



**QUANTITATIVE PHYTOCHEMICAL ANALYSIS OF DESMODIUM TRIFLORUM
FOR POTENTIAL ANTIBACTERIAL ACTIVITY AND FORMULATION OF AN
HERBAL REMEDY FOR MANAGEMENT OF RESPIRATORY TRACT INFECTIONS**

BY KIWUSO HASSAN

(BU/UP/2019/1582)

**RESEARH DISSERTATION TO BE SUBMITTED TO THE DEPARTMENT OF
CHEMISTRY FOR THE PARTIAL FULFILMENT OF THE REQUIREMENTS FOR
THE AWARD OF THE DEGREE OF BACHELOR OF SCIENCE EDUCATION OF
BUSITEMA UNIVERSITY**

APRIL 2023

DECLARATION

I Kiwuso Hassan declare that this research dissertation is my original work and has not been submitted anywhere for award of a degree where other people's work was used, this has been acknowledged and cited according to the university policy.

Signature...  Date... 31st /April/2023

KIWUSO HASSAN

1900401582

APPROVAL

This research dissertation has been submitted for examination with my approval as his university supervisor.

Signature...  Date... 31st / 05 / 2023

DR OWOR ORIKO RICHARD

Contents

DECLARATION	Error! Bookmark not defined.
APPROVAL.....	Error! Bookmark not defined.
DEDICATION	v
ACKNOWLEDGEMENT	vi
LIST OF ABBREVIATION AND ACRONYMS	vii
ABSTRACT.....	ix
Chapter 1 Introduction	1
1.1 Background	1
1.2 Problem statement	3
1.3 Objectives.....	4
1.3.1 General objective	4
1.3.2 Specific objectives.....	4
1.4 Justification	5
Chapter 2 .Literature review	6
2.1 Botany of Desmodium Triflorum	6
2.2 Traditional uses of Desmodium triflorum.....	7
2.3 Phytochemistry of Desmodium triflorum	7
2.4 Compounds isolated from Desmodium triflorum.....	8
2.5 Pharmacology/ biological activities	8
2.6 Diseases that have been tested against Desmodium triflorum.....	8
2.7 Bacteria it has been tested against.....	8
2.8 Streptococcus pneumoniae	9
2.9 The problems and effects.	9
2.10 Treatment and the failure of the treatment.....	11
2.11 How pneumonia is traditionally managed.....	11
2.11.1 Resting.....	11
2.11.2 Drink Fluids.....	11
Chapter 3 .Material and methodology	12
3.1 Plant material.....	12
3.2 Extraction	12
3.2.1 Extraction with Organic solvents (OE)	12
3.2.2 Preliminary Phytochemical analysis.....	12

3.2.3 FORMULATION OF THE HERBAL REMEDY:.....	18
Chapter 4 Results and discussion.....	19
4.1 Phytochemicals characteristics of DTE.	19
4.2 Total phytochemical content	20
4.2.1 Total alkaloid content	20
4.2.2 Total tannins	20
4.2.3 Total phenolic content	20
4.2.4 Total flavonoids.....	21
4.3 The physiochemical evaluation of the formulated herbal remedy	21
Chapter 5 Conclusion and recommendation	23
5.1 Conclusion.....	23
5.2 Recommendation.....	23
References	24

DEDICATION

This report is dedicated to my parents Mr. Womusi Micheal and Ms. Mulelengi Edinasi, who supported me spiritually, physically, emotionally and financially in the entire walk in pursuing my vision. Special thanks also to Nabukwasi Shanitah and Nabirye Sumayyah for the financial in my academic journey. Not forgetting Aunt Natozo Gorrete who showed me the plant and Nambala Oliver who worked with me from the start of this journey. This research is also dedicated to Chemistry students for their special time, support and cooperation exhibited for the success of my research

ACKNOWLEDGEMENT

I thank the almighty Allah for the gift of life, wisdom, knowledge good health and his mighty protection in the entire life both in and out of school. With great pleasure, I also thank the fraternity of Busitema University, faculty of science and education, Nagongera campus for their efforts to produce quality teachers. With great Honor to department of chemistry, and all the lecturers, explicitly Dr Kamoga Omar, Dr Richard Oriko Owor, Dr Andima Moses, Dr Egor Moses and Mr Musagala peter not forgetting the laborarotory technicians namely Dr Kigozi Moses and madam Nakijoba Lydia, for their entire commitment and time to support us from the start of the course to the end. Financial support was from Busitema University Research And Innovation Fund Award (BURIF) to Dr Owor Richard Oriko. Also special thanks to all the students of faculty of science and education for the great company during my stay in Nagongera. May the almighty God bless you all.

