

# **EVALUATING THE EFFECT OF SOUTH AFRICAN HERBAL EXTRACTS ON BREAST CANCER CELLS**

**Mpho Susan Choene**



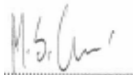
A dissertation submitted in fulfilment of the requirements for the degree of Masters in Science in  
the School of Molecular and Cell Biology, University of the Witwatersrand

Johannesburg, 2012

## DECLARATION

I declare “**EVALUATING THE EFFECT OF SOUTH AFRICAN HERBAL EXTRACTS ON BREAST CANCER CELLS**” to be my work which has not been submitted for any degree or examination at any other University and that all the sources I have used or quoted have been indicated and acknowledged by complete references.

**Mpho Susan Choene**



Signature.....

On this...19.....day of .....July.....2012

## RESEARCH OUTPUT

### Conference Outputs

#### Poster presentations

Mpho S. Choene and Lesetja R. Motadi. Evaluating the anti-tumour properties of South African herbal extracts on breast cancer. SASBMB/FASBMB congress, Drakensberg, 29-1 February, 2012

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#### Original publications and Reviews

Mpho S. Choene and Lesetja R. Motadi

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Wisdom from the forefathers' traditional medicine can induce apoptosis and inhibit cell proliferation. (Review) Biological and Biomedical Reports, 2012, 2(3), 180-189

## **DEDICATION**

This dissertation is dedicated to my mother, Ntshebo, grandmother, Alinah, grandfather, James and to my little brother, Tiisetso who have continuously supported me throughout the course of my studies. You encouraged me even when I felt like giving up.

Thank you!

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## ABSTRACT

In this research we aimed to investigate the anti-proliferative properties of three South African plants: *Kedrostis foetidissima*, *Euphorbia mauritanica* and *Elytropappus rhinocerotis* against breast cancer cells. This was done on the basis of their documented ethno-medicinal use against cancer and other ailments. The plant extracts were screened for cytotoxicity and pro-apoptotic activity against two breast cancer cell lines MCF-7 and YMB-1. With an  $IC_{50} \sim 100 \mu\text{g/ml}$ , *K. foetidissima* was the only extract that exhibited significant cytotoxicity on both cell lines, whilst *E. mauritanica* was cytotoxic to MCF-7 cells only. The cytotoxicity assay was followed by the Annexin-V detection assay to evaluate the occurrence of apoptosis. The results observed suggested that *K. foetidissima* was inducing significant apoptosis on both YMB-1 and MCF-7 cells, whilst *E. mauritanica* was inducing significant apoptosis on MCF-7 cells.

Since both *K. foetidissima* and *E. mauritanica* crude extracts induced apoptosis to MCF-7 cells, they were selected for gene expression studies on MCF-7 using real-time PCR. This was done with the aim of investigating if these extracts were having an effect on the tumour suppressors p53 and RBBP6, which were shown in previous studies to be deregulated in up to 50% of cancers. From the real-time PCR data we observed no changes in the expression levels of these genes following treatment with the herbal extracts. This may suggest that these plants have an effect on other components of the apoptotic pathway other than the tumour suppressors p53 and RBBP6.

The antiproliferative activity observed whilst treating these particular cell lines with *K. foetidissima* and *E. mauritanica* suggests that these South African herbal plants present

themselves as potential future cancer therapeutic agents; however, further studies on these herbal plants need to be performed to validate these results.

**KEYWORDS:** Apoptosis  
Breast cancer  
*Euphorbia mauritanica*  
*Kedrostis foetidissima*  
p53

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## **CHAPTER 1 INTRODUCTION AND LITERATURE REVIEW**

### **1.1 Traditional medicine**

For generations, mankind has believed and still believes in the healing powers of plants, using them as medicine to treat various common infectious diseases. Some of these plants have been used for centuries and are still used even today as a primary source of healthcare. In spite of the increasing usage of modern medicines in some areas, there are still countries that don't have regular access to essential drugs. The use of traditional plants as a great source of medicine has been greatly documented in African countries and other developing countries such as India and China. According to the World Health Organisation (WHO), 70-80% of the world's population use herbal sources as medicine (Chan 2003).

It is only in the 20<sup>th</sup> century that there has been increasing interest in the use of natural resources, especially herbal medicine, as an alternative to orthodox medicine in industrialized countries. This has thus increased interest from pharmaceuticals, and phytotherapeutic research. Scientists started developing an interest in medicinal plants due to their documented antibacterial, antifungal, antiviral and even anti-amoebic properties (McGaw *et al.* 2000). Today, up to 50% of the world drugs are made from natural products or their derivatives (Gurib-Fakim 2006). Some of these drugs include aspirin, quinine and morphine amongst others (Singh and Barrett 2006). The recent surge in natural plants research has also led to production of drugs such as taxol, camptothecin and podofilox which are now being used for chemotherapy. These drugs were derived from parts of various plants (Itokawa *et al.* 2008; Pezzuto 1997). This has been a remarkable progress in elucidating the possible anti-apoptotic properties of plants. As researchers

worldwide continue to explore plants for their anti-cancer properties, breast cancer should also be at the forefront of medicinal plant research, since it is now the leading cause of cancer in females.

## **1.2 Breast cancer**

### **1.2.1 Statistics**

Breast cancer is a malignant tumour that starts in the breast tissue; it usually starts in the ducts or in lobules. It can occur in both males and females; however, male breast cancer is rare. It is the second leading cancer in the world following lung cancer, with an estimated 1.15 million cases a year in 2002 (Parkin *et al.* 2005). The American Cancer Society estimated 182 460 cases (which accounted for 26% of all new cancer cases) for 2008 in the U.S.A alone, resulting in 40 480 deaths (Jemal *et al.* 2008). Breast cancer is primarily a postmenopausal women's disease, with the American National Cancer Institute's Cancer Review estimating the average age of women at risk to be 61 years of age. Approximately 0 cases have being reported for women below 20, but the figures are gradually increasing with every decade (Altekruse *et al.* 2009).

In South Africa, between 1986 and 1992, cervical cancer was the most leading cause of cancer amongst women, but recently breast cancer seems to have overtaken and is now the most common cancer in females (Vorobiof *et al.* 2001; Siegel *et al.* 2011). Compared to the rest of the world, breast cancer cases are relatively low in Africa, with an estimated 65 000 new cases being reported in 2002. The highest incidence was observed in South Africa with about 6474 (or 35 per 10<sup>5</sup>) cases being reported (Parkin *et al.* 2008).

### 1.2.2 Epidemiology

The lifetime risk of an individual developing breast cancer cannot be attributed to one factor alone. A number of variables play a role in the development of breast malignancies. Even though it is fairly difficult to pinpoint one cause as the main risk factor for breast cancer, studies have shown there are certain genetic, reproductive and lifestyle factors that play a major role in its induction and progression. One of these factors is age, with studies finding that the incidence rates double with every 10 years up until menopause, where the rates start to dramatically decline (McPherson *et al.* 2000). Exposure to radiation and use of oral contraceptives may be associated with induction of breast cancer (Ronckers *et al.* 2005; Urban *et al.* 2012).

Very small fraction of breast cancers could be attributed to preventable lifestyle behaviour such as obesity, alcohol consumption, lack of physical activity and long term use of hormone replacement therapy. In several studies conducted to determine the role these factors have on breast cancer incidence, the data appears to be inconsistent, hence it is hard to establish the severity these factors play in breast cancer induction that is if they even do at all (Chen *et al.* 2006; Lorincz and Sukumar 2006; Zhang *et al.* 2007). In developed countries, incidence rates are higher relative to developing countries. However, incidence rates are steadily increasing in developing countries (Key *et al.* 2001). This increase may be attributed to the adoption of a more westernized lifestyle which encompasses a more sedentary lifestyle, delayed childbearing, high alcohol use and a diet high in fat (Bray *et al.* 2004; Porter 2008).

In a lot of breast malignancies, hormones play a major role. It is known that estrogen starts breast tissue proliferation, and studies have now shown that the longer a woman is exposed to estrogen, the greater the risk of developing breast cancer. Hence, women with an early age at menarche

(<12 years), were found to be 2.2 times more likely to develop breast cancer in the future than those with late menarche (Peeters *et al.* 1994). In a different study, women who started menopause late i.e. at the age of 55, were found to have 44% higher breast cancer incidence rates than women who started menopause at 45. This may be due to a 10 year increase in the amount of estrogen the women are exposed to. Exposure to excessive exogenous estrogen plays a role in hyperestrogenia. Hyperestrogenia could trigger breast cancer in two ways: either by breast epithelium tumour promotion or by DNA damage which could be caused by estrogen metabolites directly (Butler *et al.* 2000; Imyanitov and Hanson 2004).

Another important factor which may increase susceptibility to breast cancer is inheritance of susceptibility genes, it has been estimated that 7-10% of breast cancers arise due to inherited mutated genes. The major predisposing factors to breast cancer are breast susceptibility genes, known as BRCA1 and BRCA 2, which are important for early onset of breast cancer (Hedau *et al.* 2004; Imyanitov and Hanson 2004). An initial germline mutation in one of these genes followed by a second mutation leading to the loss of function of the second allele is required for induction of tumorigenesis (Powell and Kachnic 2003). These genetic factors mostly show an autosomal dominant pattern of inheritance, with germline mutations rendering susceptibility to the following generation (Loubser *et al.* 2008; Ponzzone and Baum 1998).

### **1.3 BRCA 1 and 2**

The BRCA genes are thought to be tumour suppressors since the wild type allele is often lost in tumours of heterozygous people (Tutt and Ashworth 2002). It has been suggested that BRCA 1 and BRCA2 genes play a role in a common pathway which controls DNA damage repair and DNA replication fidelity (Powell and Kachnic 2003). They are thought to be involved in repair

of damaged DNA associated with replication by homologous recombination. This they do by interacting with a recombinational repair protein *Rad51*, thereby maintaining genomic integrity (Tutt and Ashworth 2002; Welsch and King 2001). In accordance with their role in DNA repair; a cell with a mutant BRCA gene will not be able to have a functional protein product thereby leading to defects in DNA repair mechanisms. It has been speculated that germline mutations in cancer susceptibility genes BRCA1 and BRCA2 led to an increased risk of developing breast cancer.

Breast cancer patients with mutant BRCA 1 were found to contain up to 2 to 3 fold more chromosomal rearrangements than sporadic cancers (Thangaraju 2000). Abnormalities that arise due to mutant BRCA2 include broken chromosomes and chromatids as well as formation of abnormal structures which mark an abnormal mitotic recombination which has been linked to predisposition to diseases such as Blooms syndrome and also leaves the person with high susceptibility to various types of cancers (Venkitaraman 2002). Ford *et al.* (1998) showed that up to 84% of hereditary breast cancers were linked to mutations in either BRCA1 or BRCA2 whilst only 16% of breast cancer cases were not linked to BRCA mutations.

BRCA1 tumour suppressor gene encodes an 1863-amino acid and is located at 17q21. BRCA1 is implicated in many cellular pathways including transcription, cell cycle, checkpoint control, apoptosis and DNA repair (Venkitaraman 2002). BRCA2 is a breast cancer susceptibility gene located at 13q12-q13 which the product is thought to be involved in monitoring genome integrity and cell cycle progression. BRCA 2-null mice have been reported to have a defect in embryonic cellular proliferation and die in the uterus (Cheung *et al.* 2002).

Though BRCA mutations have been implicated in a lot of breast cancer cases, there are still breast cancers which occur due to deregulation of other genes. A substantial proportion of susceptibility to breast cancer can be attributed to mutations in cell cycle regulation and apoptosis genes.

## **1.4 Cell cycle**

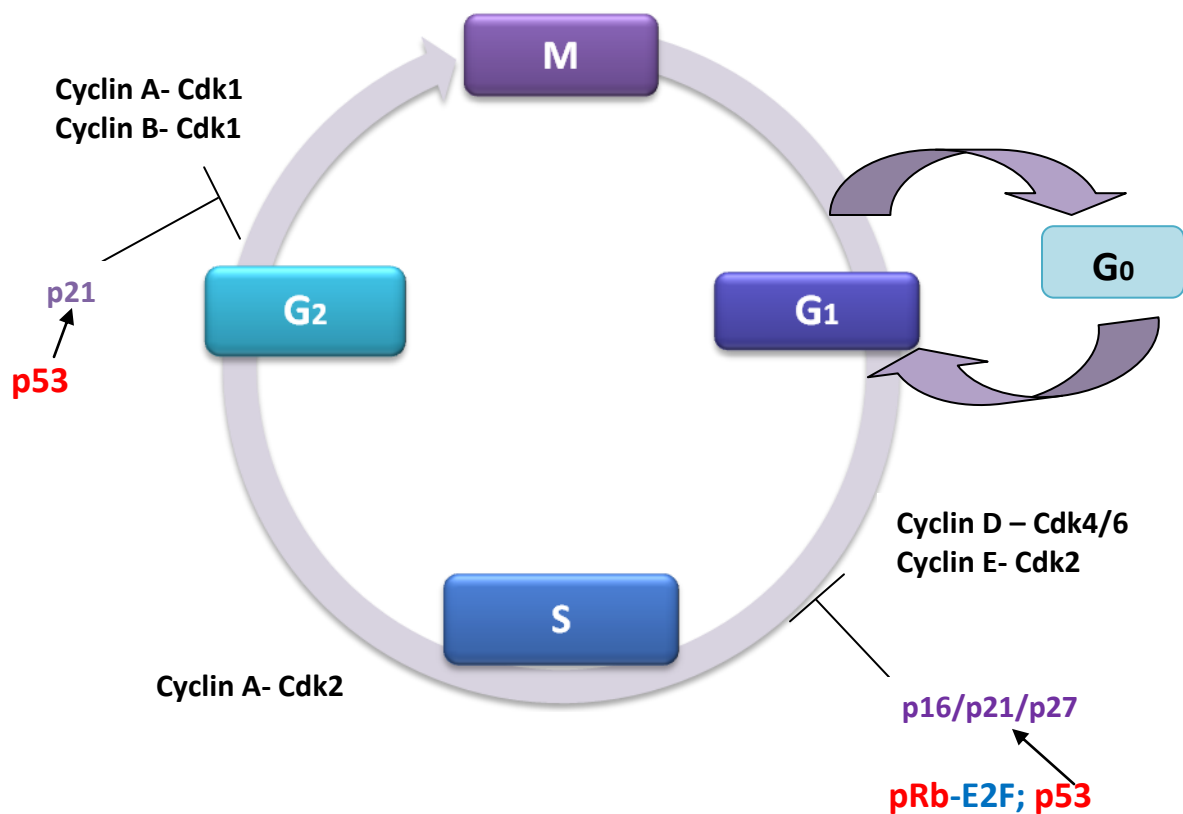
The cell cycle is an important regulator of DNA replication, cell growth and cell division (Schwartz and Shah 2005). It entails a series of highly coordinated steps, these are divided into 4 distinct stages: G1-phase - gap phase, it represents the stage of entry into the cell cycle whereby cells prepare for the S-phase; S-phase- DNA synthesis; G2- the second gap phase, a period of rapid cell growth where cells prepare for mitosis and M phase which represents mitosis, which entails cell division resulting in the formation of 2 daughter cells from each cell (Caldon *et al.* 2006; Schwartz and Shah 2005). It is essential for a cell to complete one phase before proceeding to another to ensure orderly transitions.

One of the most important things is for cells to ensure repair of damaged DNA and prevention of uncontrolled proliferation whilst going through the cell cycle. The most important mechanisms that mammalian cells have developed to deal with constant DNA damage is the development of cell cycle control checkpoints. This is a mechanism in which cell progression through the cell cycle is arrested, to prevent cells with damaged DNA from undergoing DNA replication or mitosis and this is mainly mediated by cyclin-dependent kinases (CDKs) (Kastan and Bartek 2004; Malumbres and Barbacid 2005). CDKs require specific regulatory proteins known as cyclins to be positively regulated and they are negatively regulated by CDK inhibitors (Schwartz

and Shah 2005). Depending on which stage of the cell cycle the cell is in will be the determining factor on which particular CDK will be expressed by its respective cyclin.

During the G1 phase, a signal will induce synthesis of D-cyclins which will interact with CDK4/6, these will form complexes and phosphorylate the tumour suppressor retinoblastoma (*Rb*) (Malumbres and Barbacid 2005; Potten and Wilson 2004). To bring about G1 cell cycle arrest, *Rb* protein interacts with E2F family of transcription factors (Weinberg 1995). Association between CDK2 and cyclin E drives the cell to the S-phase. Cyclin A's interaction with CDK2 will drive the cell through the S-phase. Whilst the cyclin A-CDK1 and cyclin B-CDK1 will get the cells through the S-phase to mitosis transition (Schwartz and Shah 2005) (see Figure 1.1). The two most important checkpoints being the G1 and the G2 checkpoints, the G1 checkpoint ensures that cells with a damaged DNA do not undergo DNA replication and the G2 checkpoint that unreplicated and damaged DNA do not undergo mitosis.

There has been increasing evidence that implicates cell cycle control disruption as a common pathway in the development of breast tumourigenesis. The fidelity of the cell cycle requires a coordinated complex between DNA repair, metabolic adjustments, cell death, chromatin remodeling which are controlled by regulatory surveillance systems (Elledge 1996). Mutations in genes involved in DNA damage responses and mitotic checkpoints pathways, may allow continued proliferation of genes with aberrant DNA, thus leading to increased chances of cancer (Kastan and Bartek 2004). The two most important genes that control the cell cycle that have been implicated in tumourigenesis being the tumour suppressors p53 and pRb, and these will be discussed further in the following section.



**Figure 1.1: Schematic representation of cell cycle regulation indicating G1, S and G2 checkpoints.**

The phase specific cyclin-dependent kinases (CDKs) and their regulatory proteins known as cyclins are shown. The negative regulators of the CDKs are p16, p21 and p27. The tumour suppressor protein pRb in its phosphorylated form is required for progression through the G1 phase whilst p53 is essential for both the G1 and G2 checkpoints (Stewart *et al.* 2001).

## 1.5 Apoptosis

Apoptosis is a fundamental process that occurs throughout the lives of most organisms, from humans to simple organisms such as hydra, to insects, amphibian and fish (Potten and Wilson 2004). It is an active form of programmed cell death which is important for maintaining

homeostasis by ensuring a balance between proliferation of normal cells and non-inflammatory death of damaged cells in adult multicellular organisms (Kerr *et al.* 1972). Deregulation of apoptosis could lead to induction of certain human disease including neurodegenerative disorders such as Alzheimer's and Parkinson's diseases, autoimmune disorders e.g., multiple sclerosis and rheumatoid arthritis and various types of cancers (Fadeel *et al.* 1999). It also plays a crucial role in embryonic development (Kerr *et al.* 1972). It is important for mammalian development and patterning of structures such as organs (brains, eyes) and limbs. Failure of apoptosis regulation could lead to an observable phenotype characterized by various defects (Lang 1997; Haydar *et al.* 1999; Potten and Wilson 2004).

