

**CHARACTERISATION OF LOCAL AND EXOTIC  
ACCESSIONS OF MORINGA  
(*MORINGA OLEIFERA* LAMARCK)**

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## DECLARATION

This is to certify that this thesis is the result of research undertaken by KENNETH OPARE-OBUOBI towards the award of Master Philosophy degree in the department of Crop Science of the of Agricultural and Consumer Sciences, University of Ghana.



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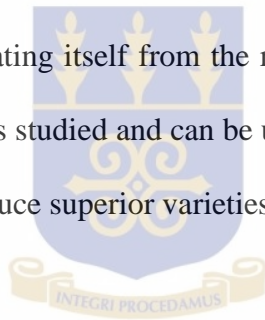
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## ABSTRACT

Knowledge of existing diversity especially at the genotypic level in plant population is crucial for the effective breeding, conservation, management, and efficient utilization of plant genetic resources (PGR) (Mondini *et al.*, 2009). However, Moringa (*Moringa oleifera* Lamarck), a common and popular plant growing in almost every part of the country with several uses and benefits still remains underutilized tropical crop in Ghana. This is due to the fact that the crop has either received little or no attention with respect to research and breeding programmes. *Moringa oleifera* seeds were collected from ten provenances with eight from Ghana and one each from United State of America and India. A completely randomised complete block design and modification of AVRDC-GRSU record sheet was used to evaluate morphological characteristics of thirty seven accessions from these provenances. Eight out of twelve traits observed revealed variability among the accessions studied. For quantitative characters studied; tripinnatum leaf length ranged between 25.2cm and 61.6cm, tripinnatum leaf width ranged between 17.8cm and 48.5cm, tripinnatum leaflet length ranged from 10.8cm to 25.9cm and tripinnatum leaflet width was from 5.7cm and 17.6cm. There were also significant differences among the accessions of *Moringa oleifera* for the vegetative features. Dry leaf samples of Moringa from eight provenances with 40 accessions were used for both biochemical and molecular analysis. The dried leaf samples were found to contain 0.0014 - 0.0736% (Fe), 0.0006 – 0.0097% Mn, and 0.0005 – 0.111% Zn. The accessions also contained 40.3 – 79.3 mg/kg (Ca), 19.23 - 48.13 mg/kg (Mg), 87.5 -253.1 mg/kg (Na), 1549 – 2272 mg/kg (K) and 0.179 – 0.438 mg/kg (P). There were significant differences in the nutritional content of the dried Moringa leaf samples except for potassium (K). The heavy metal Pb ranged from 0.0005 -0.0058%. Percentage dry matter, crude protein and nitrogen

ranged from 14.63 – 25.37%, 12.9 – 26.3% and 2.064 – 4.208% respectively. Eleven distinct bands with a monomorphic locus at a distance of 1.2cm for the forty accessions were observed in gel electrophoresis of total protein from the accessions. The bands ranged from a length of 0.1cm to 3.9cm with two accessions, E40 and E41 producing all eleven bands. Forty genotypes from different ecological zones were assayed for polymorphism using twelve decamer RAPD markers. Ten primers amplified and showed distinct DNA fragments. A high polymorphic level was recorded with 95.1% of the fragments produced being polymorphic (4.9% monomorphic). Cluster analysis showed a separation of BNR 4 accession from the rest. Accessions E41 and UE49 could not be resolved by the RAPD marker. Cluster analysis of the combined data of total protein and RAPD produced two major clusters with accession BNR4 separating itself from the rest. In conclusion, enough variation existed among the accessions studied and can be used in the establishment of Moringa breeding programme to produce superior varieties.



## DEDICATION

There are millions of excuses for not taking action, but to achieve, you have to live above all excuses. I dedicate this work to God, Almighty whose guidance and protection has brought me this far. I also dedicate this work to the fond memories of my beloved sister DIANA AWO AYIBEA OPARE-OBUIBI, (late).



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## LIST OF ABBREVIATIONS, ACRONYMS AND UNITS

### Abbreviations and Acronyms

|            |  |
|------------|--|
| Acc.       | Accession  |
| ANOVA      | Analysis of Variance   |
| ARS        | Agricultural Research Station  |
| AVRDC-GRSU | Asian Vegetable Research and Development Centre - Genetic Resource and Seed Unit |
| BNARI      | Biotechnology and Nuclear Agriculture Research Institute                         |
| CSRPM      | Centre for Scientific Research into Plant Medicine                               |
| FAO        | Food and Agriculture Organisation  |
| FDB        | Food and Drugs Board   |
| Fig        | Figure   |
| GSD        | Ghana Standards Board  |
| KNUST      | Kwame Nkrumah University of Science and Technology                               |
| NTsys      | Numerical Taxonomy System Programme  |
| RDI        | Recommended Daily Intake   |
| SARI       | Savanna Agriculture Research Institute   |
| SAHN       | Sequential, Agglomerative, Hierarchical and Nested                               |
| SRI        | Soil Research Institute  |
| UPGMA      | Unweighted Pair Group Method with Arithmetic Averages                            |
| USDA       | United States Department of Agriculture  |
| WHO        | World Health Organisation  |

**Units**

|     |                        |
|-----|------------------------|
| %   | Percentage             |
| °C  | Degree celcius         |
| cm  | Centimetre             |
| g   | Gram                   |
| mg  | Milligram              |
| kg  | Kilogram               |
| ppm | Parts per million      |
| TM  | Trade mark             |
| mm  | Millimetre             |
| ml  | Millilitre             |
| rpm | Revolutions per minute |
| *g  | Times gravity          |
| μl  | Micro litre            |
| μg  | Microgram              |

## CHAPTER ONE

### INTRODUCTION

Moringa is the monogeneric name given to the species of trees and shrubs belonging to the family *Moringaceae* Martinov of the order *Capparales* Viales (Oslon, 2001; Hawthorne and Jongking, 2006). Of the thirteen species, *M. oleifera* is the most widely utilized and well-known species and is indigenous to Sub-Himalayan regions of northwest India. It is also the most researched species of the family and has been subjected to selection through breeding methods to produce new varieties in its country of origin. It is the most common species cultivated throughout the tropics, fast growing and not very demanding with regard to climate and soil quality (Muluvi *et al.*, 1999; Tsaknis *et al.*, 1999; Vlahov *et al.*, 2002; Manzoor *et al.*, 2007). *M. stenopetala*, an African species native to Ethiopia and Northern Kenya (Lalas *et al.*, 2003), is also widely grown, but to a much lesser extent than *M. oleifera*. *M. oleifera* is known by several common names, usually dependent on the country and tribe but in English it is commonly referred to as “Horseradish tree” (due to the flavour of its root), “Drumstick tree”, “Never Die tree”, “West Indian Ben tree”, or “Radish tree” (Ramachandran *et al.*, 1980). In Ghana, the species is called by names such as Yevuti, Kpokpoti, or Yevutsi by the Ewe community (Irwine, 1969), Zingeridende in Hausa, Obnukuo, Ornyyukuo, or Zangala in Dagari community, Kpokpotsor in Damgbe, Nasadua in Krusaal and Bimoba in Gambadua (Wormadey, 2007).

*Moringa oleifera* is a fast growing plant with a high fruit production rate per acre. The leading producer of Moringa fruits is India; with an annual production rate of 1.1 to 1.3 million tonnes from 38000ha of land (Rajangam *et al.*, 2001). Small scale farmers get a net income of about USD 1500 per hectare from the cultivation of

*Moringa per annum* (Rajangam *et al.*, 2001). In Ghana the plant is grown mainly for its leaves, which are usually consumed as beverage either fresh or dried. They are also grown and used as living fence posts to demarcate lands. There is little or no information on large scale production or income generated by producers, processors and retailers of *Moringa* products in Ghana, though significant amount of revenue is obtained from these operations.

The plant is an important food commodity which serves as a natural nourishment for the people of the tropics and sub-tropics. It is rich in both macronutrients and micronutrients which are essential for human nutrition for the daily development of the human body (Anwar *et al.*, 2007). The leaves and fruits of *Moringa oleifera* are commonly eaten as vegetables mostly in the eastern part of Africa. Nutritional composition of these parts compares satisfactorily with other crops such as beans, cowpea, turnip, cassava leaves, amaranthus leaves and pumpkin leaves and even essential nutrients from other non-plant sources (CSIR, 1962; Palmer and Pitman, 1972; Maroyi, 2006). The essential amino acids, including the sulfur-containing amino acids in the *Moringa* leaves are higher (Makkar and Becker, 1997) than in foodstuff recommended by the Food and Agriculture Organization (FAO) (WHO, 1985), and with concentrations similar to those of soybean seeds.

The plant, for centuries has been used as medicine for many ailments for millions of people especially those of the tropics and subtropics (Caceres *et al.*, 1991; Jahn, 1991). Virtually every part of the plant including the flowers, seed, roots and bark has some pharmacologically active compounds useful for the treatment of diseases such as nervous debility, catarrhal infections, rheumatism, intestinal worms, ascites,

dyspepsia, venomous bites, sore throat, diuretics, skin disease, and cures for scurvy and various bladder and prostate ailments (Irvine, 1961; von Maydell, 1986; Duke, 1987; Morton, 1991). The plant can also be used as an antiseptic, cardiac and circulatory stimulant (ECHO, 2008).

For years, the seeds of *Moringa oleifera* have been shown to be one of the most effective primary coagulants for water treatment especially in rural communities (Folkard *et al.*, 1993; Doer, 2005; Onwuliri and Dawang, 2006). This is better because plant extracts possessing both coagulating and antimicrobial properties are safer for human health when used in water treatment, rather than the commonly used inorganic coagulant (alum) (Muyibi and Okuofu, 1995; Okuda *et al.*, 2001; Ali *et al.*, 2004; Akinnibosun *et al.*, 2008 and 2009) which has the tendency to induce Alzheimer's disease and has strong carcinogenic properties because of the presence of aluminium in the compound (Crapper *et al.*, 1973; Malleavialle *et al.*, 1984; Najm *et al.*, 1998). The plant's usefulness as a non-food product has also been extensively described, including being used as lumber, fencing, charcoal and as lubricating oil (Oliveiera *et al.*, 1999; Ghasi *et al.*, 2000; Kalogo and Verstraete, 2000; Saleem and Meinwald, 2000; Jahn 2001). The biomass of the plant can also be used as soil amendment material to improve poor soils as well as plant growth enhancer (Foidl *et al.*, 2001).

The late Major Courage Quashigah (former health minister of Ghana) knowing the numerous benefits of the plant called for appropriate modalities to be fashioned out into the research, processing and packaging of the Moringa plant for use as an alternative to imported vitamin and nutrient supplements administered in health

facilities (Ghanaweb, 2007). The Moringa Association of Ghana with backing from the Centre for Development of Enterprises and GSB has now completed a food certification process, inspection manual as well as code of practice documents for Moringa leaf and products (GSB, 2009a, b and c). Professor Agyeman Badu Akosa, former Director General of Ghana Health Service, also believed that if Ghanaians had good drinking water; immunized their children under five years, with Moringa taking care of the nutrition problems, the country's health burden could be lessened (Ghanaweb, 2007). The FDB of Ghana, at present recognises and accepts the use of Moringa products as food just like the Royale Cocoa Powder beverage because of the nutritional and medicinal benefits (Sumalia, 2008, personal communication).

In spite of the numerous benefits and acceptance of *M. Oleifera* by both the Ghanaian public and state institutions like GSA and the FDB, the plant has received little attention with regard to breeding programmes to produce varieties with superior agro-morphological traits. For an efficient breeding programme, there is the need to characterise and estimate the genetic relationship that exist among the local accessions via the use of biochemical and molecular markers so that elite accessions that show genetic divergence may be crossed to generate the necessary variability from which improved varieties with better morpho-agronomic characters can be developed.

Biochemical and molecular markers have been recognised as useful markers for characterisation of germplasm with higher efficiency of distinguishing among accessions than using morpho-agronomic traits (Krapovickas, 1973; Smith and Smith, 1989). These markers have proven to be reliable method for fingerprinting accessions

in germplasm collection (Liu and Furniers, 1993; Bachmann, 1994). Biochemical markers such as total protein which refers to the protein accumulated in the vegetative parts of a crop or plant has helped differentiate between accessions (Powell, 1992). These biochemical markers are among the quickest and cheapest marker systems in use. They have been used to study large numbers of samples and remain an excellent choice for projects that only need to identify low levels of genetic variation. Molecular markers further extend and compliment characterisation based on morphological or biochemical description, providing more accurate and detailed information than classical phenotypic data (Karp *et al.*, 1997).

Despite its wide distribution and fast growing the plant still, to a large extent, remains an underexploited crop or under-utilised species (NUS) in the world and specifically in Ghana (PROTA Foundation, 2008; Oliver-Bever, 2009). For maximum utilisation of the crop, proper characterisation especially at the biochemical and molecular level is very important for the identification of accessions with superior agromorphological traits. The gene pool of local accessions of the crop has been shaped and maintained mainly through natural selection. This is because accessions in the country have not been subjected to any form of selection at the genotypic level for use in breeding programmes especially when wide variation is observed in morphological characters like fruit shape, bark stem colour, petiole pigmentation of the leaves and growth habit of the plant (Owusu-Ansah, 2010). Also, in Ghana, not much research on characterisation of Moringa accessions via the use of either biochemical or molecular markers has been documented. Following the potential increase in the use of Moringa products resulting from the recent publicising of the nutritional and medicinal benefits of the plant, there is therefore the need to identify

and characterise the various accessions existing locally as well as investigate the presence or absence of harmful chemicals in the crop using the technologies that are available such as AAS.

The objectives of the study were therefore to;

1. Characterise the accessions of Moringa in Ghana based on their morphological features and biochemical characteristics.
2. Determine the level of variability and the relationship between the accessions.
3. Analyse the biochemical characteristic of the Moringa leaves and determine the lead (Pb) levels in the leaves.

## CHAPTER TWO

### LITERATURE REVIEW

#### 2.1 General Botany

*Moringa* (Adans) belongs to the order *Capparales* and is the sole genus of family of shrubs and trees called *Moringaceae* (Ramachandran *et al.*, 1980; Jahn *et al.*, 1986; Olson, 2002). *M. oleifera* the most researched of all the species is a diploid ( $2n = 28$ ) and the mating pattern is estimated to be 26% selfing and 74% outcrossing (Muluvi *et al.*, 2004). Pollination of the plant is usually by bees and other insects, as well as by birds (World Agroforestry Centre database; Morton, 1991; Jyothi *et al.*, 2004).

*Moringa* grows well in the humid tropics or hot dry lands, survives destitute soils, and has the ability to tolerate drought (Morton, 1991; Mahmood *et al.*, 2010). It also grows under wide range of rainfall with minimum annual rainfall requirements estimated at 250 mm and maximum at over 3000 mm and a pH range of 5.0–9.0 (Palada and Changl, 2003). The plant is a fast growing perennial tree which can reach a maximum height of 7-12 m and a diameter of 0.2-0.4 m at chest height with the stem usually straight but occasionally poorly formed (Fahey, 2005; Mahmood *et al.*, 2010). It reaches a height of 1.5-2 m before it begins branching but can reach up to 3.0 m (Foidl *et al.*, 2001). Its tripinnate compound leaves are feathery with green to dark green elliptical leaflets. Conspicuous, lightly fragrant flowers are borne on inflorescences 10–25 cm long, and are generally white to cream coloured, although they can be tinged with pink in some varieties. The fully grown dehiscent capsules, mature about 3 months after flowering, and remain on the tree for several months, releasing the seeds that are dispersed mainly by wind and water, but probably also by

seed-eating animals. Seeds do not retain their viability in storage at ambient temperatures for longer than 2 months (Sharma, 1982) germination percentages of 60, 48 and 7.5 % were reported for seeds after 1, 2, and 3 months respectively in India (Morton, 1991). The fruits are tri-lobed capsules, and are frequently referred to as “pods.” Immature pods are green and in some varieties have some reddish colour. The ripe fruit turns brown and on the average contains 10–12 seeds (Anwar and Bhangar 2003). Seeds are large with three papery wings. Seed hulls are generally brown to black, but can be white if kernels are of low viability. Virtually every part of the plant is useful with the leaves, flowers, immature pods (which are called long green pods), and roots being edible for both man and farm animals respectively (Foidl *et al.*, 2001; Reyes, 2006).

Propagation of *Moringa oleifera* is either by seeds or cuttings with fruit production beginning as early as 6 to 8 months when planting is done from stem and branch cuttings (Ramachandran *et al.*, 1980). Plants raised from seeds, are reportedly slower to flower with inferior quality of fruit produced (Ramachandran *et al.*, 1980). Fruit yields in the first 2 years are generally low, but from the third year onward, a single tree can yield between 600 and 1,600 fruits each year (Ramachandran *et al.*, 1980; Booth *et al.*, 1988; Morton, 1991).

## **2.2 Origin, Domestication and Distribution**

*Moringa oleifera* is a deciduous tree which is truly wild only in the western Himalayas and the eastern Punjab (Dastur, 1951). It was cultivated in India and distributed throughout the tropical Asia and Oceania in prehistoric times (Irvine, 1961). The tree was then introduced to other parts of the world mostly by

missionaries and voyagers. In the case of Netherlands New Guinea (West Papua), Indonesians took the plant there (Massal and Barrau, 1956) whilst the British brought it to Africa as an ornamental plant from India (Berger *et al.*, 1984). The British as well sent the plant to Jamaica in 1784 (Fawcett and Rendle, 1914). It is currently cultivated in several parts of the world including most countries in the Caribbean islands West Indies, Bahamas (Britton and Millspaugh, 1920) and Bermuda (Britton, 1918). In Latin America, the plant is also grown particularly in Cuba, Haiti, Jamaica, Puerto Rico, all of the Virgin Islands, Barbados, Trinidad, the Netherlands Antilles, Panama, El Salvador, Honduras, Costa Rica, Belize, Mexico, Brazil, and Venezuela (Arnoldo- Broeders, 1967, 1971; Adams, 1972; Correll and Correll, 1982; Proctor, 1984). Moringa was introduced to the United State of America by the Office of Foreign Seed and Plant Introduction of the United States Department of Agriculture from Cuba in 1915 (*Bur. Pl. Indu*, 1915)

*M. oleifera* is now cultivated and has become naturalized in other parts of the world especially in countries located in the tropical and subtropical region. These include countries like Pakistan, India, and Nepal, as well as in Afghanistan, Bangladesh, Sri Lanka, Southeast Asia, West Asia, the Arabian peninsula, East and West Africa, throughout the West Indies and southern Florida, in Central and South America from Mexico to Peru, as well as in Brazil and Paraguay (Ramachandran *et al.*, 1980; Jahn *et al.*, 1986; Lahjie and Siebert, 1987; Vivien, 1990; Francis and Liogier, 1991).

## **2.3 Species of Moringa and their distribution**

According to Olson (2002), the thirteen species of Moringa fall into three groups that reflect life form and geography, namely the bottle trees, slender tree and trees, shrubs and herbs of north-east Africa.

### **2.3.1 Bottle trees**

These species of Moringa are huge trees with bloated water-storing trunks and small radially symmetrical flowers. Three out of the thirteen species belong to this group and occur in the southern hemisphere. Namibia and Angola in south-western Africa are home to *M. ovalifolia*, while *M. drouhardii* and *M. hildebrandtii* are endemic to Madagascar (Olson, 2002).

### **2.3.2 Slender trees**

This group of Moringa trees have a tuberous juvenile stage and cream to pink slightly bilaterally symmetrical flowers. The group has three species that occur in the red sea area, Arabia, and the Indian subcontinent. The group also includes the family's best-known and most economically valuable species, *M. oleifera* which is now cultivated in all the countries of the tropics. *M. oleifera* and *M. concanensis* are mainly from India and Pakistan, barely reaching Bangladesh. *M. peregrina* the third species in the group has the widest range of all, growing from the Dead Sea area sporadically along the Red Sea coasts to northern Somalia and around the Arabian Peninsula to the mouth of the Persian Gulf which seems to be native to sub-Himalayan India.

### 2.3.3 Trees, shrubs and herbs of north-east Africa

The group consists of eight *Moringa* species found in northeast Africa. All but *M. peregrina* are endemic to northeast Africa. These species are tuberous adults or tuberous juveniles maturing to fleshy-rooted adults; colourful, bilaterally symmetrical flowers. The eight species in addition to *M. ovalifolia* occur in a geographical location termed as the Horn of Africa. The Horn of Africa is the centre of *Moringa* diversity and includes a variety of life forms. *M. peregrina* is found in northern Somalia, Arabia, and the Red Sea coasts north to the Dead Sea. The remaining seven species occur in North-Eastern Africa mainly in Kenya, Somalia and Ethiopia and include *M. arborea*, *M. borziana*, *M. longituba*, *M. pygmaea*, *M. rivaie*, *M. ruspoliana* and *M. stenopetala*. The densest concentration of *Moringa* species is in Mandera District, in the extreme northeast of Kenya, where *M. arborea*, *M. longituba*, *M. rivaie*, and *M. ruspoliana* can be found, though the species are spatially isolated from one another.

## 2.4 Uses of Moringa

### 2.4.1 Moringa as food

*Moringa oleifera* can be found in the wild but domestically it is cultivated throughout the plains, especially as hedges and in house yards, thrives best under the tropical insular climate, and is plentiful near the sandy beds of rivers and streams (The Wealth of India, 1962; Qaiser, 1973). The plant is an essential food commodity which has had myriad attention as the ‘natural nutrition of the tropics’. Parts of the plant such as the leaves, fruits, flowers and immatured pods, also called long green pods, are used in many countries especially in India, Pakistan, Philippines, Hawaii and many parts of

Africa as a highly nutritive vegetable and nutritional supplementation (D'souza and Klukarni, 1993; Anwar and Bhanger, 2003; Anwar *et al.*, 2005). The tender leaves taste like watercress and, along with the flowers, is eaten cooked or raw. The green pods with seeds have a similar flavour to that of asparagus when cooked. The seeds from ripe fruits have a peanut flavour when fried. In certain parts of the world, these parts are eaten as fresh vegetables and can be frozen or canned. Combined with curries, the parts of the plant are commonly prepared with chicken or sea food as a soup (Anwar and Bhanger 2003).

