

**Effects of 25-hydroxyvitamin D₃ on cell growth and cell death *via*
regulation of the vitamin D metabolising system in a cervical
adenocarcinoma cell line**

by

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Chemical Pathology

in the

Faculty of Health Sciences

at the

University of Pretoria

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Yours sincerely



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Study Output

Conference presentation and published abstract in conference proceedings (Appendix H)

Conference: Microscopy Society of Southern Africa (MSSA), Johannesburg, South Africa

Title: 25-Hydroxyvitamin D3 induces biochemical and morphological apoptosis in the HeLa cell line.

Authors: Zhou E.V.M., Bhoora S., Pillay T.S. and Punchoo R.

Date: 5-8 December 2022

Conference proceedings: ISSN 0250-0418 and ISBN 978-0-6398435-0-6

Abstract

Cervical cancer is the leading cause of cancer-related mortalities in South African women. Vitamin D and its metabolites exert anti-cancer actions in various cancers; however, few studies have explored the effects of vitamin D in cervical cancer.

The vitamin D metabolising system (VDMS) is responsible for the activation and breakdown of the vitamin. Vitamin D is activated by two hydroxylation steps facilitated by 25-hydroxylase (CYP27A1/CYP2R1) and 25-hydroxyvitamin D-1 α -hydroxylase (CYP27B1). The resulting active hormone, 1 α ,25-dihydroxyvitamin D₃ (1,25(OH)₂D₃), binds vitamin D receptor (VDR) in target cells to exert its effects. Both 1,25(OH)₂D₃ and its precursor 25(OH)D₃ are catabolised by 24-hydroxylase (CYP24A1). This study aimed to use the cervical cancer cell line, HeLa treated with 25(OH)D₃, to investigate the cell health parameters (cell count, viability, and cell cycle), cell death (apoptosis, autophagy and necrosis) by brightfield and transmission electron microscopy, and flow cytometry, and regulation of expression of genes and proteins in the VDMS by qPCR and Western blot.

Results indicated that treatment with 5000 nM 25(OH)D₃ significantly decreased cell count and viability and increased the sub-G1 population without inducing cell cycle arrest. Treatment-induced intrinsic apoptosis through disruption of mitochondrial membrane potential, externalisation of phosphatidylserines and activation of executioner caspases -3 and -7. Classic morphological features of apoptosis were also observed with brightfield and transmission electron microscopy. In addition, high-dose treatment with vitamin D increased CYP27B1, CYP24A1 and VDR gene and protein expression. Interestingly, multinucleated cells were observed in all cultures, but the cells were significantly larger following high dose 25(OH)D₃ treatment.

Collectively, the findings indicate that the VDMS acts in an autocrine manner to mediate the effects of supraphysiological doses of 25(OH)D₃ on the growth and death of HeLa cells. Based on the data, it can be inferred that 25(OH)D₃ holds promise as a therapeutic agent for cervical cancer treatment, and further clinical studies are warranted to explore this potential.

Keywords: Cervical cancer; 25-hydroxyvitaminD₃; apoptosis; 1 α -hydroxylase (CYP27B1); 24-hydroxylase (CYP24A1); vitamin D receptor (VDR); multinucleated cells (MNC); polyploid giant cancer cells (PGCC)

List of Abbreviations

1 α OHase	1 α -hydroxylase
1,25(OH) ₂ D ₃	1 α ,25-dihydroxyvitamin D ₃
24OHase	24-hydroxylase
25(OH)D ₃	25-hydroxyvitamin D ₃
7-AAD	7-Aminoactinomycin D
ANOVA	Analysis of variance
Atg	Autophagy-related protein
Bax	Bcl-2 Associated X-protein
BCA	Bicinchoninic acid
Bcl-2	B-cell lymphoma 2
BSA	Bovine Serum Albumin
Ca ²⁺	Calcium ions
CDK	Cyclin-dependent kinase
cDNA	Complementary deoxyribonucleic acid
CO ₂	Carbon dioxide
CYP	Cytochrome P-450 mixed-function oxidases
CYP24A1	24-hydroxylase
CYP2R1/CYP27A1	25-Hydroxylase
CYP27B1	25-Hydroxyvitamin D-1 α -hydroxylase
DEVD	Amino acid sequence Asp-Glu-Val-Asp
DMEM	Dulbecco's Modified Eagle Medium
DNA	Deoxyribonucleic acid
ER	Endoplasmic reticulum

ERSP57	Endoplasmic reticulum stress protein 57
FGF-23	Fibroblast growth factor 23
GI ₅₀	50% growth inhibition
GMP	Cyclic guanosine monophosphate
H&E	Haematoxylin and Eosin
HBSS	Hank's Balanced Salt Solution
HPV	Human Papilloma Virus
IOM	Institute of Medicine
IGFBP-3	Insulin growth factor binding protein 3
INT	2-p-iodophenyl-3-p-nitrophenyl-5-phenyl tetrazolium
LC3	Microtubule-associated protein 1A/1B-light chain 3
LC3-I	Microtubule-associated protein 1A/1B-light chain 3-1
LC3-II	Microtubule-associated protein 1A/1B-light chain 3-2
LDH	Lactate dehydrogenase
LDS	Lithium dodecyl sulphate
M	Molar
MAP	Mitogen-activated protein kinase
MARRS	Membrane-associated rapid-response steroid-binding protein
mL	millilitre
MNC	Multinucleated cell
M-PER	Mammalian Protein Extraction Reagent
MPT	Mitochondrial permeability transition
mRNA	Messenger ribonucleic acid
NADH/NAD ⁺	Nicotinamide adenine dinucleotide

NCCD	Nomenclature Committee of Cell Death
ng/mL	Nanograms per millilitre
nM	Nanomolar
OD	Optical density
p21	Cyclin-dependent kinase inhibitor protein 21
p27	Cyclin-dependent kinase inhibitor protein 27
p53	Tumour protein p53
PBS	Phosphate-buffered saline
PCD	Programmed cell death
PGCC	Polyploid giant cancer cell
PI	Propidium iodide
PMS	N-methylphenazonium methyl sulphate
PTH	Parathyroid hormone
RB	Retinoblastoma protein
PVDF	Polyvinylidene fluoride
REDOX	reduction-oxidation
RNA	Ribonucleic acid
RT-qPCR	Reverse transcription quantitative polymerase chain reaction
RXR	Retinoic X receptor
SDS	Sodium dodecyl sulphate
SEM	Standard error of mean
TEM	Transmission electron microscopy
TMRE	Tetramethylrhodamine ethyl ester
µg	micrograms

μL	microlitres
UVB	Ultraviolet radiation B
VDBP	Vitamin D binding protein
VDR	Vitamin D receptor
VDRE	Vitamin D receptor response element
WHO	World Health Organisation
WR	Working reagent
x g	Times gravity

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Chapter 1

Literature review

1.1 Introduction

Cervical cancer is the leading cause of cancer-related mortalities in South African women¹. It affects approximately 1 in 40 women and is the second most diagnosed malignancy among women in the country. Vitamin D and its metabolites exert anti-cancer actions in breast, colon, prostate, skin, and lung cancers. Cervical tissues express the vitamin D metabolising system (VDMS), but only a few studies have explored the relationship between the VDMS, its metabolites and cervical cancer.

1.2 Vitamin D: History, Metabolism, and Intracellular Signalling

1.2.1 Discovery of Vitamin D

Vitamins are trace nutrients obtained from the diet are essential for specific physiological processes². Professor Elmer McCollum's research provided evidence that cod liver oil contained a curative or preventative agent for rickets, though the exact nature of this agent was not yet identified³. Sir Edward Mellanby's experiments with dogs in 1920 aimed to uncover the precise component of cod liver oil responsible for treating or preventing rickets⁴. These studies ultimately led to the discovery and classification of vitamin D²⁻³. Huldshinsky and Chick *et al.* conducted independent studies that demonstrated how exposing children to sunlight or ultraviolet light could prevent or cure rickets². Their findings prompted many animal studies, which proved that sunlight plays a critical role in activating vitamin D and maintaining calcium balance²⁻³.