Even though apoptosis may be triggered by various stimuli, it is initiated by two major pathways the intrinsic and the extrinsic pathways (Boatright and Salvesen 2003).

#### (i) *Intrinsic pathway*

The intrinsic pathway is regulated by pro- and anti-apoptotic members of the Bcl-2 family, which will either induce or prevent permeabilization of the outer mitochondrial membrane. Once an apoptotic signal is received, it will lead to activities of signal inducing pro-apoptotic members of the Bcl-2 family leading to permeabilization of the outer mitochondrial membrane (Green and Kroemer 2004). This will lead to release of cytochrome *c* into the cell cytoplasm, which had been sequestered in the mitochondrial intermembrane space (Schuler *et al.* 2000). Cytochrome *c* will then bind to the protein apoptosis-protease-activating-factor 1 (Apaf-1), this complex will also require the binding of dATP as a cofactor to form the apoptosome complex which will be responsible for activation of caspase-9 (Purring-Koch and McLendon 2000). Active caspase-9

will lead to protease cascade which will eventually lead to cleavage of caspase-3 causing apoptosis (Li *et al.* 1997)

#### (ii) *Extrinsic pathway*

The extrinsic pathway involves cell members of the tumour necrosis factor-alpha (TNF- $\alpha$ ) superfamily. These are receptors found on the cell surface, several members have been identified so far and they include TRAIL-1 and TRAIL-2, TNFR-1, CD-95, Fas, DR6 and NGFR (Chen and Wang 2002; Lavrik *et al.* 2005). An appropriate ligand has to bind to one of the receptors on the cell surface; this will lead to recruitment and activation of caspase-8 and -10 which are part of a proteinase family called caspases to form the death inducing signalling complex (Boatright and Salvesen 2003). This will then lead to activation of effector caspases or indirectly activate downstream caspases which will then lead to apoptosis (Schuler and Green 2001).

There are a number of regulatory proteins that have been implicated in influencing cells to undergo apoptosis. However, there are two key regulatory proteins that have been largely conserved in species; they exist as multigene families with multiple homologues. They are known as Bcl-2 (family of inhibitors and promoters of apoptosis) and the p53 tumour suppressor gene and both have been extensively researched in breast cancer (Parton *et al.* 2009). DNA damage could activate p53 which may act as an inducer of apoptosis (Kopper and Petak 2008).

### **1.6 Tumour suppressor genes**

The occurrence of cancer is usually owed to gain of function mutation of proto-oncogenes, converting them to oncogenes, or loss of function mutations which occur in tumour suppressor genes (Garg 2007). The first to be discovered was *RB1*, whose mutations were found to be the

main cause of retinoblastoma (Friend *et al.* 1986). Another one which has been highly studied is p53, both these will be discussed in the following sections.

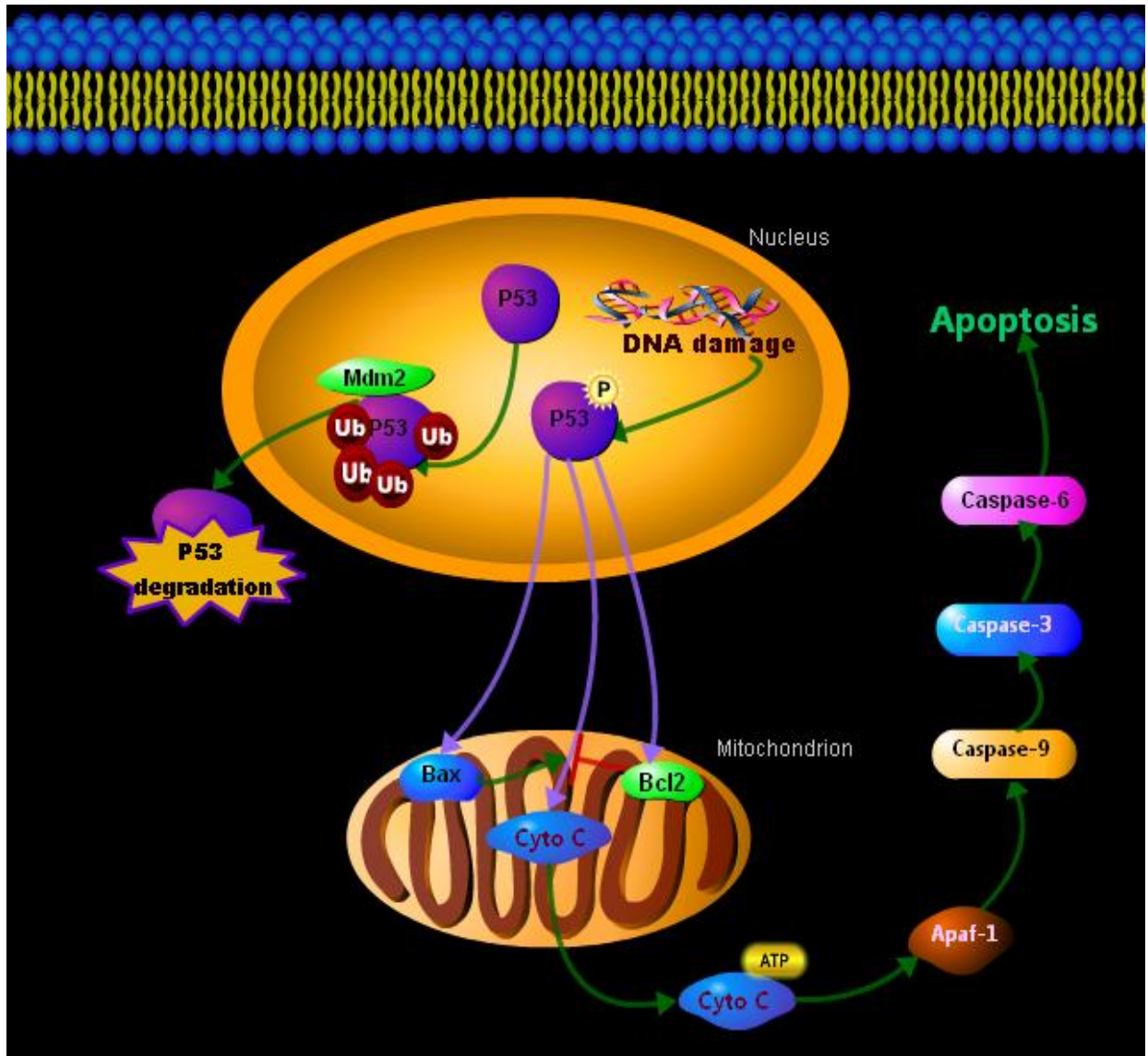
### 1.6.1 p53

p53, one of the most highly studied tumour suppressor, is important for preventing proliferation of cancer cells thereby playing an important role in maintaining genomic integrity. It is due to this crucial role that it is also known as the guardian of the genome. This role is vital in preventing development of various cancers (Barnes and Camplejohn 1996; Potten and Wilson 2004). The gene *p53* encodes a protein p53; it is this protein that plays an important role in cellular processes involved in DNA damage recognition and repair, thereby preventing proliferation of abnormal cells. The p53 protein is normally present in low concentrations and in some cells; it may even be latent or inactive. This is mediated by Mdm2 which binds to p53 and decreases its stability and half-life through the ubiquitin-proteasome pathway, requiring some kind of signal for it to be active (Lee and Lozano 2006; Levine 1997).

Following DNA damage, expression of activated oncogenes or other stressful events in normal cells, p53 is rapidly activated in high amounts which are proportional to the severity of the DNA damage and acts as a checkpoint factor, allowing cells to either undergo apoptosis or cell cycle arrest at the G1-phase (Gasco *et al.* 2002; Giono and Manfredi 2006). p53 may also act as an effector protein, allowing expression of genes involved in cell damage control and repair by interacting with proteins involved in DNA damage recognition and sending signals to those repair genes (Potten and Wilson 2004). The importance of having a functional p53 protein was further shown in mice with a null mutation that was introduced in the *p53* gene and it was found

that mice with a homozygous null mutation in *p53* had a high incidence of spontaneous tumours from a very early age (Donehower *et al.* 1992).

Of over 2500 human cancer tumours that have been analyzed, *p53* has been found to be mutated in over 50% of them (Potten and Wilson 2004). Humans with a mutant germ line *p53* gene develop Li-Fraumeni syndrome and are more easily prone to developing cancer (Hainaut and Hollstein 2000).



(Constructed using protein lounge)

**Figure 1.2: A simplified diagrammatic representation of the p53 pathway.**

In normal cells, p53-mdm2 autoregulatory feedback loop regulates amounts of p53 and keeps them low. Upon exposure to DNA damage, mdm2 is inhibited. This triggers p53, leading to transcription of its downstream targets, e.g. Bax, Bcl<sub>2</sub> and cytochrome c, eventually resulting in apoptosis.

### 1.6.2 *Retinoblastoma tumour suppressor gene (RB)*

Retinoblastoma tumour suppressor gene (*RB*) encodes the retinoblastoma protein (pRb or Rb) which is dysfunctional in many types of cancer. It belongs to a “protein pocket” family whose members possess a highly conserved sequence in that pocket domain; they interact with specific cellular proteins to mediate certain effects (Poznic 2009). The main function of the pRb is believed to be a signal transducer between the cell cycle and its transcriptional machinery (Zhu 2005). This protein regulates the G1 to S transition by binding to E2F and inhibiting promoters of expression genes required for transition through this phase (Poznic 2009). It functions to inhibit cell proliferation and is important for regulation of transcriptional factors that play a role in cellular differentiation, and ensures permanent withdrawal from the cell cycle once cell is differentiated (Poznic 2009; Weinberg 1995).

The protein Rb is thought to be a tumour suppressor; however, the mechanism of how it controls apoptosis is not yet clear. By genetically engineered mouse models, it has been speculated that it may trigger apoptosis by different pathways, the most studied being that as a regulator of E2F (Lee 2002; Zhu 2005). E2F transcriptional factors not only have a role in the cell cycle, but also regulate expression of pro-apoptotic proteins, then due to Rb’s ability to repress E2F, and then it can repress expression of these pro-apoptotic proteins such as cyclin E, CDC6 and CDK1 (MacLeod 2010).

Mutations in *RB* were first established to be the sole cause of retinoblastoma. Mutations in *RB* gene leads to production of a dysfunctional protein which may lead to increased uncontrolled proliferation and cancer. It has been further studied and has been found to be mutated in a third

of human cancers (Lee 2002). The Rb tumour suppressor pathway has been found to be one of the most common disrupted pathways in hepatocellular carcinoma (Mayhew *et al.* 2007).

### 1.6.3 RBBP6

Retinoblastoma binding protein (RBBP6), also known as potential related protein (P2P-R) or p53-associated cellular protein-testes derived (PACT) is a 250-kDa multidomain protein (Chibi *et al.* 2008; Li *et al.* 2007). The tumour suppressors p53 and Rb require specific protein to protein interactions to mediate their respective functions. RBBP 6 protein is one of very few proteins known to interact with both p53 and Rb tumour suppressors (Ntwasa 2008; Simons *et al.* 1997). RBBP6 consists of a conserved N-terminal domain known as the DWNN, a zinc finger and a conserved ring finger like motif (Pugh *et al.* 2006). It was identified by human genomic libraries for its binding properties to the tumour suppressor Rb. It binds to unphosphorylated Rb but not phosphorylated Rb on a region near the C-terminus (Sakai *et al.* 1995).

Even though the physiological role of p53-RBBP6 interaction has not been clearly understood, recent studies have given better insight on this interaction. Li *et al.* (2007) looked at the p53-RBBP6 interaction in mice. They found that RBBP6 is a negative regulator of p53, and it does this through physical interaction with mdm2, its primary regulator leading to p53 ubiquitination. Overall, they found that RBBP6 could inhibit p53 by promoting its degradation.

## 1.7 Plant extracts as anti-cancer drugs

Over the years, research on plants as anti-cancer agents has involved collection and identification of plants, which is done by a botanist, and then crude extracts are prepared from the plant material for biological screening done by phytochemists, followed by isolation of the

active compound or compounds (Balunas and Kinghorn 2005). Even though interest in medicinal plants may only have been for a couple of decades, a number of important drugs have stemmed from the different researchers and today a lot of these drugs made from natural products are widely used worldwide. Of over 50 FDA approved drugs that are used for chemotherapy, many are derived from natural sources e.g., Taxol, Camptothecin, Podofilox, Oncovin and Velban etc. which are derived from the plants *Taxus brevifolia*, *Camptotheca acuminata*, *Podophyllum peltatum* and *Catharanthus roseus*, respectively (Itokawa *et al.* 2008; Pezzuto 1997).

#### 1.7.1 Taxol

Taxol, a diterpene alkaloid was first discovered and isolated in the 1960s from the bark of a tree *Taxus brevifolia* (also known as Pacific yew or Western yew) (Guchelaar *et al.* 1994; Nicolaou *et al.* 1994). It became highly talked about in the 1980s and 1990s in America because it demonstrated efficacy as an anti-cancer agent. It has been approved for use in the treatment against metastatic breast and ovarian cancer in most countries around the world and is currently the best-selling anti-cancer drug. In America, it has also been approved for use in treating Kaposi's sarcoma and non-small lung cancer, its potential to treat others cancers continues to grow (Goodman and Walsh 2001).

Taxol's antitumour mechanism derives from its activity on microtubules. Microtubules are crucial in the cell since they play an important role in the cell's mitotic functions, cellular motility and cell shape. Taxol acts by binding to the microtubules promoting their assembly and stabilising them (Rowinsky *et al.* 1990). The stabilising of the microtubules prevents them from

depolymerising, and thus, this blocks the G1 or M stage of the cell cycle, preventing cells from undergoing cell division (Arnal and Wade 1995).

### 1.7.2 *Camptothecin*

Camptothecin is an alkaloid that can be isolated from the leaves, bark and fruit of *Camptotheca acuminata* tree (also known as Nyssaceae) (van Hengel *et al.* 1992). This plant is native to India but was introduced into the U.S.A. in the 1950s to test for potential anti-tumour, anti-viral and antibiotic properties, it was later found to have high anti-tumour properties (Potmesil 1994; Wall and Wani 1996). Camptothecin acts by binding to and strongly inhibiting DNA topoisomerase I, an enzyme important in DNA replication, thus, preventing DNA and RNA synthesis (Kessel *et al.* 1972; Kjeldsen 1992). The problem with this drug is that it is poorly soluble. It has shown anti-tumour activity on colon and gastric tumours, but it also showed hematological toxic effects. This led to an interest in producing camptothecin analogues which are more soluble. To date, two camptothecin analogues have been approved by the FDA. Topotecan is one such analogue, it is water soluble and has been found to be effective against many tumours including ovarian and colon tumours. The second is irinotecan, which is a more potent camptothecin analogue; it is primarily used in the treatment of colorectal cancers (Hind 2008; Pizzolato and Saltz 2003).

### 1.7.3 *Podofilox*

Podofilox (podophyllotoxin), and its derivatives etoposide and teniposide are extracted from rhizomes of Berberidaceae family *Podophyllum peltatum*, *Podophyllum emodi* or *Podophyllum hexandrum* (Giri and Narasu 2000). Podofilox has shown to be effective in the treatment of genital warts (*Condyloma acuminatum*). These warts are as a result of a sexually transmitted

disease caused by the human papillomavirus (HPV), which has strains that have the potential to cause cervical and genital cancer (Beutner *et al.* 1989; Bonnez 1994). Etoposide is used to treat leukemia, ovarian cancer and Hodgkin's disease. Teniposide is used to treat leukemia, brain and bladder tumours (Giri and Narasu 2000). Podophyllotoxin acts by inhibiting microtubule assembly at the beginning of metaphase thereby preventing cell division. The derivatives act in a different manner, they prevent DNA replication at the S-phase and they also bind to and inhibit DNA topoisomerase II (Giri and Narasu 2000).

#### 1.7.4 Velban (*vinblastine*), *vincristine*, *vinorelbine* and *vindesine*

Vincristine, and vinorelbine and their synthetic derivatives vindesine and vinblastine are vinca alkaloids isolated from the periwinkle plant *Catharanthus roseus* (Hait *et al.* 2007; Ngan *et al.* 2000). These vinca alkaloids are potent inhibitors of cell proliferation and have been found to exhibit different cytotoxicities to different cancers. Vinblastine is used to treat non-Hodgkin's lymphoma, testicular and breast cancer. Vincristine is effective against Hodgkin's disease and paediatric cancers. Vinorelbine is effective against metastatic breast cancer and ovarian cancer and has been approved for use against non-small lung cancer. (Aniszewski 2007; Hait *et al.* 2007). Vindesine has been found to be effective against metastatic breast cancer (Hansen and Brickner, 1984). Vinca alkaloids act by binding to depolymerized mitotic spindle microtubules, preventing their assembly (Hait *et al.* 2007; Ngan *et al.* 2000). This then brings about mitotic arrest at early metaphase (Weaver and Cleveland 2005).

## 1.8 AIMS AND OBJECTIVES

The aim of this study was to investigate the anti-tumour properties of 3 southern African plants on breast cancer cell lines. For those that exhibit a significant level of cytotoxicity, their mechanism of death induction to the cells was evaluated, to determine whether death was occurring via apoptosis or necrosis. Since apoptosis is the ideal method of death induction, we aimed to see if these plant extracts had an effect on some of the major genes involved in apoptosis being p53 and RBBP6. This was done with the aim of assisting in developing future targeted therapeutic strategies that prevent deregulation of these genes, thereby allowing their regulated expression to induce apoptosis in potential breast cancer cells.

### Objectives of the study

- To evaluate the antiproliferative activity of three crude plant extracts: *Kedrostis foetidissima*, *Euphorbia mauritanica* and *Elytropappus rhinocerotis* against two human breast cancer cell lines namely: MCF-7 and YMB-1.
- To determine whether apoptosis is the induced mode of cell death by the extracts on the breast cancer cell lines.
- To determine if the herbal extracts that exhibit cytotoxicity on the breast cancer cell lines affect the apoptotic genes p53 and RBBP6.

## **CHAPTER 2 MATERIALS AND METHODS**

### **2.1 Materials**

#### *2.1.1 Breast cancer cell lines*

The following breast cancer cell lines were used in this study: YMB-1 (human breast carcinoma, polygonal epithelial cell line) and MCF-7 (human breast adenocarcinoma, mammary epithelial cell line). YMB-1 cells are aggressive squamous cells whilst MCF-7 cells are less aggressive adenocarcinoma, both cell lines were obtained from Japan Health Resource Centre. The two cell lines were included to test if the herbal extracts would induce the same effect on different cell lines.