#### **2.4.2 Moringa as animal feed and plant growth enhancer**

Moringa leaves are also fed to livestock as well as pigs, chickens and rabbits and can be fed to fishes as food. A study in Fiji reports significant weight gain over traditional fodder when 50% of the fodder contained Moringa (Aregheore, 2002). Olugbemi (2010) reported that the inclusion of *Moringa oleifera* in cassava based broiler diets up to 5% is possible without negatively affecting productivity or haematological indices. Foidl and Reyes reported that adding Moringa leaves to cattle feed increased their daily weight gain by 32% (Trees for Life, 2005). The milk production of cattle increased by 43% after being supplemented with 15-17 kg of fresh Moringa leaves and milk production increased further by 58% and 65% when cows were fed with 2kg and 3kg of dry matter of Moringa as supplement per day respectively (Trees for Life, 2005). The adoption of Moringa as animal feed especially in third world nations will greatly improve the weight of animals as well as the production of animal milk which can boost the dietary protein supply in the population. Juice extract from green matter of the plant has been tested and is effective as a plant growth enhancer in the production of crops like soybeans, corn, turnips, black beans, red beans, white beans,

cowpeas, bell peppers, chia, sunflowers, mung beans, onions, coffee, tea, chilli peppers, melons and sorghum (Fahey 2005). Crops sprayed with green matter juice extract of the plant comparatively resulted in increased crop size. The juice extract was discovered to stimulate the growth of the plant's roots enhancing the uptake of soil nutrient as demonstrated in sugarcane production (Trees for Life, 2005). Leaf extracts have been found to increase *Rhizobium* root nodulation, nodule weight, and nitrogenase activity in mung bean (*Vigna mungo* (L.) HEPPEL) when applied to seeds or as a root dressing (Bandana *et al.*, 1987).

#### **2.4.3 Medicinal uses of Moringa**

Myriad of medicinal properties have been attributed to parts of the plant. The high quality protein of the leaves has led to its widespread use by doctors, healers, nutritionists and community leaders, to treat malnutrition and a variety of illnesses. The seeds of *Moringa* are considered to be antipyretic, acrid, bitter (Oliveira *et al.*, 1999) and reported to show antimicrobial activity (The Wealth of India, 1962). The plant also contains numerous phytochemicals, some of which are of high interest due to their medicinal properties. Current studies demonstrate that isothiocyanates have antitumor activity in cancers of the lung, breast, skin, oesophagus, and pancreas (Kalkunte *et al.*, 2006; Satyan *et al.*, 2006) and this particular plant family is rich in a fairly unique group of glycoside compounds called glucosinolates and isothiocyanates. Small proteins/ peptides were isolated from the leaves of *Moringa oleifera* possessing antifungal and antibacterial activity (Dahot, 1998). Methanolic extract from the roots has moringine and moringinine which have been reported to possess analgesic and anticonvulsive properties (Grupe *et al.*, 1999). Below are some medicinal uses of the various parts of the Moringa plant.

**Table 2.1:** Tissues of Moringa and their medicinal uses

| Plant Part | Medicinal Uses   |
|------------|--|
| Root       | Antilithic, rubefacient, vesicant, carminative, antifertility, anti-inflammatory, stimulant in paralytic afflictions; act as a cardiac/circulatory tonic, used as a laxative, abortifacient, treating rheumatism, inflammations, articular pains, lower back or kidney pain and constipation (The Wealth of India, 1962; Dahot, 1988; Padmarao <i>et al.</i> , 1996; Ruckmani <i>et al.</i> , 1998; Anwar <i>et al.</i> , 2007)  |
| Leaves     | Purgative, applied as poultice to sores, rubbed on the temples for headaches, used for piles, fevers, sore throat, bronchitis, eye and ear infections, scurvy and catarrh; leaf juice is believed to control glucose levels, applied to reduce glandular swelling. It also acts as an antiulcer and antihyperlipidemic agent (The Wealth of India, 1962; Dahot, 1988; Morton, 1991; Pal <i>et al.</i> , 1995; Makonnen <i>et al.</i> , 1997; Ghasi <i>et al.</i> , 2000; Fuglie, 2001; Patel <i>et al.</i> , 2010)                                   |
| Stem Bark  | Rubefacient, vesicant and used to cure eye diseases and for the treatment of delirious patients, prevents enlargement of the spleen and formation of tuberculous glands of the neck, to destroy tumors and to heal ulcers. The juice from the root bark is put into ears to relieve earaches and also placed in a tooth cavity as a pain killer, and has anti-tubercular activity (Bhatnagar <i>et al.</i> , 1961; Pal <i>et al.</i> , 1995; Siddhuraju and Becker, 2003; Anwar <i>et al.</i> , 2007)  |
| Gum        | Used for dental caries, and is astringent and rubefacient; Gum, mixed with sesame oil, is used to relieve headaches, fevers, intestinal complaints, dysentery, asthma and sometimes used as an abortifacient, and to treat syphilis and rheumatism (Fuglie, 2001).   |
| Flowers    | High medicinal value as a stimulant, aphrodisiac, abortifacient, cholagogue; used to cure inflammations, muscle diseases, hysteria, tumors, and enlargement of the spleen; lower the serum cholesterol, phospholipid, triglyceride, VLDL, LDL cholesterol to phospholipid ratio and atherogenic index; decrease lipid profile of liver, heart and aorta in hypercholesterolaemic rabbits and increased the excretion of faecal cholesterol (Bhattacharya <i>et al.</i> , 1982; Dahot, 1998; Mehta <i>et al.</i> , 2003; Siddhuraju and Becker, 2003) |
| Seed       | Seed extract exerts its protective effect by decreasing liver lipid peroxides, antihypertensive compounds thiocarbamate and isothiocyanate glycosids have been isolated from the acetate phase of the ethanolic extract of Moringa pods. The seeds also possesses anti-inflammatory, antioxidant, antimicrobial and anticancer properties (Caceres <i>et al.</i> , 1992; Faizi <i>et al.</i> , 1998; Lalas and Tsaknis, 2002; Patel <i>et al.</i> , 2010)  |

#### **2.4.4 Moringa as oil**

The seeds contain 35-40% non-drying oil, and the remaining seed cake after extraction is reported to be very high in crude protein (nearly 60%), making it a desirable source of animal fodder (Duke, 1983). Known commercially as “ben oil”, its oil qualities are similar to that of olive oil and are rich in palmitic, stearic, behmic, and oleic acids (Verma *et al.*, 1976; Nautiyal and Venhataraman, 1987) and is used for human consumption, and in cosmetics and soaps (Delaveau and Boiteau, 1980; Ramachandran *et al.*, 1980; Szolnokim, 1985). The degummed oil has favourable characteristics for use as a replacement for petroleum diesel (Tsaknis, *et al.*, 1999). The oil is highly valued by perfumers for its power of absorbing and retaining odours, and by watchmakers as a lubricant (Ramachandran *et al.*, 1980)

#### **2.4.5 Moringa seeds as water purifying agent**

*Moringa* seeds are one of the best natural coagulants discovered so far (Ndabigengesere and Narasiah, 1998) and crushed seeds are a viable replacement of synthetic coagulants (Kalogo *et al.*, 2000). The seeds are effective for high turbidity water and show similar coagulation effects to alum (Muyibi and Evison, 1995b). Muyibi and Evison, (1995b) also reported that coagulation effectiveness of *Moringa* seeds could reduce turbidity by between 92% and 99% though its effectiveness depended on the initial turbidity of the water. Rural women in Sudan, use *Moringa* seed extract to treat the high turbid Nile water instead of alum due to fear of alum causing gastrointestinal disturbances and Alzheimer’s disease (Crapper *et al.*, 1973; Miller *et al.*, 1984; Martyn *et al.*, 1989; Muyibi, 1994). *Moringa* seeds also possess antimicrobial properties (Madsen *et al.*, 1987; Olsen, 1987) making it possible for it to be used as an antiseptic in the treatment of drinking water (Obioma and Adikwu,

1997). Broin *et al.*, (2002) reported that a recombinant protein in the seed is able to flocculate Gram-positive and Gram-negative bacteria cells, giving the seed of the plant an anti-microbial activity. *Moringa* seeds could be used as a less expensive biosorbent for the removal of cadmium (Cd) from aqueous media (Sharma *et al.*, 2006).

#### **2.4.6 Other uses**

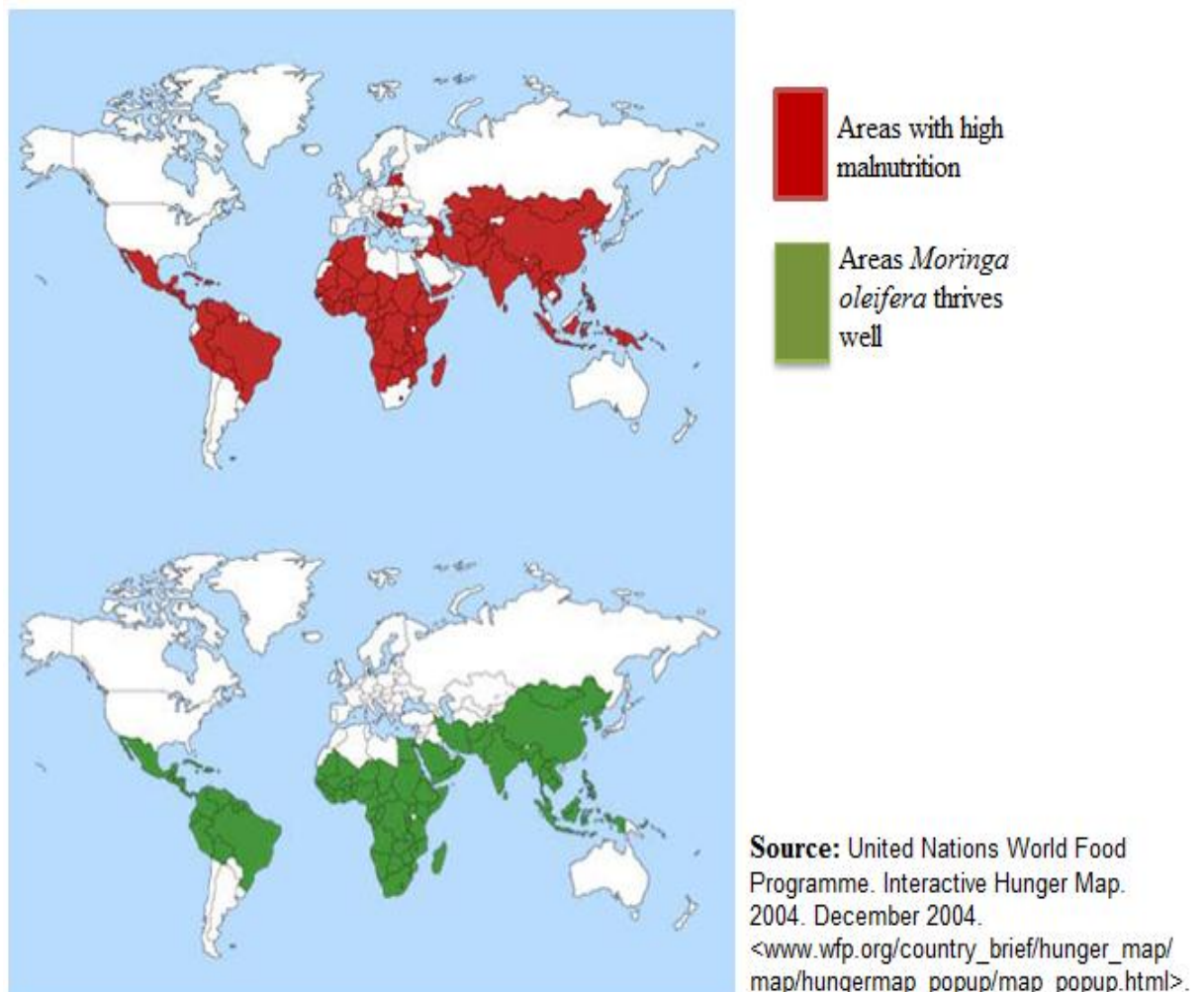
*Moringa* plants are used in alley cropping because it has a large tap root and few lateral roots, it does not compete for nutrients in the subsoil with other crops (ECHO, 2008). It also adds to the nutrients available as it produces many protein rich leaves. They grow very quickly but do not provide too much shade due to the structure of their leaves (ECHO, 2008). They are also very good for reclaiming marginal land. The plant is used in the textile industry for shuttles and picking-sticks and is also suitable for pulp production for newsprint (Singh *et al.*, 1983; Guha *et al.*, 1968), cellophane, and textiles (Nautiyal and Venhataraman, 1987; Mahajan and Sharma, 1984). The corky bark yields a coarse fibre, which is utilized in making mats, paper, and cordage. The stem exudes a mucilaginous gum that is used in leather tanning and calico printing (Ramachandran *et al.*, 1980; Nautiyal and Venhataraman, 1987). The plant is also used as a natural herbicide. This is usually done by digging *Moringa* leaves into the soil before planting; damping off disease (*Pythium debaryanum*) can be prevented among seedlings. In some parts of Nigeria, crushed leaves of the plant are used as domestic cleaning agent for cleaning cooking utensils or even walls (Fuglie, 2001).

## 2.5 Nutritional content and benefits

The leaves of Moringa have been reported to contain high amount of  $\beta$ -carotene, protein, vitamin C, calcium and potassium and serve as a good source of natural antioxidant and thus enhance the shelf-life of fat containing foods due to the presence of various types of antioxidant compounds such as ascorbic acid, flavonoids, phenolics and carotenoids (Dillard and German, 2000; Siddhuraju and Becker, 2003). “Gram for gram, fresh leaves of *M. oleifera* have four times the vitamin A of carrots, seven times the vitamin C of oranges, four times the calcium of milk, three times the potassium of bananas, three quarters the iron of spinach, and two times the protein of yogurt” (Trees for Life, 2005). The plant is referred to as “mother’s best friend” in the Philippines simply because of its ability to increase a nursing mother’s milk production when consumed (Estrella *et al.*, 2000) and is at times prescribed for anaemia (Siddhuraju and Becker, 2003). Fuglie (2005) reported that serving 8g of dried Moringa leaf powder will satisfy a child within ages 1-3 with 14% of the protein, 40% of the calcium, 23% of the iron, and nearly all the vitamin A that the child needs in a day.

The numerous nutritional benefits of the plant complemented by the ability of the plant to thrive in most areas in the tropics and subtropics gives it the potential of curbing the malnutrition and hunger problems of the world. The areas with high malnutrition of the world have the plant thriving there as shown in figure 2.1. Health workers in countries like Senegal and Haiti have adopted the use of Moringa powder to combat malnutrition especially in small children, pregnant and nursing mothers (Price, 1985a). Zija™, a commercially-promoted *Moringa oleifera* containing product sold in the United States of America is an important nutritional supplement

with several medicinal properties and contains low calories, quality protein dietary supplementation (containing 19 of the 20 most common amino acids), low carbohydrate, lower-fat alternative to common liquid dietary supplements, and may serve to reverse the effects of many nutrient-depleting prescription and non-prescription drugs (Johnson, 2005).



**Fig 2.1:** Map showing areas of malnutrition and areas where the *Moringa oleifera* plant grows

## **2.6 Characterisation and Diversity**

Characterisation of biological organisms deals with identifying or accessing the variations that exist between the organisms either at the phenotypic or genotypic level (Mondini *et al.*, 2009). Efficiently managed and effectively utilized germplasm collection must first and foremost be well-characterized. Historically, characterization of germplasm collections has been carried out at several levels, from descriptions of taxonomy (Hilu, 1989), biogeography (Lyman, 1984; Nabhan, 1985; Brush, 1989; Zimmerer *et al.*, 1991), and morphology and agronomic characters (Chapman, 1989) to biochemical analysis (Doebley, 1989; Gepts, 1990) and the study of molecular traits and markers (Clegg, 1990; Gepts, 1995). Diversity originally was assessed by measuring variations that occur at the phenotypic traits (i.e. morphological variation) of crop plants such as growth habit, flower colour, or quantitative agronomic traits like yield potential but soon became inadequate because they are subjected to environmental variations. Due to these limitations, molecular tools were then developed to measure the variations that occur at the genetic level between species, accessions or varieties. The assessment of genetic diversity within and between populations is routinely performed at the molecular level using various laboratory-based techniques such as allozyme, total protein or DNA based analysis, which measure levels of variation directly. Genetic diversity may also be estimated using morphological, and biochemical characterization and evaluation.

### **2.6.1 Importance of Characterisation**

The assessment of genetic variability and relatedness in any germplasm collection is very vital for effective utilisation of the germplasm for breeding programmes, identification of conservation priorities (Bekele, 1983; Demissie and Bojórnsland,

1996) and/or elimination of duplicates in the gene stock and establishment of core collections (Nisar *et al.*, 2007). A practical application of the knowledge of genetic diversity is its use in sorting of populations for genome mapping experiments (Kaga *et al.*, 1996). Diversity studies especially at the molecular level of cultivated plants and their wild relatives bring important information for establishment of appropriate breeding strategies, especially when interspecific crosses are necessary for mapping purposes or for the incorporation of new features (Benko-Iseppon, 2001).

### **2.6.2 Morphological characterisation**

This form of characterisation deals with variations that exist at the phenotypic level, in other words visually observed characters. It often does not require expensive technology but large tracts of land are frequently required for these experiments, making it sometimes more expensive than molecular assessment. The traits measured often show phenotypic plasticity; this allows assessment of diversity in the presence of environmental variation. Oslon (2002) in a study on both morphological features such as leaf form, leaf glands life form, woody anatomy, gum duct (articulated laticifers), flowers and fruit and testa anatomy; molecular data on chloroplast genome (cpDNA), gene sequence of ribulose -1-5-2 biphosphate carboxylase / oxygenase (rbcL) revealed that Caricaceae and Moringaceae are sister taxa. Resmi *et al.*, (2005) observed variability among drumstick (*Moringa oleifera* Lam.) accessions from central and southern Kerala based on the observation that accessions from Thiruvananthapuram (MO 13, 24 and 26) had three distinct flowering peaks in a year, while others showed one or two peaks. Moringa also exhibits considerable variability in weight of cotyledons, seed size or yield, and phenology, and especially in the length, appearance, and quality of pods (Duke 1987; Jahn 1989; Morton 1991).

### **2.6.3 Biochemical characterisation**

Biochemical analysis is based on the separation of proteins into specific banding patterns through the use of biochemical markers (Mondini *et al.*, 2009). The use of biochemical markers involves the analysis of total protein, seed storage proteins and isozymes. The isozymes technique uses enzymatic functions and is comparatively inexpensive yet powerful method of measuring allele frequencies for specific genes. It is a fast method which requires only small amounts of biological material. However, only a limited number of enzymes are available and thus, the resolution of diversity is limited.

Characterization of germplasm using biochemical fingerprinting has got special attention due to its increased use in crop improvement and the selection of desirable genotypes for breeding crops (Mondini *et al.*, 2009). Genetic markers and protein profiling are also successfully used to resolve the taxonomic and evolutionary problems of several crop plants (Khan, 1990; Murphy *et al.*, 1990; Nakajima, 1994; Das and Mukarjee, 1995; Ghafoor *et al.*, 2002) It is also known that variation in protein bands elaborate the relationship among the collection from various geographical regions (Satija, 2002; Asghar *et al.*, 2003; Ghafoor *et al.*, 2003).

### **2.6.4 Molecular characterisation**

Molecular form of diversity is based on a large variety of DNA molecular markers, which can be employed for analysis of variation. Different markers have different genetic qualities. They may be dominant or co-dominant can amplify anonymous or well characterized loci, can contain expressed or non-expressed sequences, etc. Molecular markers work by highlighting differences (polymorphisms) within a

nucleic sequence between different individuals. These differences include insertions, deletions, translocations, duplications and point mutations. However, they cannot determine the activity of specific genes in the genome of the organism. Currently, molecular markers complement the use of morphological and biochemical markers and also introduce many new possibilities to increase our understanding of the genetic constitution of plants (Tanksley, 1993; Karp *et al.*, 1997).

The development of new PCR-based molecular markers, such as Randomly Amplified Polymorphic DNA (RAPDs), Simple Sequence Repeats (SSRs), Amplified Fragment Length Polymorphisms (AFLPs) and Single Nucleotide Polymorphism (SNP) has created the opportunity for fine-scale genetic characterization of germplasm collections that were previously impossible. The PCR-based markers generate relatively large amounts of data per unit time (Powell *et al.*, 1996) due to the high polymorphic nature and the ease with which they are processed. The high levels of variation detected with these molecular markers has allowed germplasm managers, plant breeders, and geneticists to pursue with new vigour, important questions relating to crops and their genetic resources (Bretting and Widrlechner, 1995).

#### **2.6.4.1 RAPDs**

Random Amplified Polymorphic DNAs were the first PCR-based molecular markers to be employed in genetic variation analyses (Welsh *et al.*, 1990; Williams *et al.*, 1991). This class of molecular markers are generated through the random amplification of genomic DNA using short primers (decamers or hexamers), separation of the obtained fragments on agarose gel in the presence of ethidium bromide and finally, visualization under ultraviolet light. The use of short primers is

necessary to increase the probability that, although the sequences are random, they are able to find homologous sequences suitable for annealing. DNA polymorphisms are then produced by “rearrangements or deletions at or between oligonucleotide primer binding sites in the genome” (Williams *et al.*, 1991). The main advantages of the RAPD technology are suitability for the work on anonymous genome, application to the problem where only limited amount of DNA is available and its efficiency and cost effectiveness (Hadrys *et al.*, 1992; Tingey and Tufo, 1993; Rabouam *et al.*, 1999). A study of genetic variability using RAPDs in evaluating 75 accessions of *Moringa oleifera* from the Sudan and Guinea savannah zone in Nigeria produced high degree of polymorphism (74% polymorphic) (Abubakar *et al.*, 2011). Beyene, (2005) studied the genetic variation in 19 accessions with 71 individuals of *Moringa stenopetala* germplasm from Ethiopia by using RAPD as genetic marker. He reported a mean diversity range estimate of Shannon’s index from 0.00 for Lasho and Berber to 0.3623 for Konso Gamolle within populations. The entire population was found to have an index of 0.3124. He concluded that there was a low genetic variability among the accessions studied. Muluvi *et al.*, (1999) analysed the genetic variation in *M. oleifera* Lam using four pairs of amplified fragment length polymorphism (AFLP) primers which revealed polymorphism between and within populations and reported significant differences between populations from different regions, even though outcrossing perennial plants are expected to maintain most variation within populations. Muluvi *et al.*, (1999) further suggested that provenance source is an important factor in the conservation and exploitation of *M. oleifera* genetic resources.