LIST OF ABBREVIATION AND ACRONYMS

TCM: Traditional China Medicine

CAP: Community-Acquired Pneumonia

HAP: Hospital-Acquired Pneumonia

VAP: Ventilator-Acquired Pneumonia

DNA: Deoxyribonucleic acid

Vgg16: Visual Geometry Group

PCR: Polymerase Chain reaction

MV: Mechanically ventilated

PSB: Protected Specimen brush

DT: Desmodium triflorum

AE: Aqueous extract

OE: Organic extract

V/V: Volume by Volume

ml: milliliter

Conc H₂SO₄: Concentrated sulphuric acid

FeCl₃: Ferric Chloride

DTE: Desmodium triflorum extract

Na₂CO₃: Sodium carbonate

mg: Milligram

M: Molar

HCl: Hydrochloric acid

LIST OF TABLES

Table 1. Formulation of the herbal remedy

Table 2. Phytochemical characteristics of DTE

Table 3. Total phytochemical content of DTE

ABSTRACT

Objective: To quantitatively analyze the phytochemistry of *Desmodium triflorum* for potential antibacterial activity and formulation of an herbal remedy for management of respiratory tract infections.

Materials and methods: Phytochemical analysis of *Desmodium triflorum* belongs to order Fabales in Fabaceae Family, sub family desmodieae, Genus *Desmodium* and Species: *D. triflorum* was examined using organic extracts and aqueous extracts of the whole plant. The aqueous extract was used in the formulation of the herbal remedy.

Results: The phytochemical analysis of aqueous and organic extracts confirmed the presence of Alkaloids, tannins, saponins, phenolics, glycosides, Quinones and flavonoids which show high antibacterial activity.

Conclusion: Although the active components were not isolated but antibacterial active plant principles such as Alkaloids, tannins, saponins, phenolics, glycosides, Quinones and flavonoids were observed in the extract.

Chapter 1 Introduction

1.1 Background

Medicinal plants are a source of great economic value in the African continent (Dzoyem, Tshikalange, & Kuete, 2013). Ancient knowledge has been the basis of modern medicine and will remain as one important source of future medicine and healing (Sharma, Parashar, & Kabra, 2013). The future of natural products drug discovery is a multi-dimensional problem requiring several parameters such as safety and efficacy of the compound to be evaluated during drug selection (Ahuja & Sharma, 2014). The advent of latest technologies that enhance drug design hypotheses such as Artificial intelligence involving docking techniques means that technology has become part of drug discovery (Bender & Cortés-Ciriano, 2021). This has resulted in increased speed in drug discovery and evaluation of the safety (Bender & Cortés-Ciriano, 2021; Bleicher, Böhm, Müller, & Alanine, 2003).

Recent advances in analytical and computational techniques have opened new avenues to process complex natural products and to use their structures to derive new and innovative drugs (Thomford et al., 2018). Nature has bestowed a very rich botanical wealth and a large number of diverse types of plants grow in different parts of the world. From the very beginning of human existence (Issazadeh et al., 2012; Parekh, Jadeja, & Chanda, 2005), man has familiarized himself with plants and used them a number of ways up to date (Лисицын, 2004). Primitive man in search of food and to cope with human suffering began to distinguish those plants suitable for medicinal purpose from others with definitive pharmacological action. The relationship between plants and man has grown (Shakya, 2016) and many plants are used as medicines for example anti-malarial, quinine (1) from bark of *Cinchona* species, anti-bacterial, phellandrene (2) from *Eucalyptus phellandra* and anticancer, taxol (3) from the bark of *yew tree*.

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