Subsequent experiments proved the existence of two variations of vitamin D. Vitamin D₂ (ergocalciferol), sourced from plants, is derived from the provitamin ergosterol (Figure 1.1a). Vitamin D₃ (cholecalciferol) is obtained from animals (Figure 1.1b)⁴⁻⁵. In addition, vitamin D₃ can be synthesised in the skin upon exposure to solar ultraviolet B (UVB) radiation. Ultraviolet radiation cleaves 7-dehydrocholesterol in the skin to pre-vitamin D₃. Pre-vitamin D₃ thermally isomerises to vitamin D₃ (cholecalciferol)⁶. Both vitamin D₂ and D₃ can be sourced from the diet; however, very few foods naturally contain or are fortified with vitamin D⁵.

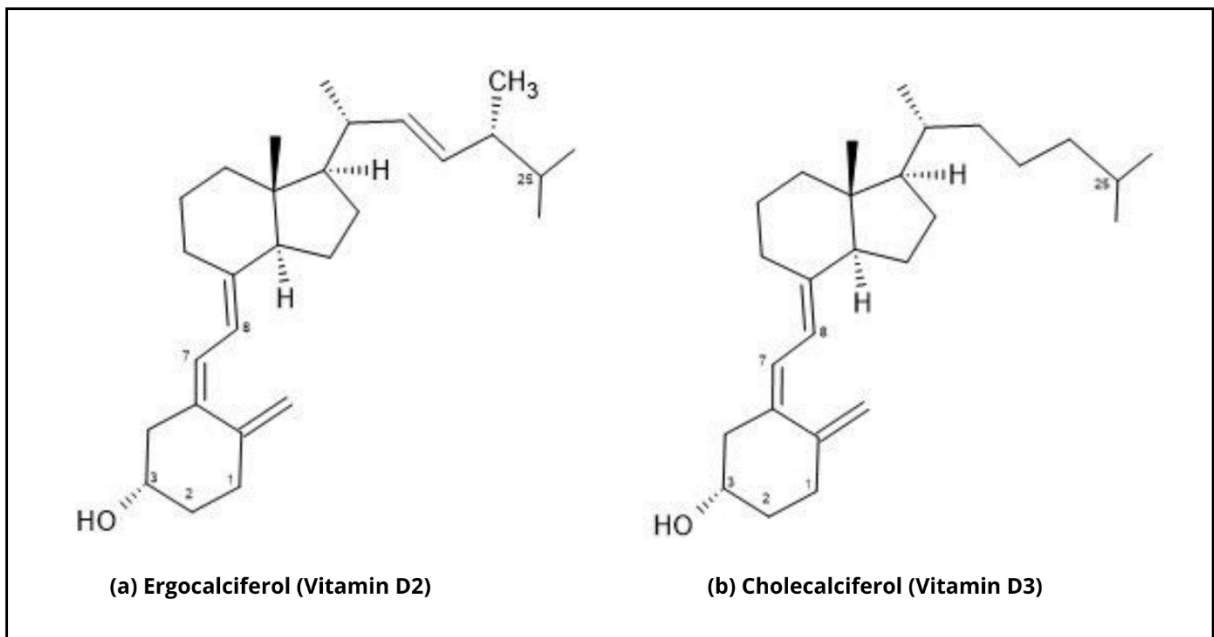


Figure 1.1 Chemical structures of forms of Vitamin D. Structures of vitamin D₂ and D₃ drawn using ACD/ChemSketch (Freeware) 2021.1.0. These two compounds undergo two sequential hydroxylation steps to produce the active metabolite, 1,25-dihydroxyvitamin D (calcitriol).

1.2.2 Vitamin D metabolism

Cytochrome P-450 mixed function oxidases (CYPs) activate vitamin D₂ and D₃ obtained from the diet and skin through two hydroxylation steps. The initial hydroxylation, which takes place on carbon-25 (C25), is carried out by 25-hydroxylase (CYP2R1/CYP27A1) in the liver. The resulting metabolite, 25-hydroxyvitamin D₃ (25(OH)D₃/calcidiol), circulates in blood bound to the vitamin D binding protein (VDBP) or albumin. In the kidneys, 25-hydroxyvitamin D-1 α -hydroxylase (CYP27B1, 1 α OHase) mediates the second hydroxylation step by adding a hydroxyl group to the alpha position of carbon 1 on 25(OH)D₃⁷. This process produces 1 α ,25-dihydroxyvitamin D₃ (1,25(OH)₂D₃/calcitriol), which is the potent form of vitamin D₃ (Figure 1.2)⁴. Both 25(OH)D₃ and 1,25(OH)₂D₃ are catabolised by 24-hydroxylase (CYP24A1, 24OHase), resulting in an inactive water-soluble compound called calcitroic acid that is excreted in bile^{5,7}.