## 2.7 Biochemical (Physico-chemical) Analysis

The food we consume is basically needed to provide the body with the minimum requirements of the six groups of nutrients namely carbohydrates, fats, proteins, mineral elements, vitamins and water (Alfred and Patrick, 1985). One food group that contains nutrients which can be absorbed by the body to be used as energy sources, body building, regulatory and protective material is vegetables. Leafy vegetables are a category of vegetables that is widely eaten throughout the world and are comparatively rich in fibre while cereals, root vegetables and other foodstuff are relatively poor sources (Brain and Allan, 1986). The low calories and negligible quantities of utilizable energy of vegetables make them ideal for obese people who can satisfy their appetite without consuming too much carbohydrate (Oke and Ojofeitmi, 1988). Generally vegetables have low amounts of protein content but Moringa leaves are exceptional and there is increasing awareness of the importance of it in maintaining health, particularly in areas where animal protein is scarce (Baker and Griffin, 1967). Pallavi and Dipika, (2010) after assessing the effect of different methods of drying (sun, shade and oven drying) on the nutritive value of the *Moringa oleifera* leaf with its fresh counterparts reported that there was significant increase ( $p < 0.01$ ) in all the nutrients in the dried samples of the leaves making them a concentrated source of nutrients. Shade dried samples had the highest nutrient retention followed by sun drying and oven dried samples but the differences were not statistically significant ( $p > 0.05$ ).

Oduro *et al.*, (2008) evaluated nutritional potential of *M. oleifera* and seven varieties of sweet potato (*Ipomea batatas*) leaves. They reported that the *M. oleifera* leaves contain higher levels of crude protein, crude fibre, iron and calcium making it a very

rich source of dietary nutrient compared to *Ipomea batatas*. Kwenin *et al.*, (2011) also evaluated the nutritional value of some African indigenous green leafy vegetables in Ghana. They concluded that *Moringa oleifera* was high in protein content with value of 6.60% compared to other locally consumed vegetables such as *Xanthosoma sagittifolia* (Kontomire) and *Talinum triangulare* (Bokoboko) with protein content of 4.65% and 5.10% respectively. Fuglie, 2005 also reported that 8g serving of dried leaf powder will satisfy a child within the ages 1-3 with 14% of the protein, 40% of the calcium, 23% of the iron, and nearly all the vitamin A that the child needs in a day. Also a 100g portion of leaves could provide a woman with over a third of her daily need of calcium and give her important quantities of iron, protein, copper, sulphur, and B-vitamins. Reddy *et al.*, (2005) after utilizing extracts of three plant foods, namely amla (gooseberry), drumstick leaves (*M. oleifera*) and raisins as sources of natural antioxidants in the preparation of biscuits, reported that even though all the above three extracts exhibited a high percentage of antioxidant activity compared with the effect of butylatedhydroxyanisole, extracts from drumstick leaves and amla were more effective than raisins in controlling lipid oxidation during storage of biscuits. The leaves of the plant had the richest amount of protein source with an adequate profile of amino acids and ash, while the immature pods show a high content of dietary fibre and low lipid content (Sánchez-Machado *et al.*, 2009). Abdulkarim *et al.*, 2005 also reported high levels of total proteins (383.0 SD=13.0g/kg dry matter) in *Moringa oleifera*, which were greater than the protein level (18-25%) for important leguminous seeds used for human nutrition and nearly double the contents of cereals (Singh and Singh, 1992). The seed lipid content (412.0 SD= 22.2g/kg dry matter) reported by Oliveira *et al.*, (1999) is greater than that of some soybean varieties (149-220g/ kg meal) (Vasconcelos *et al.*, 2001).

## 2.8 Presence of Lead (Pb) in Plants

The toxicity of heavy metals like lead is a problem for ecological, evolutionary and environmental reasons (Nagajyoti *et al.*, 2008). The metal is a highly toxic pollutant as it is introduced into the atmosphere through automobile exhausts (Lagerwerff and Specht, 1970). Dense traffic releases detrimental exhaust gases and toxic pollutants like unburnt and partially burnt hydrocarbons, lead compounds and other elements that are contained in petrol polluting the city environment (Iqbal *et al.*, 2001). In the soil, it may be caused by broken-down lead paint, residues from lead-containing gasoline or pesticides used in the past, contaminated landfills, or from nearby industries such as foundries or smelters (Woolf *et al.*, 2007). Much of the global lead contamination has occurred as a result of mining and iron smelting activities (Huang *et al.*, 1996). According to FAO/WHO standard (Codex Alimentarius Commission, 2001) the safe level of lead in vegetables is 2 mg/kg above which it becomes toxic for human consumption. The soil being a major source of heavy metals, consciousness of the critical contents or limit on safe crop production needs be observed. The following critical levels of heavy metals for arable land have been established; Zn – 300 ppm; Cu – 100 ppm; Cr – 100 ppm; Pb – 100 ppm; Ni – 50 ppm; Cd – 3 ppm and Hg – 2 ppm (Mengel and Kirby 1987). In plants, the critical levels established for heavy metals are Zn (150-200 ppm); Cu (15-20 ppm); Cr (1-2 ppm); Cd (5-10 ppm); Pb (10-20 ppm); Hg (2-5 ppm); Ni (20-30 ppm) and Co (20-30 ppm). Higher contents than these are very likely to be toxic (Sauerbeck 1982).

Exposure of the human body to lead may cause a decrease in lifespan and have health effects in the long term (Needleman, 2004). Death rates from a variety of causes have been found to be higher in people with elevated blood lead levels; these include

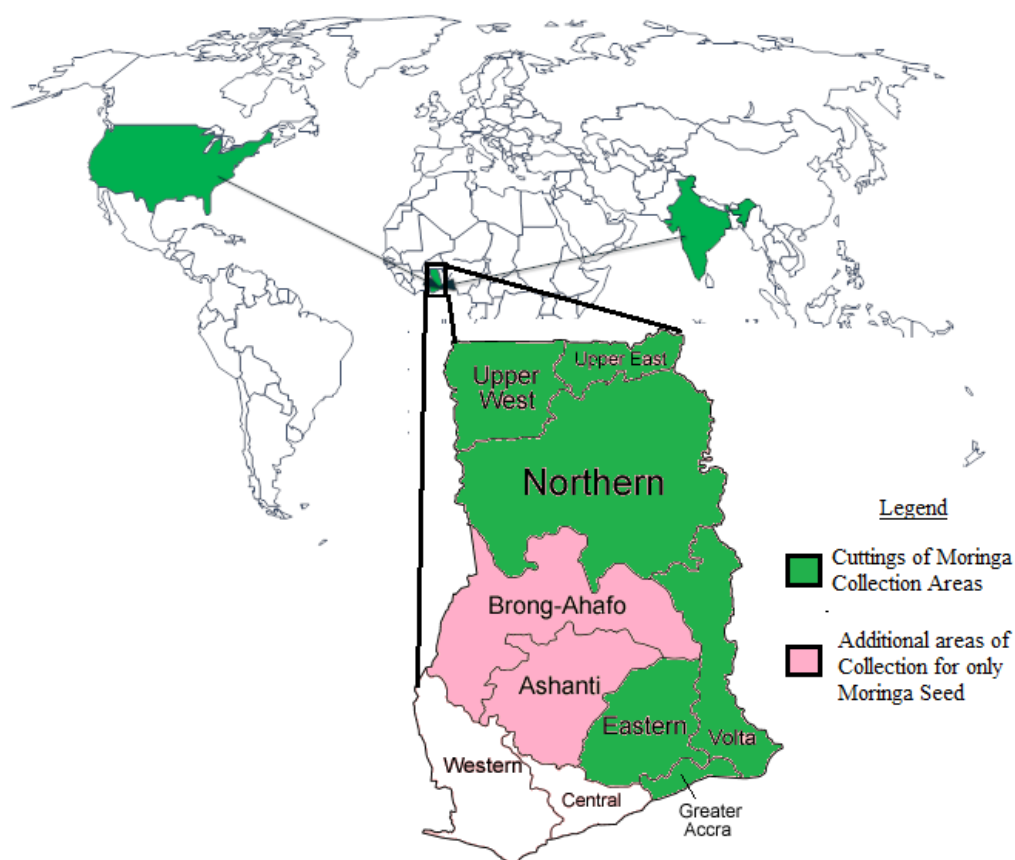
cancer, stroke, and heart disease, and general death rates from all causes (Rossi, 2008). The metal is considered a possible human carcinogen based on evidence from animal studies (Merrill *et al.*, 2007). The metal also has an adverse effect on plants as it inhibit germination and retard plant growth as commonly observed at toxicity level (Lerda, 1992; Wierzbicka and Obidzinsca, 1998; Antosiewicz and Wierzbicka, 1999; Shaukat *et al.*, 1999; Iqbal and Shazia, 2004). The toxicity of lead negatively impacts on seed germination and seedling growth of some tree species (Shafiq and Iqbal, 2005). Studies on concentrations of lead in some leafy vegetables of Makarfi, Nigeria revealed 321 mg/kg of lead in the dry weight of Moringa leaves studied (Garba *et al.*, 2010). This far exceeds FAO/WHO standard (Codex Alimentarius Commission, 2001) the safe level of lead in vegetables. Magat *et al.*, (2009) reported lead level of 90 mg/kg in some Moringa accessions studied in the Philippines giving enough reason that the plant can pick and accumulate lead concentrations like it has been reported in Chinese cabbage (Wong *et al.*, 1996; Chove *et al.*, 2006). Foliar application of lead affected growth and yield of wheat (Rashid and Mukhirji, 1993). High significant effects of lead on shoot, root lengths and seedling was observed in dry biomass of *Lythrum salicaria* (Joseph *et al.*, 2002).

## CHAPTER THREE

### MATERIALS AND METHODS

#### 3.1 Collection of Moringa Accessions

Moringa seeds and cuttings were collected from seven regions of Ghana, namely Greater Accra, Central, Western, Eastern, Ashanti, Volta, Northern, Upper East and Upper West representing the different ecological zones of the country. Fourteen additional collections from the Biotechnology and Nuclear Agriculture Research Institute (BNARI) of Ghana Atomic Energy Commission (GAEC) were included of which two accessions were collected from United States of America (USA) and India as shown in figure 3.1.



**Fig 3.1:** Origins of *Moringa oleifera* accessions used in the study

### 3.2 Morphological Characterisation

The accessions were planted at the University of Ghana farm, Legon on the soil belonging to the Nyigbenya-Hatso series characterised by well drained, red, sandy clay loam to clay with abundant rough stone concretions and quartz gravels classified by the soil research institute (FAO/UNESCO, 1990). A randomised complete block design was used for the experiment with three replicates. Seeds were nursed, pricked out and transplanted onto the permanent plot after four weeks. Planting was done at a distance of 2.5m within rows and 2.5m between rows. A total land area of 656.25m<sup>2</sup> was used for the study with seven plants per row. A Moringa record sheet with modification from AVRDC-GRSU was used for the morphological characterisation. The passport data of the various accessions used for the morphological studies are shown in appendix 1.

#### 3.2.1. Data Collection

Quantitative and qualitative data were taken on the vegetative and morphology of the plant. Mean values were computed by dividing the sum of the parameters measured by the number of samples selected from the observation unit. Data was taken six months after planting. Data collected were as follows;

1. *Tripinnatum leaf length*: This is the average length of the leaf measured from the petiole attached to the stem to the tip or apex of the leaf.
2. *Tripinnatum leaf width*: This measured the average width recorded from the tip of the longest opposite leaflets in the leaf.
3. *Tripinnatum leaflet length*: This parameter measured the average length of a leaflet from the tip of the leaflet to the point of attachment to the midrib of the leaf.

4. *Tripinnatum leaflet width*: This was measured from the furthest points of the leaflet across the midrib.
5. *Growth habit*: Plants were classified as exhibiting compact, intermediate or tall (erect)
6. *Leaf type*: leaf was scored either as simple or compound for the various accessions in study.
7. *Leaf arrangement*: Plant data for the accessions on this parameter was recorded as opposite, alternate, pinnate or palmate.
8. *Leaflet arrangement*: Data for the accessions with regard to leaflet arrangement was recorded as opposite, alternate, pinnate or palmate.
9. *Leaflet shape*: The leaflet shape of the plants was recorded as simple or compound.
10. *Leaf colour*: For each accession the colour of the leaf was recorded as *light green*, green or dark green.
11. *Petiole pigmentation*: the petiole pigmentation was observed as green, dark green, purple or dark purple.
12. *Colour of the stem bark*: The colour of the woody stem observed and recorded as whitish, green, brown or dark brown.

### **3.3 Biochemical Studies**

#### **3.3.1 Experimental material**

Dried leaves of plants established from stem cuttings of *Moringa oleifera* from eight different locations were used for both biochemical analysis and molecular

characterisation. The passport data of the various accessions used for both the biochemical and molecular studies are shown in appendix 2.

### 3.3.2 Dry matter determination

The hot empty crucible was weighed ( $W_2$ ) and 1.0 g ( $W_1$ ) of air dried sample transferred into it. The crucible with the sample in it was placed in an oven at 105 °C overnight. The hot crucible with the dried sample was weighed ( $W_3$ ) and the percentage dry matter content computed.

$$\text{Dry matter \%} = \frac{(W_3 + C_2) - (W_2 + C_1) * 100}{W_1}$$

Where  $W_1$  = weight of air dried sample

$W_2$  = weight of empty crucible

$W_3$  = weight of crucible + oven dried sample

$C_1$  = correction for  $W_2$  read from the balance (due to hot weighing)

$C_2$  = correction for  $W_3$  read from the balance (due to hot weighing)

### 3.3.3 Crude protein determination

0.3 g ground dried leaf sample was weighed into a digestion tube. 1 g catalyst mixture of  $K_2SO_4$  and anhydrous  $CuSO_4$  was added in the ratio 10:1.5 ml conc.  $H_2SO_4$  was then added and the tube placed in the digester with the temperature raised to 350 °C. The sample was digested for about an hour and a half or until the colour changed to light blue. The tube was removed from the block and allowed to cool. 30 ml of distilled water was carefully added to the digestion tube. The steam distillation procedure was then used to determine the ammonium in the digestion mixture. This

was done by first placing the digestion tube onto the Tecator steam distillation apparatus. The distiller was then set according to manufacturer's instruction. The digestion tube was inserted into the system and 150 ml of distillate collected into the receiver flask containing 25 ml 4% boric acid. The distillate was then titrated against a standard acid (0.1N HCl). The colour change was from green to steel-blue then pink with the end-point occurring when a drop of acid resulted in a colour change from steel-blue to light pink. The percentage nitrogen content computed and multiplied by 6.25 to give the percentage crude protein.

$$\% \text{ N} = \frac{14.01*(V-B)*n}{W*10*DM\%}$$

$$\% \text{ Crude Protein} = \text{N} * 6.25$$

Where:

N = Nitrogen

V = volume of HCl consumed

B = blank titration

n = normality of HCl

W = weight of sample taken

DM% = Dry matter of sample

### 3.3.4 Determination of macronutrients

1.0 g of the plant sample was weighed into a silica crucible and heated to ash in a muffle furnace for 5 hours at 500–600 °C. The ash residue was dissolved in dilute HNO<sub>3</sub> and filtered through acid-washed filter paper in a 100 ml volumetric flask and the volume made up to the mark with distilled water. A blank digest were prepared of the same amounts of reagents to account for any contamination through the acids used in the digestion. The estimation of calcium (Ca) magnesium (Mg) and potassium

(K) was done using the atomic absorbance spectrometer (AAS). For sodium (Na) the flame photometer was used.

#### 3.3.4.1 Determination of Ca, Mg and K using AAS

5 ml aliquot of the above extract was transferred into a 100 ml volumetric flask and made up to the volume. The sample solution was then atomised and the absorbance observed. The corresponding concentration for the absorbance recorded represented the content of Ca and Mg in the sample solution. The percentage calcium, magnesium or potassium was computed.

$$\text{Ca/Mg/ K (g) in 100g (\%Ca/ Mg/ K)} = \frac{C * df * 100}{1000000} = \frac{C * 2000 * 100}{1000000} = \frac{C}{5}$$

Where C = concentration of Ca ( $\mu\text{g/ml}$ ) as read from the standard curve;

df = dilution factor, which is  $100 \times 20 = 2\,000$ , as calculated below:

1 g of sample made to 100 ml (100 times);

5 ml of sample solution made to 100 ml (20 times).

1 000 000 = factor for converting  $\mu\text{g}$  to g.

#### 3.3.4.2 Determination of phosphorus using spectrophotometric vanadium phosphomolybdate method

5 ml of the digest was transferred into a 50-ml volumetric flask, and 10 ml of vanadomolybdate reagent added. The volume was made up to the mark with distilled water, and thoroughly mixed. It was then kept for 30 minutes for a yellow colour to develop and read on spectrophotometer at 420 nm. The percentage phosphorus was then computed as follows:

$$\text{P content (g) in 100g (\% P)} = \frac{C * df * 100}{1000000} = \frac{C * 1000 * 100}{1000000} = \frac{C}{10}$$

Where  $C$  = concentration of P ( $\mu\text{g/ml}$ ) as read from the standard curve;

$df$  = dilution factor, which is  $100 \times 10 = 1\,000$ , as calculated below:

1 g of sample made to 100 ml (100 times);

5 ml of sample solution made to 50 ml (10 times).

1 000 000 = factor for converting  $\mu\text{g}$  to g.

### 3.3.5 Determination of micronutrients and lead (Pb)

0.3 g of dry Moringa leaf sample was weighed into a clean 125 ml 'pyrex' conical flask. 4 ml of concentrated  $\text{H}_2\text{SO}_4$  was then added and the flask swirled carefully to ensure that the entire sample is wetted. The flask and its content were heated in a fume hood chamber on an electric sand bath plate set at medium heating. The flask was removed, cooled and about 10 drops of  $\text{H}_2\text{O}_2$  added to the solution. 3-4 drops of  $\text{H}_2\text{O}_2$  were added at a time again slowly to avoid vigorous reaction of the contents. The flask was swirled with the contents kept at the bottom of the flask and reheated avoiding excessive heating that causes spattering. The flask and its content were cooled and 6 drops of  $\text{H}_2\text{O}_2$  added before reheating. Cooling and adding drops of  $\text{H}_2\text{O}_2$  were done until there was a change in colour from black to dark brown. The temperature of the heater was then increased on the hot electronic sand-bath plate and cooling continued while adding 6 drops of  $\text{H}_2\text{O}_2$  and heating. When the solution stayed colourless upon cooling, peroxide was added and left for the last time on the burner at high temperature for 10-15 minutes. The solution was cooled and contents transferred into a 100 ml volumetric flask and brought to the mark with distilled water, cooled and mixed thoroughly. The solution was then used to determine iron, manganese, zinc and lead levels. Two blank digests were prepared of the same

amounts of the reagents (i.e. H<sub>2</sub>SO<sub>4</sub> and H<sub>2</sub>O<sub>2</sub>). All data were corrected against the mean blank value and the percentage metal content computed.

$$\% \text{ Metal} = \frac{(A-B) * V * 100}{1000 * 1000 * W}$$

Where; A = AAS reading

B = Blank reading

V = Volume of extract

W = Weight of sample

**Table 3.1:** The wavelength, slit and current at which the various metals read.

| <b>Metal</b> | <b>Wavelength (nm)</b> | <b>Slit</b> | <b>Current (mA)</b> |
|--------------|------------------------|-------------|---------------------|
| Fe           | 248.33                 | 1.8/1.5     | 30                  |
| Zn           | 213.86                 | 2.7/1.8     | 15                  |
| Mn           | 279.48                 | 1.8/0.6     | 20                  |
| Pb           | 283.31                 | 2.7/1.05    | 10                  |

### 3.4 Molecular Characterisation

#### 3.4.1 Total protein studies

##### 3.4.1.1 Total protein extraction

A modified protocol of Bio-Rad Laboratories (Philadelphia, USA) was used for electrophoretic studies of total protein variation at 10% sodium dodecyl sulphate (SDS) polyacrylamide gel. The collected accessions were screened for variations in the total protein fingerprints using fresh leaf samples. About 0.2 g of Moringa leaves

were washed with distilled water. The washed leaves were then macerated using ceramic mortar and pestle with 500  $\mu$ l of Tris-citric buffer of pH 7.0 (extraction buffer). The macerated leaves were transferred into eppendorf tubes and centrifuged at 15000 rpm for six minutes. The supernatant were then collected and transferred into new eppendorf tubes.

#### **3.4.1.2 Preparation of gel for electrophoresis**

Polyacrylamide gel electrophoresis was used to separate the DNA samples studied. Two glass plates, a comb and Perspex spacers were prewashed in 6M HCl and rinsed several times in distilled water. The plate sets were blocked with plasticine to prevent the gel from leaking away until it solidified. The comb was inserted and 1cm below the comb marked with a marker. The plate set was then filled with a prepared separating gel (appendix 4g) to about 2 cm to the brim, topped with a few drops of n-butanol and allowed to polymerise in about 40 minutes. The n-butanol was then washed off with distilled water and the space left filled with the prepared stacking gel (figure 4d) almost to the brim. The comb was immediately inserted in the stacking gel and the gel allowed to polymerise for about 30 minutes.

#### **3.4.1.3 Loading and running the electrophoresis**

After the gel had set, the plate sets were removed from the plasticine and the gel assembly set placed in the electrophoresis tank. The electrode buffer pH 8.3 was poured into the bottom and the top compartments to make contact with the ends of the gel and form a circuit. The comb was removed and each well loaded with 15  $\mu$ l of total protein extracts of an accession. The power leads were then connected to make an anionic contact (black lead) to the top and cathode contact (red lead) to the bottom.

The system was then run at constant voltage of 50V until the front/tracking dye moved into the separating gel. The voltage was adjusted to 80V and allowed to run until the front/tracking dye moved within 2mm to the bottom of the separating gel.