In a study of this nature, it would have been ideal to include a normal breast cell line. However, this was not possible for the following reasons. Normal breast cells undergo a limited number of cell divisions and readily enter senescence under *in vitro* conditions. This clearly poses a problem for i) the maintenance of large quantities of uniform cell populations and ii) when the primary goal of the analyses is to assess programmed cell death. In attempting to address this issue, some researchers have made use of oncogenic (viral/cellular) or carcinogenic e.g., benzo[a]pyrene (B[a]P) transformation to give the cells “extended life”. It can then be argued, however, that this defeats the whole purpose of keeping the cells “normal” (Wazer *et al.* 1995). Nevertheless, the herbal extracts were screened for cytotoxicity on MRC5 fibroblast cells in the lab by Thafeni (unpublished) and results are shown in appendix D.

### 2.1.2 Plant collection

The three herbal extracts used in this study: *Kedrostis foetidissima*, *Euphorbia mauritanica* and *Elytropappus rhinocerotis* were collected from various locations around South Africa. The specimens were obtained from the School of Molecular and Life Sciences, University of Limpopo.

### 2.1.3 Plant extraction

The stems and leaves were collected, washed and frozen. The frozen plant material was ground to a fine powder in liquid nitrogen using a warring blender. Once ground, the plant material was weighed and extracted using absolute methanol (1 g/10 ml, w/v) at room temperature for 24 hours. The resulting extract was filtered through a Whatman filter paper, and then the filtrate was dried at 40°C under low pressure using a Büchi rotavapor R-205 (Büchi Labortechnik AG, Switzerland). Once dried, the extract was weighed and dissolved in 100% dimethyl sulfoxide (DMSO, Sigma) to the desired concentration and stored as a stock solution in an airtight container at -20°C until use.

### 2.1.4 Cell culture expansion

MCF-7 breast cancer cells were routinely cultured in 25 cm<sup>2</sup> tissue culture flasks with minimum essential media (MEM with L-glutamine, Sigma) supplemented with 10% fetal bovine serum (FBS, Sigma) and 1% penicillin/streptomycin (Sigma). YMB-1 cells were cultured in 25 cm<sup>2</sup> tissue culture flasks with RPMI-1640 (Sigma) media supplemented with 10% FBS and 1% penicillin/streptomycin. The cells were incubated at 37°C, 95% humidity and 5% CO<sub>2</sub>.

## 2.2 Experimental assays

### 2.2.1 MTT assay

The MTT {3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide} assay is a simple colorimetric assay used to measure cell cytotoxicity, proliferation or viability. MTT is a pale yellow substrate which enters live cells and passes into the mitochondria to produce a dark blue insoluble formazan. It is converted to formazan by cleavage of the tetrazolium ring by dehydrogenase enzymes of viable cells. The formazan requires an organic solvent to be solubilised; the solubilised formazan can then be measured on a spectrophotometer (Mosmann, 1983). Since MTT can only be reduced in metabolically active cells, it can thus be used to determine the amount of viable cells versus the dead ones. For this study, this assay was used to measure the effect of liquid plant extract on breast cancer cell viability.

Ninety six well tissue culture plates were used to culture cells for the MTT assay. Into each well,  $5 \times 10^3$  cells in 90  $\mu$ l of media were seeded. These were then incubated overnight. Cells were treated with varying concentrations of the herbal extracts (10, 30, 50 and 100  $\mu$ g/ ml, w/v); these were prepared from 40 ng/ml stock solution by diluting with either supplemented MEM or RPMI-1640 media. A non-treatment control (cells, media and later MTT) and a blank (only cells and media) were included. Cells were treated over a period of 24 hours. This was followed by addition of 10  $\mu$ l of MTT (Sigma) (prepared to 5 mg/ml 1x PBS at 37°C) into each well except for the blank, these were further incubated for 4 hours. To dissolve the formazan crystals formed, 90  $\mu$ l of DMSO was added into each well and the plate was read using Bio-Rad Microplate reader at an absorbance of 570 nm.

Percentage Cell Viability was calculated as follows:

$$\% \text{ Cell viability} = \frac{\text{Absorbance of treated cells} - \text{Absorbance of blank}}{\text{Absorbance of untreated cells} - \text{Absorbance of blank}} \times 100$$

$$\text{Absorbance of untreated cells} - \text{Absorbance of blank}$$

### 2.2.2 Flow cytometry

The flow cytometer is highly sophisticated equipment used to analyse individual cells in heterogeneous populations. It allows thousands of cells to pass through a light beam every second (Potten and Wilson 2004), and then it can distinguish these cells based on size and health amongst other parameters. This technique was employed following the MTT assay to investigate whether the herbal extracts were inducing cytotoxicity to the cells by either necrosis or apoptosis. Annexin V (Ambion Annexin V-FITC Apoptosis Detection Kit) was used to quantitate the percentage of cells undergoing apoptosis, necrosis and viable cells. Annexin V has high calcium dependent affinity to phosphatidylserine residues; these residues are normally embedded in the cytoplasmic plasma membrane in healthy cells but are translocated to the surface of the cells during apoptosis. Therefore, Annexin V can bind to these residues acting as a probe to detect and measure apoptosis.

MCF-7 and YMB-1 cells ( $1 \times 10^5$ ) were incubated overnight. They were then treated with plant extract at a concentration that induced ~50% cell death and incubated for 24 hours. Suspended YMB-1 cells were collected by centrifugation whilst attached MCF-7 cells were detached by trypsinization (Trypsin-EDTA, Sigma). MCF-7 cells were rinsed twice with 2 ml MEM and YMB-1 with RPMI-1640 at 37°C. The cells were resuspended in 500 µl of 1x Annexin V

Binding Buffer. This was followed by addition of 5 µl of Annexin V-FITC and 5 µl propidium iodide and 5 minutes incubation. The cells were then analysed by flow cytometry.

### 2.2.3 RNA extraction

Since RNA is highly susceptible to degradation, safety measures were taken throughout the processing steps to minimize its degradation. Working surfaces, glassware, pipettors and gloves were sprayed with RNaseZAP (Sigma) to eliminate RNase. RNA was isolated from treated and untreated cells using the High Pure RNA Isolation Kit (Roche). This kit is designed to extract total RNA from cultured cells; it includes a DNase digestion step which aims to remove all contaminating DNA from the RNA. Before RNA extraction, cultured cells were rinsed twice with 2 ml 1x PBS (Sigma) at 37°C then RNA was extracted using RNA Isolation Kit according to manufacturer's specifications, (see Appendix A1). Resulting RNA was quantified using a nanodrop (NanoDrop technologies, USA), reading was taken at 260A, A260/A280 ratio of ~2.0 was regarded as pure.

### 2.2.4 Reverse transcription

For one step Real time PCR (RT-PCR), mRNA had to first be reverse transcribed to cDNA. This process was conducted following RNA extraction and it employs the use of specific primers with sequence complementary to the mRNA site where reverse transcription is meant to initiate. Extracted RNA was reverse transcribed using the ImProm-II Reverse Transcription System (Promega). Sequence specific primers were selected for use to generate specific cDNAs from the RNA target. The reverse transcription cocktail (see Appendix A2) was prepared in PCR tubes to a final volume of 20 µl. The tubes with the reverse transcription cocktail were briefly vortexed

then placed into the Multigene Gradient Thermal cycler system (Labnet International, Inc). The RNA was reverse transcribed under the conditions explained in Appendix A3. The resulting cDNA was used for one-step Real Time (RT)-PCR.

#### 2.2.5 *Real Time PCR (RT-PCR)*

Real time polymerase chain reaction is the most sensitive technique for simultaneous mRNA detection and quantification. It has been the preferred method of choice for quantitating changes in gene expression patterns. It can be used to quantify mRNA levels from much smaller samples, such as mRNA from a single cell.

To monitor cDNA amplification in RT-PCR, SYBR Green JumpStart Taq Ready Mix (Sigma) was used in our study. It contains JumpStart Taq DNA polymerase and only requires addition of template, primers and water. It is a fluorescent intercalating dye which presents the simplest and cheapest way of detecting PCR product in real time. The dye will only fluoresce when bound to double stranded DNA, thus allowing for fluorescence to increase as the number of double stranded DNA increases, this allows for DNA concentration to be quantified at each cycle. The real-time PCR cocktail is explained in Appendix A4.

Quantitative analysis of the mRNA transcripts was carried out using the Roche lightcycler. The parameters on the machine were set as follows:

**Table 2.1: Real Time PCR parameters.**

| Analysis          | Cycle  | Segment              | Temperature in °C | Hold time  |
|-------------------|--------|----------------------|-------------------|------------|
| Enzyme activation | 1      | 1                    | 95                | 2 minutes  |
| Quantification    | 35- 45 | <u>Amplification</u> | 95                | 45 seconds |
|                   |        | Annealing            | 55                | 45 seconds |
|                   |        | Extension            | 72                | 45 seconds |
| None              | 1      | Final extension      | 72                | 10 minutes |
| Melting           | 1      | <u>Melting curve</u> |                   |            |
|                   |        | Denaturation         | 95                | 10 seconds |
|                   |        | Annealing            | 65                | 15 seconds |
|                   |        | Extension            | 95                | 0.1°C/Sec  |
| None              | 1      | Cooling              | 40                | 30 seconds |

## **CHAPTER 3 SCREENING INDIGENOUS SOUTH AFRICAN PLANTS FOR CYTOTOXICITY ON BREAST CANCER**

### **3.1 Introduction**

South Africa has a remarkably rich flora biodiversity, comprising of about 8% (20 000) of the world's plant species. Of these, about 20% (3689) are used as medicine, and from these, about 350 are commonly used and traded medicinal plants (Cherry 2005; Van Wyk *et al.* 2008). A great proportion of black South Africans rely on traditional medicine, with about 70-80% consulting with traditional healers as a source of primary healthcare. Traditional healers use various traditional indigenous plants to heal different ailments. They either use the roots, stems or leaves of individual plants or combinations of a number of these plants.

However, a great number of people harvest their own plants or obtain them from local *muthi* vendors. This vast usage of medicinal plants stems from the fact that these plants are easily accessible or can be obtained at a very affordable price. Indigenous knowledge on the use of these medicinal plants has been passed down from generation to generation, providing great expertise on their ethnobotanical use. This has allowed for these natural plant medicines to have been tested for efficacy and side effects throughout human history. The obstacle faced however, is little or no knowledge on the molecular mechanisms of active compounds and the chemical profiles of these plants (Lee *et al.* 2010). Hence, it is imperative for those plants which have been used traditionally as medicine for treatment of various diseases, that they undergo pharmacological and biological studies to test for their activities and efficacy.

### **3.2 *Kedrostis foetidissima***

#### *3.2.1 Classification*

*Kedrostis foetidissima*, also called *Utuvishe* in Xhosa belongs to a group of plants known as cucurbitacins. Cucurbitacins are a group of bitter tasting plants mostly found in the plant family Curcubitaceae. There are however, a few other cucurbitacins that may be found in other plant kingdoms (Miro 1995). Cucurbitacins are a fairly large family comprising about 130 genera and 900 species (Jeffrey 1980). The characteristic constituent of this family of plants are cucurbitacins, these are tetracyclic triterpenoids derived from the skeleton of these plants. Cucurbitacins are all named after successive letters of the alphabet from A to R. Cucurbitacins B and D are the most common in the plant kingdom. A previous study identified cucurbitacins B, D, E and I as present in *K. foetidissima* (Miro 1995).

#### *3.2.2 Botanical description*

*Kedrostis foetidissima* is a leguminous herb. It is a succulent perennial climber with a very unpleasant smell. It grows to a height of between 0, 3-3 metres (Figure3.1).

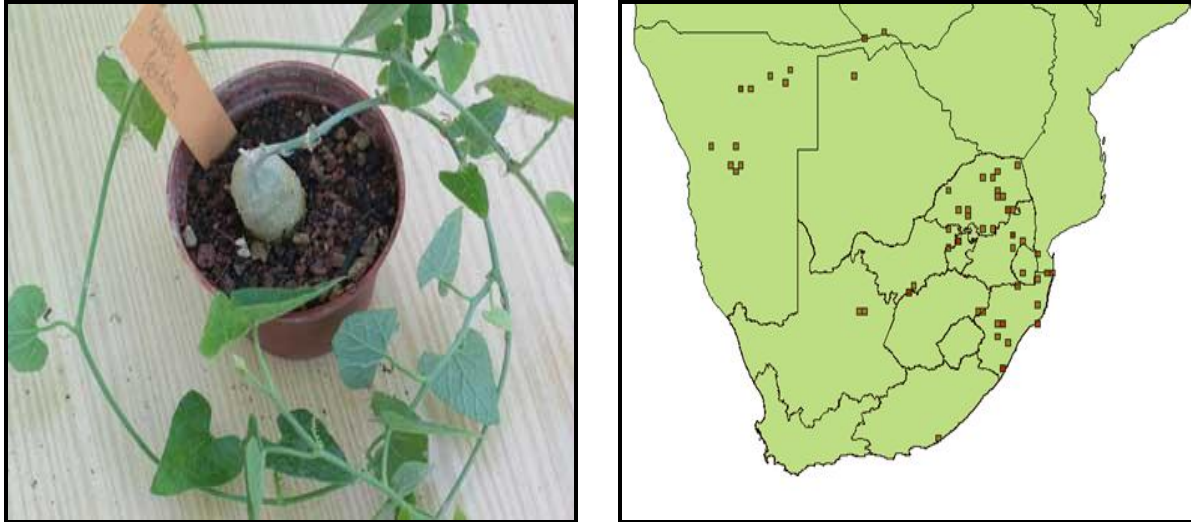
#### *3.2.3 Geographical distribution*

The distribution of *K. foetidissima* stems in the southern African region including Namibia, Botswana through to Gauteng, KwaZulu-Natal, Limpopo, Mpumalanga, Northwest and Northern Cape [South African National Biodiversity Institute (SANBI)].

#### 3.2.4 Traditional medicinal use

Fruits belonging to the cucurbitacins family have been used for centuries due to the wide range of biological activities they exhibit. They are useful for human health due to their ability to act as natural laxatives, to treat liver diseases and inflammatory response such as rheumatism amongst others. Other species have been found to exhibit insecticidal, antifungal and antibacterial properties. These plants have been of great interest in scientific research and a number of compounds from this group have been investigated for antitumour activities and positive effects on cardiovascular health (Miro 1995; Rahman 2008).

*Kedrostis foetidissima* has been used by tribes throughout Africa to treat various human and livestock ailments. It is used by the Masai people in Kenya to treat cattle suffering from bloat; crushed plant is feed to the ailing cattle; they also use it to treat children suffering from measles, the crushed plant is added to bath water of the sick children (Ole-Miaron 2003). The Zay people in Ethiopia use it to treat chest pains (Giday *et al.* 2003). Tanzanians use it to treat measles (Otieno *et al.* 2007). In South Africa, especially in the Xhosa culture, an infusion of the tuber is used as a ritual wash to bring luck, a piece can also be put under the tongue during times of trouble e.g., in court cases (Cocks and Dold 2006)



**Figure 3.1: *Kedrostis foetidissima* (Cucurbitaceae).**

A perennial climber with a simple tendril (left). Geographical distribution of *K. foetidissima* in Southern Africa (right). (<http://www.bihrmann.com/audiciforms/SUBS/ked-foe-sub.asp> and SANBI)

### **3.3 *Euphorbia mauritanica***

#### **3.3.1 Classification**

*Euphorbia mauritanica* belongs to a genus called *Euphorbia*. This genus belongs to a family of plants known Euphorbiaceae. Euphorbiaceae is the most diverse genera with over 2000 species. Of these 2000 species over 200 can be found in South Africa alone (Pienaar 1984; Webster 1994; Zimmermann 2010).

#### **3.3.2 Geographical distribution**

The *Euphorbia* genus can be found all over the world, e.g., Arabia, Morocco, Canary Islands and South Africa amongst others (Jacobsen 1946). They occur mostly in temperate regions, with

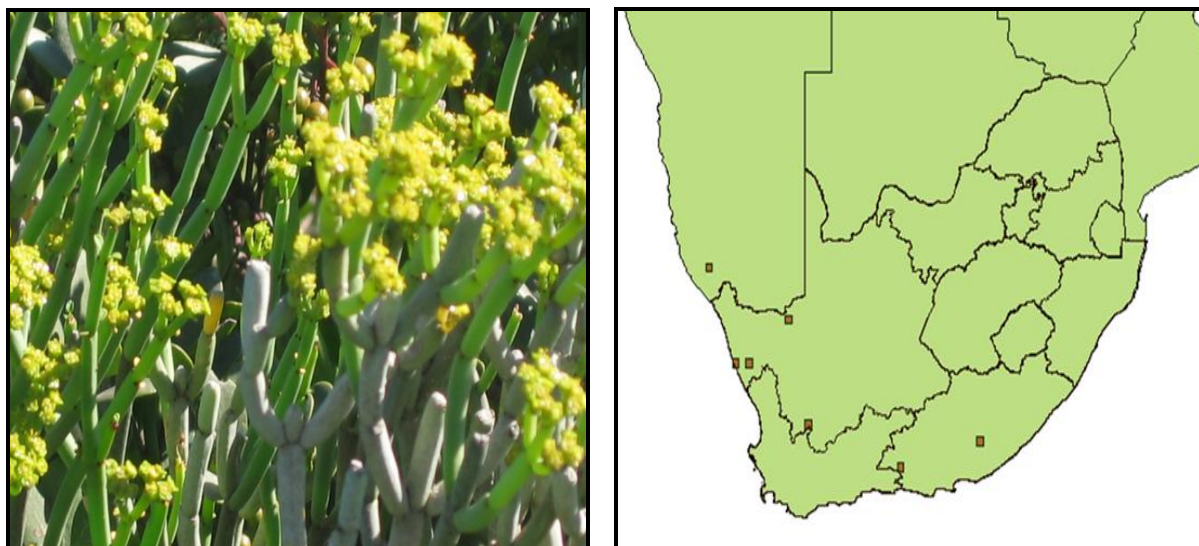
most of the succulent species being indigenous to South Africa. *Euphorbia mauritanica* is mostly dominant in KwaZulu-Natal, Northern Cape, Western Cape, Eastern Cape, Free State as well as Namibia and Lesotho (Pienaar 1984).