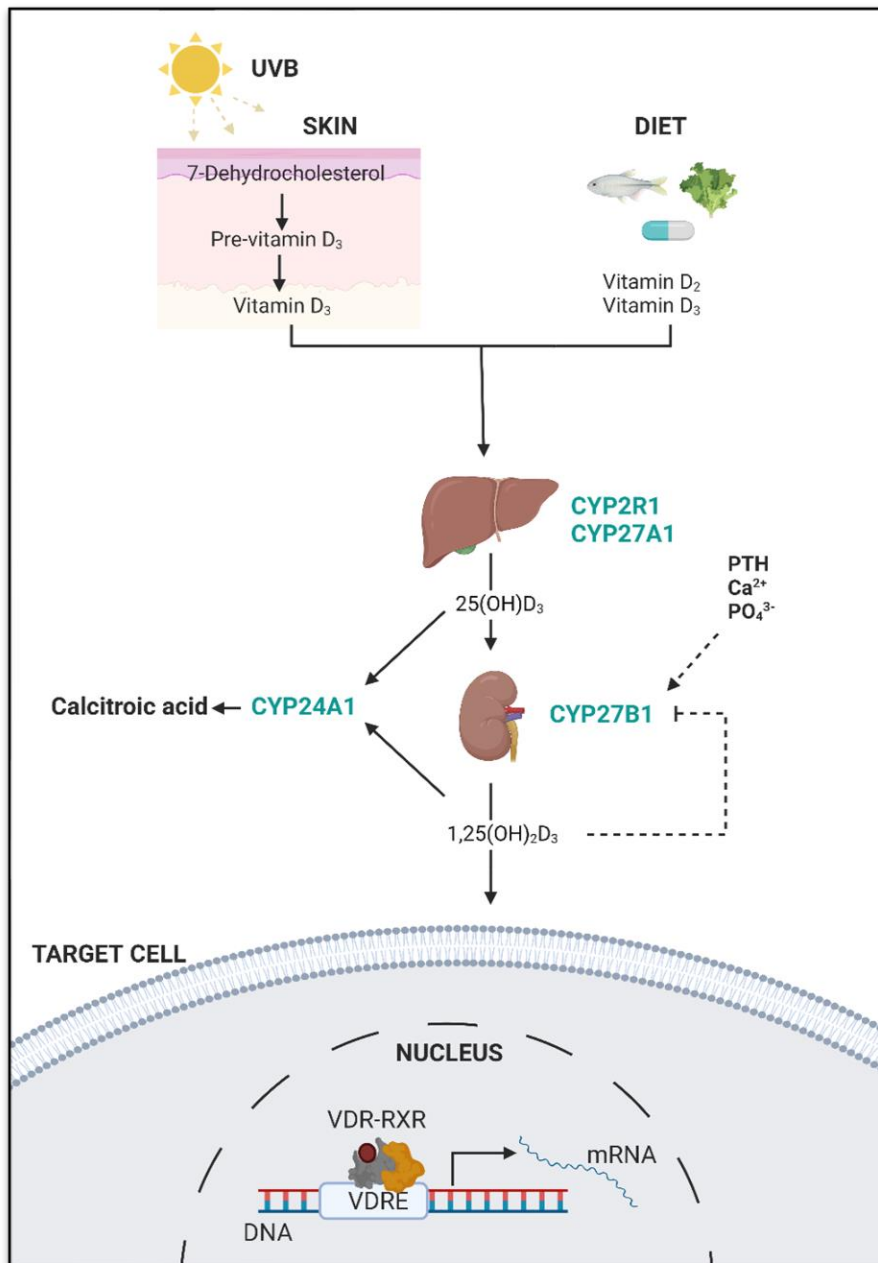


Figure 1.2 Vitamin D metabolism and cell signalling. Vitamin D₃ is synthesised upon exposure to ultraviolet B radiation (UVB) or is taken up in the diet. It is transported to the liver, where it undergoes hydroxylation via CYP2R1 or CYP27A1. The product, 25(OH)D₃, is transported to the kidneys, where it is hydroxylated to produce 1,25(OH)₂D₃ via CYP27B1. This 1,25(OH)₂D₃ can inhibit CYP27B1 expression through a negative feedback mechanism. CYP27B1 expression can be increased by parathyroid hormone (PTH) and calcium (Ca²⁺), and phosphate ions (PO₄³⁻). The 1,25(OH)₂D₃ binds the vitamin D receptor (VDR), which then dimerises with the retinoid x receptor and initiates transcription in the target cell. Both 25(OH)D₃ and 1,25(OH)₂D₃ are broken down by CYP24A1 to produce calcitriol. Created with BioRender.com

The cytochrome P450 oxidases involved in vitamin D metabolism are located in either the endoplasmic reticulum (ER) (CYP2R1) or in the mitochondria (CYP27A1, CYP27B1 and

CYP24A1) of a cell⁸. The activity of these enzymes is not solely limited to the liver and kidney but has been observed in several other tissues such as the lungs, adrenal glands, intestines, colon, breast, and prostate⁶. Therefore, 1,25(OH)₂D₃ can exert its effects in either an endocrine or intracrine, autocrine, or paracrine mode of action⁷.

1.2.3 Genomic effects of 1,25(OH)₂D₃

The most biologically active metabolite of vitamin D, 1,25(OH)₂D₃, is hormonally active. The binding of 1,25(OH)₂D₃ to the vitamin D receptor (VDR) in target cells is utilised in signal transduction pathways^{4,9}. The VDR is a member of the steroid-thyroid-retinoid receptor superfamily of ligand-activated transcription factors (nuclear receptors) and can interact with other transcription factors⁷. The retinoid X receptor (RXR) is the best-studied transcription factor to bind to VDR. The liganded VDR heterodimerises with RXR and then binds to regulatory vitamin D response elements (VDRE) in promoter regions of target genes¹⁰⁻¹¹. As a result, coregulators are recruited, and transcription is initiated.

The VDR is present in several tissues, and 1,25(OH)₂D₃ regulates 3-5% of genes. Therefore, vitamin D can stimulate upregulation or downregulation of gene expression in multiple physiological systems and tissues^{7,12}. One such gene is CYP24A1 which deactivates 1,25(OH)₂D₃. This autoregulation renders excess 1,25(OH)₂D₃ inactive¹³⁻¹⁴.

1.2.4 Non-genomic effects of vitamin D₃

The activity of 1,25(OH)₂D₃ has been observed in non-genomic pathways. The non-genomic actions of vitamin D influence various signalling pathways by acting through receptors such as the membrane-associated rapid-response steroid-binding protein (1,25D₃-MARRS) receptor¹⁵, a membrane-associated receptor. The downstream regulated pathways operate through phosphoinositide, calcium ions (Ca²⁺), cyclic guanosine monophosphate (GMP), and mitogen-activated protein (MAP) kinase⁹. The MARRS receptor is identical to the endoplasmic reticulum stress protein 57 (ERSP57)¹⁵, which, when associated with vitamin D, has a protective effect against skin cancer and sunlight-induced deoxyribonucleic acid (DNA) damage⁷. However, the non-genomic actions in most cells are not clear.

Although all the actions and mechanisms of vitamin D metabolites are not completely understood, some mechanisms have been clarified by studying various diseases such as cancer.

1.2.5 Vitamin D deficiency

Vitamin D is best known for stimulating calcium and phosphorus absorption, vital for healthy bone mineralisation. Vitamin D deficiency is associated with osteomalacia, osteoporosis, osteopenia, heart disease, diabetes, hypertension, asthma, and cancer⁵. Serum 25(OH)D₃ clinically establishes vitamin D status. The Institute of Medicine (IOM) and the Endocrine society define the cut-off for deficiency at 20 and 30 ng/mL (50 and 75 nmol/L), respectively⁷. Studies have shown that serum levels above 40 ng/mL (100 nmol/L) correlate with reduced risk of several cancers¹⁴. Dietary supplementation at amounts ranging from 2000-4000 international units daily are required to maintain serum 25(OH)D levels above 100 nmol/L¹⁴.

1.3 Vitamin D and cancer

The World Health Organisation (WHO) classifies cancer as a leading cause of death worldwide¹⁶. The term cancer broadly describes the uncontrolled growth and division of abnormal cells. More specifically, cancer originates from genomic mutations triggered by environmental exposure, oncoviruses, and replication errors, leading to alterations in the epigenetics, heterotypic interactions, and the number and arrangement of chromosomes in a cell¹⁷. These changes, known as the hallmarks of cancer, undergo evolutionary and clonal selection during malignancy.

There are believed to be about eight defined hallmarks of cancer¹⁷. These are evading growth suppressors, sustaining proliferative signalling, inducing angiogenesis, resisting cell death, avoiding immune destruction, enabling replicative immortality, deregulating cellular metabolism, and triggering invasion and metastasis¹⁸, as shown in Figure 1.3. Vitamin D, combined with enzymes and other proteins, has been demonstrated to negatively affect cell proliferation, angiogenesis, and metastasis^{7,9} in prostate, breast, and colon cancers.