#### **3.4.1.4 Staining and destaining the gel**

The gel was removed from the plate and placed in a plastic tray and stained with Coomassie brilliant blue (appendix 3b) for about 2 hours. The gel was then destained by washing in destaining solution as (appendix 3c) until the background became clear with only the protein bands taking up the stain. The gel was then stored in 20% glycerol solution, photographed and the bands scored. The buffer systems, running conditions and staining procedures for total protein studies are shown in appendix 3 and 4.

### **3.4.2 Molecular genotyping using Random Amplified Polymorphic DNA (RAPD)**

#### **3.4.2.1 DNA extraction (D'neasy plant mini-kit)**

About 0.2 g of plant material was ground under liquid nitrogen to a fine powder using mortar and pestle. The tissue powder and the liquid nitrogen were transferred into an eppendorf tube and the liquid nitrogen allowed to evaporate before the sample thawed. 400 µl of buffer AP1 and 4 µl of RNase A stock solution (100 mg/ml) was added to a maximum of 100 mg of ground plant tissue and vigorously vortexed. The mixture was incubated for 10 minutes at 65°C and was mixed 2-3 times during incubation by inverting tube. 130 µl of buffer AP2 was added to the lysate, mixed and incubated for 5 minutes on ice. The lysate in a 2 ml collection tube was centrifuged

for 2 minutes at a speed of 8000 rpm. The supernatant was carefully collected to a new tube without disturbing the cell-debris pellet. 255  $\mu$ l buffer AP3 and 450  $\mu$ l of ethanol (96-100%) was added to the 450  $\mu$ l of clear lysate and mixed by pipetting. 650  $\mu$ l of the mixture was then transferred to a new tube and centrifuged for a minute at  $\geq 6000$  g (i.e.  $\geq 8000$  rpm) and the supernatant collected and discarded. The process was repeated and the supernatant discarded. 500  $\mu$ l of buffer AW was added to the precipitate and centrifuged for 1 minute at a speed  $\geq 6000$  x g (i.e.  $\geq 8000$  rpm) and the supernatant collected and discarded. 100  $\mu$ l of preheated (65°C) buffer AE was pipetted and added to the precipitate. The mixture was incubated for 5 minutes at room temperature and then centrifuged for 1 minute at  $\geq 6000$  g (i.e.  $\geq 8000$  rpm) to elute. The elution process was repeated.

#### **3.4.2.2 DNA quantification**

DNA concentration and its purity were determined by measuring the optical density using the UV absorbance spectrophotometer. The quality of DNA was assessed on a 0.8 % (w/v) agarose gel electrophoresis that contains 0.5X tris borate EDTA (TBE) buffer. The DNA concentration was estimated with a Hoefer TKO 100 mini fluorometer. The intensity of orange fluorescence was also used for rough estimation of the concentration of DNA together with UV absorbance. The purity was measured by spectrophotometry.

#### **3.4.2.3 Polymerase chain reaction and gel electrophoresis**

PCR amplification conditions were: pre-denaturation at 94 °C for 4 mins, 45 cycles of denaturation at 94 for 1 min, annealing at 34 °C for 1 min, elongation step at 72 °C for

2 mins. Final cycle at 72 °C for 5 mins and final product was then cooled at 4 °C. RAPD-PCR products were then analysed by electrophoresis on a 1.2% agarose gel with 1XTAE buffer and stained with 5 µl ethidium bromide. Power was supplied at 70V for 2 hours. DNA banding patterns were then visualized using Benchtop 3UV<sup>TM</sup> transilluminator (Cambridge, UK).

**Table 3.2:** List of the RAPD primers and their nucleotide sequences used in the study

| Primer No. | Primer Code | Nucleotide Sequence (5' - 3') |
|------------|-------------|-------------------------------|
| 1          | OPC-08      | TGGACCGGTA                    |
| 2          | OPC-04      | CCGCATCTAC                    |
| 3          | OPC-07      | CTCCCGACGA                    |
| 4          | OPC-16      | CACACTCCAG                    |
| 5          | OPC-06      | GAACGGACTC                    |
| 6          | OPC-13      | AAGCCTCGTC                    |
| 7          | OPC-18      | TGAGTGGGTG                    |
| 8          | OPC-15      | GACGGATCAG                    |
| 9          | OPC-01      | TTCGAGCCAG                    |
| 10         | OPC-19      | GTTGCCAGCC                    |

### 3.5 Data Analysis

#### 3.5.1 Morphological data

Quantitative data obtained were analysed using the analysis of variance (ANOVA).

All statistical comparisons were done at 5% significance level. Where the analysis of

variance indicated significant differences among treatments, least significance difference (L.S.D) was used to compare the treatment means. The standard errors and ranges were calculated from the mean values for each accession. Qualitative data obtained for the accessions were grouped based on their descriptor state. Qualitative data were scored on presence (1) and absence (0) basis. The data obtained were entered onto an excel sheet where characters were given codes using numbers and used to compute the similarity coefficient and to generate a dendrogram to reflect their relatedness using the NTSYS-pc version 2.2i, (Applied Biostatistics Inc, USA) and to generate a matrix determining similarity among samples according to Dice's coefficients. The similarity matrix was then used to draw a dendrogram with the clustering method UPGMA via the SAHN module in the software.

### **3.5.2 Biochemical Data**

Biochemical data obtained were analysed using the analysis of variance (ANOVA) with all statistical comparisons done at 5% level of significance. Where the analysis of variance indicated significant differences among treatments, least significance difference (L.S.D) was used to compare the treatment means. The standard errors and ranges were calculated from the mean values for each accession. Graphs were also used to represent analysed data.

### **3.5.3 Molecular data**

#### **3.5.3.1 Total protein**

For analysis of total protein data, the samples were genotyped for the presence and absence of bands as 1 and 0, respectively. The data were then used to draw a

dendrogram. NTSYS-pc version 2.2i, (Applied Biostatistics Inc, USA) was used to generate a matrix determining similarity among samples according to Dice's method. The similarity matrix was then used to draw a dendrogram with the clustering method UPGMA via the SAHN module in the software. Both total protein and RAPD data were combined and used to draw a dendrogram.

### 3.5.3.2 RAPD

Amplification products were scored for presence (1) or absence (0) of bands in a binary matrix at the same molecular weights. Where a PCR product was not obtained, data for specific locus and genotype were treated as missing data. Only polymorphic bands were included in the binary data set, and similarities were calculated using Dice coefficient (Dice 1945; Nei and Li, 1979)

$$\text{Dice's coefficient } (D_{ij}) = \frac{2a}{(2a+b+c)}$$

Where  $D_{ij}$  = similarity between two individuals  $i$  and  $j$

$a$  = number of bands present in both  $i$  and  $j$ ,

$b$  = present in  $i$  but not in  $j$ ,

$c$  = present in  $j$  but absent in  $i$ .

Cluster analyses using the UPGMA (unweighted pair-group method with arithmetic averages; Sokal and Michener, 1958) were carried out on the similarity matrices phenograms (dendrogram) constructed via the SAHN module using the software NTSYS-pc version 2.2i, (Applied Biostatistics Inc, USA).

## CHAPTER FOUR

### RESULTS

#### 4.1 Morphological analysis

##### 4.1.1 Variation in vegetative characteristics of the *Moringa* accessions

At six months after planting (MAP), mean tripinnatum leaf length ranged between 25.2 cm and 61.6 cm whilst the tripinnatum leaf width ranged between 17.8 cm and 48.5 cm (Table 4.1). For both parameters the minimum values were recorded on accessions collected from Ashanti region of Ghana while the maximum was recorded on accessions from the Upper East region. Tripinnatum leaflet length and tripinnatum leaflet width ranged from 10.8 cm to 25.9 cm and 5.7 cm and 17.6 cm respectively (Table 4.1). Accessions from Ashanti region had the lowest tripinnatum leaflet length and tripinnatum leaflet width with values of 10.8 cm and 5.7 cm respectively (Table 4.1). Accessions collected from Upper East had the highest tripinnatum leaflet length of 25.9 cm whilst those assembled from Greater Accra with a value of 17.6 cm had the maximum tripinnatum leaflet width. There were significant differences among the accessions of *Moringa oleifera* for the vegetative features determined in this study ( $P \leq 0.05$ , Appendices 6a, 6b, 6c and 6d).

**Table 4.1:** Vegetative characteristics of *Moringa oleifera* accessions.

| Origin of accession     | Mean vegetative features at 6 MAP $\pm$ SE (cm) |                                   |                                   |                                   |
|-------------------------|---|-----------------------------------|-----------------------------------|-----------------------------------|
|                         | Tripinnatum leaf length                         | Tripinnatum leaf width            | Tripinnatum leaflet length        | Tripinnatum leaflet width         |
| USA                     | 54.4 $\pm$ 6.55                                 | 43.1 $\pm$ 5.37                   | 23.9 $\pm$ 3.10                   | 13.9 $\pm$ 1.55                   |
| AR                      | <u>25.2 <math>\pm</math> 2.70</u>               | <u>17.8 <math>\pm</math> 0.99</u> | <u>10.8 <math>\pm</math> 0.58</u> | <u>5.7 <math>\pm</math> 0.27</u>  |
| BA                      | 37.8 $\pm$ 3.16                                 | 23.5 $\pm$ 2.40                   | 14.4 $\pm$ 2.23                   | 8.9 $\pm$ 1.14                    |
| ER                      | 41.6 $\pm$ 6.91                                 | 29.9 $\pm$ 5.65                   | 16.8 $\pm$ 2.99                   | 9.2 $\pm$ 1.43                    |
| GA                      | 50.3 $\pm$ 1.69                                 | 38.2 $\pm$ 3.97                   | 20.1 $\pm$ 1.30                   | <b>17.6 <math>\pm</math> 5.88</b> |
| INDIA                   | 43.8 $\pm$ 1.79                                 | 33.0 $\pm$ 2.35                   | 17.9 $\pm$ 1.29                   | 9.6 $\pm$ 0.77                    |
| NR                      | 34.7 $\pm$ 1.67                                 | 23.5 $\pm$ 2.58                   | 14.9 $\pm$ 1.13                   | 8.8 $\pm$ 0.46                    |
| UER                     | <b>61.6 <math>\pm</math> 3.11</b>               | <b>48.5 <math>\pm</math> 5.90</b> | <b>25.9 <math>\pm</math> 1.78</b> | 13.3 $\pm$ 1.25                   |
| UWR                     | 44.5 $\pm$ 2.83                                 | 34.4 $\pm$ 3.70                   | 20.6 $\pm$ 1.74                   | 12.6 $\pm$ 1.81                   |
| VR                      | 37.3 $\pm$ 5.35                                 | 27.5 $\pm$ 4.88                   | 14.9 $\pm$ 2.64                   | 9.0 $\pm$ 1.16                    |
| <b>L.s.d p&lt; 0.05</b> | <b>11.59</b>                                    | <b>11.93</b>                      | <b>5.99</b>                       | <b>6.28</b>                       |
| <b>%CV</b>              | <b>21.0</b>                                     | <b>29.1</b>                       | <b>25.9</b>                       | <b>45.1</b>                       |

**Note;** SE = standard error of the mean, minimum values are underlined whilst maximum values are bolded. AR= Ashanti region, BA = Brong Ahafo region, ER = Eastern region, GA = Greater Accra region, NR = Northern region, UER = Upper East region, UWR = Upper West region, VR = Volta region and USA = United States of America.

#### **4.1.2. Qualitative attributes of the *Moringa oleifera* accessions**

There was variation in the morphological traits in the *Moringa oleifera* accessions studied (Table 4.2). Some of the variations observed were in traits such as colour of leaf, petiole pigmentation and stem bark colour of the accessions. Leaf colour varied from light green to dark green. Most of the accessions had purple petioles, others petioles were dark purple and a few were green in colour. Four accessions showed a growth habit of intermediate between erect and compact with two accessions exhibiting compactness. The remaining accessions exhibited erect growth form. Variations in the stem bark colour of the accessions comprised whitish/ silvery, green and dark brown. Three of the accessions had a whitish bark colour with an accession exhibiting a brown colour. In the case of leaf type, leaf arrangement, leaflet shape and leaflet arrangement the accessions appeared to be monomorphic for the traits.

**Table 4.2:** Qualitative attribute of the *Moringa oleifera* accessions

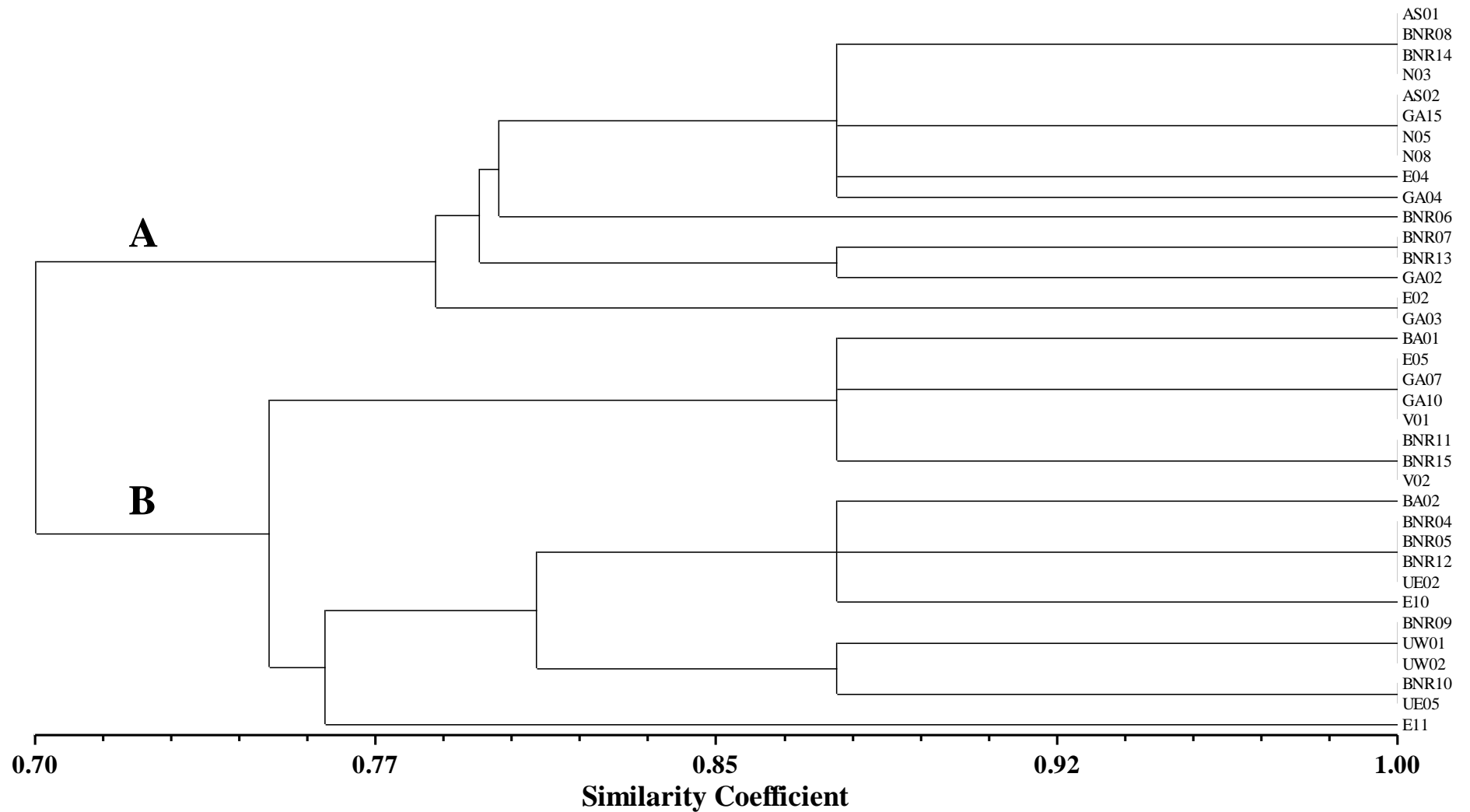
| <b>ID</b> | <b>Leaf colour</b> | <b>Leaf type</b> | <b>Leaf arrangement</b> | <b>Leaflet shape</b> | <b>Leaflet arrangement</b> | <b>Petiole pigmentation</b> | <b>Growth habit</b> | <b>Bark Colour</b> |
|-----------|--------------------|------------------|-------------------------|----------------------|----------------------------|-----------------------------|---------------------|--------------------|
| AS01      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| AS02      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| BA01      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Whitish            |
| BA02      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Intermediate        | Green              |
| BNR04     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| BNR05     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| BNR06     | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Intermediate        | Green              |
| BNR07     | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Green              |
| BNR08     | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| BNR09     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| BNR10     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Intermediate        | Dark brown         |
| BNR11     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Green              |
| BNR12     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| BNR13     | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Green              |
| BNR14     | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| BNR15     | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Green              |
| E02       | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Green                       | Erect               | Dark brown         |
| E04       | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Brown              |
| E05       | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Dark brown         |
| E10       | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Compact             | Green              |
| E11       | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Compact             | Whitish            |

**Table 4.2:** Continued qualitative attribute of the *Moringa oleifera* accessions

| <b>ID</b> | <b>Leaf colour</b> | <b>Leaf type</b> | <b>Leaf arrangement</b> | <b>Leaflet shape</b> | <b>Leaflet arrangement</b> | <b>Petiole pigmentation</b> | <b>Growth habit</b> | <b>Bark Colour</b> |
|-----------|--------------------|------------------|-------------------------|----------------------|----------------------------|-----------------------------|---------------------|--------------------|
| GA02      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Green                       | Erect               | Green              |
| GA03      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Green                       | Erect               | Dark brown         |
| GA04      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Whitish            |
| GA07      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Dark brown         |
| GA10      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Dark brown         |
| GA15      | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| N03       | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| N05       | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| N08       | Light green        | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| UE02      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Green              |
| UE05      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Intermediate        | Dark brown         |
| UW01      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| UW02      | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Purple                      | Erect               | Dark brown         |
| V01       | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Dark brown         |
| V02       | Dark green         | Compound         | Alternate               | Compound             | Opposite                   | Dark Purple                 | Erect               | Green              |

#### **4.1.3. Cluster analysis for Qualitative attributes of the *Moringa oleifera* accessions**

Cluster analysis resolved the qualitative data for the *Moringa oleifera* accessions into two major clusters A and B at a similarity level of 70% (figure 4.1). Major cluster A contained sixteen accessions; AS01, BNR 08, BNR 14, N03, AS02 GA15, N05, N08, E04, GA04, BNR06, BNR07, BNR13, GA02, E02 and GA03. Cluster A further separated into two sub-clusters at 78% similarity level with accession E02 and GA03 separating themselves from the rest of the accessions in major cluster A. Major cluster B contained; BA01, E05, GA07, GA10, V01, BNR11, BNR15, C05, V02, BA02, BNR04, BNR05, BNR12, UE02, E10, BNR09, UW01, UW02, BNR10, UE05 and E11. It further separated at 75% similarity level into two sub-clusters containing BA01, E05, GA07, GA10, V01, BNR11, BNR15 and V02 in one group and BA02, BNR04, BNR05, BNR12, UE02, E10, BNR09, UW01, UW02, BNR10, UE05 and E11 in the other group. Accessions AS01, BNR 08, BNR 14 and N03; AS02 GA15, N05 and N08; BNR07 and BNR13; E02 and GA03 in major cluster A and E05, GA07, GA10 and V01; BNR11, BNR15 and V02; BNR04, BNR05, BNR12 and UE02; BNR09, UW01 and UW02; BNR10 and UE05 in major cluster B could not separate at 100% similarity level suggesting that they are duplicates or they are the same based on the traits studied.



**Fig 4.1:** Dendrogram generated by UPGMA cluster analysis showing the relationship among Moringa accessions based on Dice similarity from qualitative data.

## 4.2. Biochemical analysis

### 4.2.1 Concentration of micro minerals in the dried leaf powder of *Moringa oleifera* accessions

Table 4.3 shows the levels of micro-minerals such as iron (Fe), manganese (Mn) and zinc (Zn) in the dried leaf samples of *Moringa oleifera* accessions collected from different locations. Fe content in the dried leaf samples ranged from 0.0014% to 0.0736%. The accessions from Upper West region had the lowest Fe content and those from India had the highest. The concentration of Mn detected in the samples varied from 0.0006% to 0.0097%. Accessions from Upper East region had the minimum amount of Mn whereas those from India gave the highest concentration of the mineral element. The accessions collected from Northern region had the lowest concentration of Zn with a value of 0.0005% and the highest quantity (0.0111%) obtained from accessions assembled from India. There were significant differences among the accessions of *Moringa oleifera* studied for concentration of micro-minerals ( $P \leq 0.05$ , Appendices 7a, 7b and 7c). The quantity of heavy metal lead (Pb) detected in the dried leaf samples ranged from 0.0005% to 0.0056%. Accessions from USA had the lowest Pb concentration whilst those from India had the highest concentration of the metal. There was a significant difference among the accessions examined ( $P \leq 0.05$ , Appendix 7d).