### 3.3.3 Botanical description

Euphorbiaceae display a wide range of growth morphologies ranging from annual and perennial herbs, succulents to trees and shrubs. *Euphorbia mauritanica*, also known as *Yellow milk bush*, *Gifmelkbos* and *Jakkalskos* is a freely branched succulent shrub that produces yellow flowers in spring. It can grow to a height of between 0,07 -2,74 metres (Figure 3.2). The leaves are up to 1,3 cm long, which are only on young growing stems. It has smooth green branches which produce white milky latex. It produces fruit with a three chambered capsule, with each capsule containing a seed (Pienaar 1984; Webster 1994; Botha and Venter 2002; Zimmermann, 2010).

### 3.3.4 Traditional medicinal use

Plants of the Euphorbiaceae family have a long history of use as traditional medicine. In Africa and Fiji, a decoction or maceration of some Euphorbiaceae plants are used to treat asthma, cough, diabetes, cure ear-ache, bowel complaints and infertility (Singh 1986; Igoli *et al.* 2005). Some have been used for centuries as traditional medicine against cancers and tumours (Rizk 1987). In countries like Egypt, *Euphorbia* species are widely used in the treatment of stomach, uterus, liver and kidney cancer (Sayed 1980). The root of *E. mauritanica* is infused in boiling water and used to treat toothache. *Euphorbia mauritanica*'s latex is used as treatment for warts (van Wyk *et al.* 2008).



**Figure 3.2: *Euphorbia mauritanica* (Euphorbiaceae).**

A shrub producing yellow flowers in spring (left). Geographical distribution of *E. mauritanica* in southern Africa (right) ([http://sophy.u-3mrs.fr/Afriqsud/Photo-cpAFS /E/](http://sophy.u-3mrs.fr/Afriqsud/Photo-cpAFS/E/)

*Euphorbia\_mauritanica* and SANBI)

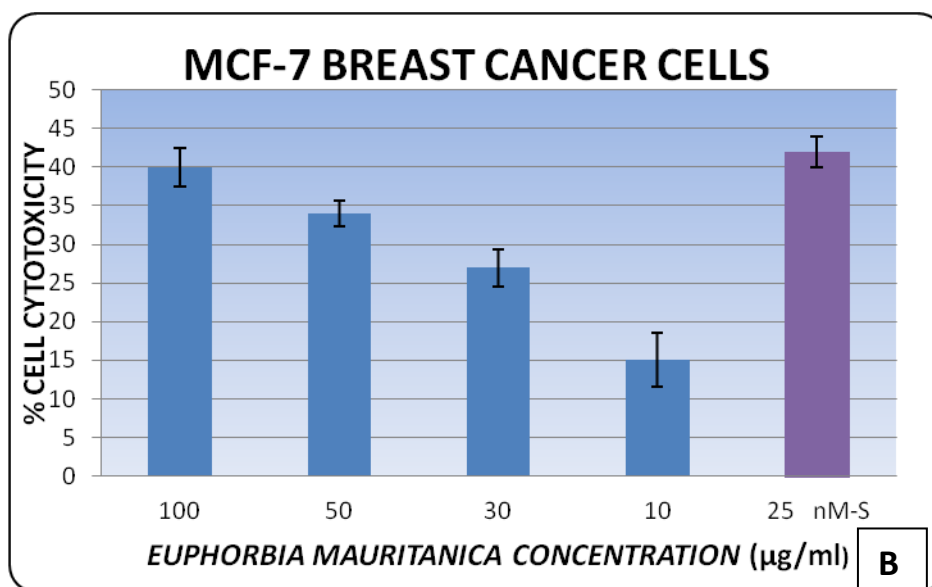
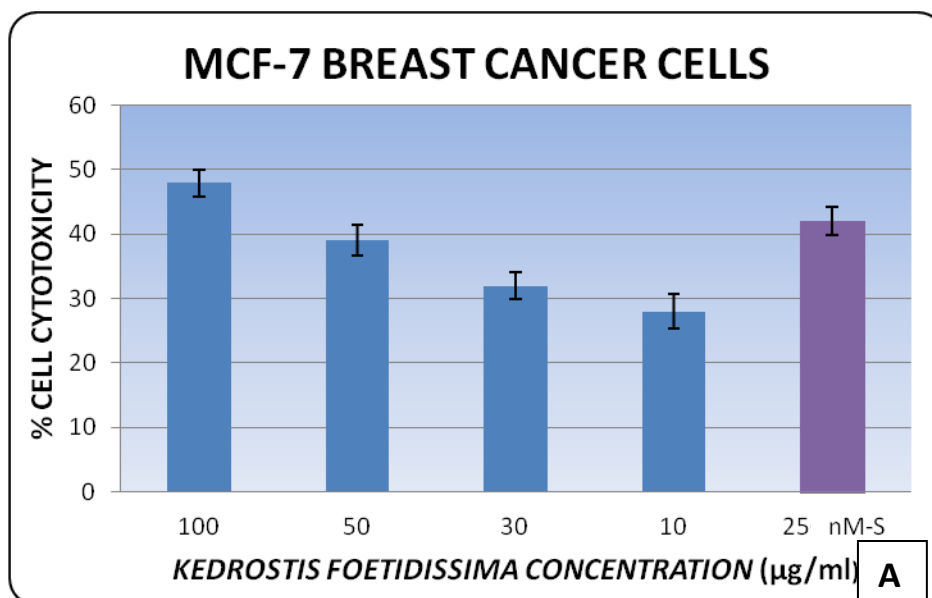
From section 1.5, it is evident that plants have shown great potential in anti-cancer therapeutics. South Africa with its huge biodiversity is also playing a role in production of medicine used around the world. Some of the most popular medicine such as Cape aloe (*Aloe ferox*) and devil's claw (*Harpagophytum procumbens*) are indigenous to South Africa (van Wyk 2008). It is now up to scientists to explore other indigenous plants, especially those that have been used for centuries by South African people to treat different ailments for potential anti-cancer activities. For this study, we aimed to investigate the possible antiproliferative properties of the 3 southern African herbal extracts, namely: *Kedrostis foetidissima*, *Euphorbia mauritanica* and *Elytropappus rhinocerotis* against breast cancer cells. This was done by conducting an MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide) assay, and the IC<sub>50</sub> was

determined. All experiments were repeated at least five times (five assays with eight replicates) to ensure accuracy of the results.

### **3.4 Results and discussion**

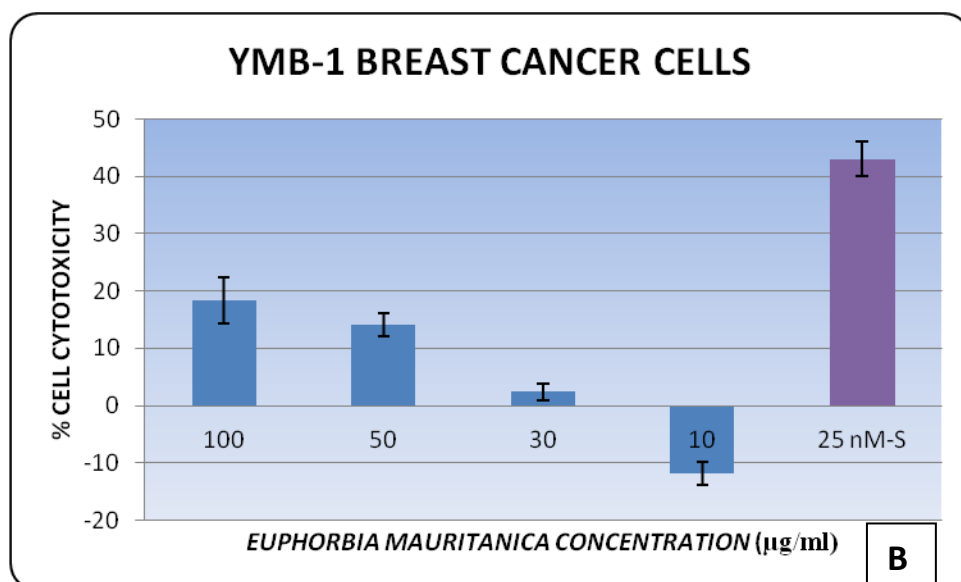
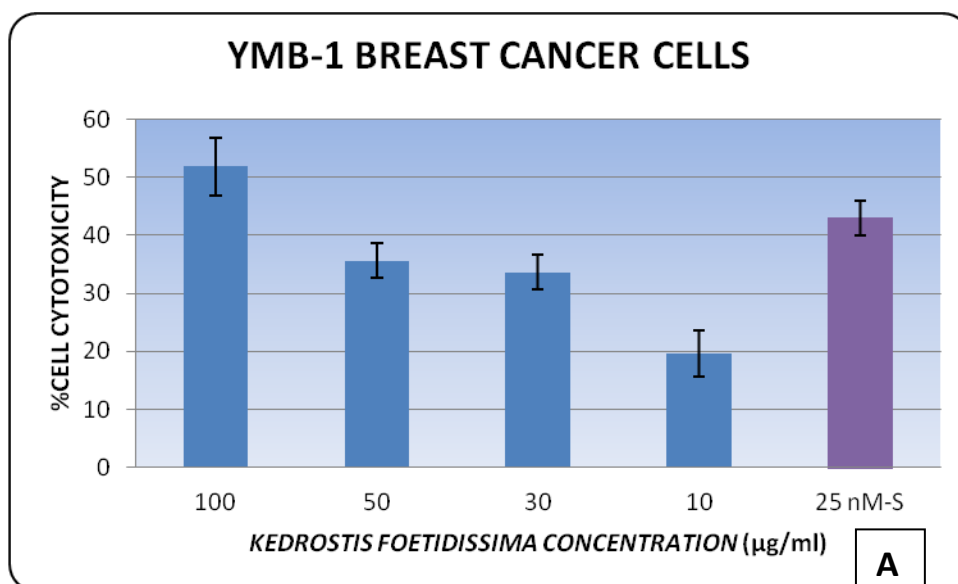
MTT assay is a cell cytotoxicity or viability assay. The three herbal extracts were screened for their possible antiproliferative effect against YMB-1 and MCF-7 breast cancer cell lines. The MTT assay method used is described in Chapter 2, Section 2.2.1. Results obtained suggest the crude extracts of the plant *Elytropappus rhinocerotis* did not exhibit any antiproliferative activity against either MCF-7 or YMB-1 breast cancer cell lines, showing very minimal activity at the highest concentration tested of 100 µg/ml, following 24 hour treatment (see Appendix A).

Results observed for *E. rhinocerotis* in this study correlate with those obtained in a study by Essack (2006) whereby the cytotoxicity of *E. rhinocerotis* on MCF-7 cells was tested; the results obtained in that study suggested that the herbal extract exhibited insignificant antiproliferative activity. At a concentration of 5 mg/ml, cell death of less than 20% would be considered negligible (Essack 2006). It is interesting to note that they studied the herbal extract at high concentrations of between 0,625-5 mg/ml whilst in our study the concentrations ranged between 10-100 µg/ml. This may suggest that *E. rhinocerotis* generally does not have any antiproliferative properties on MCF-7 and YMB-1 cell lines; however, further studies testing much higher concentrations of this plant extract may be required to completely rule out its cytotoxicity against MCF-7 and YMB-1 breast cancer cells.



**Figure 3.3: Cytotoxicity effects of the herbal extracts *Kedrostis foetidissima* A) and *Euphorbia mauritanica* B) on MCF-7 breast cancer cells.**

Cells were treated for 24 hours at increasing concentrations of 10, 30, 50 and 100 µg/ml using the plant aqueous extracts. 25 nM-S represents the positive control, staurosporine, at a concentration of 25 nM. n = 5, p < 0.05.



**Figure 3.4: Cytotoxicity effects of the herbal extracts *Kedrostis foetidisima* A) and *Euphorbia mauritanica* B) on YMB-1 breast cancer cells.**

Cells were treated for 24 hours at increasing concentrations of 10, 30, 50 and 100 µg/ml using the plant aqueous extracts. A staurosporine control was included. 25 nM-S represents the positive control, staurosporine, at a concentration of 25 nM.  $n = 5$ ,  $p < 0.05$

The antiproliferative activity of the herbal plant extracts on the breast cancer cell lines was measured using the  $IC_{50}$  value principle, which is a principle based on the concentration of the plant extract that causes 50% cell death. The lower the  $IC_{50}$  value of an extract on a cell line, the more potent it is considered to be. The aqueous extracts of both the herbal plants *Kedrostis foetidissima* and *Euphorbia mauritanica* exhibited considerable levels of antiproliferative activity against MCF-7 breast cancer cells presented in Figure 3.3.

Treatment with *K. foetidissima* at a concentration of 100  $\mu\text{g/ml}$  induced 48% cell death ( $IC_{50} > 100 \mu\text{g/ml}$ ) (Figure 3.3 A). *Euphorbia mauritanica* induced 40% cell death at a concentration of 100  $\mu\text{g/ml}$  ( $IC_{50} > 100 \mu\text{g/ml}$ ) (Figure 3.3. B). The positive control, staurosporine induced 43% cell death at a concentration of 25 nM. In a previous study aimed at investigating the possible antiproliferative properties of *E. mauritanica* on MCF-7 cells (Essack 2006), no significant cytotoxicity could be observed, with less than 10% cell death being observed even at a high concentration of 5 mg/ml. Results were contrary to what was observed in this study with 40% cell death resulting from a much lower concentration of 100  $\mu\text{g/ml}$ .

In addition to the cell cytotoxicity assay for *K. foetidissima* and *E. mauritanica* on MCF-7 breast cancer cells, we aimed to investigate the possible antiproliferative effects of these herbal extracts on a different breast cancer cell line. Hence their cytotoxicity was tested on YMB-1 cell lines. The positive control, staurosporine, induced 42% cell death at a concentration of 25nM on the YMB-1 cells. On comparing to the cytotoxicity of *K. foetidissima* on YMB-1 to MCF-7, it was observed that *K. foetidissima* extract exhibited a slightly higher cytotoxicity on YMB-1 cells, with 53% cell cytotoxicity being observed at a concentration of 100  $\mu\text{g/ml}$  and 37% cytotoxicity at a concentration of 50  $\mu\text{g/ml}$ , meaning  $50 \mu\text{g/ml} < IC_{50} > 100 \mu\text{g/ml}$  (Figure 4.2. A). For MCF-7

cells, an  $IC_{50} > 100$   $\mu\text{g/ml}$  was observed. The second herbal extract *E. mauritanica* however, exhibited an insignificant amount of cytotoxicity on YMB-1 cells, with less than 20% cell death occurring at a concentration of 100  $\mu\text{g/ml}$  whilst 40% cell death was observed for MCF-7 at the same concentration. These results indicate that the induced cytotoxicity is cell line specific; hence the two cell lines exhibited different levels of sensitivity to *E. mauritanica* extract.

Most plants which have been used as traditional medicine for the treatment of cancer and inflammation are high in cucurbitacins (Rios *et al.* 2005). Research into antiproliferative properties of various cucurbitacins has led to interesting discoveries, where cucurbitacins B, D and R, isolated from *Begonia heracleifolia* were found to exhibit cytotoxicity to prostate and nasopharyngeal cancer at  $IC_{50}$  ranging between 0,003 and 3, 81  $\mu\text{g/ml}$  (Rios *et al.* 2005). These exhibited a high level of cytotoxicity. For our study, a significant level of cytotoxicity was still induced by *K. foetidissima* on both MCF-7 and YMB-1 cell lines, whose  $IC_{50}$  was at concentrations of between 50 and 100  $\mu\text{g/ml}$ . Another study observed that cucurbitacins B, D, E, I, J, K and L exhibited high cytotoxicity to Hela and KB human cell cultures with  $IC_{50}$  of 0.005-1  $\mu\text{g/ml}$  (Konopa *et al.* 1974). This research and other previous studies are all indicating promising results in the use of cucurbitacins as potential future therapeutics not only of breast cancer but other cancers as well.

Plants of the Euphorbiaceae family have been used for centuries as traditional medicine against cancers and tumours (Rizk 1987). However, a large number of *Euphorbia* have not been scientifically screened for anti-cancer activity. Hence, we thus tested the cytotoxicity of the crude extracts of *Euphorbia mauritanica* on two breast cancer cell lines: MCF-7 and YMB-1. A survey on the chemical data isolated from over 120 Euphorbiaceae identified a large diversity of

structurally unique chemical compounds. These include methyl esters, their derivatives and diterpene polyesters. Other compounds identified include alkaloids, phenolic compounds, cyanogenic glucosides and glucosinolates (Rizk 1987). Some of these chemical compounds have been shown to exhibit antiproliferative activity against various cancer cells. A study by Engi *et al.* (2007) looked at the effect of diterpenes isolated from methanol extracts of nine *Euphorbia* species on human colon cancer cells and observed a moderate antiproliferative effect of  $IC_{50} < 3.5 \mu\text{g/ml}$ -  $17.46 \mu\text{g/ml}$ .

Studies by Whelan and Ryan (2003), investigated the effect of ethanolic extracts of *Euphorbia* species: *E. grandidens*, *E. grandicorni*, *E. latea*, *E. coerulescens*, *E. trigona*<sup>1</sup>, *E. istigy*, *E. candelabrum*, *E. pentagona*, *E. triangularis* against HEP-2 (Human epidermoid cancer) cells at concentrations of  $8.53 \mu\text{g/ml}$ ,  $85.3 \mu\text{g/ml}$  and  $853.9 \mu\text{g/ml}$ . They found that of the nine plant extracts tested, four extracts resulted in reduced cell viability at concentrations of  $85.3 \mu\text{g/ml}$  and  $853.9 \mu\text{g/ml}$ , with the most potent extract causing  $IC_{50}$  at  $57 \mu\text{g/ml}$ . Interestingly, for our study, we observed increase in proliferation for YMB-1 cells that were treated with *E. mauritanica* at a concentration of  $10 \mu\text{g/ml}$  (Figure 3.4. B), in their study, they observed a similar trend with seven of their extracts also promoting cellular proliferation at a concentration of  $8.53 \mu\text{g/ml}$ . This may indicate that some *Euphorbia* species, including *E. mauritanica*, may actually induce cell proliferation to some cancers at low doses.

It was evident from our study that the crude extracts of the two plants *E. mauritanica* and *K. foetidissima* were exhibiting antiproliferative properties on the breast cancer cells, but the next important issue was to determine if the observed cytotoxicity was as a result of apoptosis or necrosis.