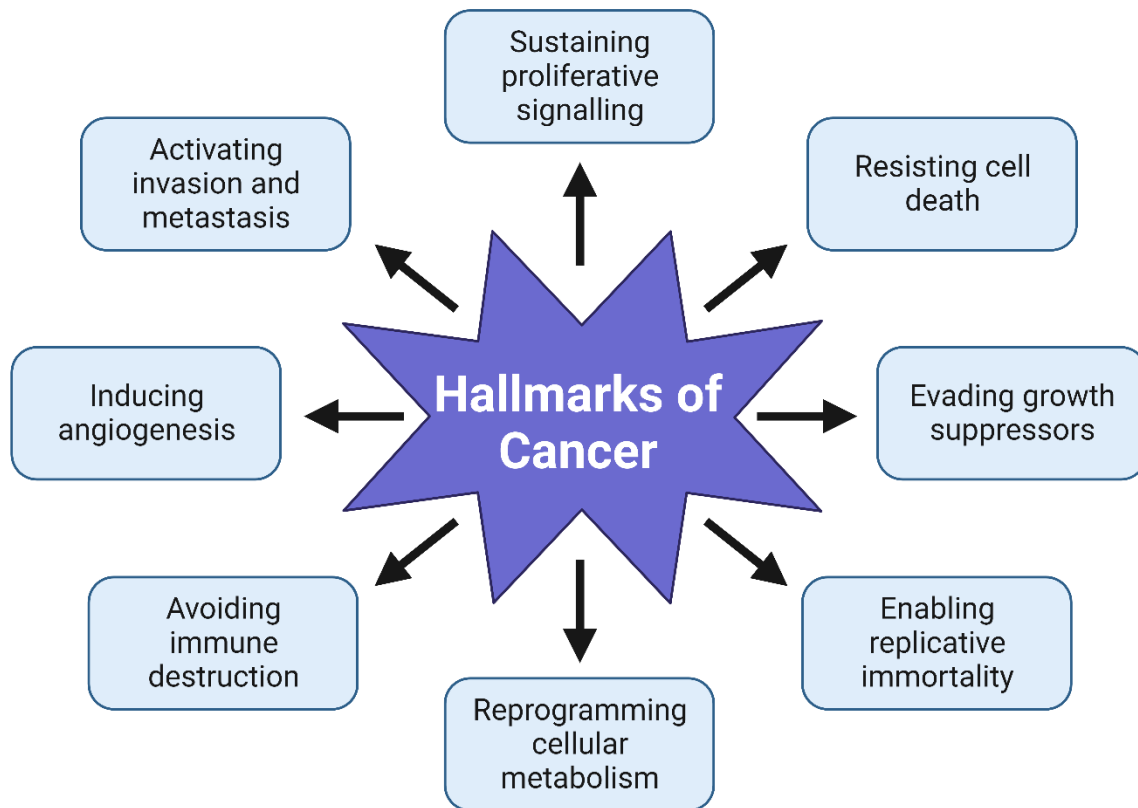


Figure 1.3 Hallmarks of cancer. The eight hallmarks of cancer were proposed by Hanahan and Weinburg in 2000. These hallmarks are believed to be how cells are selected during malignancy. The hallmarks are: sustaining proliferative signalling, resisting cell death, evading growth suppressors, enabling replicative immortality, reprogramming cellular metabolism, avoiding immune destruction, inducing angiogenesis, and activating invasion and metastasis.¹⁸

1.3.1 Anti-proliferation

Cell proliferation is the process by which cells grow and replicate. Normal cells divide continuously until they reach a differentiated state, after which proliferation is arrested. Carcinogenesis results from the deregulation of proliferation¹⁹. Growth factors and receptors are usually altered in carcinogenesis, resulting in abnormal proliferation. This can be counteracted when the cell cycle is tightly regulated¹⁷. Deregulation of the cell cycle enhances the development of malignancy and the overall survival of the abnormal cells.

Vitamin D metabolites inhibit cell proliferation in both normal and malignant cells through multiple cell-specific mechanisms²⁰⁻²¹. They regulate growth factors, the cell cycle, and signalling pathways, demonstrating anti-proliferative properties. Vitamin D treatment increases the expression of insulin-like growth factor-binding protein 3 (IGFBP-3) and cyclin-dependent kinase (CDK) inhibitors such as cyclin-dependent kinase inhibitor protein -21 and -27 (p21 and

p27, respectively), thereby inhibiting cell proliferation. Calcitriol also inhibits telomerase activity and the Wnt/ β -catenin signalling pathway, essential for adult tissue homeostasis²¹⁻²².

1.3.2 Cell cycle arrest

The cell cycle is an ordered process necessary for normal cell growth and the maintenance of tissue homeostasis^{17,23}. The cycle is divided into five phases: quiescence or resting phase (G_0), Gap 1 (G_1), DNA synthesis (S), Gap 2 (G_2), and mitosis (M)²⁴. During the S phase, DNA is replicated to produce two copies of genetic material²⁵. The genetic material is evenly distributed between two daughter cells during M phase. The two gap phases (G_1 and G_2) are required for cell growth (Figure 1.4).

The progression of the cycle is tightly regulated by cyclins, cyclin-dependent protein kinases (CDKs), and tumour suppressors²⁶. For a cell to divide, DNA replication must occur to produce daughter cells that are identical²⁴. Progression through the G_1/S checkpoint is regulated by Cyclin D-CDK4 and cyclin E-CDK2. These cyclin complexes phosphorylate the retinoblastoma (RB) protein leading to the release of E2F transcription factors, enhancing the expression of genes required for DNA replication^{25,27}. Cyclin A-CDK2 phosphorylates cdc6 and E2F1, and drives cells out of the S phase into the G_2 phase of the cell cycle once DNA replication has occurred, Progression through the G_2/M phase is mediated by cyclin B-CDK1 which triggers mitosis²⁸.

If significant DNA damage is detected in the G_1 phase, tumour protein p53 stimulates the transcription of cyclin-dependent kinase inhibitor 1 (p21), inhibiting all cyclin-dependent kinase complexes. This leads to cell cycle arrest until the damage is resolved²⁶. Tumour suppressors such as p53 prevent uninhibited cell proliferation, characteristic of cancer²⁶. Cancer cells differ because the signalling pathway has one or more alterations giving them a proliferative and growth advantage¹⁷.

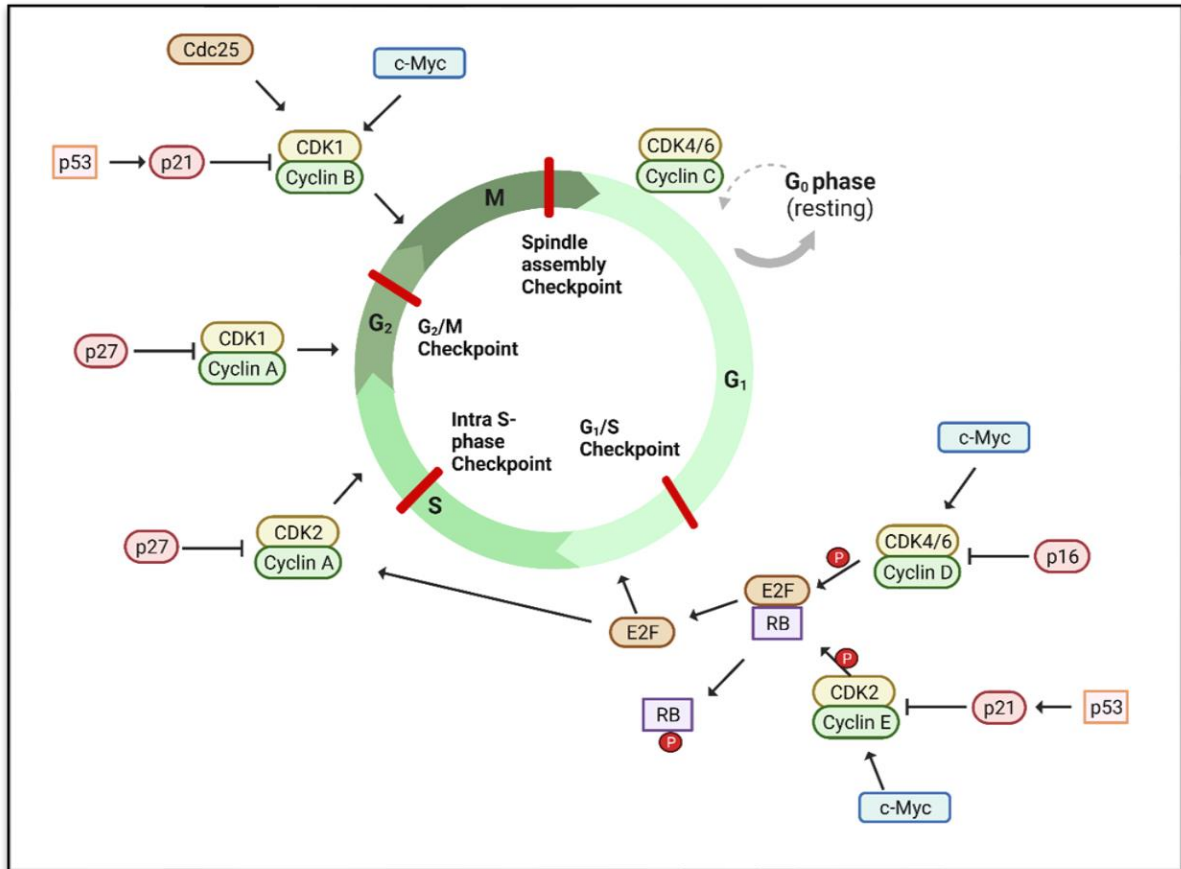


Figure 1.4 The cell cycle. The cell cycle consists of four phases: (1) the G_1/G_0 phase, (2) the S phase, (3) the G₂ phase and (4) the M phase. Cyclins and cyclin-dependent kinases regulate progression through the cycle. Cyclin-CDKs are inhibited by the p16, p21 and p27 proteins. Created with BioRender.com