**Table 4.3:** Micro mineral composition in the dried leaf of *Moringa oleifera* accessions

| Origin of<br>accession    | Mean Micro Mineral Composition (%) $\pm$ SE |  |  |  |
|---------------------------|---|--|--|--|
|                           | Fe (%) $\pm$ SE                             | Mn (%) $\pm$ SE                        | Zn (%) $\pm$ SE                        | Pb (%) $\pm$ SE                        |
| USA                       | 0.0033 $\pm$ 0.00017                        | 0.0011 $\pm$ 0.00031                   | 0.0006 $\pm$ 0.00006                   | <u>0.0005 <math>\pm</math> 0.00006</u> |
| ER                        | 0.0039 $\pm$ 0.00054                        | 0.0007 $\pm$ 0.00006                   | 0.0031 $\pm$ 0.00245                   | 0.0019 $\pm$ 0.00020                   |
| GR                        | 0.0044 $\pm$ 0.00012                        | 0.0008 $\pm$ 0.00025                   | 0.0006 $\pm$ 0.00009                   | 0.0016 $\pm$ 0.00009                   |
| India                     | <b>0.0736 <math>\pm</math> 0.00202</b>      | <b>0.0097 <math>\pm</math> 0.00088</b> | <b>0.0111 <math>\pm</math> 0.00065</b> | <b>0.0056 <math>\pm</math> 0.0007</b>  |
| NR                        | 0.0027 $\pm$ 0.00043                        | 0.0007 $\pm$ 0.00003                   | <u>0.0005 <math>\pm</math> 0.00010</u> | 0.0013 $\pm$ 0.00015                   |
| UER                       | 0.0044 $\pm$ 0.00123                        | <u>0.0006 <math>\pm</math> 0.00006</u> | 0.0007 $\pm$ 0.00003                   | 0.0010 $\pm$ 0.00007                   |
| UWR                       | <u>0.0014 <math>\pm</math> 0.00003</u>      | 0.0007 $\pm$ 0.00036                   | 0.0005 $\pm$ 0.00003                   | 0.0008 $\pm$ 0.00006                   |
| VR                        | 0.0026 $\pm$ 0.00026                        | 0.0010 $\pm$ 0.00019                   | 0.0008 $\pm$ 0.00006                   | 0.0027 $\pm$ 0.00031                   |
| <b>L.s.d (p&lt; 0.05)</b> | <b>0.0025</b>                               | <b>0.0010</b>                          | <b>0.0027</b>                          | <b>0.0008</b>                          |
| <b>%CV</b>                | <b>12</b>                                   | <b>31</b>                              | <b>68.9</b>                            | <b>23.7</b>                            |

**Note:** SE = standard error of the mean, minimum values are underlined whilst maximum values are bolded. ER = Eastern region, GA = Greater Accra region, NR = Northern region, UER = Upper East region, UWR = Upper West region, VR = Volta region and USA = United States of America. Conversion rate: 1% = 10000 ppm; 1ppm =1 mg/kg

#### **4.2.2. Concentration of macro minerals in the dried leaf powder of *Moringa oleifera* accessions**

Table 4.4 shows the mean macro mineral composition of dried leaf of *Moringa oleifera* accessions collected from the various provenances. Samples from USA had the lowest concentration of Ca (40 mg/kg) whilst those accessions from Northern region of Ghana had the highest Ca concentration of 79.3 mg/kg. The lowest quantity of Mg detected was from Volta region (19.23 mg/kg) accessions whilst accessions from Northern region had the highest quantity of Mg of 48.13 mg/kg. The Na content of the *Moringa oleifera* samples ranged from 87.5 mg/kg to 253.1 mg/kg (Table 4.4). Accessions from Volta region had the lowest Na concentration whilst those from Greater Accra region had the highest level of Na in the dried leaves of Moringa. The accessions from the Upper East had the lowest concentration of 1549 mg/kg of potassium (K) in the dried leaves sample of Moringa whilst Upper West region had the highest K concentration of 2272 mg/kg. The phosphorus concentration in the accessions ranged from 0.179 mg/kg recorded in the accessions from India to 0.438 mg/kg in accessions from Greater Accra region as shown in Table 4.4. There were significant differences between the accessions for the macro minerals studied except potassium concentration as shown in appendices 8a, 8b 8c, 8d and 8e.

**Table 4.4:** Macro mineral composition of dried leaf of *Moringa oleifera* accessions

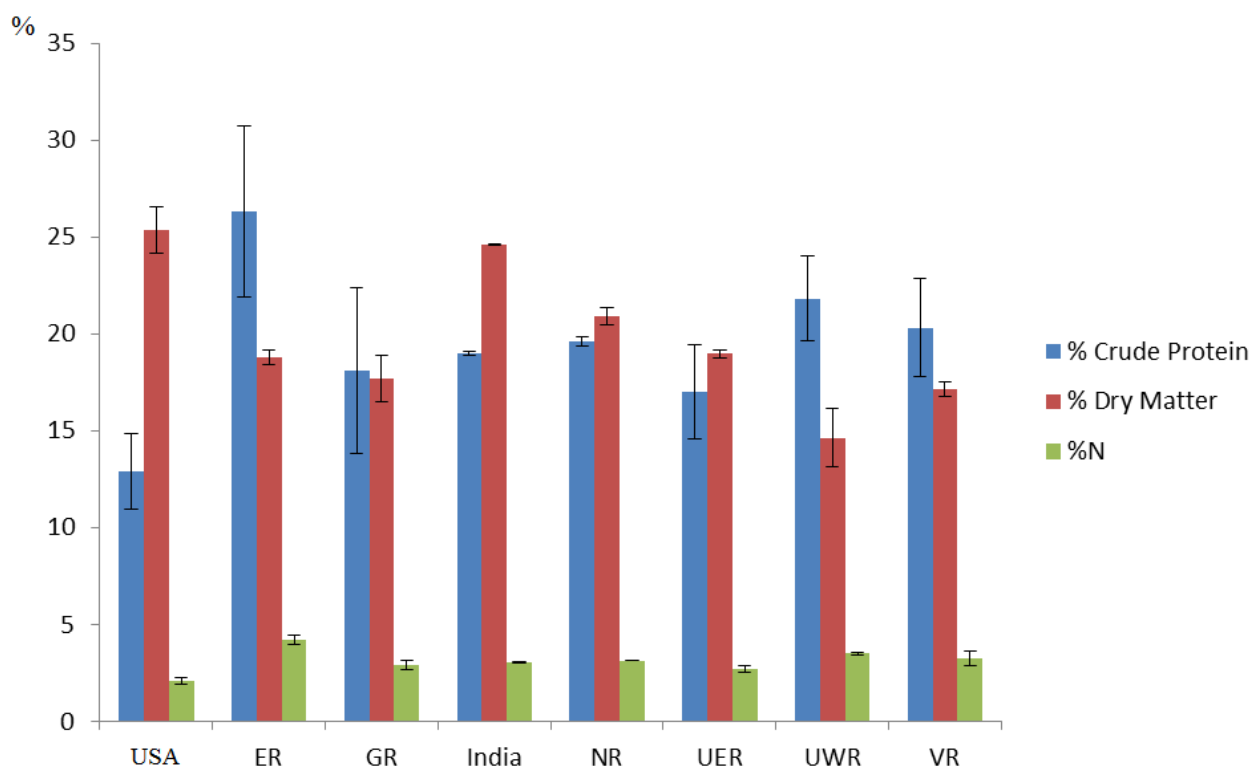
| Origin of accession       | Mean Macro Mineral Composition (mg/kg DM) $\pm$ SE |                                    |                                     |                                     |                                     |
|---------------------------|--|------------------------------------|-------------------------------------|-------------------------------------|-------------------------------------|
|                           | Ca   | Mg                                 | Na                                  | K                                   | P                                   |
| USA                       | <u>40.3 <math>\pm</math> 3.23</u>                  | 22.67 $\pm$ 1.19                   | 185.9 $\pm$ 8.88                    | 2155 $\pm$ 175.21                   | 0.183 $\pm$ 0.017                   |
| ER                        | 44.5 $\pm$ 4.08                                    | 26.87 $\pm$ 2.12                   | 128.1 $\pm$ 3.25                    | 1868 $\pm$ 194.13                   | 0.396 $\pm$ 0.006                   |
| GR                        | 48.2 $\pm$ 2.39                                    | 26.70 $\pm$ 1.21                   | <b>253.1 <math>\pm</math> 14.06</b> | 2131 $\pm$ 279.31                   | <b>0.438 <math>\pm</math> 0.033</b> |
| India                     | 44.2 $\pm$ 2.14                                    | 20.60 $\pm$ 1.69                   | 191.7 $\pm$ 1.59                    | 2074 $\pm$ 62.65                    | <u>0.179 <math>\pm</math> 0.004</u> |
| NR                        | <b>79.3 <math>\pm</math> 1.98</b>                  | <b>48.13 <math>\pm</math> 1.85</b> | 184.4 $\pm$ 10.17                   | 1700 $\pm$ 211.81                   | 0.271 $\pm$ 0.039                   |
| UER                       | 53.8 $\pm$ 4.89                                    | 23.57 $\pm$ 1.17                   | 192.2 $\pm$ 3.13                    | <u>1549 <math>\pm</math> 35.04</u>  | 0.304 $\pm$ 0.009                   |
| UWR                       | 41.3 $\pm$ 1.21                                    | 20.8 $\pm$ 1.65                    | 167.2 $\pm$ 13.10                   | <b>2272 <math>\pm</math> 394.42</b> | 0.183 $\pm$ 0.010                   |
| VR                        | 50.3 $\pm$ 4.27                                    | <u>19.23 <math>\pm</math> 1.04</u> | <u>87.5 <math>\pm</math> 6.51</u>   | 1162 $\pm$ 93.72                    | 0.267 $\pm$ 0.024                   |
| <b>L.s.d (p&lt; 0.05)</b> | <b>9.84</b>  | <b>4.88</b>                        | <b>39.12</b>                        | <b>ns</b>                           | <b>0.1174</b>                       |
| <b>%CV</b>                | <b>11.2</b>  | <b>10.7</b>                        | <b>12.9</b>                         |                                     | <b>24.1</b>                         |

**Note:** SE = standard error of the mean, ns = not significant, minimum values are underlined whilst maximum values are bolded. ER = Eastern region, GA = Greater Accra region, NR = Northern region, UER = Upper East region, UWR = Upper West region, VR = Volta region and USA = United States of America.

#### 4.2.3 Percentage crude protein and percentage dry matter of dried leaf samples of *Moringa oleifera* accessions.

Percentage nitrogen of *Moringa oleifera* accessions studied ranged from 2.06% to 4.21%. The lowest percentage nitrogen was observed in the accession from USA and the highest in the accessions from Eastern region (figure 4.2). Significant differences existed between the provenances studied for percentage nitrogen (appendix 9a). The percentage crude protein of *Moringa oleifera* accessions studied showed that

accessions from the eastern region contained the highest crude protein value of 26.3% whilst the lowest value of 12.9% occurred in accessions from USA (figure 4.2). However, the accessions were not significantly different from each other with regard to percentage crude protein (appendix 9b). The percentage dry matter content of the accessions ranged from 14.63% to 25.37% (figure 4.2). The lowest was recorded in accessions from the Upper West region of Ghana and the greatest from the accessions from USA. There were significant differences in percentage dry matter of the accessions studied (appendix 9c).



**Fig 4.2:** Percentage crude protein and dry matter of dried leaves of *M. oleifera* accessions.

**Table 4.5: Nutrient composition per 100 g of dried Moringa leaves and USDA-RDI**

| Origin of Accession | Nutrient per 100g of dried Moringa leaves (mg) |      |       |       |       |        |        |        |
|---------------------|--|------|-------|-------|-------|--------|--------|--------|
|                     | Fe   | Mg   | Mn    | P     | Zn    | K      | Ca     |        |
| USA                 | 3.30   | 2.27 | 1.10  | 0.018 | 0.60  | 215.50 | 4.03   |        |
| ER                  | 3.90   | 2.69 | 0.70  | 0.040 | 3.10  | 186.80 | 4.45   |        |
| GR                  | 4.40   | 2.67 | 0.80  | 0.044 | 0.60  | 213.10 | 4.82   |        |
| India               | 73.60  | 2.06 | 9.70  | 0.018 | 11.10 | 207.40 | 4.42   |        |
| NR                  | 2.70   | 4.81 | 0.70  | 0.027 | 0.50  | 170.00 | 7.93   |        |
| UER                 | 4.40   | 2.36 | 0.60  | 0.030 | 0.70  | 154.90 | 5.38   |        |
| UWR                 | 1.40   | 1.92 | 0.70  | 0.018 | 0.50  | 227.20 | 4.13   |        |
| VR                  | 2.60   | 2.08 | 1.00  | 0.027 | 0.80  | 116.20 | 5.03   |        |
| Mean conc.          | 12.04  | 2.61 | 1.91  | 0.028 | 2.24  | 186.39 | 5.02   |        |
| RDI                 | Male (mg/d)                                    | 8.0  | 400.0 | 2.3   | 700.0 | 11.0   | 4700.0 | 1000.0 |
|                     | Female (mg/d)                                  | 18.0 | 310.0 | 1.8   | 700.0 | 8.0    | 4700.0 | 1000.0 |

Source: USDA National Nutrient Database for Standard Reference, Release 24

**Note:** ER = Eastern region, GA = Greater Accra region, NR = Northern region,

UER = Upper East region, UWR = Upper West region, VR = Volta region and

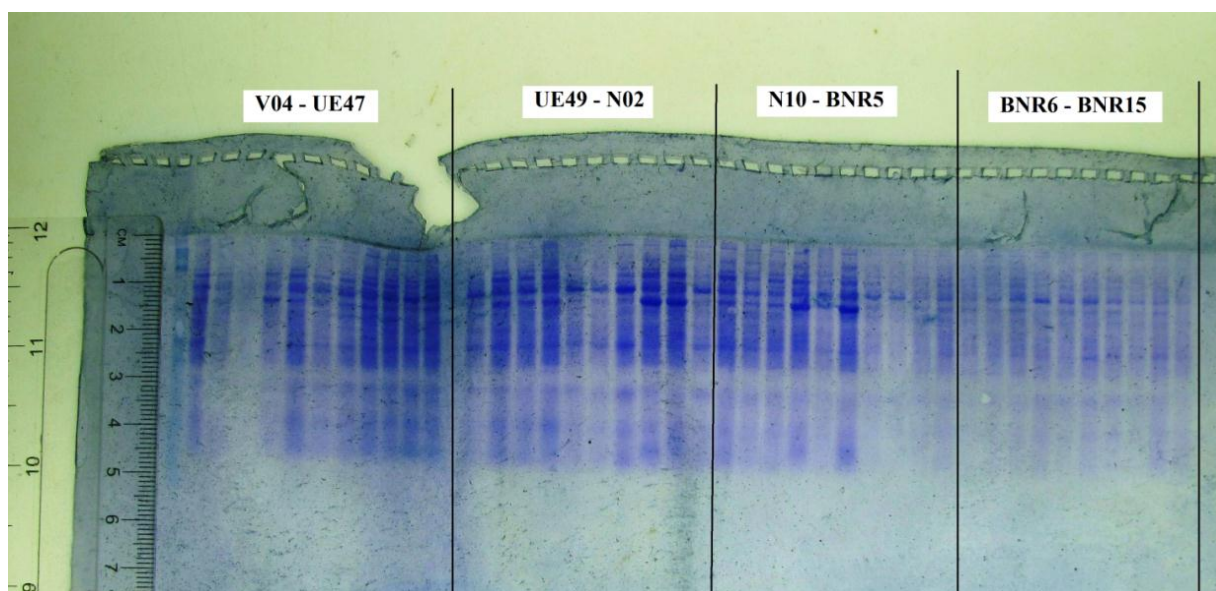
USA = United States of America.

### 4.3 Molecular characterisation of *Moringa oleifera* leaf samples

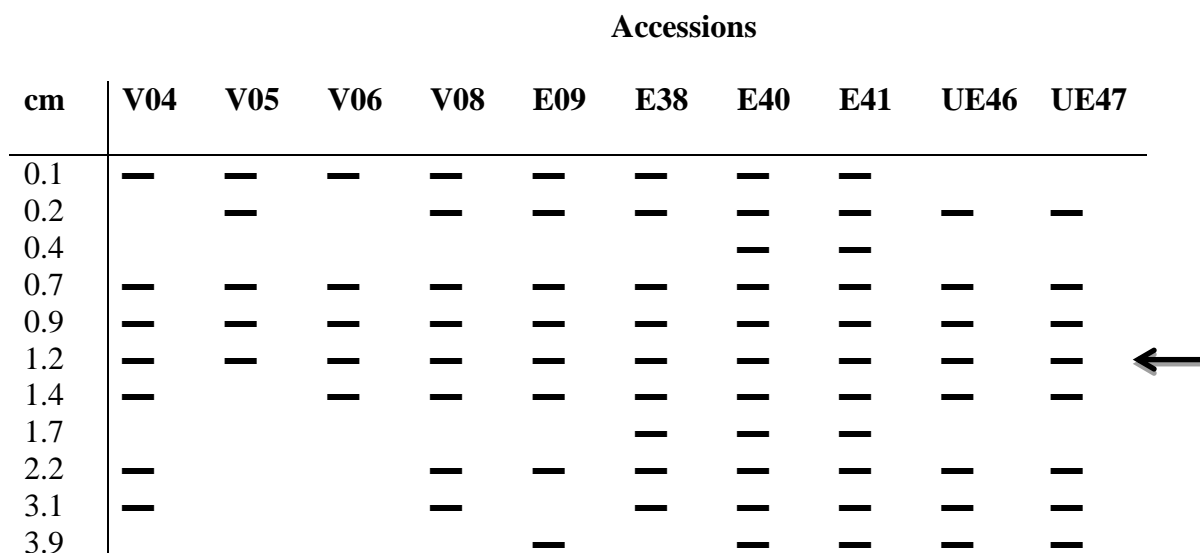
Forty (40) *Moringa oleifera* accessions from Upper East, Upper West, Northern, Eastern, Volta and Greater Accra regions of Ghana, United States of America (USA) and India were electrophoresed to determine the variation existing at the biochemical and genotypic level. Variations were observed in the accessions studied for both total protein and RAPDs markers.

#### 4.3.1 Total Protein

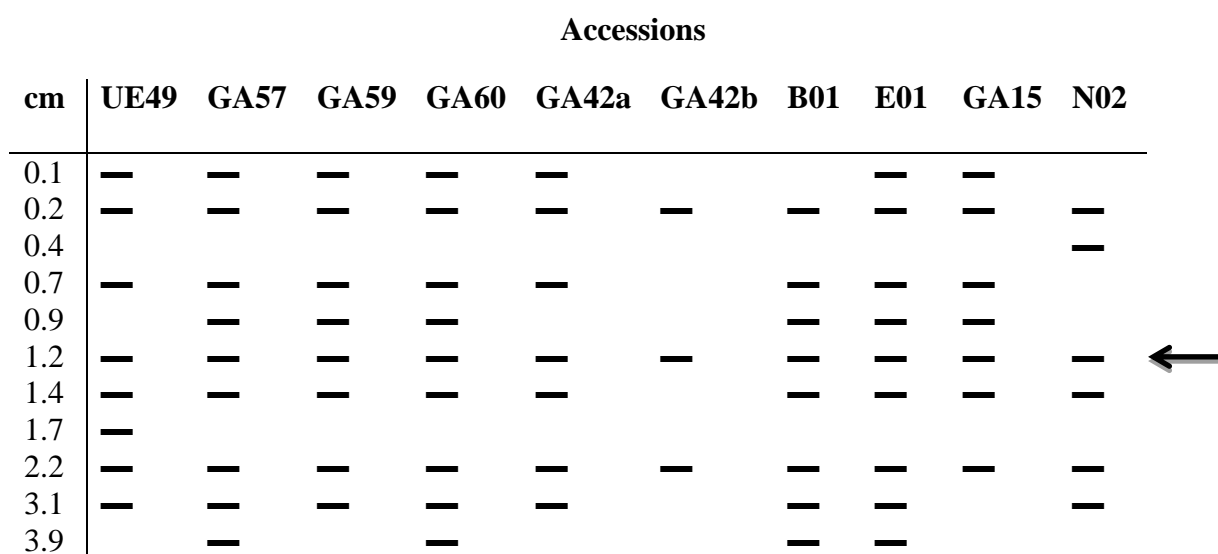
A total of eleven (11) prominent bands were resolved. Two out of the forty accessions, namely E40 and E41 showed all the eleven bands. The bands ranged from a distance of 0.1cm to 3.9cm with a single band at distance of 1.2 cm which was common to all the forty (40) accessions electrophoresed (Figures 4.3a, 4.3b, 4.3c, 4.3d). A sample photograph of the gel is shown in plate 1.



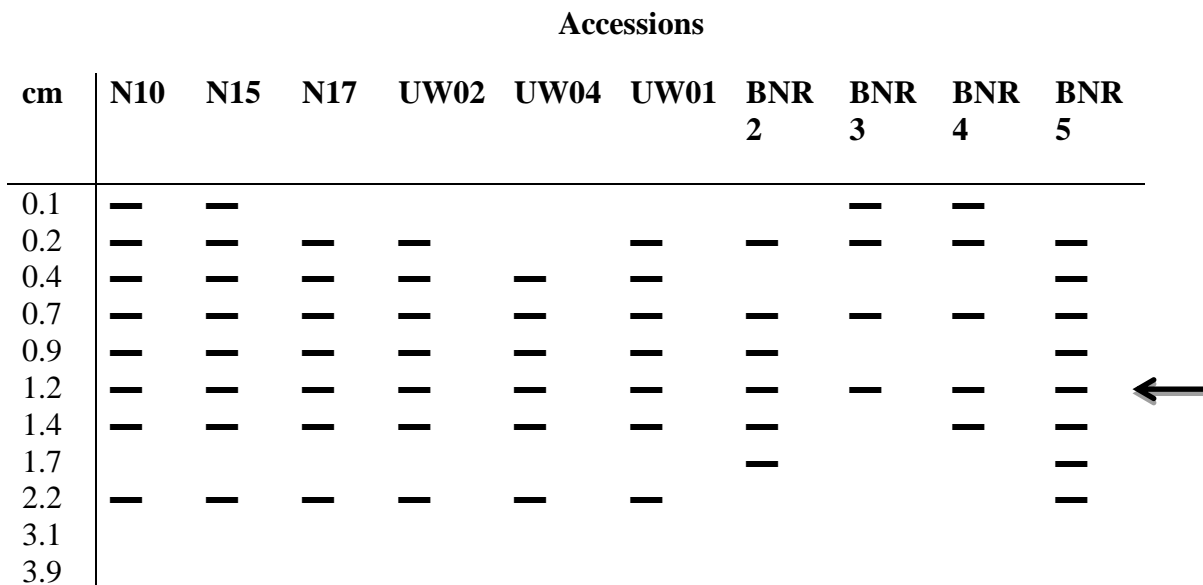
**Plate 1:** Total Protein Gel of 40 *Moringa oleifera* accessions collected from different parts of the country.