## **CHAPTER 4 SCREENING OF HERBAL EXTRACTS FOR MODE OF DEATH INDUCTION (APOPTOSIS OR NECROSIS)**

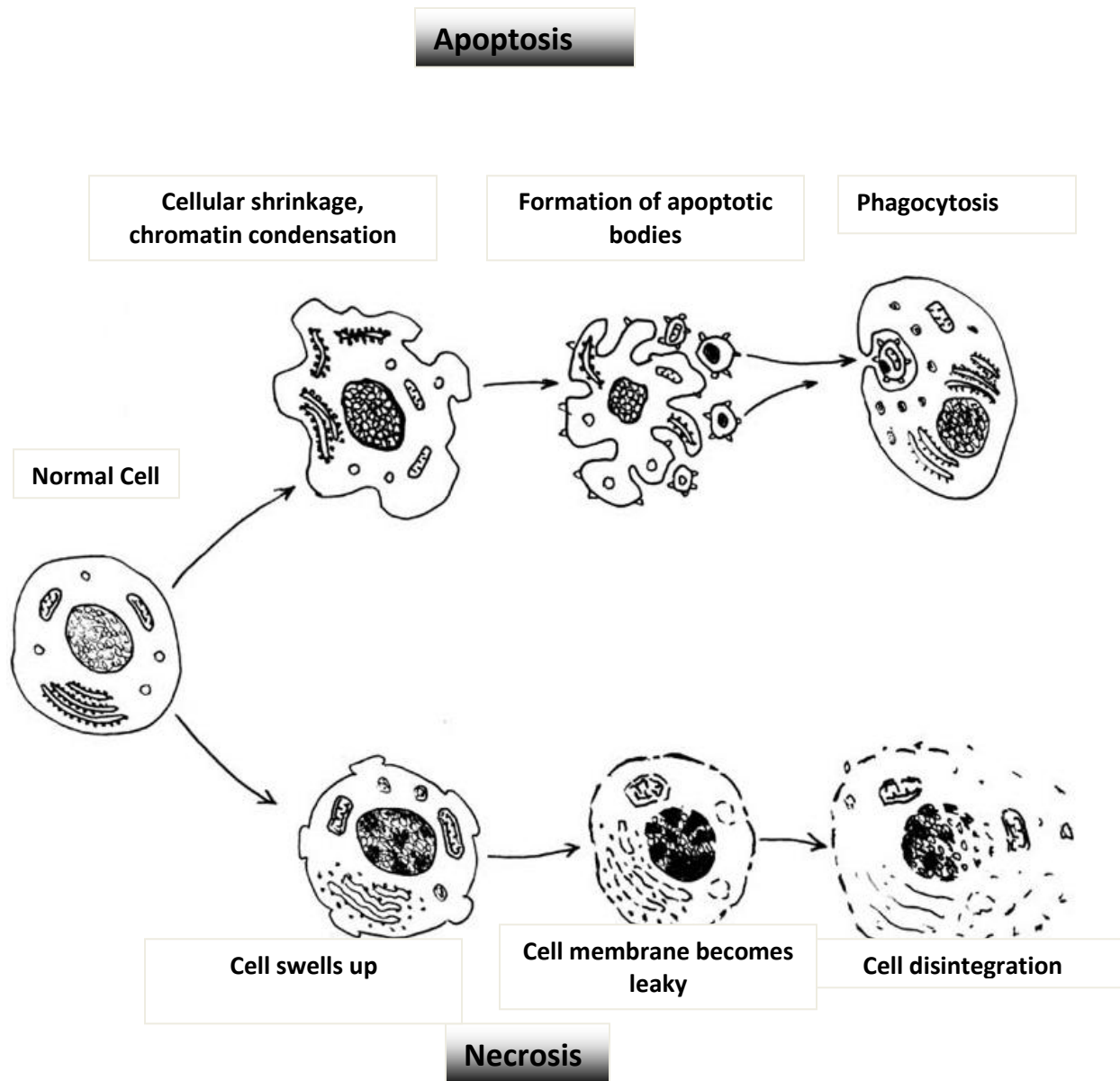
### **4.1 Introduction**

Apoptosis and necrosis are two different types of cell death characterised by different modes or morphologies of death (Figure 4.1). Apoptosis is a form of programmed cell death, whereby cells undergo orderly typical morphological changes whereas necrosis involves rapid lysis of a cell (Kumar 1998; Raffray and Cohen 1997). Necrosis is generally not the ideal death pathway since it leads to leakage of damaged internal components of the dying cell into surrounding tissue, which may be harmful to the surrounding cells, and leads to induction of an inflammatory response (Potten and Wilson 2004).

**Table 4.1: Differences between apoptotic and necrotic cellular death pathways**

| <b>Apoptosis</b>                                                                                                                                                                                   | <b>Necrosis</b>                                                                |
|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|--------------------------------------------------------------------------------|
| <b>Morphological features</b>                                                                                                                                                                      |                                                                                |
| Early loss of cell to cell contact and involves single cells (d)                                                                                                                                   | Late loss of cell to cell contact and involves clusters (d)                    |
| Anoikis- detachment from the surface                                                                                                                                                               | No anoikis                                                                     |
| Cell shrinkage (e)                                                                                                                                                                                 | Cell swelling (e)                                                              |
| Mitochondrial condensation, no swelling (a)                                                                                                                                                        | Mitochondria swelling (a)                                                      |
| DNA fragmentation- nuclear DNA breaks down in large, then small nucleosomal fragments (b)                                                                                                          | Nuclear swelling (e)                                                           |
| Formation of apoptotic bodies                                                                                                                                                                      | No apoptotic bodies                                                            |
| Rapid recognition, uptake and degradation of apoptotic bodies by phagocytes (c)                                                                                                                    | No cellular uptake by phagocytes, attraction of inflammatory cells (c,e)       |
| No leakage of cell contents (a)                                                                                                                                                                    | Cells swell up and suddenly collapse, leading to spilling of cell contents (b) |
| <b>Apoptosis</b>                                                                                                                                                                                   | <b>Necrosis</b>                                                                |
| <b>Biochemical and molecular features</b>                                                                                                                                                          |                                                                                |
| ATP levels are preserved (e)                                                                                                                                                                       | Rapid intracellular ATP depletion (e)                                          |
| Oxidative stress (e)                                                                                                                                                                               | —                                                                              |
| Cytochrome C released from mitochondria (e)                                                                                                                                                        | —                                                                              |
| Protein degradation mediated by specific proteolytic enzymes known as caspases (a)                                                                                                                 | No regulated protein degradation (a)                                           |
| Plasma membrane remains intact almost until the end, then losses asymmetry and integrity. Phosphatidyl serine residues located in the inner plasma membrane translocate to the cell surface. (a,b) | Plasma membrane losses integrity rapidly (b)                                   |

Key: a) Kroemer *et al.* 1998; b) Fiers *et al.* 1999; c) Savill *et al.* 1993; d) Potten and Wilson 2004; e) Wlodkowic *et al.* 2009



**Figure 4.1: Characteristic features that distinguish between apoptotic and necrotic cell death (van der Meer *et al.* 2010).**

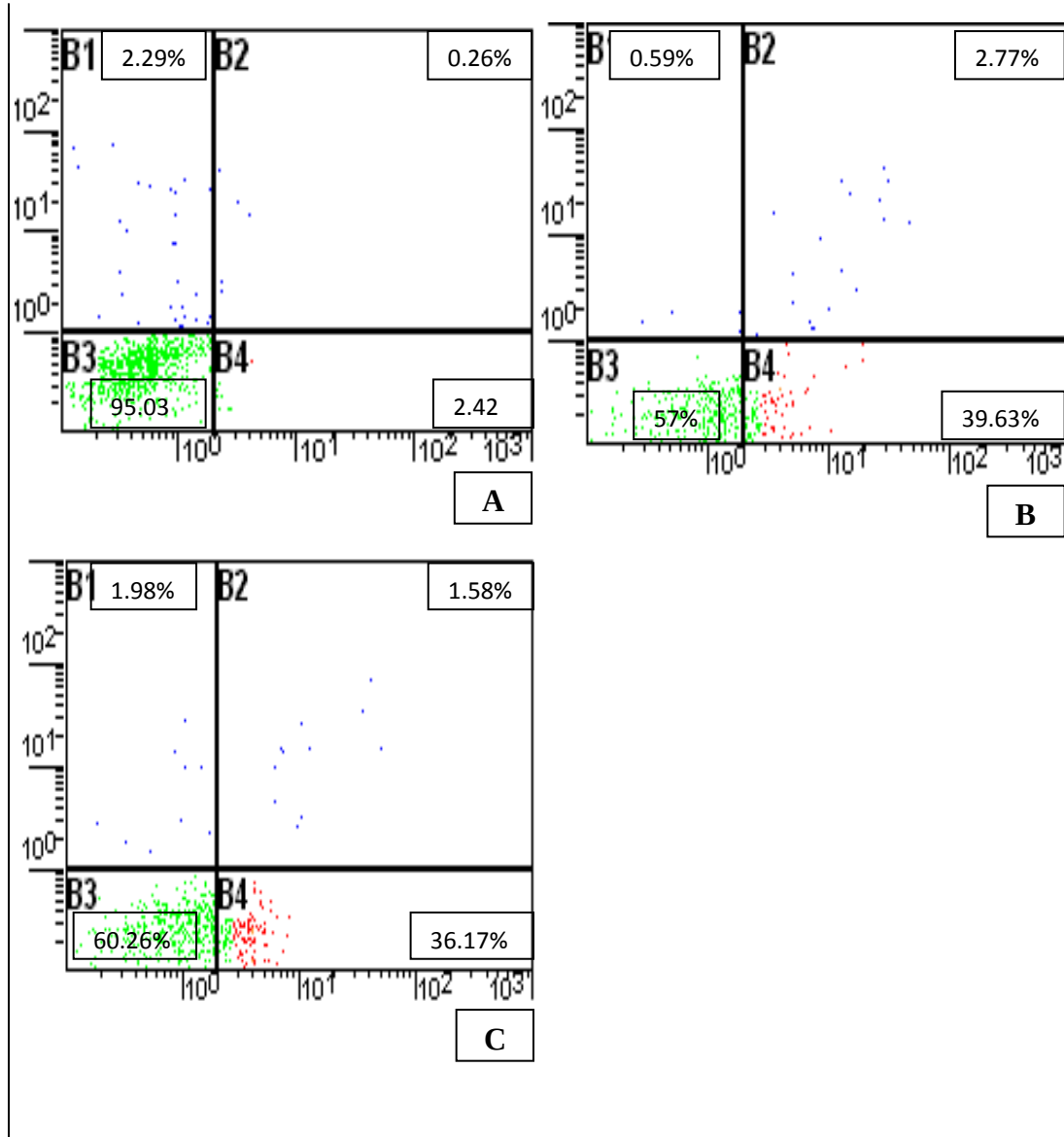
Necrotic cells are characterised by swelling up of the cell, loss of membrane integrity and eventually the cells lyse releasing their internal components into the surrounding tissue. Apoptotic cells shrink, chromatin condenses, nuclear DNA breaks down and apoptotic bodies form which are then phagocytosized by either macrophages or surrounding cells (van der Meer *et al.* 2010).

When working with potential therapeutic agents, it is important that they induce high cytotoxicity to their target cells whilst showing none or very minimal toxicity to the surrounding unaffected healthy tissue. From Table 4.1, it is clear that apoptosis would be the desired method of death since it is more regulated, considering the fact that dying cells are taken up by phagocytes and do not lyse to release their internal components to neighbouring cells, which would otherwise prove to be catastrophic.

The purpose of this section was to screen the mode of death induction of the crude extracts of *K. foetidissima* on MCF-7 and YMB-1 cells and *E. mauritanica* on MCF-7 cells since they showed significant cytotoxicity on the cells. MCF-7 and YMB-1 breast cancer cells, treated with the herbal extract concentrations that induced about 50% cell death were incubated with Annexin V according to manufacturer's specifications (see section 2.2.2). A negative control of untreated cells and positive control of cells treated with 25 nM of staurosporine were included. Staurosporine is a known inducer of apoptosis in many different cancer cell types (Motadi 2009). Flow cytometry was employed to analyse the percentage of cells that underwent apoptosis compared to the percentage of necrotic and live cells following treatment with the herbal extracts.

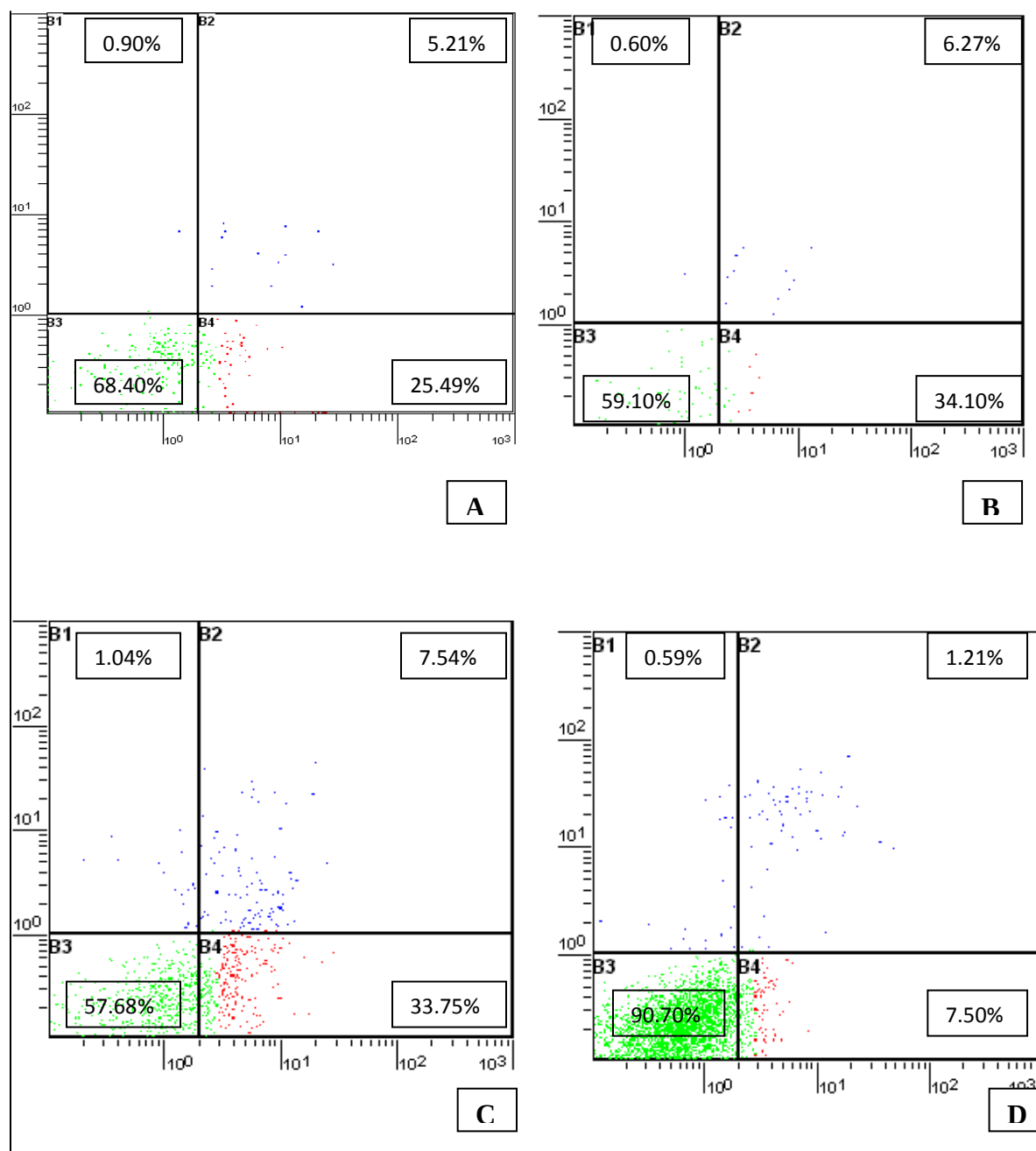
## 4.2 Results and discussion

The graphs below are representative of the flow cytometry apoptosis analysis



**Figure 4.2: Scatter analysis diagrams for YMB-1 cells.**

A) Untreated cells, B) Staurosporine (25 nM) treated cells and C) *Kedrostis foetidissima* (100  $\mu\text{g/ml}$ ) treated cells.



**Figure 4.3: Scatter analysis diagrams for MCF-7 cells.**

Cells were treated with A) *Kedrostis foetidissima* (100 µg/ml), B) *Euphorbia mauritanica* (100 µg/ml), C) Staurosporine (25 nM) and D) Untreated.

The cells were scattered into one of four quadrants: Quadrant B1 contains cells undergoing necrosis, quadrant B2 contains cells undergoing late apoptosis, quadrant B4 contains cells undergoing early apoptosis and quadrant B3 contains live cells (Figures 4.2 & 4.3).

Since *K. foetidissima* seemed to be the only extract that exhibited cytotoxicity on YMB-1 cells, it was then further assessed for possible pro-apoptotic activity on this cell line. The YMB-1 cells were treated with *K. foetidissima* at a concentration of 100 µg/ml for 24 hours and tested for markers of apoptosis. On observing results for the positive control, staurosporine (Figure 4.2 B) 40% of the cells stained positive for externalised phosphatidylserine residues, indicating that 40% of the cells underwent apoptosis whilst 57% remained alive. The cells treated with *K. foetidissima* extract demonstrated a similar percentage of phosphatidylserine externalisation to that of staurosporine, with about 40% of cells undergoing apoptosis and about 60% remaining alive (Figure 4.2 C). A negative control of untreated cells was included which from flow cytometer analysis indicated that 95% of the cells were alive and less than 5% were dying naturally either via necrosis or apoptosis (Figure 4.2 A). The flow cytometer results suggested that the herbal extract *K. foetidissima* mainly induced cell death to YMB-1 cells via an apoptotic pathway.

MCF-7 cells exhibited significant sensitivity to both herbal extracts *E. mauritanica* and *K. foetidissima* hence we then aimed to investigate if these herbal extracts were inducing death to the cells via necrosis or apoptosis. Similarly to the previous experiments, staurosporine (25 nM) was used as a positive control, and the flow cytometry results suggested that over 40% of cells stained positive for externalised phosphatidylserine. MCF-7 cells were treated with both *K. foetidissima* and *E. mauritanica* at concentrations of 100 µg/ml. Flow cytometry analysis

indicated that treatment with *K. foetidissima* resulted in about 30% cells staining positive for externalised phosphatidylserine residues i.e. undergoing apoptosis and with almost 70% remaining alive. *Euphorbia mauritanica* treatment resulted in about 40% cells undergoing apoptosis whilst about 60% remained viable (Figure 4.6). A no treatment control was included which showed that over 90% of the cells were still alive after 24 hours and the rest of the cells had died mainly due to apoptosis (Figure 4.3).