Vitamin D derivatives have been reported to arrest the cell cycle at the G_0/G_1 phase²⁹ through the upregulation of p21³⁰. This occurs via the transcription factor actions of liganded of VDR at the p21 promoter region³⁰. As previously mentioned, the transcription of p21 is also regulated by p53. Studies have demonstrated that p53 can bind to VDR to regulate the transactivation of p21 messenger ribonucleic acid (mRNA). Activated vitamin D has also been reported to cause cell cycle arrest at the G₂/M phase in ovarian cancer cells via upregulation of the growth arrest and DNA-Damage-Inducible alpha (GADD45 α) protein; however, further studies are needed to explore the mechanism.

1.3.3 Cell death

Cell death is part of a regenerative process by which a cell ceases to perform its function. Under normal circumstances, old cells naturally degrade in an organised manner, defined as programmed cell death (PCD)³¹. This creates an equilibrium between cell death and the

survival of normal cells. PCD determines the fate of the cell when the balance is disrupted³². The Nomenclature Committee of Cell Death (NCCD) categorizes programmed cell death into apoptosis, autophagy, and necrosis based on morphological differences^{31,33-34}.

1.3.3.1 Apoptosis

Cells primarily eliminate themselves through apoptosis, classified as type 1 PCD. Biochemical and morphological changes are used to differentiate apoptosis from autophagy and necrosis^{31-32,35}. Biochemical changes include alterations in mitochondrial membrane potential³⁶, the externalization of phosphatidylserines on the cell membrane³⁷, activation of terminal caspases³⁸⁻³⁹, and DNA fragmentation^{32,40}. Morphological changes include cell shrinkage, chromatin condensation, and plasma membrane blebbing⁴¹. The resulting apoptotic bodies are phagocytosed, degraded, and removed without inciting an inflammatory response^{35,41}. Moreover, apoptosis enables organisms to regulate their cell numbers and tissue size and confers the ability to protect against foreign invasion⁴⁰.

Apoptosis can occur via the intrinsic pathway or the extrinsic pathway. In the intrinsic pathway, stimuli cause the mitochondrial permeability transition (MPT) pore to open. This disrupts the mitochondrial membrane potential and leads to the release of pro-apoptotic proteins into the cell. These proteins activate caspases which in turn fragment DNA. The extrinsic pathway involves binding a ligand to death receptors in the target cell membrane, thereby activating caspase activity and inducing apoptotic cell death^{34,40,42}.

Cancer cells are characterised distinctly by their abnormal control of apoptotic pathways⁴³. Studies have shown that $1\alpha,25(\text{OH})_2\text{D}_3$ suppresses pro-survival proteins B-cell lymphoma 2 (Bcl-2) and telomerase reverse transcriptase while simultaneously activating pro-apoptotic protein Bcl-2 associated X-protein (BAX), thereby inducing apoptosis^{7,12}.

1.3.3.2 Autophagy

Autophagy-dependent cell death is classified as type 11 PCD³¹, and in normal cellular homeostasis, it is essential in recycling cellular content^{31,44}. As its name suggests, autophagy is a self-digestion process aiding in the degradation of damaged organelles⁴⁴⁻⁴⁵. The process of autophagy can, therefore, relieve acute cellular stress or lead to cell death^{32,46}. Liu et al. (2010) stipulated that autophagy regulates multiple physiological processes, such as cell starvation, differentiation, survival and death⁴⁵.

During autophagy, stressor molecules trigger the expression of signal molecules on the surface of damaged organelles. This promotes the formation of a double-membrane vesicle around the organelle known as the autophagosome⁴². The autophagosome will fuse with a lysosome containing hydrolytic enzymes that degrade the engulfed organelles. The autophagy-related (Atg) genes regulates the multi-step process. The most common molecular detection of autophagy is the antibody-based detection of the Microtubule-associated protein 1A/1B-light chain 3 (LC3) gene. This gene is expressed in most cells and is the LC3 protein is cleaved to LC3-I and LC3-II. The latter is localised in the autophagosome membrane and can be detected through immunoblotting to identify autophagy⁴⁶⁻⁴⁷.

Vitamin D induces autophagy in HL-60 human myeloid leukaemia cells and MCF-7 breast cancer cells by upregulating proteins that inhibit mTOR^{11,48}. However, because autophagy can lead to cell death or cell survival, studies must be conducted at varying stages of tumour progression or under varying *in vitro* conditions to conclude how vitamin D treatment can be used in tumour treatments.

1.3.3.3 Necrosis

Necrosis, categorised as type III PCD³¹, is defined as the accidental death of cells resulting in disruption of the plasma membrane causing spillage of intracellular proteins, consequently activating host immune response⁴⁷. In contrast to apoptosis, necrosis is a random, uncontrolled and passive process³¹.

Necrosis can be detected in cells by assessing the lactate dehydrogenase (LDH) activity⁴⁹. The LDH enzyme is involved in the interconversion of pyruvate and lactate, simultaneously with the oxidation and reduction of nicotinamide adenine dinucleotide (NADH/NAD⁺)⁴⁹. When necrosis occurs, LDH leaks into the cytosol. Necrosis is an adaptive form of PCD as it can eliminate cell survival and induce an immune response in target cells⁴³.

1.3.4 Vitamin D metabolising enzymes and receptor in cancer

1.3.4.1 VDR

The vitamin D receptor mediates many effects of 1,25(OH)₂D₃. The VDR has a short N-terminal domain, followed by a DNA-binding domain with two zinc fingers and an RXR-binding domain. Within these domains lie the ligand and coactivator binding domains¹².

Research findings have shown that the impact of cancer on VDR differs vastly across tissues. It has been hypothesised that VDR expression in cancerous tissues is lower than in healthy tissues. However, treatment with vitamin D metabolites has been shown to increase the expression of VDR in colon, breast and lung cancer cell lines²².

1.3.4.2 CYP27B1

Vitamin D metabolising system features both classification of cytochrome P450 enzymes (CYPs), microsomal and mitochondrial. These CYP subtypes do not function independently and are part of the electron transport chain⁵⁰. The CYP27B1 is classified as a mitochondrial vitamin D related CYP and is situated on the inner mitochondrial membrane.

Several factors regulate the expression of both renal and extrarenal CYP27B1. Increased parathyroid hormone (PTH), pro-inflammatory cytokines and low calcium upregulate CYP27B1 expression. In contrast, fibroblast growth factor 23 (FGF-23), hyperphosphataemia, and 1,25(OH)₂D₃ downregulate CYP27B1 gene expression via VDR-mediated action⁵⁰.

Although 1,25(OH)₂D₃ is chiefly synthesised in the kidney, biochemical investigations have shown its expression in extrarenal and cancer tissues. Numerous in vitro studies have shown that the activity of CYP27B1 is lost with increased cancer severity²². Hsu et al. demonstrated that 25(OH)D₃ treatment decreased CYP27B1 activity and growth inhibition in normal and benign prostatic primary cultures⁵¹. It is suggested that the level CYP27B1 determines the anti-cancer protection action of 25(OH)D₃²².