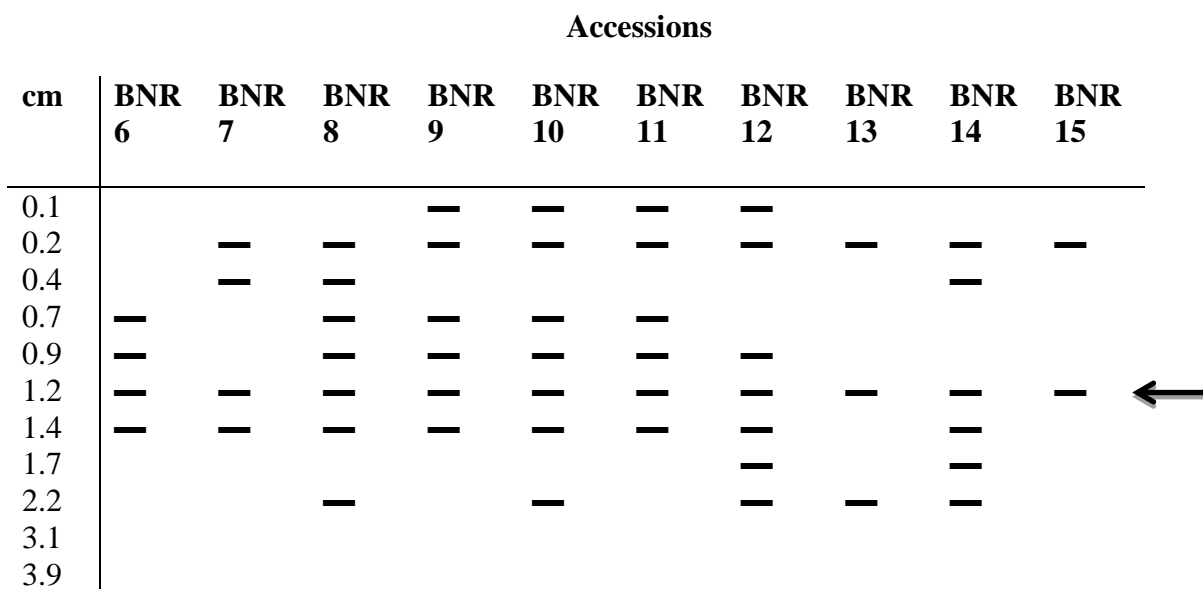
**Fig 4.3a:** Schematic diagram showing the total protein banding pattern of the 10 *Moringa oleifera* accessions from Volta, Eastern and Upper east region.



**Fig 4.3b:** Schematic diagram showing the total protein banding pattern of the 10 *Moringa oleifera* accessions from Upper East, Greater Accra, Brong Ahafo, Eastern and Northern region.



**Fig 4.3c:** Schematic diagram showing the total protein banding pattern of the 10 *Moringa oleifera* accessions from Northern region, Upper West region and BNARI.



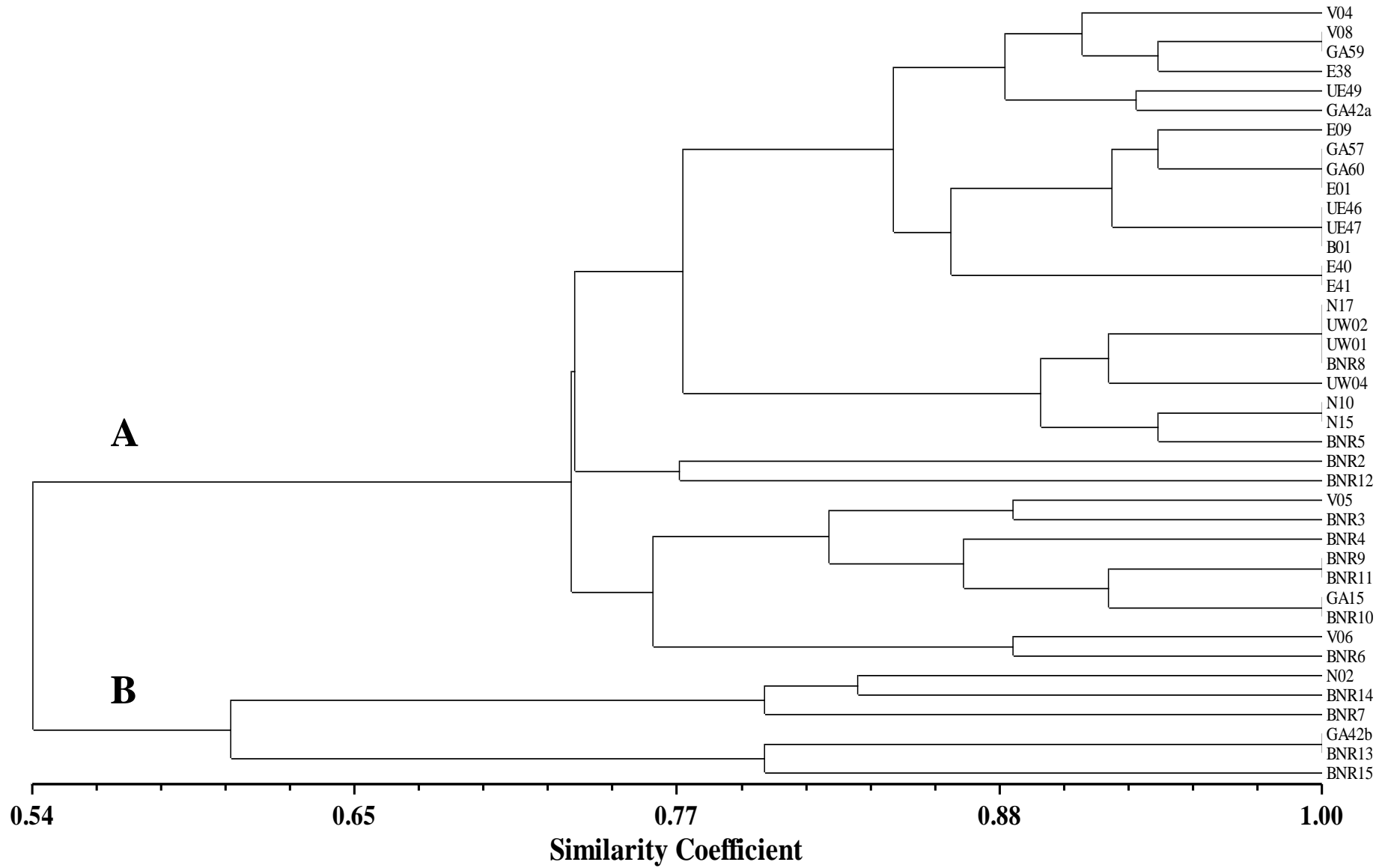
**Fig 4.3d:** Schematic diagram showing the total protein banding pattern of the 10 *Moringa oleifera* accessions from BNARI.

← Monomorphic locus

### 4.3.2 Total protein cluster analysis

The dendrogram generated from total protein profile for the forty (40) *Moringa oleifera* accessions is shown in figure 4.4. The cluster analysis resolved the forty accessions into two major clusters (i.e. A and B) at a similarity level of 54%. Major cluster A contained thirty four (34) accessions; V04, V08, GA59, E38, UE49, GA42a, E09, GA57, GA60, E01, UE46, UE47, B01, E40, E41, N17, UW02, UW01, BNR8, UW04, N10, N15, BNR5, BNR2, BNR12, V05, BNR3, BNR4, BNR9, BNR11, GA15, BNR10, V06 and BNR6. Cluster A further separated into two sub-clusters at 73% similarity level with accessions V04, V08, GA59, E38, UE49, GA42a, E09, GA57, GA60, E01, UE46, UE47, B01, E40, E41, N17, UW02, UW01, BNR8, UW04, N10, N15, BNR5, BNR2 and BNR12 in one set and V05, BNR3, BNR4, BNR9, BNR11, GA15, BNR10, V06 and BNR6 in the other set.

Major cluster B contained; N02, BNR14, BNR7, GA42b, BNR13 and BNR15. It further separated at 61% similarity level into two sub-clusters containing N02, BNR14 and BNR7 in one group and GA42b, BNR13 and BNR15 in the other. The following groups of accessions from major cluster A could not be separated from each other at 100% similarity V08 and GA59; GA57, GA60 and E01; UE46, UE47 and B01; E40 and E41; N17, UW02, UW01 and BNR 8; N10 and N15; BNR 9 and BNR 11; GA15 and BNR10. Similarly accessions GA42b and BNR 13 in major cluster B could not separate at 100% similarity level suggesting that they are very closely related or duplicate accessions.



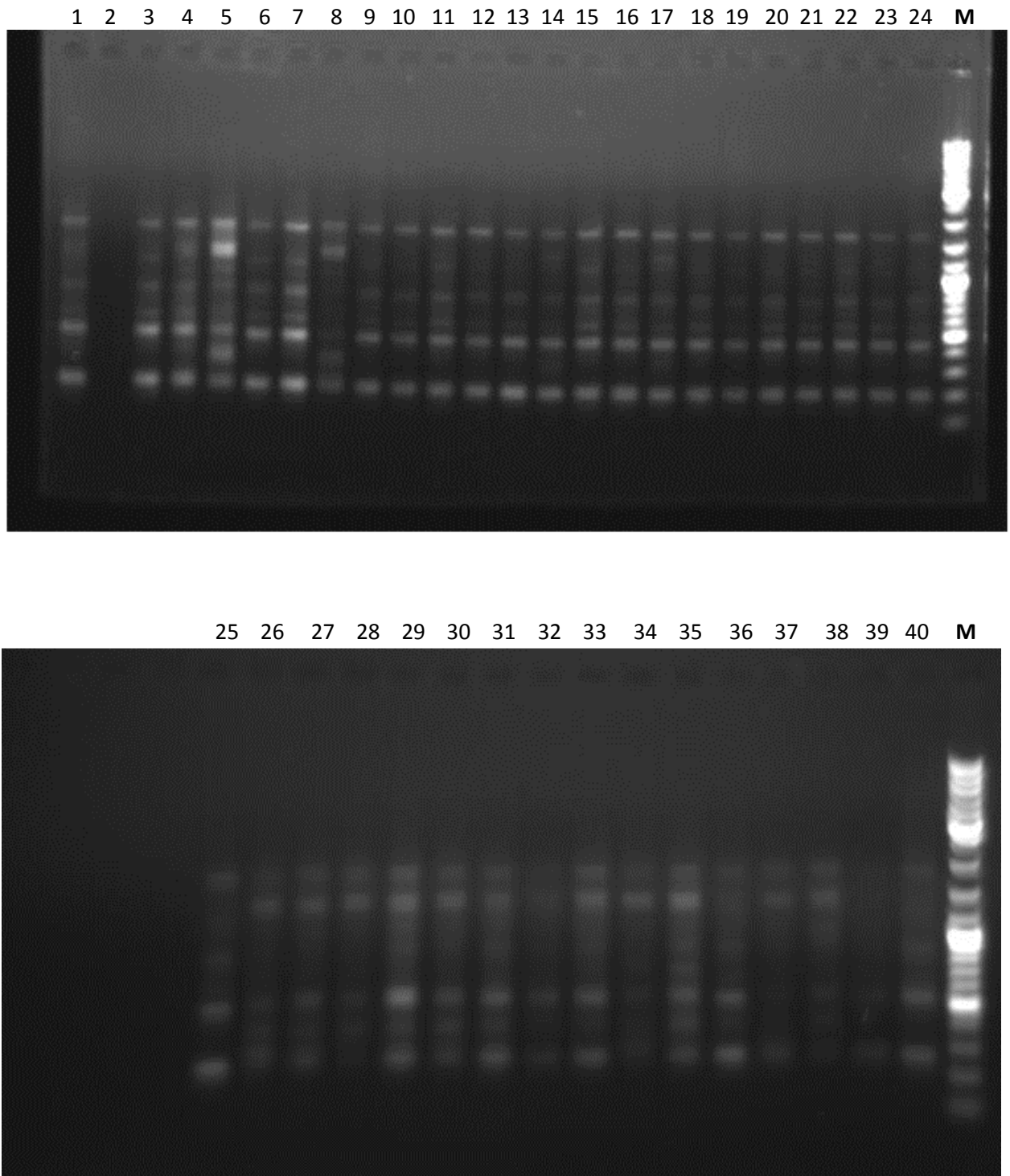
**Fig 4.4:** Dendrogram generated by UPGMA cluster analysis showing the relationship among forty (40) *Moringa* accessions based on Dice similarity from Total protein data

### 4.3.3 RAPD markers

10 out of the 12 RAPD markers used generated visible bands. A total of 82 prominent polymorphic RAPD fragments were generated. The number of amplified loci ranged from 7 to 10. Of the 10 primers used, the number of RAPD loci generated was highest for the primers OPC-07 and OPC-06 which amplified 10 fragments each as shown in table 4.5. The lowest number of bands was produced by the primers OPC-08, OPC-04, OPC-16 and OPC-19 with seven loci each (table 4.5). The average number of fragments and percentage polymorphic bands produced per primer were 8.2 and 95.1 respectively. A representative profile of RAPD products amplified with primer OPC-08 from all 40 *Moringa oleifera* accessions is shown in plate 2.

**Table 4.6:** List of primers and their sequences used for the RAPD analysis and corresponding number of bands generated.

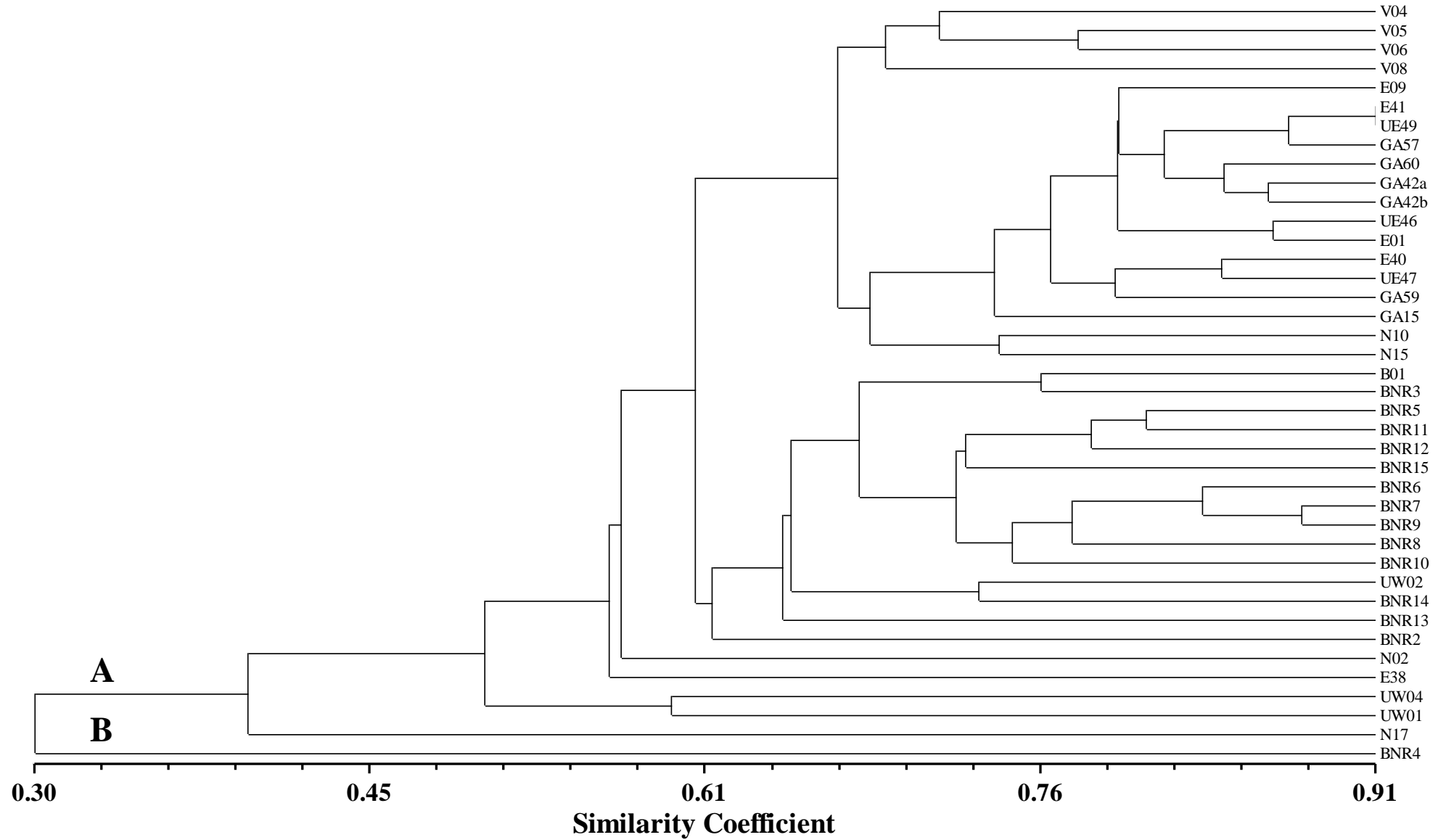
| Primer No.   | Primer code | Sequence (5'–3') | Total No. of bands | Polymorphic bands |             | Monomorphic bands |            |
|--------------|-------------|------------------|--------------------|-------------------|-------------|-------------------|------------|
|              |             |                  |                    | No.               | %           | No.               | %          |
| 1            | OPC-08      | TGGACCGGTA       | 7                  | 7                 | 100         | –                 | –          |
| 2            | OPC-04      | CCGCATCTAC       | 7                  | 7                 | 100         | –                 | –          |
| 3            | OPC-07      | CTCCCGACGA       | 10                 | 9                 | 90          | 1                 | 10         |
| 4            | OPC-16      | CACACTCCAG       | 7                  | 7                 | 100         | –                 | –          |
| 5            | OPC-06      | GAACGGACTC       | 10                 | 8                 | 80          | 2                 | 20         |
| 6            | OPC-13      | AAGCCTCGTC       | 9                  | 9                 | 100         | –                 | –          |
| 7            | OPC-18      | TGAGTGGGTG       | 8                  | 8                 | 100         | –                 | –          |
| 8            | OPC-15      | GACGGATCAG       | 8                  | 8                 | 100         | –                 | –          |
| 9            | OPC-01      | TTCGAGCCAG       | 9                  | 9                 | 100         | –                 | –          |
| 10           | OPC-19      | GTTGCCAGCC       | 7                  | 6                 | 85.7        | 1                 | 14.3       |
| <b>Total</b> | –           | –                | <b>82</b>          | <b>78</b>         | <b>95.1</b> | <b>4</b>          | <b>4.9</b> |



**Plate 2:** Sample of DNA profile generated by OPC-01; lanes 1-40 corresponding to the code of the 40 *Moringa oleifera* accessions in appendix 2. M - GeneRuler DNA ladder Mix (Fermentas)

#### **4.3.4. RAPD analysis of 40 *Moringa oleifera* accessions**

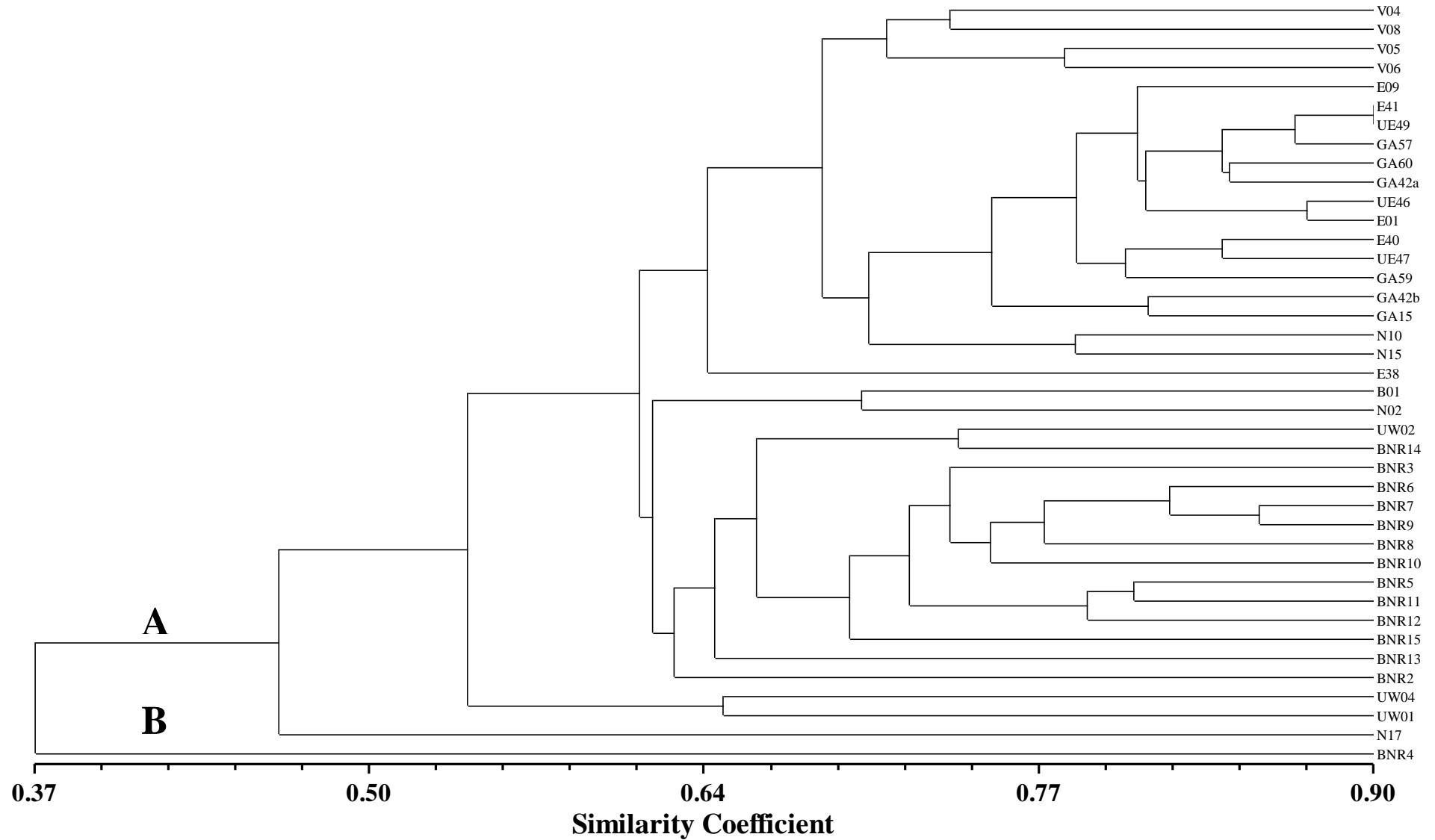
The dendrogram generated for the RAPD data for the *Moringa oleifera* accessions was resolved into two major clusters A and B at a similarity coefficient of 30% as shown in figure 4.5. Major cluster A contained all the accessions analysed except BNR 4 which was in cluster B. Major cluster A further separated into two sub-clusters at 40% similarity level with accessions N17 separating itself from the rest of the accessions. Accessions E41 and UE49 in major cluster A could not separate at 91% similarity coefficient suggesting that they may be very closely related or the same.