Extracts of *K. foetidissima* were assayed for markers of pro-apoptotic activity on MCF-7 and YMB-1 cell lines whilst *E. mauritanica* was assayed for markers of pro-apoptotic activity on MCF-7 cell lines only. On assessing cells in the second and fourth quadrant representing cells undergoing late and early apoptosis respectively, it was observed that *K. foetidissima* induced the highest apoptotic activity in YMB-1 than in MCF-7 cells (Figures 4.3C & 4.4A). This difference was also observed in the cytotoxicity assay. It is promising however, that the majority of cells were undergoing apoptosis in both cell lines. In a study by Essack (2006), contradictory results were obtained, with no phosphatidylserine residues externalisation being observed for MCF-7 cells treated with *E. mauritanica*.

In summary, the Annexin V apoptotic assay has demonstrated that the aqueous extracts of two South African herbal plants *K. foetidissima* and *E. mauritanica* induce their cytotoxic effects by an apoptotic mechanism. This indicates that these plants possess compound/s which can trigger an apoptotic pathway, which is promising in that induction of apoptosis correlates with anti-proliferation of cancer cells. Even though regulation of apoptosis is controlled by a huge network comprising a number of genes, the two most important genes which have been found to be highly deregulated in cancer are p53 and RBBP6. Following the flow cytometry apoptotic assay,

we then aimed to investigate if these herbal extracts have any effect on these two tumour suppressor genes, using real-time PCR.

## **CHAPTER 5 TUMOUR SUPPRESSOR GENES**

### **5.1 Introduction**

The majority of breast cancers arise from deregulation of genes involved in the cell cycle or apoptosis. As with all drugs that are manufactured to treat different diseases, it is important for potential natural anti-cancer agents to exhibit a certain level of specificity on their target. The FDA approved chemotherapeutic drugs as mentioned in section 1.5 all exhibit specific activity on their target at various steps of the cell cycle. Their activity ranges from inhibiting microtubule assembly, to preventing DNA and RNA synthesis whilst some work by preventing DNA replication (Giri and Narasu 2000; Kessel *et al.* 1972; Kjeldsen 1992).

Tumour suppressor genes have now become of great interest in understanding molecular pathogenesis of cancer. The tumours suppressors which have been highly found to be deregulated in breast cancer include p53 and RB genes. p53, a tumour suppressor is very crucial in the control of cell proliferation. Deregulation of p53 has been implicated in over 50% of cancers (Potten and Wilson 2004). This may stem from its role as a sequence specific DNA binding transcription factor, whose downstream targets are mostly gene products which control cell cycle arrest, DNA repair and apoptosis (Picksley and Lane 1994). Retinoblastoma (Rb), the first tumour suppressor to be discovered, acts as a crucial component of the cell cycle checkpoint. It functions to prevent excessive cell growth, by inhibiting progression of aberrant DNA along the cell cycle through G1 into the S-phase (Garg 2007; Mayhew *et al.* 2007). Rb has been found to be mutated in a third of human cancers (Lee 2002). From the extensive amount of research done on p53 and Rb deregulation, it can be said that they are arguably the most deregulated pathways in human cancer. The tumour suppressors p53 and Rb require specific

protein to protein interactions to mediate their respective functions. RBBP 6 protein is one of very few proteins known to interact with both p53 and Rb.

In this study, the cytotoxicity assay showed that MCF-7 exhibited sensitivity to both *E. mauritanica* and *K. foetidissima* whilst YMB-1 cells were sensitive to *E. mauritanica* only (Figures 3.3 & 3.4). The flow cytometric assay further indicated that death was induced by an apoptotic pathway (Figures 4.3 & 4.4). With the high rate of p53 and Rb mutations that have been shown to occur in cancer and the fundamental role these genes play in tumour suppression, we thus aimed to investigate if the herbal extracts could restore or upregulate the levels of p53 and reduce expression levels of RBBP6 in MCF-7 breast cancer cells thereby increasing apoptosis. This was done by means of real-time PCR, which can accurately measure changes in gene expression levels.

Real-Time PCR is a highly sensitive and powerful method used for quantitating small changes in gene expression. Its accuracy and the rapid quantification of results make it the ideal method of nucleic acid quantification (Klein 2002; Pfaffl 2001). Real-Time PCR was employed to (i) investigate the expression levels of the above mentioned tumour suppressors: p53 and RBBP6 in MCF-7 breast cancer cells and (ii) to investigate if the herbal extracts which were inducing apoptosis to these cells were having any effect on the expression levels of these tumour suppressors.

## **5.2 Primer efficiency assessment using standard curves**

To answer the above questions, real-time PCR specific primers were designed using NCBI for collection of gene sequences, primer3plus for designing primers and NCBI Blast for ensuring

that the primers were specific to our target gene and did not amplify non-target sequences. Prior to real-time PCR, the primers were checked for efficiency by normal PCR.

The designed primer sequences were as follows:

**GADPH:** Forward – 5' -GAG TCA ACG GAT TTG GTC GT- 3'

Reverse – 5' -TTG ATT TTG GAG GGA TCT CG- 3'

Product Size: 238 bp

**β-actin:** Forward – 5' -AGA GCT ACG AGC TGC CTG AC- 3'

Reverse – 5' -AGC ACT GTG TTG GCG TAC AG- 3'

Product Size: 184 bp

**RBBP6:** Forward – 5' -CAG CGA CGA CTA AAA GAA GAG TCT- 3'

Reverse – 5' -GGT AAT TGC GGC TCT TGC CT- 3'

Product Size: 199 bp

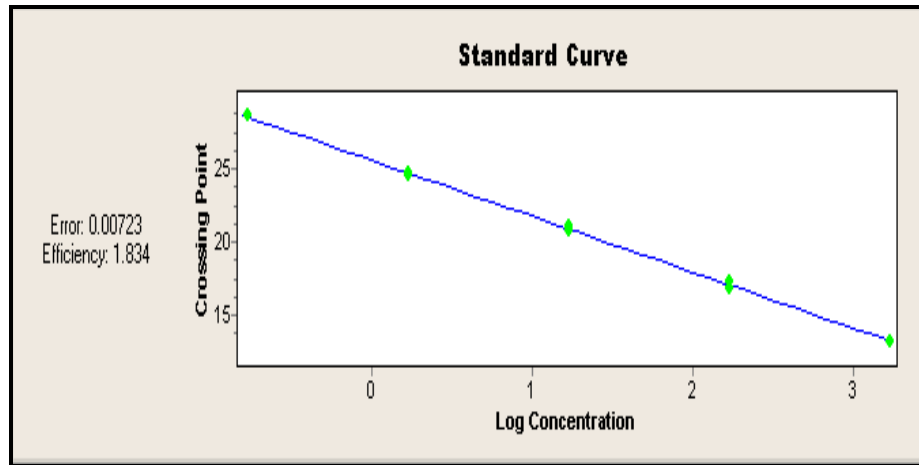
**p53:** Forward – 5' -GTT CCG AGA GCT GAA TGA GG- 3'

Reverse – 5' -TGA GTC AGG CCC TTC TGT CT -3'

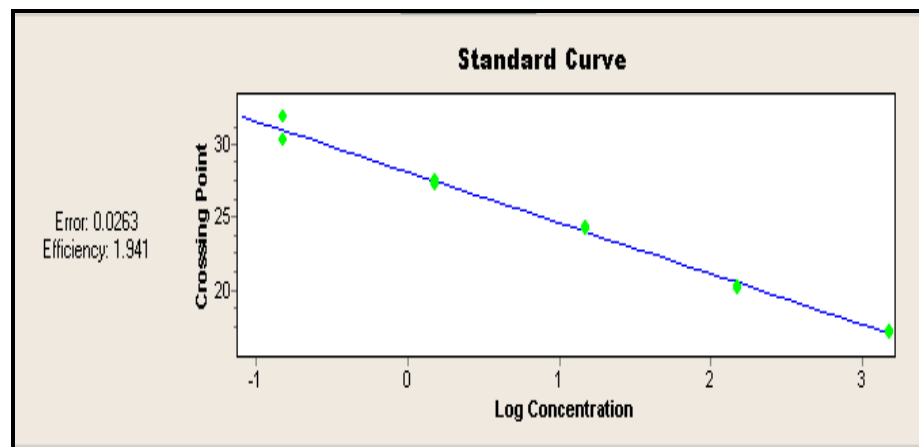
Product Size: 157 bp

Efficiency of the primers was assessed by preparing standard curves from 10 fold serial dilutions (undiluted, 1:10, 1:100, 1:1000 and 1:10000) of MCF-7 cDNA. Beta-actin and GADPH are representative of the housekeeping genes used to validate efficacy of the real-time PCR

experiments. The real-time PCR cocktail was prepared as explained in appendix A4 and the real-time PCR running parameters are explained in Table 2.1.



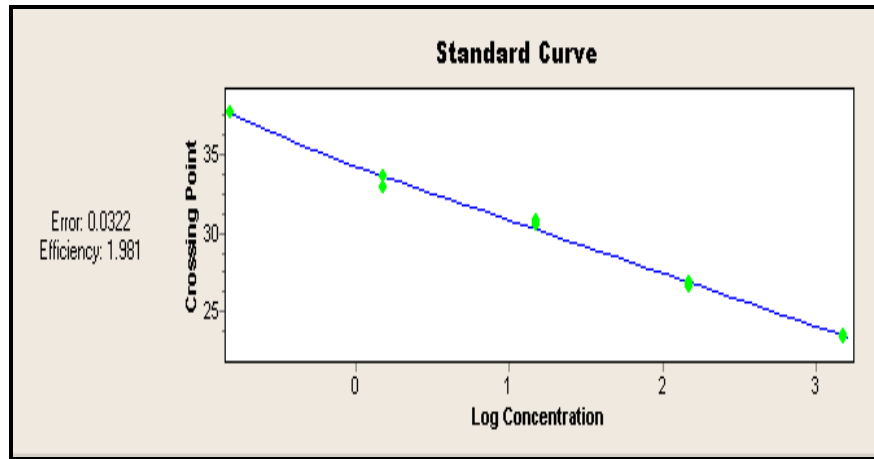
A



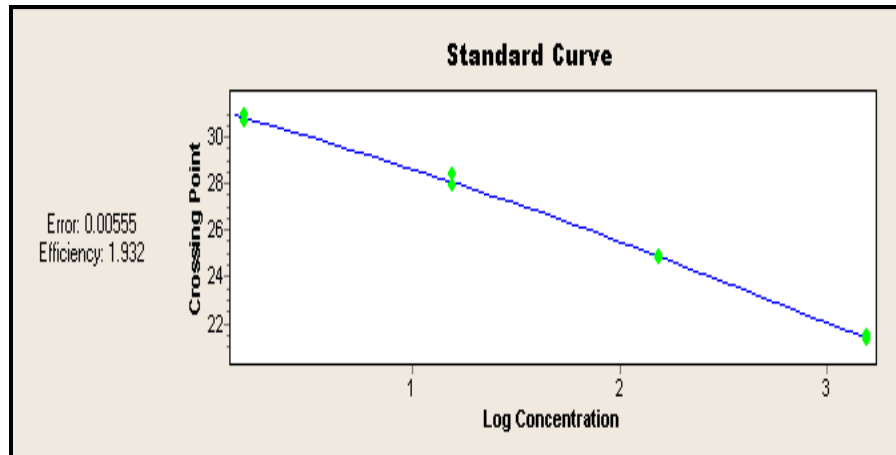
B

**Figure 5.1: Standard curves for Beta-actin (A) and GADPH (B) primers which served as housekeeping genes.**

The amplification efficiencies of the slopes were 1.834 and 1.941, respectively. n= 2



**A**



**B**

**Figure 5.2: Standard curves for the tumour suppressors RBBP6 (A) and p53 (B) primers.**

The amplification efficiencies of the slopes were 1.981 and 1.932, respectively. n= 2

### 5.3 Relative quantitative real-time polymerase chain reaction

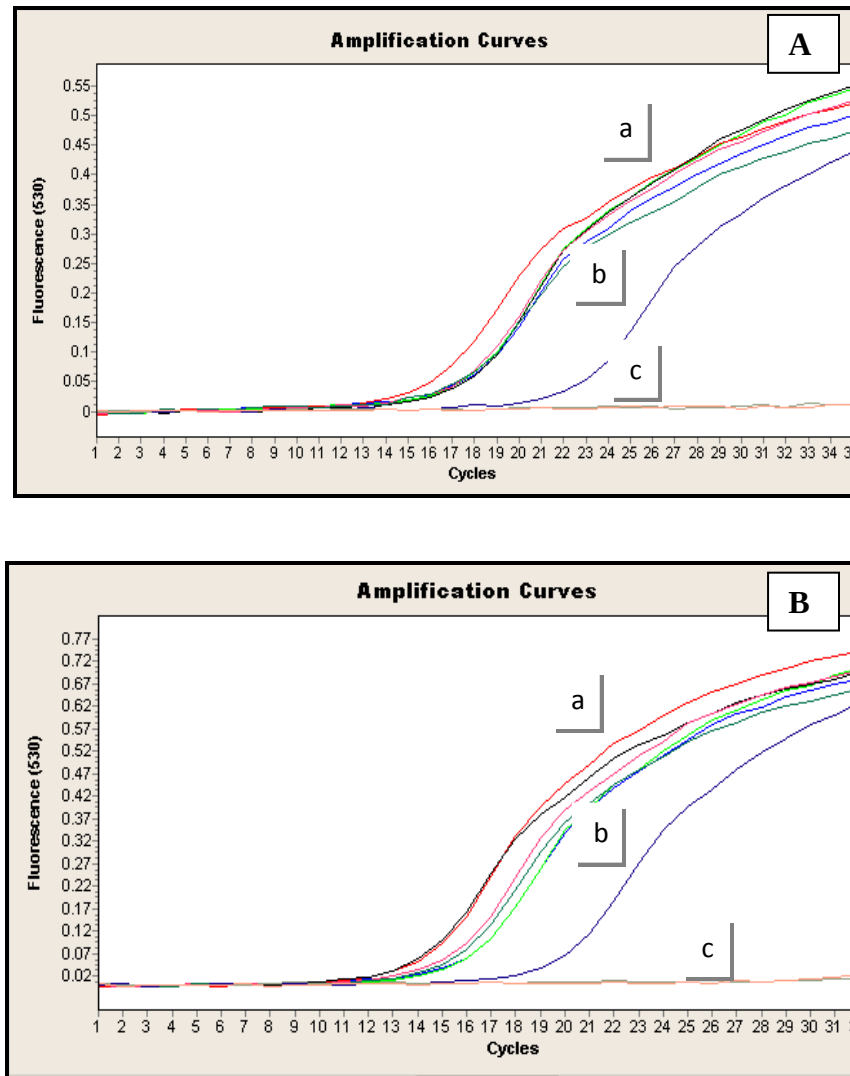
Relative quantification, which is based on the relative expression of the target gene versus that of a housekeeping gene, was the preferred mode of quantification for this study. Housekeeping genes are constitutively expressed in all nucleated cells since they are required for basic cell survival. Their mRNA is considered stable even under changing experimental conditions (Pfaffl 2001). Housekeeping genes such as glyceraldehyde-3-phosphate dehydrogenase (GADPH), actins, tubulins, cyclophilin, 18S rRNA and 28S rRNA are an absolute prerequisite for use in real-time RT-PCR experiments to ensure accurate normalisation of the experiments (Schmittgen and Zakrajsek 2000). For this study, mRNA expression levels were normalised to the levels of GADPH and  $\beta$ -actin transcripts, which served as our housekeeping or reference genes.

One-step real-time PCR requires a separate step of reverse transcription of mRNA to cDNA before running RT-PCR. This method is ideal in that the starting amount of cDNA can be equalised to ensure the difference in expression level of genes is not due to experimental error. The following mRNA templates were reverse transcribed as in section 2.2.3 in preparation of RT-PCR:

- i. Untreated MCF-7 cells
- ii. MCF-7 cells treated with *K. foetidissima*
- iii. MCF-7 cells treated with *E. mauritanica*
- iv. Standard ( Prepared from untreated cells at one of the concentrations correlating with one of the points or concentrations on the standard curve)
- v. Negative control (-ve)
- vi. No template control (NTC)

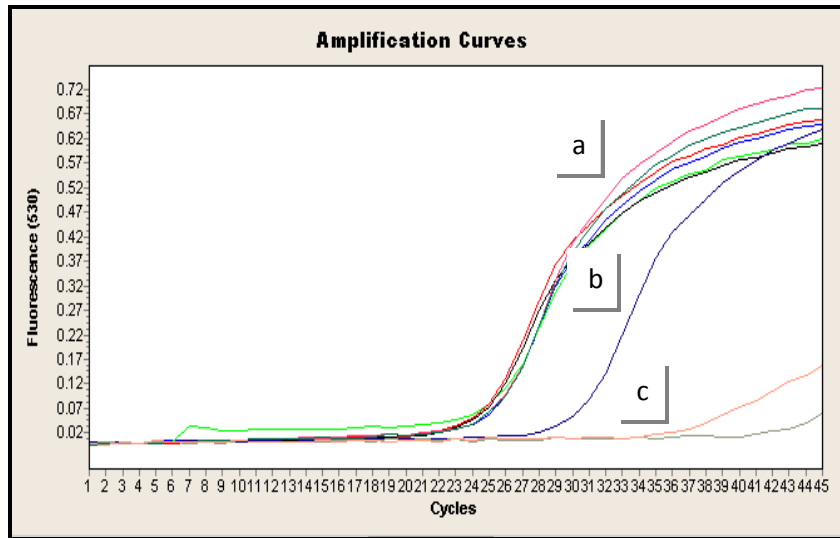
To validate if the selected housekeeping genes were the best possible for normalisation of these experiments, we had to test if they were stably expressed under changing experimental systems. Real-time PCR amplification curves were constructed using (1) GAPDH and (2)  $\beta$ -actin primers. Each run comprised of (i), (ii) and (iii), all experiments were conducted in duplicates with (iv), (v) and (vi) also included.

## 5.4 Results and discussion



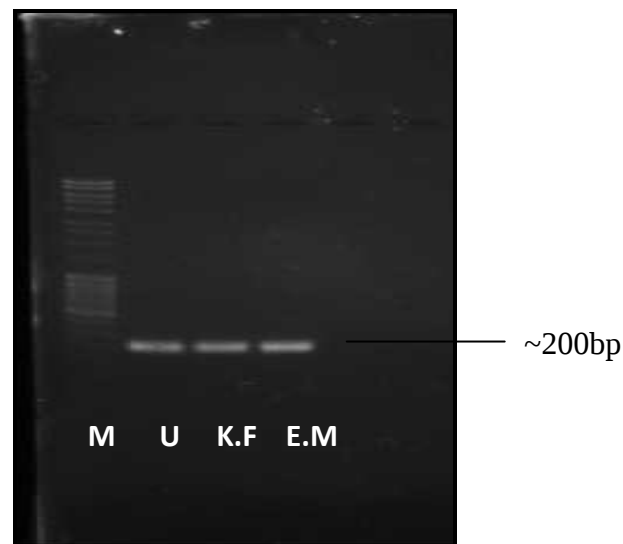
**Figure 5.3: Amplification curves for the housekeeping genes.**

A) GADPH and B)  $\beta$ -actin primers. The curves were constructed using cDNA from a) untreated MCF-7 cells, *K. foetidissima* and *E. mauritanica* treated cells. All experiments were performed in duplicates b) A standard, c) NTC and -ve control were included.