1.2.4.3 CYP24A1

The CYP24A1 enzyme is an inner mitochondrial membrane CYP⁵⁰ responsible for neutralising vitamin D activity⁵². The expression and function of CYP24A1 are regulated in a classic negative-feedback loop⁵⁰.

It has been hypothesised that CYP24A1 expression is upregulated in cancer compared to normal tissue. Upregulation of CYP24A1 expression has been shown in colorectal, breast, ovarian and lung carcinomas⁵³⁻⁵⁴ suggesting that a feature of advanced cancer is increased catabolism of intracellular 1,25(OH)₂D₃. Elevated CYP24A1 degrades high 1,25(OH)₂D₃ intracellular concentrations and thus induces cancer actions²².

1.4 Cervical cancer

1.4.1 Epidemiology and pathophysiology

Cervical cancer is the fourth most common cancer among females worldwide, with roughly 90% of cases occurring in low- and middle-income countries⁵⁵⁻⁵⁷. Approximately 600 000 new cases and over 300 000 deaths from this cancer are reported annually⁵⁶. Cervical cancer is highly treatable when detected in the early stages⁵⁸. However, prevention and management are significantly affected by socio-economic conditions, health care infrastructure and competing illnesses⁵⁵. According to a report by the World Health Organisation, only 62% of countries globally have cervical cancer screening programmes, with under half of them reaching the target populations, further increasing the burden of the disease⁵⁷.

Cervical cancer originates in the narrow, muscular opening of the uterus, known as the cervix. The cervix is divided into the ectocervix (stratified squamous epithelial cells) and the endocervix (columnar epithelial cells)⁵⁵. The two regions meet at the “transformation zone,” which is believed to be the site of premalignant transformation. The common histological subtypes of cervical cancer that arise are squamous cell carcinoma and adenocarcinomas, accounting for approximately 70% and 25% of all cervical cancers, respectively⁵⁶.

The main etiological agent in developing cervical cancer cases is high-risk Human Papillomavirus (HPV)^{56,59}. Human papillomaviruses are small double-stranded DNA viruses that can integrate into the host genome due to persistent infection⁵⁵. Invasive cervical cancers are attributed to HPV-16 and HPV-18, which contain transcriptional units E6 and E7 essential for viral replication⁵⁹.

In South Africa, cervical cancer is the most frequent cancer among women between 15 and 44 years of age⁶⁰. A cervical cancer diagnosis at an earlier age is due to the burden of HIV infections, which decreases immunity and favours HPV acquisition⁵⁶. Women infected with HIV have an increased risk of cervical cancer compared to HIV-negative women.

1.4.2 Cervical cancer and vitamin D

Unambiguous evidence has been presented on the anti-cancer effects of vitamin D₃ in various cancers; however, few studies have indicated the direct method by which vitamin D affects cervical cancer. Some patient studies have validated an inverse correlation between UVB irradiation and the incidence rate of cervical cancer and mortality from cervical cancer^{10,61}.

As previously mentioned, the vitamin D receptor and members of the vitamin D metabolising system play a role in the anti-cancer actions of vitamin D and its metabolites. Investigations by Kloss and colleagues revealed that the human cervical cancer cell line, HeLa, expresses CYP27A1, CYP27B1 and CYP24A1⁶² and that treatment with 1,25(OH)₂D₃ significantly increased gene expression of CYP24A1. Kloss *et al.* also treated HeLa cells with cholecalciferol and 25(OH)D₃, and their results indicated that both cell lines expressed VDR, which is essential for the expression of the other vitamin D metabolising enzymes. Results also showed that CYP24A1 expression increased 13-fold and 176-fold after 72-hour incubation with 100nM of 1,25(OH)₂D₃ in HeLa and OVCAR-3 cells, respectively.

Studies *in vitro* studies by our research group have demonstrated a decrease in cell count and cell viability, induction of apoptosis and upregulation of vitamin D metabolising enzymes in SiHa cells after treatment with cholecalciferol⁶³. In another study, our research group demonstrated similar results in the CaSki cervical cancer cell line³⁰.

To date, there is strong evidence for using vitamin D to prevent colon, rectal and breast cancer¹⁴. It has also been postulated that vitamin D may effectively prevent cancers associated with the bladder, brain, endometrial lining, oesophagus, gallbladder, kidney, lungs, ovaries, pancreas, prostate, and stomach¹⁴. However, the lack of clinical trials, randomized trial evidence and inconsistent observational research remain challenging to understand vitamin D's potential as an anti-cancer therapeutic, thus limiting its development and use in cancer treatment¹². One of the underlying research issues is the concern of increasing serum 25(OH)D₃ to a dangerously high level due to treatment¹². Hence, further studies are needed to elucidate the effects of vitamin D in the treatment and prevention of cancer. Furthermore, there is insufficient evidence to elucidate the effects of vitamin D on cervical cancer.

1.5 Aim and objectives

1.5.1 Aim

This study aimed to use the cervical cancer cell line, HeLa, treated with 25(OH)D₃ to investigate the cell growth, cell death and regulation of expression of genes and proteins in the vitamin D metabolising system.

1.5.2 Objectives

The objectives were:

- To determine the concentration of 25(OH)D₃ that causes 50% growth inhibition (GI₅₀) using the crystal violet assay
- To assess cell growth in 25(OH)D₃ treated cells by enumerating cells with crystal violet assay, alamarBlue® assay and the Muse® Cell Cycle assay
- To evaluate biochemical apoptosis using the Muse® Mitopotential assay, the Muse® Annexin V and Dead Cell assay, and the Muse® Caspase-3/7 assay
- To evaluate biochemical markers of autophagic and necrotic cell death using the Muse® autophagy LC3-antibody-based kit and lactate dehydrogenase assay, respectively
- To assess gross morphological and ultrastructural changes using brightfield microscopy and transmission electron microscopy (TEM)
- To evaluate the expression of CYP27B1, CYP24A1, and VDR, vitamin D metabolising mRNA and proteins using the Quantitative reverse transcription Polymerase Chain Reaction (RT-qPCR) and Western blots

Chapter 2

Materials and Methods

2.1 Cell culture

2.1.1 Materials

The human cervical cancer cell line, HeLa, was purchased from the American Type Culture Collection (ATCC® CCL-2™). Dulbecco's Modified Eagle Medium (DMEM) was purchased from Invitrogen (Carlsbad, CA, USA). Glutamine, Pen-strep and 25-hydroxycholecalciferol (Catalogue number: H4014) were purchased from Sigma-Aldrich (St. Louis, MO, USA) and foetal calf serum from Inqaba Biotechnical Industries (Pretoria, South Africa). Sodium chloride, potassium chloride, potassium dihydrogen phosphate, and sodium dihydrogen phosphate were purchased from Alfa Aesar (Haverhill, MA, USA). Disposable cell-culture flasks were purchased from Lasec (Capetown, South Africa).

2.1.2 Thawing cells

HeLa cells were thawed at room temperature and centrifuged for 5 minutes at 200 x g. Cells were resuspended in pre-warmed DMEM and transferred to a 25 mL culture flask containing 4 mL growth medium. The culture flask was incubated in a ASTEC Water Jacket Type incubator (Model: WCI-165, Japan) at 37°C and 5% carbon dioxide (CO₂) until cells reached 90% confluency, at which point seeding was conducted.

2.1.3 Cell seeding and treating

The cells were exposed to 1 mL of trypsin (0.25%) following their washing with 1X phosphate-buffered saline (PBS; pH 7.4; explained in Appendix A) three times. After incubation at 37°C for 2 minutes, detached cells were collected in 2 mL of fresh medium and counted using Trypan Blue and a haemocytometer. Cells were seeded at different concentrations for various assays and incubated overnight before treatment with 25(OH)D₃ at physiological (260 nM) and supraphysiological (5000 nM) concentrations for 72 hours. Solvent (0.5% ethanol, v/v) and medium controls were included in all experiments.