**Fig 4.5:** Dendrogram generated by UPGMA cluster analysis showing the relationship among 40 *Moringa* accessions based on Dice similarity from RAPD data

#### **4.3.5 Combination of Total protein and RAPD analysis of 40 *Moringa oleifera* accessions**

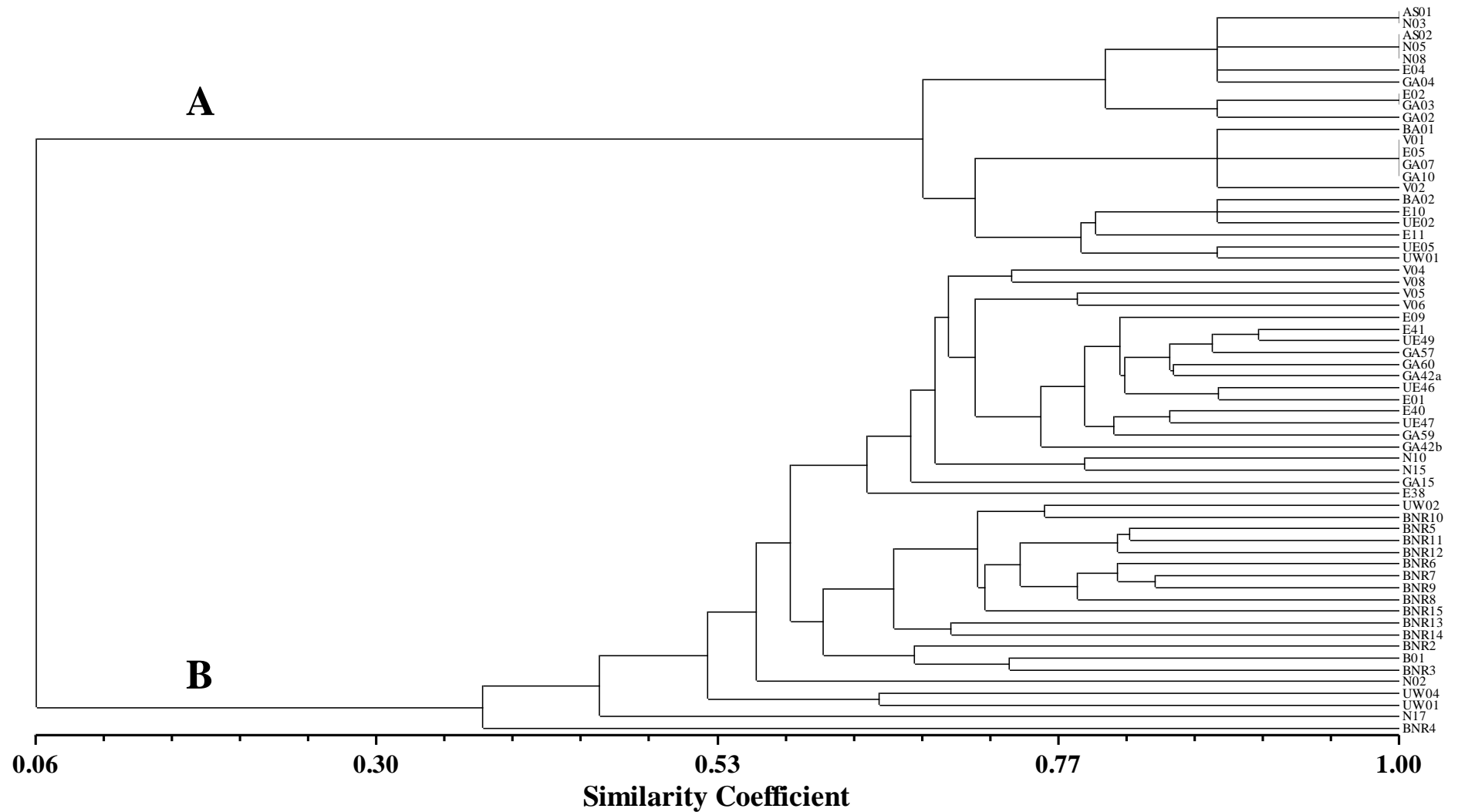
The cluster analysis generated for the combination of Total protein and RAPD data for the *Moringa oleifera* accessions was resolved into two major clusters A and B at a similarity coefficient of 37% (figure 4.6). Major cluster A comprised all the accessions apart from accession BNR 4 which was in major cluster B. Major cluster A further separated into two sub-clusters at 46% similarity level with accessions N17 separating itself from the rest of the accessions. Accessions E41 and UE49 in major cluster A could not separate at 90% similarity coefficient suggesting that they may be closely related.



**Fig 4.6:** Dendrogram generated by UPGMA cluster analysis showing the relationship among 40 Moringa accessions based on Dice similarity from the combination of Total protein and RAPD data

#### **4.3.6 Combination of Molecular and Morphological analyses of *Moringa oleifera* accessions**

Cluster analysis generated from the combination of Molecular and Morphological data for the *Moringa oleifera* accessions was resolved into two main clusters, A and B at a similarity coefficient of 6% (figure 4.7). Main cluster A consisted of AS01, N03, AS02, N05, N08, E04, GA04, E02, GA03, GA02, BA01, V01, E05, GA07, GA10, V02, BA02, E10, UE02, E11, UE05 and UW01. It further separated into two sub-clusters at 67% similarity level with accessions AS01, N03, AS02, N05, N08, E04, GA04, E02, GA03 and GA02 forming one sub-clusters and accessions BA01, V01, E05, GA07, GA10, V02, BA02, E10, UE02, E11, UE05 and UW01 making up the other half. Main cluster B comprised of V04, V08, V05, V06, E09, E41, UE49, GA57, GA60, GA42a, GA42b, UE46, E01, E40, UE47, GA59, N10, N15, GA15, E38, UW02, BNR10, BNR5, BNR11, BNR12, BNR6, BNR7, BNR9, BNR8, BNR15, BNR13, BNR14, BNR2, B01, BNR3, N02, UW04, UW01, N17 and BNR4. Main cluster B further separated into two sub-clusters at 39% similarity level with accession BNR4 separating itself from the remaining accessions. These groups of accessions AS01 and N03; AS02, N05 and N08; E02 and GA03; V01, E05, GA07 and GA10 in main cluster A could not separate at 100% similarity coefficient.



**Fig 4.7:** Dendrogram generated by UPGMA cluster analysis showing the relationship of *Moringa* accessions based on Dice similarity from the combination of molecular and morphological data

## CHAPTER FIVE

### DISCUSSION

#### 5.1 Variability among accessions for morphological characteristics

The provenances of *Moringa oleifera* Lam. from the ten different locations, eight in Ghana and two from United State of America and India, studied exhibited variations with respect to the eight morphological traits studied. These were tripinnatum leaf length (at 6MAP), tripinnatum leaf width (at 6MAP), tripinnatum leaflet length (at 6MAP), tripinnatum leaflet width (at 6MAP), leaf colour, petiole pigmentation, growth habit and stem bark colour. In germplasm management, morphological characterisations are usually employed as markers either within or between species, varieties or accessions (Acquaah, 2007). These markers called descriptors give the description of taxonomy of the crop (Hilu, 1989). The information though basic is vital and provides the breeder with the information on genetic closeness of the accessions. Information acquired from these markers aid breeders in the selection of accessions with desirable traits for both farmers and consumers (Gitonga *et al.*, 2008).

Diversity in morphological characteristics such as tripinnatum leaf length, tripinnatum leaf width, tripinnatum leaflet length and tripinnatum leaflet width give important information to *Moringa* plant breeders in selecting accessions with large leaf area. This is because the leaves are the most consumed part of the plant especially in Ghana. In this instance larger leaf area will yield larger amounts of dried leaf powder which will mean more income for farmers and business persons who process the leaves into powder for sale. Variability in the qualitative information such as leaf

colour, petiole pigmentation and stem bark colour as well as growth habit recorded in the study was desirable. Edwards *et al.*, (2000) also observed variations in these characters in another species, *Moringa stenopetala* in their study. Diversity provides the breeder with the genetic material for selection, improvement as well as conservation of the plants (Acquaah *et al.*, 1992; Brothers and Kelly, 1993). For instance, the choice of a dark coloured leaf over light coloured one will be desired by the breeder because of the high amount of chlorophyll which will enable the plant to produce more photosynthate for effective growth. Also, dark leaf colour will be preferred for beverage especially when fresh leaves are used.

## **5.2 Variations in Biochemical Analysis**

Variations in nutrient composition among *Moringa oleifera* accessions could be from genetic source, environmental source or cultivations method (Broin 2006; Dechasa, 2006). In this study, the leaf samples from cuttings of the various accessions from the different location were used. The environmental conditions as well as cultivation methods were the same for all accessions studied. Therefore the only source of variation affecting the biochemical variations namely variation in the amount of micro and macronutrients may be the genotypic differences of provenances used in the study.

Minerals are required for normal growth, activities of muscles and skeletal development (such as calcium), cellular activity and oxygen transport (copper and iron), chemical reaction in the body and intestinal absorption (magnesium), fluid balance and nerve transmission (sodium and potassium), as well as the regulation of acid-base balance (phosphorus). Iron is useful in prevention of anemia and other

related diseases (Oluyemi *et al.*, 2006). Manganese plays a role in energy production and in supporting the immune system (Muhammad *et al.*, 2011). Zinc is useful for protein synthesis, normal body development and recovery from illness (Muhammad *et al.*, 2011). The minerals determined are very essential macro minerals since they make the edible parts of Moringa useful for food or feed (Dechassa *et al.*, 2003). The average concentration of Ca observed was 50.24 mg/kg. This was lower compared to the findings of authors like Aslam *et al.*, (2005) and Oduro *et al.*, (2008) who reported 229.3 mg/kg and 200.9 mg/kg respectively. However, the value compared well with those reported by Council of Scientific and Industrial Research., (1962); Verma *et al.*, (1975) and Owusu-Ansah, (2010) viz 44 mg/kg, 44 mg/kg and 73.4 mg/kg respectively. The average amounts of Mg (26.07 mg/kg) and P (0.2776 mg/kg) obtained was lower than those reported by Owusu-Ansah, (2010) who obtained average concentrations of Mg and P of 26.6 mg/kg and 0.59 mg/kg respectively. The mean concentration of K recorded in the study was 1863.9 mg/kg. This was lower than the concentration reported by Aslam *et al.*, (2005) who had 2098.2 mg/kg. The quantity of K detected was less than the USDA-RDI (4700 mg/d) for an adult human being.

Regarding the micro-nutrients, Fe is an essential trace element for haemoglobin formation, normal functioning of the central nervous system and in the oxidation of carbohydrates, proteins and fats (Adeyeye and Otokiti, 1999). The mean value of Fe in the dried Moringa leaf samples was 0.012% (i.e. 120 mg/kg). This is similar to values reported Magat *et al.*, (2009) of 100 mg/kg but higher than that reported by Oduro *et al.*, (2008) for Fe composition in dry leaf powder of Moringa. The micronutrients Mn and Zn had 1.9 mg/kg and 2.2 mg/kg respectively. These were

lower than values obtained by Magat *et al.*, (2009) who reported 340mg/10kg for Mn and 100mg/10kg for Zn in Moringa leaf.

In plants, the critical levels of the heavy metal Pb is 10-20 ppm and higher concentrations than these are very likely to be toxic (Sauerbeck 1982). According to the FAO/WHO standard (Codex Alimentarius Commission, 2001) the safe level of lead (Pb) in vegetables is 0.2 mg/kg. The recorded average Pb level of the accessions studied was 0.0019% (i.e. 1.9 mg/kg), which is above the accepted safe level of 0.2 mg/kg in vegetables but below the critical level of 10-20 ppm in plants. The observed lead level in the accessions studied is considered toxic for human or animal consumption but not toxic to the plant. Authors like Garba *et al.*, (2010) of Nigeria and Magat *et al.*, (2009) of Philippines also reported high lead levels in Moringa species taken as vegetables in their countries. They reported 321 mg/kg and 90 mg/kg respectively. The amount of Pb in the dried leaf samples of Moringa probably may be due to the area or location of collection especially in the local collections which included collections from the road side and near filling stations. Lead (Pb) pollution has been shown to be commensurate with population and vehicular density (Afolami *et al.*, 2010). Also high levels of Pb in some plants are attributable to pollutants in irrigation water, farm soil or due to pollution from the highways traffic (Qui *et al.*, 2000). Like the Chinese cabbage which has been reported to pick up and accumulate higher Pb concentrations (Wong *et al.*, 1996; Chove *et al.*, 2006), *Moringa oleifera* may also behave likewise especially when exposed to lead pollutants.

The lowest percentage dry matter among the accessions was observed in the accessions from the Upper West, meaning they had the highest moisture content and

hence may be more prone to deterioration since foods with high moisture content are more prone to perishability (Fennema and Tannenbaum, 1996). Inferably, the accessions from USA will be less prone to deterioration since they have the least amount of moisture content. Similar observations of variability in dry matter of the leaves from different accessions have been reported by Fuglie, (2001); Magat *et al.*, 2009 and Oduro *et al.*, (2008).

With the exception of the accessions from USA which had 12.9% crude protein, all the Moringa accessions had 17% protein or more with those from the Eastern region of Ghana recording the highest amount of 26.3%. The results obtained for the range of protein levels in the accessions are similar to levels documented by Fuglie, (2001), Elkhailifa *et al.*, (2007) and Oduro *et al.*, (2008). This makes the leaves of Moringa a good source of protein since plant food that provides more than 12% of its calorific value from protein are considered good source of protein (Pearson, 1976). The crude protein levels observed in the study were higher than the levels reported in *Cucurbita moschata*, *Trianthema portulacastrum*, *Spinacia oleracea*, *Abelmoschus esculentus*, *Praecitrullus fistulosus*, *Luffa acutangula* and *Cucumis sativus* with protein percentage of  $10.77 \pm 0.23$ ,  $19.63 \pm 0.9$ ,  $20.82 \pm 0.07$ ,  $16.17 \pm 0.26$ ,  $15.34 \pm 0.30$ ,  $13.47 \pm 0.05$  and  $10.57 \pm 0.23$  respectively (Hussain *et al.*, 2010). Percentage nitrogen corresponds with percentage protein. The average nitrogen level of the accessions of 3.1% is lower than those observed by Magat *et al.*, (2009) who recorded an average of 4.737%.

The results of the study showed that all the nutrients with the exception of Fe occurred in minute quantities compared to the USDA- RDI. The average quantity of 12 mg/100g of iron detected was higher when compared to the USDA- RDI for an adult male but less

than that needed by an adult female (that is, about two thirds of the USDA- RDI of required by an adult female). This may be as a result of the drying process since drying foodstuffs under high temperature over a period of time can contribute to nutrient loss.

### 5.3 Molecular Analysis

Biochemical markers assess accurate genetic diversity index (Akhtar, 2001; Rabbani *et al.*, 2001). Protein electrophoresis is a powerful tool for population genetics (Parker *et al.*, 1998). Irwin *et al.*, (1998) reported that closely related accessions are normally located within 80–90% similarity. Crosses between accessions with similarity indices of 80–100% may, therefore, not be recommended. The results of the study on total protein marker revealed that considerable variation was available in the accessions. This is because a sizeable number of the accessions studied had a similarity index of 54 – 77% (figure 4.4). Following the recommendation of Irwin *et al.* (1998), crosses between the accessions BNR 7 and BNR 15 may provide superior genotypes from which useful agronomic types can be selected. In spite of the variation existing among the accessions, some accessions from Ghana were similar which probably may be attributable to a common parent or ancestry. This is likely because most Moringa planting materials in Ghana are collected from cuttings or seeds of plants from relatives, friends and loved ones who recommend the crop due to its numerous benefits. Large scale production from seed is not yet established.

RAPD technique is an effective method of screening for nucleotide sequence polymorphisms randomly distributed throughout the genome of an organism (Subudhi and Huang, 1999) which provides adequate information on the variability of

accessions studied. For the RAPD marker analysis, there was a high level of polymorphism (95.1%) among the accessions of *Moringa oleifera* studied. This may be due to the fact that the variations observed may be variations that exist at the genotypic level. Accession pair E41 and UE49 in cluster A could not be resolved at 91% similarity (figure 4.5) suggesting that they could be closely related or be the same. Accession BNR04 from USA was resolved from the rest of the accessions, forming cluster B whilst accession N17 was also resolved from the other accessions in cluster A showing their uniqueness among the accessions studied (figure 4.5). The observed similarity level of 30% (that is 70% genetic variability) shows that there is a high genetic variability among the 40 *Moringa* accessions analysed. This confirms the ability of RAPD markers to detect variations in the samples. The study compares well with Abubakar *et al.*, (2011) who reported high degree of polymorphism (74%) among 75 *Moringa* accessions from Sudan and Guinea savannah zones in Nigeria. Fu *et al.*, (2003) obtained 51.2% genetic differentiation coefficient in *Changnium smyrnioides* using RAPD marker suggesting high genetic diversity among the populations of these plants. The result from the present study indicates the ability of RAPD to discriminate among *Moringa oleifera* accessions.

In analysing the electrophoresis data, an assumption made is that differences in banding patterns among accession reflect differences in their genetic make-up (Shecter and de Wet, 1975). A combination of two marker systems shows much diversity or variability among the accession studied since they cover a large section of the genome. This confirms that the use of different marker systems tend to give better picture of the genetic relatedness than single marker. The combined dendrogram of total protein and RAPD markers suggest high level of genetic variability since a

similarity coefficient of 37% (that is 63% genetic variability) was observed. Accession BNR 4 was resolved from the rest of the accessions studied forming cluster B. This shows the level of distinctiveness of the accession from the rest of the accessions used in the study. Similarly in cluster A of the RAPD dendrogram, accession N17 was resolved from the rest of the accessions for the combined data of total protein and RAPD analysis. The accessions E41 and UE49 were similar at 90% similarity coefficient suggesting that they were either duplicates or samples with common parents (figure 4.6).

The combined dendrogram of morphological and molecular analysis generated very high variability among the accessions used in the study. A similarity coefficient of 6% (that is 94% variability) was observed in the study (figure 4.7). Again, accession BNR4 in major cluster B was resolved from the other accessions studied showing its uniqueness. Accession pairs AS01 and N03; AS02, N05 and N08; E02 and GA03 were 100% similar and could not be resolved. These tied accessions are the ones that were subjected to morphological analysis but not molecular analysis. From the cluster analysis of RAPD data, combined data from total protein and RAPD as well as combined data from molecular and morphological data, the accessions N17 and BNR04 are the most divergent genotypes among the accessions studied hence crosses between them will provide superior genotypes from which useful agronomic traits can be selected for any Moringa breeding programme (Irwin *et al.*, 1998).

## CHAPTER SIX

### CONCLUSION AND RECOMMENDATIONS

#### 6.1 Conclusion

From the study, it may be concluded that;

- *Moringa oleifera* Lam. accessions studied have considerable amount of variability in morphological characteristics such as leaf colour, petiole pigmentation, stem bark colour and growth habit.
- Cluster analyses of both total protein and RAPD markers revealed variations in the accessions studied.
  - The total protein profile of the accessions studied revealed a moderate level of diversity since almost half of the total accessions analysed could not be resolved at 100% similarity.
  - On the other hand, the RAPD profile distinguished between most of the accessions, resolving almost all the accessions except the pair E41 and UE49 at 91% similarity.
  - A combination cluster analysis of RAPD and total protein markers could still not resolve accessions pair E41 and UE49. These accessions are therefore closely related.
- Regarding the relationship between the accessions, it may be concluded that accessions BNR04 and N17 from United States of America and Northern region of Ghana respectively were the most distinctive accessions in the study.

- These two accessions are the most diverse. A cross between them will generate recombinants from which genotypes with superior and useful agronomic traits could be selected.
- A high variability existed in the micro as well as macronutrient composition of the leaves. This variation offers a good opportunity for exploitation in plant improvement programmes.
  - The minerals Ca, Mg, Na, K, P, Fe, Mn and Zn in Moringa accessions studied were in appreciable quantities and showed marked variation in their concentrations but were less than the USDA- RDI.
  - The crude protein content was high compared to leafy vegetable such as “Kontomire” (*Xanthosoma sagittifolia*) and “Bokoboko” (*Talinum triangulare*) which are commonly consumed locally. The high protein with a fairly high concentration of potassium and iron; make it a potential leafy vegetable that is suitable for fortification of foods.
  - Among the local accessions studied, those from Greater Accra, Northern and Upper East regions exhibited the highest levels of certain macro-nutrients and crude protein. Crosses between these have the potential to concentrate the superior characteristics in new varieties.
- The high concentration of lead detected in the accessions studied suggests that *Moringa oleifera* Lam could have high levels of the metal Lead.

## 6.2 Recommendations

- Studies on agro-morphological characteristics such as yield, 100 seed weight, pod colour, pod length, pod width and number of seeds/pod and reproductive traits such as flower colour, flower aromatic, Days to flowering (from sowing to 50% with inflorescence) should be considered in the future to further ascertain the variability that exist at the phenotypic level.
- Further investigation into lead accumulation in *Moringa oleifera* Lam. as well as the ability of the plant to pick up and accumulate other heavy metals such as cadmium, arsenic, mercury, nickel and chromium in the edible parts of the plant is very necessary and should be carried out.
- The use of more modern molecular markers such as microsatellites or single nucleotide polymorphic (SNP) that easily detect polymorphism even among close relations could be adopted to further investigate the relationships between the accessions especially E41 and UE49 at the genotypic level.
- The accessions that have not been subjected to morphological analysis including accessions E41 and UE49 should be studied to confirm the relationships between them.