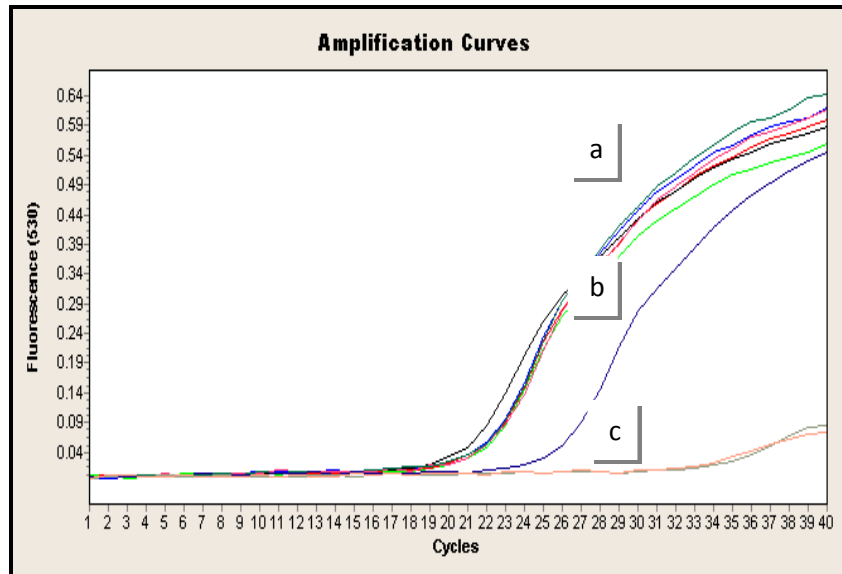
**Figure 5.4: Amplification curves for RBBP6 expression.**

a) untreated MCF-7 cells, *K. foetidissima* and *E. mauritanica* treated MCF-7 cells b) A standard, c) NTC and -ve control were included.



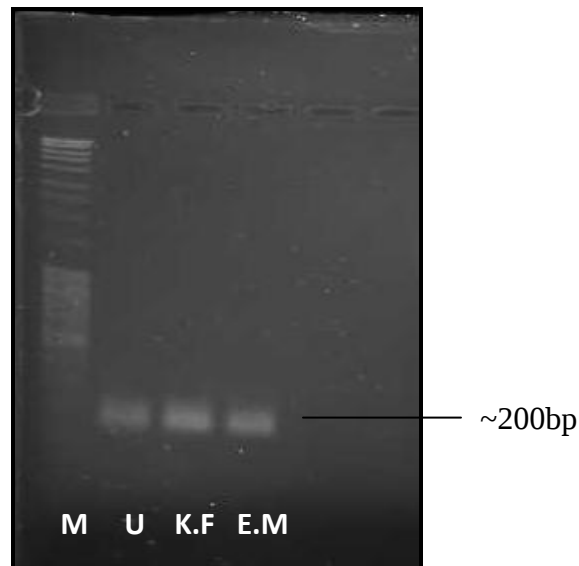
**Figure 5.5: RT-PCR products resolved on a gel following RBBP6 amplification.**

M represents the molecular weight marker (MassRuler DNA Ladder Mix, Fermentas), U represents untreated MCF-7 cells, K.F and E.M represent *K. foetidissima* and *E. mauritanica* treated MCF-7 cells, respectively.



**Figure 5.6: Amplification curves for p53 expression.**

a) Untreated MCF-7 cells, *K. foetidissima* and *E. mauritanica* treated MCF-7 cells. b) A standard, c) NTC and -ve control were included.



**Figure 5.7: RT-PCR products resolved on a gel following p53 amplification.**

M represents the molecular weight marker, U represents untreated MCF-7 cells, K.F and E.M represent *K. foetidissima* and *E. mauritanica* treated MCF-7 cells, respectively.

There are currently a number of mathematical models used to analyse real-time PCR data. A fairly new and simple method by Pfaffl (2001) was used since it seems to provide the most accurate data quantification. This method compares the ratio of the target gene to that of the reference (housekeeping) gene. The relative expression ratio of an unknown target is computed based on its real-time PCR efficiencies, the crossing point difference of the unknown target versus that of the control in comparison to that of the control (Pfaffl 2001).

$$\text{Relative expression} = \frac{(E_{\text{target}})^{\Delta C_{\text{Ptarget}}(\text{control} - \text{sample})}}{(E_{\text{ref}})^{\Delta C_{\text{Pref}}(\text{control} - \text{sample})}} \quad (\text{Pfaffl 2001})$$

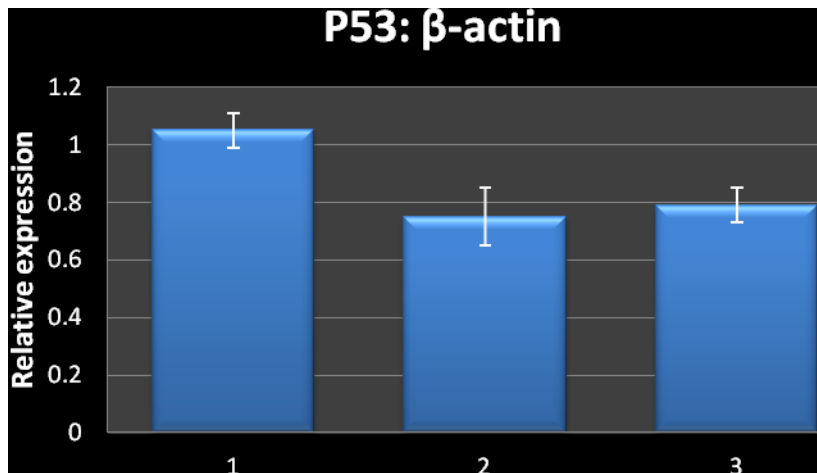
Where:

$E_{\text{target}}$  = real time PCR efficiency of target gene

$E_{\text{ref}}$  = real time PCR efficiency of reference gene

$\Delta C_{\text{Ptarget}}$  = Deviation in crossing point of control - crossing point of target gene

$\Delta C_{\text{Pref}}$  = Deviation in crossing point of control - crossing point of reference gene



**P53:  $\beta$ -actin**

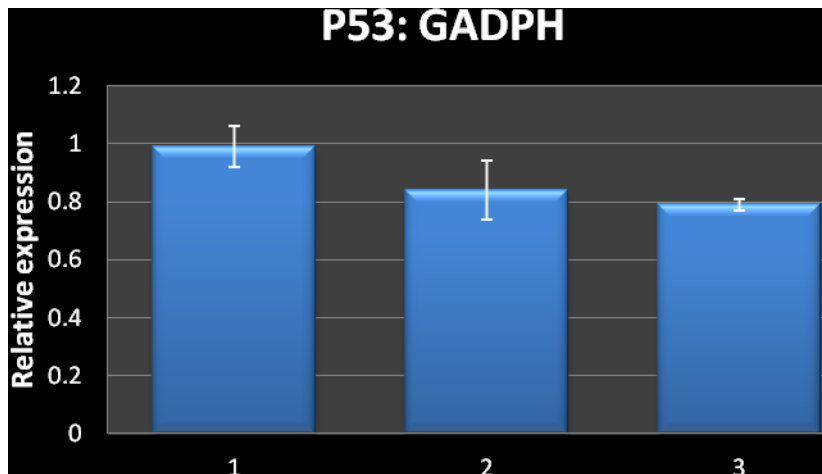
1. Untreated - 1.05

2. *K. foetidissima* - 0.75

3. *E. mauritanica* - 0.79

**Figure 5.8: Relative expression of p53 versus that of  $\beta$ -actin housekeeping gene.**

P53 expression levels for 1. Untreated MCF-7 cells, 2. *K. foetidissima* and 3. *E. mauritanica* treated cells. n=2.



### **P53: GADPH**

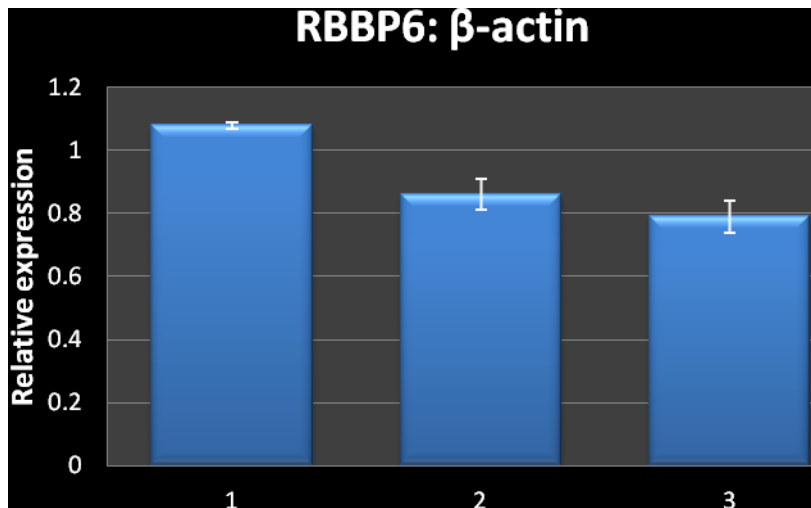
1. Untreated – 0.99

2. *K. foetidissima* - 0.84

3. *E. mauritanica* - 0.79

**Figure 5.9: Relative expression of p53 versus that of GADPH housekeeping gene.**

P53 expression levels for 1. Untreated MCF-7 cells, 2. *K. foetidissima* and 3. *E. mauritanica* treated cells. n=2.



**RBBP6:  $\beta$ -actin**

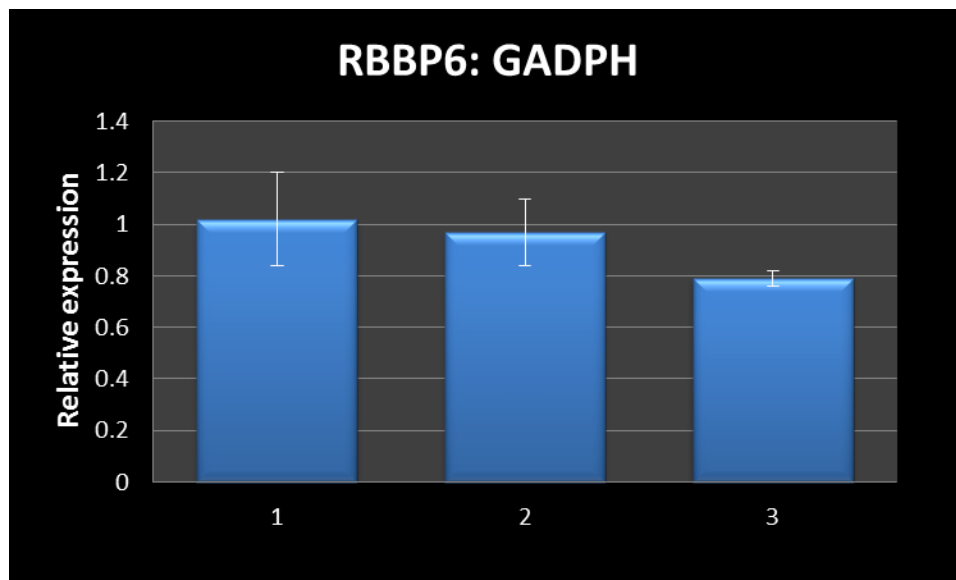
1. Untreated – 1.08

2. *K. foetidissima* - 0.86

3. *E. mauritanica* - 0.79

**Figure 5.10: Relative expression of RBBP6 versus that of  $\beta$ -actin housekeeping gene.**

RBBP6 expression levels for 1. Untreated MCF-7 cells, 2. *K. foetidissima* and 3. *E. mauritanica* treated cells. n=2.



### **RBBP6: GADPH**

1. Untreated – 1.02

2. *K. foetidissima* - 0.97

3. *E. mauritanica* - 0.79

**Figure 5.11: Relative expression of RBBP6 versus that of GADPH housekeeping gene.**

RBBP6 expression levels for 1. Untreated MCF-7 cells, 2. *K. foetidissima* and 3. *E. mauritanica* treated cells. n=2

Relative quantitative RT-PCR revealed that the levels of the RBBP6 transcript following normalisation with  $\beta$ -actin were 1.08 in the untreated MCF-7 cells (Figure 5.10). Treatment with *K. foetidissima* slightly reduced expression levels to 0.86 and with *E. mauritanica* it was reduced to 0.79. Following normalisation with GAPDH, the RBBP6 levels seemed to have been slightly reduced from 1.02 to 0.97 whilst treatment with *E. mauritanica* led to a slightly higher reduction of 0.79 (Figure 5.11).

The relative expression levels of p53 in MCF-7 cells following normalisation with  $\beta$ -actin were 1.05 before treatment (Figure 5.8). Treatment with *K. foetidissima* slightly reduced expression levels to 0.75 and with *E. mauritanica* it was reduced to 0.79. Following normalisation with GAPDH, relative expression of p53 in the untreated cells was 0.99, it was slightly reduced to 0.84 following treatment with *K. foetidissima*. Treatment with *E. mauritanica* yielded a slight reduction of 0.79 (Figure 5.9). The herbal extracts *K. foetidissima* and *E. mauritanica* exhibit minimal effect on the expression levels of the tumour suppressors p53 and RBBP6. Both extracts caused a slight reduction in the expression of both genes.

In summary, the crude extracts of *E. mauritanica* and *K. foetidissima* induced apoptosis on MCF-7 cells, but, RT-PCR analysis indicated that their mode of action is not by upregulating p53 or downregulating RBBP6. Since apoptosis is controlled by such a huge network of genes and signalling molecules, it is possible that these plant extracts control other components of the apoptotic pathway. Further research that incorporates isolation and identification of the active compounds would be required to better understand their molecular mechanism and the pathway components they affect.

## **CHAPTER 6 GENERAL DISCUSSION AND CONCLUSION**

### **6.1 General discussion**

Historically, plants, herbs and spices were a folkloric source of medicinal agents, and as modern medicine expanded, many useful drugs were developed from lead compounds discovered from medicinal plants. This approach has provided leads against various pharmacological targets, including cancer, malaria and pain, and remains an important route to new pharmaceuticals (Essack 2006). Recent advances in cytotoxic and phytochemical screening have provided scientists with insight into the bioactive properties of medicinal plants, which has led to the development of new medicines. In this study, *Kedrostis foetidissima*, *Euphorbia mauritanica* and *Elytropappus rhinocerotis* plants which are indigenous to South Africa and its neighbouring countries were screened for their possible antiproliferative and/or pro-apoptotic effect on two breast cancer cell lines: MCF-7 and YMB-1.

MTT assay was used to measure significant variations in the inhibitory activity of the 3 plant extracts on breast cancer cell lines. The best thing about MTT is that it does not have limitations and one can test as many times as possible. It was used to measure IC<sub>50</sub>, the concentration of the plant extract that induces 50% cell death on the breast cell lines. Untreated cells were included as a control and cells treated with staurosporine as a positive control. Little to no cell cytotoxicity was observed when *Elytropappus rhinocerotis* was used in these two cell lines. From our results as described in Chapter 3 (Figures 3.3 & 3.4), we observed interesting results with *Kedrostis foetidissima* being able to induce up to 48% and 53% cell death in MCF-7 and YMB-1 cells respectively, at a concentration of 100 µg/ml. Whereas *Euphorbia mauritanica* induced up to

40% cell death in MCF-7 but insignificant cell death of less than 20% in YMB-1 could be observed. In this case, only MCF-7 showed to be a cell line that can be used to assess the growth inhibition potential of *Euphorbia mauritanica*. The results obtained especially for *E. mauritanica* treatments, are similar to those observed by Kummalue *et al.* (2011) who compared inhibition of cell growth by *Sapindus rarak* water extract on two lung cancer cell lines and a breast cancer cell line; they also observed a significant growth inhibition in a dosage dependent manner, but only in the A549 cell line, and no significant growth inhibition was observed in the other two cell lines. These results therefore suggest that this plant extract might be cell line specific and cancer type dependent.

After the confirmation that the two plant extracts indeed managed to induce significant cell death in the two breast cancer cell lines tested, we had to verify if the type of cell death induced was due to apoptosis, which is genetically controlled or if it was due to necrosis. One way of verifying this was by the use of flow-cytometry staining which uses both the propidium iodide and FITC. Propidium iodide stains cells in their late apoptotic stages as well as G1 cell cycle arrest and FITC stains cells in the early stages of apoptosis. *Kedrostis foetidissima* was then further assessed for possible pro-apoptotic activity using flow cytometer on MCF-7 and YMB-1 cells, whilst *E. mauritanica* was assessed on MCF-7 cells. Similarly to the cytotoxicity assays, staurosporine was used as the positive control whilst cells that were not treated with anything but media were used as the control.

The results confirmed our earlier findings that indeed cell death had been induced following treatment with the two plant extracts, and that it was as a result of apoptosis. YMB-1 and MCF-7 cells treated with *K. foetidissima* at a concentration of 100 µg/ml demonstrated a 40% and 30%

phosphatidylserine residue externalization, respectively; whilst MCF-7 cells treated with *E. mauritanica* demonstrated a 40% phosphatidylserine residue externalization, which is an indication of induced apoptosis. Staurosporine induced just over 40% apoptosis in both cell lines. This demonstrated that treatment of MCF-7 cells with *K. foetidissima* and *E. mauritanica* and of YMB-1 cells with *K. foetidissima* resulted in a significant induction of apoptosis, while showing relatively no significant toxicity to the cells in the form of necrosis.

Research into the development of novel small-molecule plant-derived extracts and compounds, which may prevent carcinogenesis, curtail its progression, or even cure the disease is still at the forefront of cancer therapeutics (D'Archivio *et al.* 2007). With studies by Creemers *et al.* (1994) showing compounds such as irinotecan and topotecan derived from Nyssaceae *Camptotheca accuminata* as the main inducers of apoptosis against colorectal and ovarian cancer. *Kedrostis foetidissima*, which we found to be the most potent of the plant extracts tested, has been found to contain cucurbitacins B, D, E, I, J, K and L by Konopa *et al.* (1974). They further went on to show that *K. foetidissima* exhibits high cytotoxicity to Hela and KB human (Konopa *et al.* 1974), however the synergistic activity of these cucurbitacins need to be studied further to understand their exact activities on the different cancer cell lines.