2.2 Growth inhibition concentration determination

Growth inhibition of 50% (GI_{50}) refers to the concentration of the drug in the medium, which inhibits cell growth by 50% under standard cell incubation conditions⁶⁴. To determine the GI_{50} concentration of 25(OH) D_3 , HeLa cells were seeded at 25 000 cells/mL in a 96-well plate and treated with a serial dilution series of seven concentrations of 25(OH) D_3 , with a maximum concentration of 5000 nM. Solvent and medium controls were included. After 72 hours of incubation at 37°C and 5% CO_2 , the crystal violet assay (described in section 2.3) was used to determine the GI_{50} concentration, with absorbances read at 570 nm using the ELx800 Universal Microplate Reader (Bio-Tek Instruments Inc., Winooski, VT, USA).

2.3 Crystal violet

2.3.1 Principle of assay

Crystal violet is a triarylmethane dye that binds to and stains ribose-containing molecules. As cells undergo death, they detach from culture plates, which makes crystal violet a useful tool for determining the number of viable cells. The degree of staining is directly proportional to the number of viable cells present⁶⁵⁻⁶⁶.

2.3.2 Materials and overview of assay

Crystal violet indicator (Catalogue number: C0775), Glutaraldehyde (Catalogue number: G5882) and Triton X-100 (Catalogue number: X100) were purchased from Sigma-Aldrich (St. Louis, MO, USA).

A six-step process was followed in crystal violet experiments to analyse protein extracts from HeLa cell cultures (Figure 2.1).

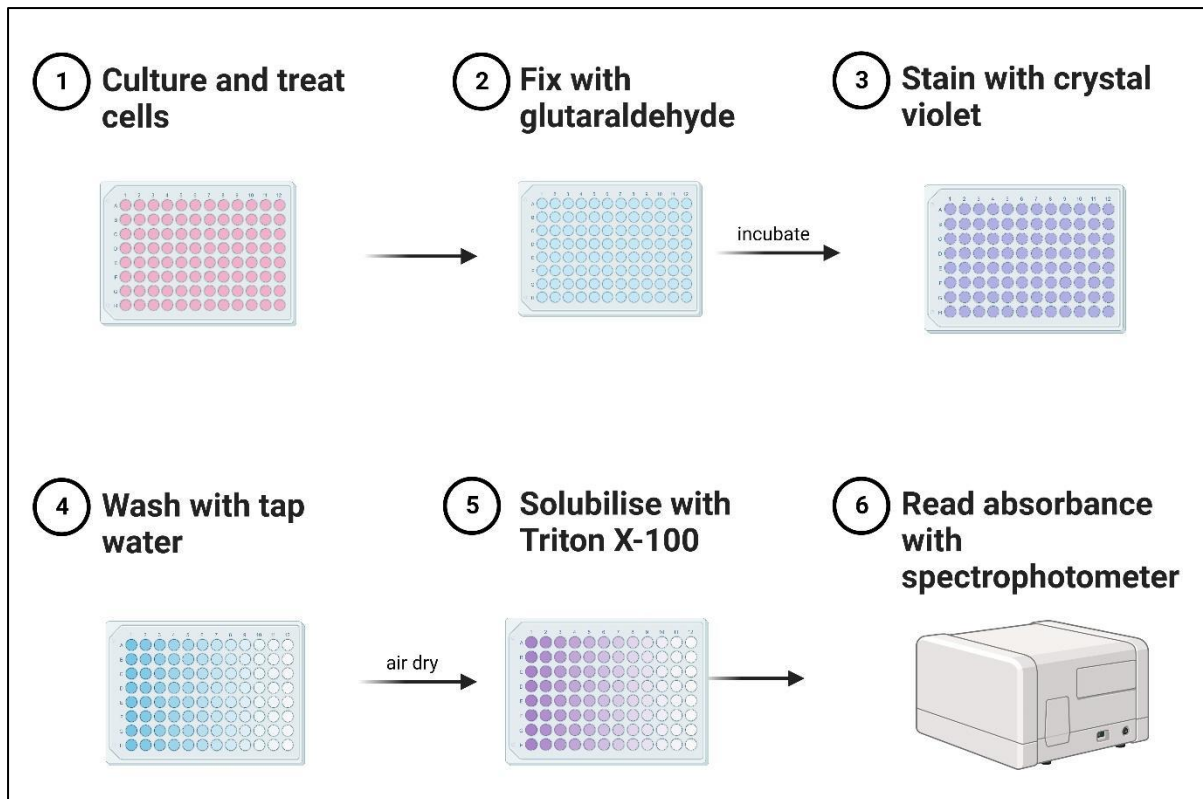


Figure 2.1 Overview of workflow for crystal violet experiments (Source: personal collection created with Biorender.com)

2.3.3 Methodology

The assay was conducted following instructions by Feoktiskova *et al.*⁶⁵. Cells were seeded at 25 000 cells/mL in a 96-well plate and treated with 25(OH)D₃ following overnight incubation. After 72 hours of incubation, cells were fixed with 100 μ L of 1% glutaraldehyde (Sigma-Aldrich) for 15 minutes at room temperature and stained with 100 μ L of 0.1% crystal violet (Sigma-Aldrich) for 30 minutes. Stained cells were washed under running water and air-dried overnight. The stain was solubilised with 200 μ L of 0.5% triton X-100 (Sigma-Aldrich). The optical density was measured at 570 nm by the ELx800 Universal Microplate Reader (Bio-Tek Instruments Inc., Winooski, VT, USA).

2.4 AlamarBlue® assay

2.4.1 Principle of the assay

The alamarBlue® reagent uses the reduction-oxidation (REDOX) indicator resazurin, a non-toxic and cell-permeable phenoxazine-3-one dye that acts as an electron acceptor in the electron transport chain⁶⁶. When reduced to resorufin, fluorescence increases and the colour of the cell culture medium changes from blue to red. The fluorescence/absorbance change is measured over 4 hours to determine the metabolic activity of the cells⁶⁷.

2.4.2 Materials and overview of assay

The alamarBlue® dye (catalogue number: DAL1025) was purchased from ThermoFisher (Waltham, MA, USA).

A three-step process was followed in alamarBlue® experiments to analyse protein extracts from HeLa cell cultures (Figure 2.2).

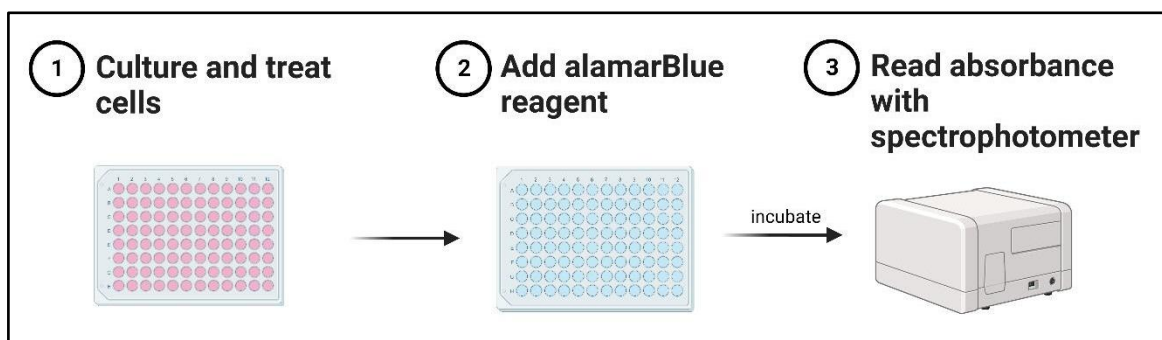


Figure 2.2 Overview of workflow for alamarBlue® experiments (Source: personal collection created with Biorender.com)

