log 1 1.2 0.941 0.841 0.855 0.931 1.001 1.1 1.20% (survers) 22.5 Log 1 1.1.010 0.8415 0.855 0.931 1.001 1.1 1.20% (survers) 22.5 Log 1 1.025 0.8815 0.8916 0.9911 1.0017 1.0019 10 1.1 1.0059 10 1.1 1.0059 10 1.2 1.005 1.0019 10 1.2 1.005 1.0019 10 1.2 1.005 1.0019 10 1.2 1.005 1.0019 1.0 1.2 1.005 1.0019 1.0019 1.0019 1.0019 1.0019 1.001 1.2 1.0019 1.0 1.2 1.0019 1.0 1.2 1.0019 1.0 1.2 1.0019 1.0019 1.0 1.2 1.0019 1.2 1.0019 1.2 | Sarces p 1.3 Plateform Bioport Absorbanc Raw FAXE 1 2 Sarces 1 2 Sarces 3 1 1 2 2 3 | Sarveur J.J. Plateform Endpoint Absorbanc Raw FASE 1 9 10 3 surveus 22.5 2069 2.16 2.014 0.847 0.931 1.338 1.386 1.473 5 1.4000 5 1.4000 22.5 2017 2.014 0.847 0.931 1.338 1.386 1.473 0.9126667 0.9056667 0.9056667 0.9056667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.91266667 0.9126667 0.9126667 0.91266667 0.9126667 0.9126667 0.9126667 0.91266667 <td>Coll plate 1,3 Plate/form Endpoint Absorbanc Raw PALE 1 9 10 11 12 200% (outure control) 22.5 0.777 0.881 0.844 0.344 0.345 0.341 0.343 0.991 11 12 200% (outure control) 22.5 0.776 0.881 0.44 0.346 0.345 0.341 0.346 0.976 0.975</td> <td>Colliptier 1.3 Plateform Endpoint Absorbanc Raw FAXE 1 P 9 10 11 E. coll Tromp21. 0001 04001 04006 0740 0400 11 12 100% coll 11 100% coll 11 12 100% coll 11 12 100% coll 11 12 100% coll 11 130% coll 10 11 12 100% coll 10 12 100% coll 10 11 12 100% coll 10 12 10 11 12 100% coll 10 12 10 11 12 10 11 12 10 11 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 11 12 10 12 11 12 10 12 11 12 10</td> <td>Colliptier 1.3 Plateform Endpoint Absorbanc Raw FAMSE 1 9 10 11 12 100% (outwee control 213. 1991 1.117 1.2453 0.0444 0.0007 0.0007 0.016 0.334 1.0078 (outwee control 1.02308 0.01 1.1 1.1076 (outwee control 1.02308 0.01 1.02308 0.01 0.01 0.01 1.02308 0.01 0.02016 0.0316 0.0316 0.01 0.02016 0.02016 0.0216</td> <td>Colliplate LibrateForm Example List <thlist< th=""> <thlist< th=""> List</thlist<></thlist<></td> | Coll plate 1,3 Plate/form Endpoint Absorbanc Raw PALE 1 9 10 11 12 200% (outure control) 22.5 0.777 0.881 0.844 0.344 0.345 0.341 0.343 0.991 11 12 200% (outure control) 22.5 0.776 0.881 0.44 0.346 0.345 0.341 0.346 0.976 0.975 | Colliptier 1.3 Plateform Endpoint Absorbanc Raw FAXE 1 P 9 10 11 E. coll Tromp21. 0001 04001 04006 0740 0400 11 12 100% coll 11 100% coll 11 12 100% coll 11 12 100% coll 11 12 100% coll 11 130% coll 10 11 12 100% coll 10 12 100% coll 10 11 12 100% coll 10 12 10 11 12 100% coll 10 12 10 11 12 10 11 12 10 11 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 10 12 11 12 10 12 11 12 10 12 11 12 10 | Colliptier 1.3 Plateform Endpoint Absorbanc Raw FAMSE 1 9 10 11 12 100% (outwee control 213. 1991 1.117 1.2453 0.0444 0.0007 0.0007 0.016 0.334 1.0078 (outwee control 1.02308 0.01 1.1 1.1076 (outwee control 1.02308 0.01 1.02308 0.01 0.01 0.01 1.02308 0.01 0.02016 0.0316 0.0316 0.01 0.02016 0.02016 0.0216 | Colliplate LibrateForm Example List List <thlist< th=""> <thlist< th=""> List</thlist<></thlist<> |
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| Mass of extract (g) | Sample | Mass of A. arra Hexane extract (| Sample | Mass of A. atra Methanol extract (g) | Sample | Mass of E. globulus Hexane extract (g) | Sample | Mass of E. globulus Methanol extract (g) |
|---------------------|--------|----------------------------------|-------------------|--|---------------------------------------|--|--------|--|
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| 0,1975 | 2 | 0,1975 | 5 | 0,8334 | œ | 0,4032 | Ħ | 2,2338 |
| 0,2015 | m | 0,2015 | 9 | 0,8881 | o | 0,4002 | 12 | 2,3621 |
| 0,9787 | | 0,219233333 | | 0,922066667 | | 0,3979 | | 2,276233333 |
| 0,8994 | | | | | | | | |
| 0,8881 | | | | | | | | |
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ADDENDUM C: Average Percentage Yield of Extractions