On the other hand, *E. mauritanica* belongs to Euphorbiaceae family, whose extracts include methyl esters, their derivatives and diterpene polyesters. Other compounds that have been discovered include alkaloids, phenolic compounds, cyanogenic glucosides and glucosinolates which have been identified as players in the induction of apoptosis in some cells (Rizk 1987). This study clearly shows promising results for this plant extract; however, further studies that

entail isolation and research into the biological activities of the active compounds from the plant *E. mauritanica* would provide better insight.

As with all potential anti-tumour agents, it is crucial to understand the molecular mechanism underlying their apoptotic activities. In this study, we concentrated on the two genes RBBP6 and p53. The involvement of p53 and apoptosis in the cucurbitacins-mediated effect has been studied in various cancer cells (Miro 1995; Rios *et al.* 2005). In response to various cellular stresses, p53 is phosphorylated on NH<sub>2</sub>-terminal residues. In particular, the phosphorylation of Ser 15 affects the interaction with the negative regulator MDM2 oncoprotein and enhances contribution to the stabilization of p53 (Chen *et al.* 2008; Suzuki *et al.* 2007). Phosphorylation of p53 is mediated by phosphoinositide-3-kinase (PI3K)-related proteins, including *Ataxia telangiectasia*-mutated (ATM), ATM- and Rad3- related kinase (ATR) and DNA-dependent protein kinases (DNA-PKs).

While p53 plays numerous roles in the cell, our experiments focused on distinguishing whether *K. foetidissima* and *E. mauritanica*-induced apoptosis was p53 dependent or independent. On the other hand RBBP6 is a negative regulator of p53 that results in the ubiquitination of p53 thereby leading to cell proliferation (Motadi *et al.* 2011). *Kedrostis foetidissima* and *Euphorbia mauritanica* treated cells revealed that p53 was not significantly increased (Tables 5.1 and 5.2) following treatment with the extracts. Similarly, RBBP6 levels did not differ among the *K. foetidissima* and *E. mauritanica* treated cells with less than 1/16 fold decrease (Tables 5.3 and 5.4). The biological activities of these plants have not been studied in depth, however, several studies have looked at the molecular mechanisms of different cucurbitacins and *Euphorbia* species on different cancer cells. Lee *et al.* (2010) reviewed the molecular mechanism of

cucurbitacins and gathered that they target different oncogenic signalling pathways that have been implicated in cancer such as the MAPK, JAK-STAT and the Akt-PKB pathways. The JAK-STAT pathway induces Signal Transducers and Activators of Transcription (STATs) and Janus-Kinases (JAKs). Cucurbitacin B was found to inhibit downstream phosphorylation of some STATs and JAKs in pancreatic tumours, thereby inhibiting this pathway affecting several of its downstream targets involved in apoptosis such as p53 and Bcl-2 (Lee *et al.* 2010). Escandell *et al.* 2008 investigated cucurbitacin R and cucurbitacin I on HCT116 cells harbouring a Ras mutant and found that they induced expression of p53 and p21. They further reported that cucurbitacin R resulted in p53 protein overexpression and Bcl-2 downregulation following 8, 12 and 18 hour treatments in RAW 264.7 macrophages.

One of the most prominent compounds that has been identified in *Euphorbia* species are jatrophone esters, which belong to *Euphorbia semiperfoliata* (Cragg and Newman 2005). This was discovered following research on chemotypes of the drug taxol (explained in section 1.7.1), which exhibited poor water solubility and side effects. Miglietta *et al.* (2002) looked at the effect of these jatrophone polyesters (250 nM) on MCF-7 cells over a period of 16 hours and observed that they highly induced expression of p53 and also abrogates Bcl-2 probably through the Raf-1 mediated signaling pathway. These results, however, are contradictory to our observations where expression of p53 was slightly reduced. This could be attributed to the diversity of compounds which have been identified from *Euphorbia* species. Hence future studies will require further isolation of the active compounds within *E. mauritanica* and molecular mechanism investigations.

In summary, the crude extracts of *K. foetidissima* induced apoptosis in both MCF-7 and YMB-1 cells whilst those of *E. mauritanica* induced apoptosis in MCF-7. RT-PCR indicated that their mode of action is in a p53 independent manner. Since apoptosis is controlled by such a huge network of genes and signalling molecules, it is possible that these plant extracts control other components of the apoptotic pathway. Further research that incorporates isolation and identification of the active compounds would be required to better understand their molecular mechanism and the molecular pathway that follows this apoptosis induction.

## **6.2 Conclusion and future perspectives**

Inducers of apoptosis are considered good agents in anti-cancer therapeutics. Small molecules and natural plant extracts have recently attracted attention to modern medical science research with their non-lethal activity. In this study, crude extracts of three indigenous South African plants were screened for their antiproliferative activity against two breast cancer cell lines: MCF-7 and YMB-1. This study revealed that not only did *K. foetidissima* and *E. mauritanica* plant extracts demonstrate antiproliferative activity to the breast cancer cells, but they induced cell death by apoptosis, rendering them good agents in anticancer drug discoveries.

It could not be determined in this research which components of the apoptotic pathway are being regulated. Hence, further research which includes isolation and characterisation of the active compounds present in these plants is required. Further research into the molecular targets of the isolated active compounds would also provide better insight on the molecular mechanisms of action of these plant extracts.

In conclusion, this study has further suggested the importance of using or targeting traditional plant extracts as potential therapy against several cancers and other illnesses. However, we

should continue to approach this with caution as most plant extracts turn to be unsuccessful once several secondary metabolites are separated.

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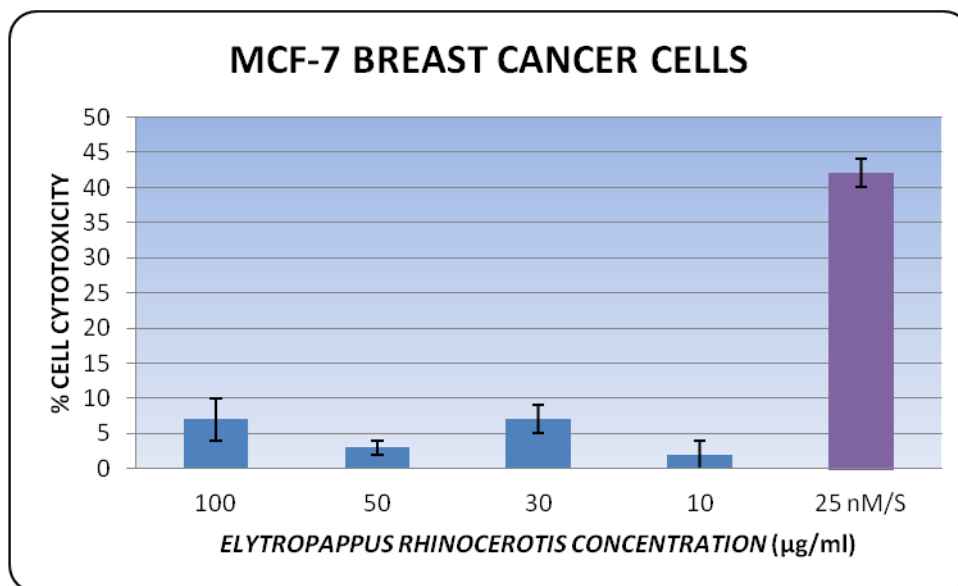
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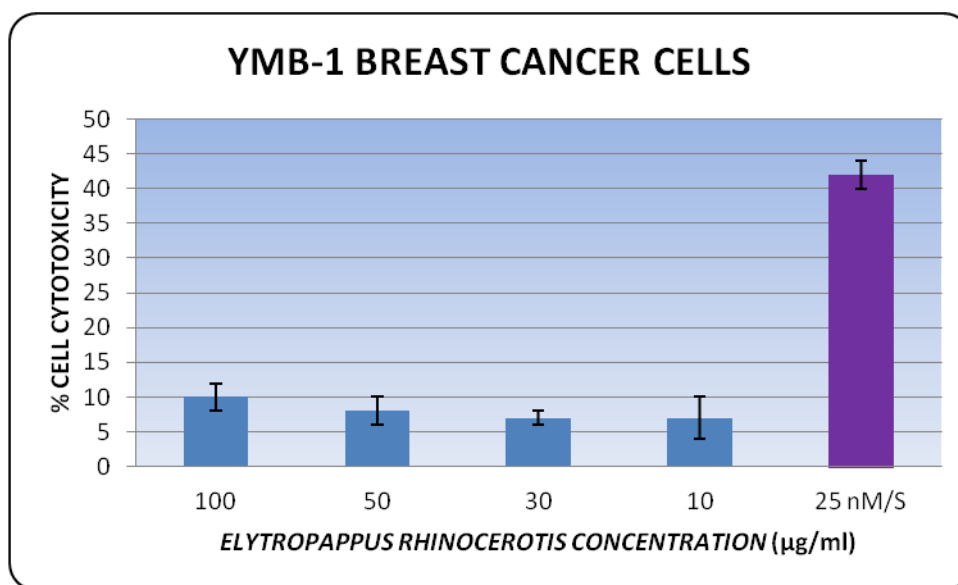
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[Date of access 11/10/2011]

## APPENDIX A



**Figure A.1: Cytotoxicity of *Elytropappus Rhinocerotis* on MCF-7 breast cancer cells.**



**Figure A.2 Cytotoxicity of *Elytropappus Rhinocerotis* on YMB-1 breast cancer cells.**

(Figure A1 and A2) Cells were treated for 24 hours at increasing concentrations of 10, 30, 50 and 100 µg/ml using the plant aqueous extracts. A staurosporine control was included. 25 nM-S represents the positive control, staurosporine, at a concentration of 25 nM. n = 5, p < 0.05

## APPENDIX B

**Table B.1: RNA extraction protocol**

|        |                                                                                                                                                              |
|--------|--------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Step 1 | Cells were harvested( $10^6$ ) and resuspended in 200 $\mu$ l PBS                                                                                            |
| Step 2 | 400 $\mu$ l Lysis/Binding buffer was added, contents were mixed well, then added to upper reservoir of the High Pure Filter Tube assembly                    |
| Step 3 | These were centrifuged for 15seconds at 8000 x g, flowthrough was discarded.                                                                                 |
| Step 4 | Each sample was incubated with a mixture of 90 $\mu$ l DNase I Incubation Buffer and 10 $\mu$ l DNase I for 15 minutes at 25°C.                              |
| Step 5 | Wash Buffer 1(500 $\mu$ l) was added to each sample, these were centrifuged for 15seconds at 8000 x g. Flowthrough was discarded.                            |
| Step 6 | Wash Buffer II (500 $\mu$ l) was added to each sample, these were centrifuged for 15seconds at 8000 x g. Flowthrough was discarded.                          |
| Step 7 | Wash Buffer II (200 $\mu$ l) was added to each sample, these were centrifuged for 2 minutes at 13000 x g. Flowthrough was discarded through collection tube. |
| Step 8 | 50 $\mu$ l of Elution Buffer was added then centrifuged for 1minute at 8000 x g                                                                              |

**Table B.2: The cocktail for cDNA synthesis**

| <b>Reagent</b>            | <b>Volume</b> |
|---------------------------|---------------|
| 10X reaction buffer       | 2.0 µl        |
| 25 mM Magnesium Chloride  | 3.0 µl        |
| Deoxyribonucleotide mix   | 2.0 µl        |
| Specific primers          | 2.5 µl        |
| RNase inhibitor           | 1.0 µl        |
| AMV reverse transcriptase | 1.0 µl        |
| sdH <sub>2</sub> O        | 7.5 µl        |
| RNA sample                | 1.0 µl        |
| Total volume              | 20 µl         |

**Table B.3: Reverse transcription conditions**

| Step   | Temperature | Time (minutes) |
|--------|-------------|----------------|
| Step 1 | 25 °C       | 10             |
| Step 2 | 42 °C       | 60             |
| Step 3 | 70 °C       | 15             |
| Step 4 | 4 °C        | ∞              |

**Table B.4: Real-time PCR cocktail**

| Reagent                                                           | Volume |
|-------------------------------------------------------------------|--------|
| SyBr Green                                                        | 10 µl  |
| Forward primer                                                    | 1 µl   |
| Reverse primer                                                    | 1 µl   |
| H <sub>2</sub> O                                                  | 7 µl   |
| 1µl of the respective cDNAs were added lastly to the capillaries. |        |
| Total                                                             | 20 µl  |

## APPENDIX C

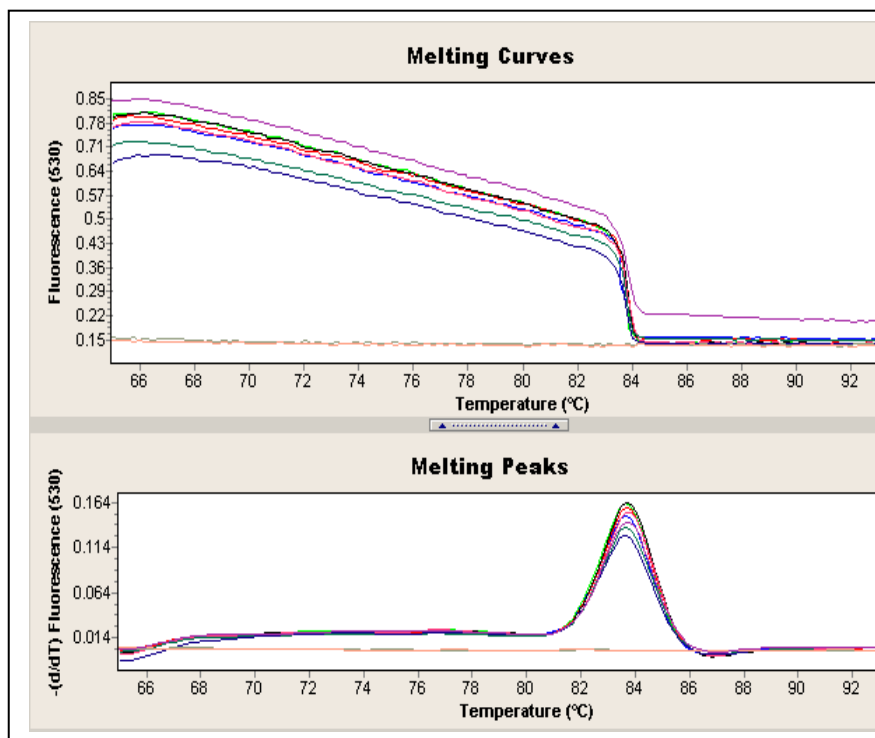
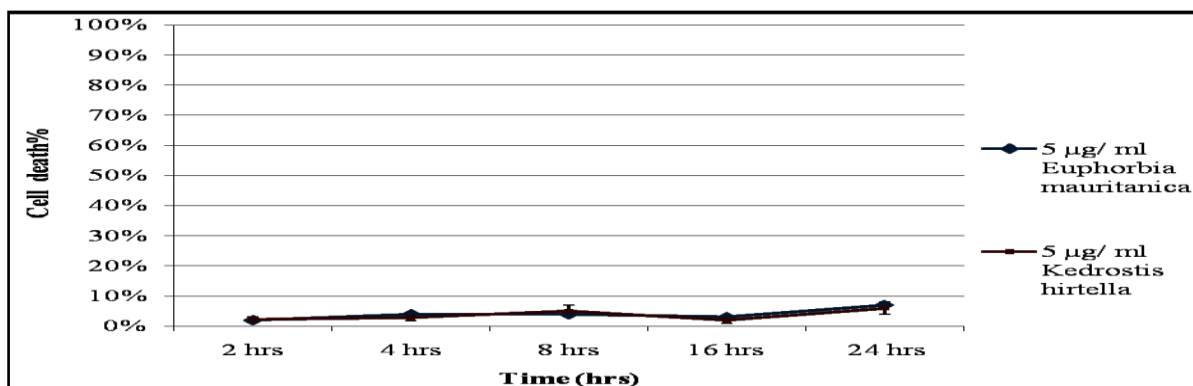


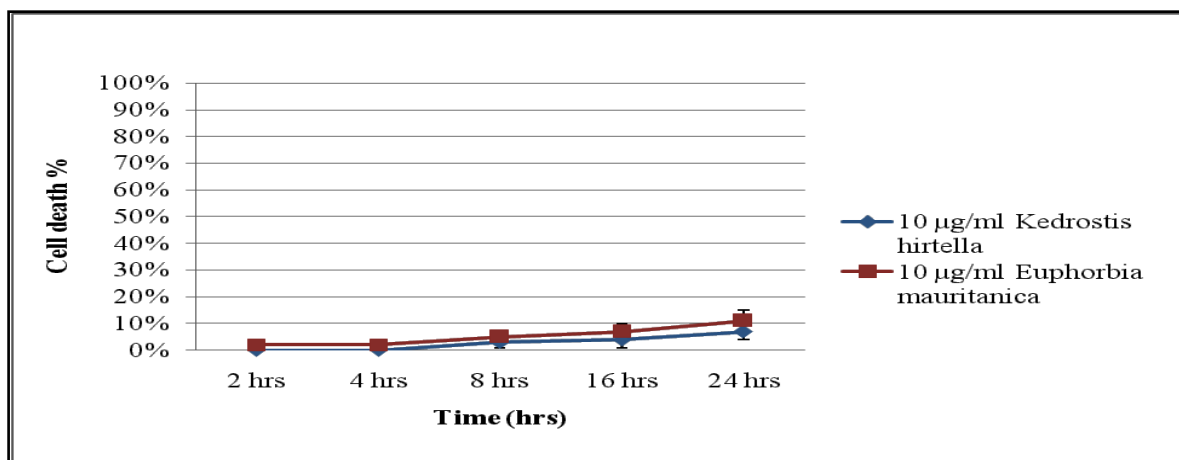
Figure C.1: Melting curves for GADPH

## APPENDIX D



**Figure D.1: Cytotoxicity of *Euphorbia mauritanica* and *Kedrostis hirtella* extracts.**

Cells were treated with herbal extracts at a concentration of 5 µg/ml on MRC5 primary human fibroblast cells, minimal growth inhibition was observed at low concentrations (Thafeni unpublished).



**Figure D.2 : Cytotoxicity of *Euphorbia mauritanica* and *Kedrostis hirtella* extracts.**

Cells were treated with herbal extracts at a concentration of 10 µg/ml on MRC5 primary human fibroblast cells, minimal growth inhibition was observed at minimal concentrations (Thafeni unpublished).

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