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## APPENDICES

**Appendix 1:** Passport data of 36 accessions (seeds) of *M. oleifera* used for morphological studies

| Acc. Code | Town Collected                              | Region Collected | Acc. Collector                      |
|-----------|---|------------------|-------------------------------------|
| AS01      | Kwadaso                                     | Ashanti          | Mr. K. Opare-Obuobi                 |
| AS02      | Adiebra TUC                                 | Ashanti          | Mr. K. Opare-Obuobi                 |
| BA01      | Bomaa Ward B                                | Brong Ahafo      | Mr. E. W. Anyomi                    |
| BA02      | Akrudwa                                     | Brong Ahafo      | Mr. D. Opare-Obuobi                 |
| E02       | Akusa                                       | Eastern          | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| E04       | Mampong                                     | Eastern          | Mr. K. Opare-Obuobi                 |
| E05       | Nsawam prison                               | Eastern          | Mr. A. Bajo                         |
| E10       | Kpong                                       | Eastern          | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| E11       |   | Eastern          | Mr. K. Opare-Obuobi & Mr. Ankara    |
| GA02      | Amasaman                                    | Greater Accra    | Mr. K. Opare-Obuobi & Mr. Ankara    |
| GA03      | Dzorwulu (Methodist church)                 | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA04      | Tifa Daavi                                  | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA07      | Association sch. (Airport residential area) | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA10      | Patrice lumuber road (Opp. Ent. Insurance)  | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA15      | ARS- Nungua                                 | Greater Accra    | Mr. K. Opare-Obuobi                 |
| N03       | Kpalsogu (House no. 23)<br>Tolon Kumbungu   | Northern         |                                     |
| N05       | Shishegu (House no. 861)<br>Tamale metro    | Northern         |                                     |
| N08       | Shishegu (House no. 916)<br>Tamale metro    | Northern         |                                     |
| UE02      | Manga (CSIR- SARI station)                  | Upper East       | Mr. F. Kusi                         |
| UE05      | Bawku central                               | Upper East       | Mr. F. Kusi                         |
| UW01      | Tumu (Agric village)                        | Upper West       | Mr. A. Haruna                       |
| UW02      | Sissala east dist. -Tumu (Market square)    | Upper West       | Mr. A. Haruna                       |
| V01       | Juapong                                     | Volta            | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| V02       | Dzodze                                      | Volta            | Mr. E. Sapei                        |
| BNR04     | USA   | Abroad           | Department of Horticulture, KNUST   |
| BNR05     | Wa  | Upper West       | Mr. Arhin, ADRA-Ghana               |
| BNR06     | USA   | Abroad           | Department of Horticulture, KNUST   |
| BNR07     | South Volta                                 | Volta            | Mr. Nii Marcus Arday                |
| BNR08     | Wa  | Upper West       | Mrs. Rebecca Gyedu Karikari         |

|       |                            |            |                           |
|-------|----------------------------|------------|---------------------------|
| BNR09 | South Volta                | Volta      | Miss Peace Gbecko-Kove    |
| BNR10 | Bolgatanga                 | Upper East | Mrs Grace Tsema           |
| BNR11 | Bolgatanga                 | Upper East | Mrs Grace Tsema           |
| BNR12 | Bolgatanga                 | Upper East | Mrs Grace Tsema           |
| BNR13 | Bolgatanga                 | Upper East | Mrs Grace Tsema           |
| BNR14 | India                      | Abroad     | Mr. Pascal/ Amaglo, India |
| BNR15 | CSRPM- Mampong-<br>Akwapim | Eastern    | CSRPM- Mampong-Akwapem    |

**Appendix 2:** Passport data of 40 accessions (stem cutting) of *M. oleifera* used for biochemical and molecular studies

| Acc. Code | Town Collected                                   | Region Collected | Acc. Collector                      |
|-----------|--|------------------|-------------------------------------|
| E01       | Koforidua  | Eastern          | Mr. K. Opare-Obuobi & Mr. D. Djan   |
| E09       | Asikuma  | Eastern          | Mr. K. Opare-Obuobi & Mr. Ankarah   |
| E38       | Aburi  | Eastern          | Mr. K. Opare-Obuobi & Mr. Ankarah   |
| E40       | Yaw Adepa  | Eastern          | Mr. Opare-Obuobi & Mr. E. O. Ankrah |
| E41       | Ahwodjo  | Eastern          | Mr. Opare-Obuobi & Mr. E. O. Ankrah |
| GA15      | ARS Nungua                                       | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA42a     | Pokuase 1  | Greater Accra    | Mr. A. Bajo                         |
| GA42b     | Pokuase 2  | Greater Accra    | Mr. A. Bajo                         |
| *GA54     | Dzorwulu (Mehodist Church)                       | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA59      | Tifa   | Greater Accra    | Mr. K. Opare-Obuobi                 |
| GA60      | Airport Res. Area                                | Greater Accra    | Mr. K. Opare-Obuobi                 |
| N02       | Kasalga Tamale metro (Shea butter village)       | Northern         | Mr. Agyare                          |
| N10       | Nyankpala (SARI quarters 8) Tolon-Kumbungu       | Northern         | Mr. Agyare                          |
| N15       | Kpalsogu (house no. 27) Tolon-Kumbungu           | Northern         | Mr. Agyare                          |
| N17       | Kpalsogu (house no. 9) Tolon-Kumbungu            | Northern         | Mr. Agyare                          |
| UE46      | Nyorego  | Upper East       | Mr. F. Kusi                         |
| UE47      | Manga-SARI station                               | Upper East       | Mr. F. Kusi                         |
| UE49      | Bolga- Kumbosko                                  | Upper East       | Mr. F. Kusi                         |
| UW01      | Sissala West dist. Pulma village                 | Upper West       | Mr. A. Haruna                       |
| UW02      | Sorbelle village Dam site                        | Upper West       | Mr. A. Haruna                       |
| *UW03     | Sorbelle village - Near Techno-serve farm stores | Upper West       | Mr. A. Haruna                       |

|        |                            |               |                                     |
|--------|----------------------------|---------------|-------------------------------------|
| UW04   | Funsi- Near Police station | Upper West    | Mr. A. Haruna                       |
| V04    | Kpong                      | Volta         | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| V05    | Sokode Gbogame             | Volta         | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| V06    | Sokode Lokoe               | Volta         | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| V08    | Kpeve                      | Volta         | Mr. K. Opare-Obuobi & Mr. P. Mireku |
| BNR 2  | USA                        | Abroad        | Department of Horticulture, KNUST   |
| BNR 3  | Legon                      | Greater Accra | Agric. Science Research farm, UG    |
| BNR 4  | USA                        | Abroad        | Department of Horticulture, KNUST   |
| BNR 5  | Wa                         | Upper West    | Mr. Arhin, ADRA-Ghana               |
| BNR 6  | USA                        | Abroad        | Department of Horticulture, KNUST   |
| BNR 7  | South Volta                | Volta         | Mr. Nii Marcus Arday                |
| BNR 8  | Wa                         | Upper West    | Mrs. Rebecca Gyedu Karikari         |
| BNR 9  | South Volta                | Volta         | Miss Peace Gbecko-Kove              |
| BNR 10 | Bolgatanga                 | Upper East    | Mrs Grace Tsema                     |
| BNR 11 | Bolgatanga                 | Upper East    | Mrs Grace Tsema                     |
| BNR 12 | Bolgatanga                 | Upper East    | Mrs Grace Tsema                     |
| BNR 13 | Bolgatanga                 | Upper East    | Mrs Grace Tsema                     |
| BNR 14 | India                      | Abroad        | Mr. Pascal/ Amaglo, India           |
| BNR 15 | CSRPM - Mampong-Akwapem    | Eastern       | CSRPM- Mampong-Akwapem              |

\* Accession not included in Total Protein studies

### Appendix 3: Buffer Systems and Electrophoresis Solutions

#### a. Sample Buffer (SDS reducing buffer)

| Chemical Composition         | Quantity (ml) |
|------------------------------|---------------|
| Deionised water              | 3.55          |
| 0.5M Tris-Citric HCl, pH 6.8 | 1.25          |
| Glycerol                     | 2.5           |
| 10% (w/v) SDS                | 2.0           |
| 0.5% (w/v) bromophenol blue  | 0.2           |
| *β-mercaptoethanol           | 0.5           |
| <b>Total volume</b>          | <b>9.5</b>    |

\*added to solution prior to use.

## b. Staining Solution

| <b>Chemical Composition</b> | <b>Quantity (ml)</b> |
|-----------------------------|----------------------|
| Coomassie blue              | 0.25g                |
| 50% Methanol                | 90ml                 |
| Glacial acetic acid         | 10ml                 |

## c. Destaining Solution

| <b>Chemical Composition</b> | <b>Quantity (ml)</b> |
|-----------------------------|----------------------|
| Methanol                    | 150ml                |
| Glacial acetic acid         | 50ml                 |

**APPENDIX 4:** Composition of solutions for Gel Preparation

## Stock Solutions and Buffers

## a. Acrylamide/Bis (30% T, 2.6% C)

| <b>Chemical Composition</b>                   | <b>Quantity</b> |
|---|-----------------|
| Acrylamide (29.2g/100ml)                      | 87.6g           |
| N'N'-bis-methylene-acrylamide<br>(0.8g/100ml) | 2.4g            |

## b. 10% (w/v) SDS

10g SDS dissolved in 90ml distilled water and topped up to the 100ml mark.

## c. Gel buffer; 1.5M Tris-HCl, pH 8.8

| <b>Chemical Composition</b> | <b>Quantity</b> |
|-----------------------------|-----------------|
| Tris base (18.15g/100ml)    | 27.23g          |
| Deionised water             | 80ml            |

- d. Stacking gel buffer; 0.5M Tris-HCl, pH 6.8

| <b>Chemical Composition</b> | <b>Quantity</b> |
|-----------------------------|-----------------|
| Tris base                   | 6g              |
| Deionised water             | 60ml            |

- e. 10x Electrode (running) buffer, pH 8.3

| <b>Chemical Composition</b> | <b>Quantity</b> |
|-----------------------------|-----------------|
| Tris base                   | 30.3g           |
| Glycine                     | 144.0g          |
| SDS                         | 10.0g           |

- f. 10% Ammonium persulfate  
100mg of ammonium persulfate dissolved in 1ml of deionised water

- g. Separating gel Preparation (10%)

| <b>Chemical Composition</b>       | <b>Quantity</b> |
|-----------------------------------|-----------------|
| Distilled H <sub>2</sub> O        | 4.1ml           |
| 30% Acrylamide stock              | 3.3ml           |
| Gel buffer stock (pH 8.8)         | 2.5ml           |
| 10% Ammonium per Sulphate (fresh) | 50µl            |
| TEMED                             | 10µl            |

- h. Stacking gel Preparation

| <b>Chemical Composition</b>       | <b>Quantity</b> |
|-----------------------------------|-----------------|
| Distilled H <sub>2</sub> O        | 5.7ml           |
| 30% Acrylamide stock              | 1.7ml           |
| Gel buffer stock                  | 2.5ml           |
| 10% SDS                           | 0.1ml           |
| 10% Ammonium per Sulphate (fresh) | 50µl            |
| TEMED                             | 5µl             |

**APPENDIX 5: Microprep buffer**

| <b>Chemical Composition</b> | <b>Quantity</b> |
|-----------------------------|-----------------|
| DNA extraxct buffer         |                 |
| Nuclei lysis buffer         |                 |
| 5% Sarkosyl                 |                 |
| Sodium Bisulfite*           | 0.4g/100mls     |

\*added prior to use.

**Appendix 6a: ANOVA for tripinnatum leaf length of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|-------|---------|--------|------|-------|
| Rep_I stratum       | 4     | 323.41  | 80.85  | 0.99 |       |
| Region              | 9     | 4901.47 | 544.61 | 6.67 | <.001 |
| Residual            | 36    | 2939.41 | 81.65  |      |       |
| Total               | 49    | 8164.30 |        |      |       |
| L.s.d p< 0.05       | 11.59 |         |        |      |       |

**Appendix 6b: ANOVA for tripinnatum leaf width of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|-------|---------|--------|------|-------|
| Rep_I stratum       | 4     | 243.53  | 60.88  | 0.70 |       |
| Region              | 9     | 4048.98 | 449.89 | 5.20 | <.001 |
| Residual            | 36    | 3112.04 | 86.45  |      |       |
| Total               | 49    | 7404.56 |        |      |       |
| L.s.d p< 0.05       | 11.93 |         |        |      |       |

**Appendix 6c: ANOVA for tripinnatum leaflet length of *Moringa oleifera* accessions**

| Source of variation | d.f. | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|------|---------|--------|------|-------|
| Rep_I stratum       | 4    | 46.50   | 11.62  | 0.53 |       |
| Region              | 9    | 972.56  | 108.06 | 4.96 | <.001 |
| Residual            | 36   | 785.11  | 21.81  |      |       |
| Total               | 49   | 1804.17 |        |      |       |
| L.s.d p< 0.05       | 5.99 |         |        |      |       |

**Appendix 6d: ANOVA for tripinnatum leaflet width of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.    | m.s.  | v.r. | F pr. |
|---------------------|-------|---------|-------|------|-------|
| Rep_I stratum       | 4     | 84.55   | 21.14 | 0.88 |       |
| Region              | 9     | 528.07  | 58.67 | 2.45 | 0.027 |
| Residual            | 36    | 863.57  | 23.99 |      |       |
| Total               | 49    | 1476.18 |       |      |       |
| L.s.d p< 0.05       | 6.282 |         |       |      |       |

**Appendix 7a: ANOVA for iron composition of dried leaf powder of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.      | m.s.      | v.r.   | F pr. |
|---------------------|--------|-----------|-----------|--------|-------|
| Rep_I stratum       | 2      | 7.702E-06 | 3.851E-06 | 1.84   |       |
| Region              | 7      | 1.304E-02 | 1.862E-03 | 890.30 | <.001 |
| Residual            | 14     | 2.928E-05 | 2.092E-06 |        |       |
| Total               | 23     | 1.307E-02 |           |        |       |
| L.s.d p< 0.05       | 0.0025 |           |           |        |       |

**Appendix 7b: ANOVA for manganese composition of dried leaf powder of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.      | m.s.      | v.r.  | F pr. |
|---------------------|--------|-----------|-----------|-------|-------|
| Rep_I stratum       | 2      | 1.703E-06 | 8.517E-07 | 2.41  |       |
| Region              | 7      | 2.102E-04 | 3.002E-05 | 84.80 | <.001 |
| Residual            | 14     | 4.957E-06 | 3.540E-07 |       |       |
| Total               | 23     | 2.168E-04 |           |       |       |
| L.s.d p< 0.05       | 0.0010 |           |           |       |       |

**Appendix 7c: ANOVA for zinc composition of dried leaf powder of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.      | m.s.      | v.r.  | F pr. |
|---------------------|--------|-----------|-----------|-------|-------|
| Rep_I stratum       | 2      | 5.403E-06 | 2.701E-06 | 1.14  |       |
| Region              | 7      | 2.832E-04 | 4.046E-05 | 17.01 | <.001 |
| Residual            | 14     | 3.330E-05 | 2.379E-06 |       |       |
| Total               | 23     | 3.219E-04 |           |       |       |
| L.s.d p< 0.05       | 0.0027 |           |           |       |       |

**Appendix 7d: ANOVA for lead composition of dried leaf powder of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.      | m.s.      | v.r.  | F pr. |
|---------------------|--------|-----------|-----------|-------|-------|
| Rep_I stratum       | 2      | 1.351E-06 | 6.754E-07 | 3.26  |       |
| Region              | 7      | 5.555E-05 | 7.935E-06 | 38.27 | <.001 |
| Residual            | 14     | 2.903E-06 | 2.073E-07 |       |       |
| Total               | 23     | 5.980E-05 |           |       |       |
| L.s.d p< 0.05       | 0.0008 |           |           |       |       |

**Appendix 8a: ANOVA for calcium composition of dried leaf powder leaf of *Moringa oleifera* accessions**

| Source of variation | d.f. | s.s.    | m.s.   | v.r.  | F pr. |
|---------------------|------|---------|--------|-------|-------|
| Rep_I stratum       | 2    | 67.29   | 33.64  | 1.06  |       |
| Region              | 7    | 3334.71 | 476.39 | 15.08 | <.001 |
| Residual            | 14   | 442.37  | 31.60  |       |       |
| Total               | 23   | 3844.36 |        |       |       |
| L.s.d p< 0.05       | 9.84 |         |        |       |       |

**Appendix 8b: ANOVA for magnesium composition of dried powder of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.     | m.s.    | v.r.  | F pr. |
|---------------------|-------|----------|---------|-------|-------|
| Rep_I stratum       | 2     | 4.331    | 2.165   | 0.28  |       |
| Region              | 7     | 1830.316 | 261.474 | 33.72 | <.001 |
| Residual            | 14    | 108.563  | 7.754   |       |       |
| Total               | 23    | 1943.21  |         |       |       |
| L.s.d p< 0.05       | 4.877 |          |         |       |       |

**Appendix 8c: ANOVA for sodium composition of dried leaf powder leaf of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.    | m.s.   | v.r.  | F pr. |
|---------------------|-------|---------|--------|-------|-------|
| Rep_I stratum       | 2     | 4119.0  | 2059.5 | 4.13  |       |
| Region              | 7     | 50364.6 | 7194.9 | 14.42 | <.001 |
| Residual            | 14    | 6985.8  | 499.0  |       |       |
| Total               | 23    | 61469.4 |        |       |       |
| L.s.d p< 0.05       | 39.12 |         |        |       |       |

**Appendix 8d: ANOVA for phosphorus composition of dried powder of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.     | m.s.     | v.r. | F pr. |
|---------------------|--------|----------|----------|------|-------|
| Rep_I stratum       | 2      | 0.002461 | 0.001230 | 0.27 |       |
| Region              | 7      | 0.203541 | 0.029077 | 6.47 | 0.002 |
| Residual            | 14     | 0.062880 | 0.004491 |      |       |
| Total               | 23     | 0.268882 |          |      |       |
| L.s.d p< 0.05       | 0.1174 |          |          |      |       |

**Appendix 8e: ANOVA for potassium composition of dried leaf powder leaf of *Moringa oleifera* accessions**

| Source of variation | d.f.   | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|--------|---------|--------|------|-------|
| Rep_I stratum       | 2      | 149113  | 74557  | 0.16 |       |
| Region              | 7      | 2959814 | 422831 | 0.93 | 0.512 |
| Residual            | 14     | 6344706 | 453193 |      |       |
| Total               | 23     | 9453634 |        |      |       |
| L.s.d p< 0.05       | 1178.9 |         |        |      |       |

**Appendix 9a: ANOVA for nitrogen composition of dried powder of *Moringa oleifera* accessions**

| Source of variation | d.f.  | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|-------|---------|--------|------|-------|
| Rep_I stratum       | 2     | 0.9884  | 0.4942 | 1.13 |       |
| Region              | 7     | 27.8557 | 3.9794 | 9.13 | <.001 |
| Residual            | 14    | 6.0990  | 0.4356 |      |       |
| Total               | 23    | 34.9431 |        |      |       |
| L.s.d p< 0.05       | 1.156 |         |        |      |       |

**Appendix 9b: ANOVA for crude protein composition of dried leaf leaf**powder of *Moringa oleifera* accessions

| Source of variation | d.f. | s.s.    | m.s.  | v.r. | F pr. |
|---------------------|------|---------|-------|------|-------|
| Rep_I stratum       | 2    | 147.20  | 73.60 | 0.86 |       |
| Region              | 7    | 312.63  | 44.66 | 0.52 | 0.805 |
| Residual            | 14   | 1202.88 | 85.92 |      |       |
| Total               | 23   | 1662.71 |       |      |       |
| L.s.d p< 0.05       | ns   |         |       |      |       |

**Appendix 9c: ANOVA for dry matter composition of dried**powder of *Moringa oleifera* accessions

| Source of variation | d.f.  | s.s.    | m.s.   | v.r. | F pr. |
|---------------------|-------|---------|--------|------|-------|
| Rep_I stratum       | 2     | 0.133   | 0.067  | 0.01 |       |
| Region              | 7     | 285.200 | 40.743 | 5.26 | 0.004 |
| Residual            | 14    | 108.454 | 7.747  |      |       |
| Total               | 23    | 393.787 |        |      |       |
| L.s.d p< 0.05       | 4.874 |         |        |      |       |

**Appendix 10: AVRDC- GRSU CHARACTERIZATION RECORD SHEET**

Crop: **Moringa** Accession No. \_\_\_\_\_  
 Sowing Date \_\_\_\_\_ Plot No. \_\_\_\_\_  
 Transplanting Date \_\_\_\_\_ Name \_\_\_\_\_  
 Location \_\_\_\_\_ Species \_\_\_\_\_  
 Origin \_\_\_\_\_

**SEEDLING DATA**

1. Germination period (no. of days from sowing to first germination)
2. Cotyledon Colour  
 1=light green    3=green    5=purplish green    7= purple    9=dark purple
3. Hypocotyl colour  
 1=light green    3=green    5=purplish green    7= purple

**VEGETATIVE DATA**

4. Growth habit  
 1= compact    2= intermediate    3= tall (erect)
5. Plant height (cm) (at flowering stage)
6. Tripinnatum leaf length (cm)
7. Tripinnatum leaf width (cm)
8. Leaf type  
 1= simple    2= compound
9. Leaf arrangement  
 1= opposite    2= alternate    3=pinnate    4=palmate

## 10. Leaflet shape

1=simple      2=compound

## 11. Leaflet arrangement

1= opposite      2= alternate      3=pinnate      4=palmate

## 12. Number of secondary rachis

## 13. Leaf Colour

1=Light green      3=Green      5= Dark green

## 14. Leaf flavour

1=bitter      3=normal      5=strong

## 15. Tripinnatum leaflet length (cm)

## 16. Tripinnatum leaflet width (cm)

## 17. Petiole pigmentation

1=Green      3=Dark Green      5= Purple      7=Dark Purple

## 18. Stem bark colour

1=Whitish      3=Green      5= Brown      7=Dark Brown

**INFLORESCENCE DATA**

## 19. Days to flowering (from sowing to 50% with inflorescence)

## 20. Flower Colour

1= white      2= cream

## 21. Flower aromatic

1=weak      5=Intermediate      7=strong

## 22. No. of Petals

## 23. Petal length

## 24. Petal width

25. No. of Sepals
26. Sepal length
27. Sepal width
28. Height of Pistil
29. Height of Stamen

### **FRUIT DATA**

30. Pod Colour  
1= green      2=greenish brown      3=brown
31. Pod length (cm)
32. Pod width (cm)
33. Number of seeds/pod

### **SEED DATA**

34. Seed Colour  
1=brown      2=dark brown
35. 1000-seed weight (g)
36. Pollen fertility