2.4.3 Methodology

Following the Invitrogen protocol⁶⁷, HeLa cells were seeded at a concentration of 25 000 cells/mL in a 96-well plate and treated with 25(OH)D₃ following overnight incubation. After a 72-hour incubation with treatment, the culture medium was replaced with 100 µL of fresh medium and 10 µL alamarBlue®. The 96-well plate was covered with foil and incubated for 4 hours at 37°C and 5% CO₂. Absorbance was measured at 570 nm and 600 nm using the ELx800 Universal Microplate Reader (Bio-Tek Instruments Inc., Winooski, VT, USA). Percentage cytotoxicity was calculated using the following equation:

$$\% \text{ cytotoxicity} = \left(\frac{\text{Average 600 nm absorbance} - 570 \text{ nm absorbance}}{\text{Average (average 600 nm absorbance} - 570 \text{ nm absorbance of solvent)}} \right) \times 100$$

2.5 Cell cycle assay

2.5.1 Principle of the assay

The cell cycle is the fundamental mechanism through which cells grow and divide. To distinguish cells at different stages of the cell cycle, a flow cytometric assay utilising a unique combination of propidium iodide (PI) and RNase A is employed. Propidium iodide binds to DNA and indicates DNA content. The assay distinguishes cells based on their DNA content, with fluorescence intensity from PI increasing as DNA replication takes place⁶⁸.

2.5.2 Materials and overview of assay

Muse™ Cell Cycle Kit (Catalogue Number: MCH100106) was purchased from Merck (Modderfontein, South Africa). Micro pipettor tips and 1.5 mL microcentrifuge tubes were purchased from Lasec (Cape Town, South Africa). Ethanol (96%) was purchased from Sigma-Aldrich (St. Louis, MO, USA).

A six-step process was followed in cell cycle experiments to analyse protein extracts from HeLa cell cultures (Figure 2.3).

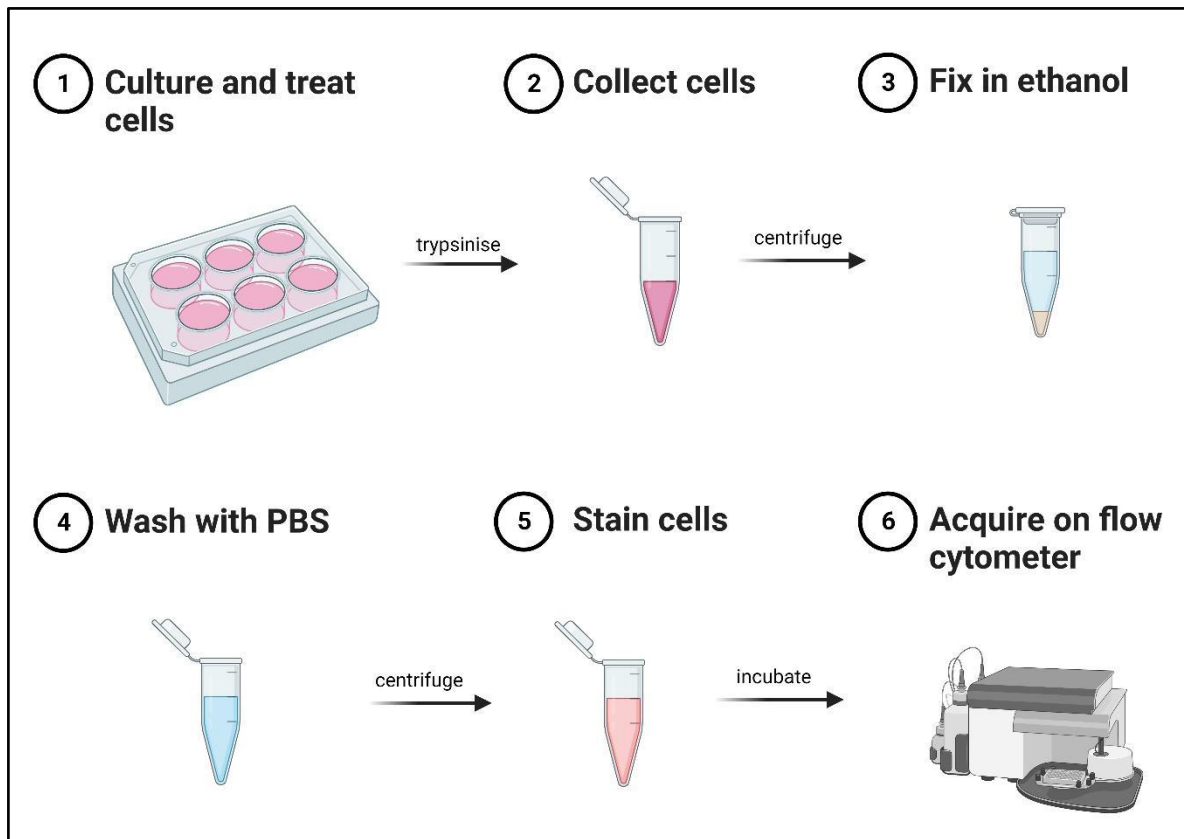


Figure 2.3 Overview of workflow for cell cycle experiments (Source: personal collection created with Biorender.com)

2.5.3 Methodology

Following the manufacturer's instructions⁶⁸, experimental and control cells were seeded at a concentration of 1×10^6 cells/mL. After a 72-hour incubation period with the treatment, cultures were transferred to 50 mL conical tubes and centrifuged at $300 \times g$ for 5 minutes. The supernatants were discarded, and the pellets were re-suspended in 1X PBS (pH 7.4) and gently vortexed. The cells were centrifuged again at $300 \times g$ for 5 minutes, and the supernatant was removed, leaving approximately 50 μ L of 1X PBS (pH 7.4) per 1×10^6 cells. The cells were re-suspended in the remaining PBS by gentle vortexing and fixed by adding the cell suspension to 1 mL of ice-cold 70% ethanol drop-wise while vortexing at medium speed. The samples were then frozen at -20°C for 3 hours.

To stain the experimental and control cell cultures, 200 μ L of the ethanol-fixed cells were pipetted into microcentrifuge tubes, and the cells were centrifuged at $300 \times g$ for 5 minutes. The supernatant was discarded, and the pellets were washed by re-suspending in 1X PBS (pH 7.4) and centrifuging again. The supernatant was discarded, and the cells were re-suspended in 200 μ L of Muse™ Cell Cycle Reagent. The samples were incubated at room temperature for

30 minutes and protected from light. The samples were then analysed using the Guava® Muse™ Cell Analyser (Luminex; Austin, TX, USA). The Muse™ software v1.8.0.3 (Windows, Luminex Corporation, Austin, TX, USA) plotted DNA content index vs cell count. The cell populations in the cell cycle's G0/G1, S and G2/M phases were analysed.

2.6 Mitopotential assay

2.6.1 Principle of the assay

This assay aims to identify apoptosis by evaluating modifications in mitochondrial membrane potential. The MitoPotential dye (tetramethylrhodamine ethyl ester, TMRE) discriminates depolarised and non-depolarised mitochondria, whereas the 7-aminoactinomycin D (7-AAD) dye distinguishes live and dead cells^{40,69}. The dot plots show the distribution of four cell populations: live, live and depolarised, dead and depolarised, and dead cells.

2.6.2 Materials and overview of assay

A Muse™ MitoPotential Kit (Catalogue Number: MCH100110) containing Muse™ MitoPotential Dye (Part No.4700-1580), Muse™ MitoPotential 7-AAD (Part No. 4700-1585) and 1X Assay Buffer (Part No. 4700-1330) was purchased from Merck (Modderfontein, South Africa). Micropipette tips and microcentrifuge tubes were purchased from Lasec (Cape Town, South Africa).

A six-step process was followed in mitopotential experiments to analyse protein extracts from HeLa cell cultures (Figure 2.4).

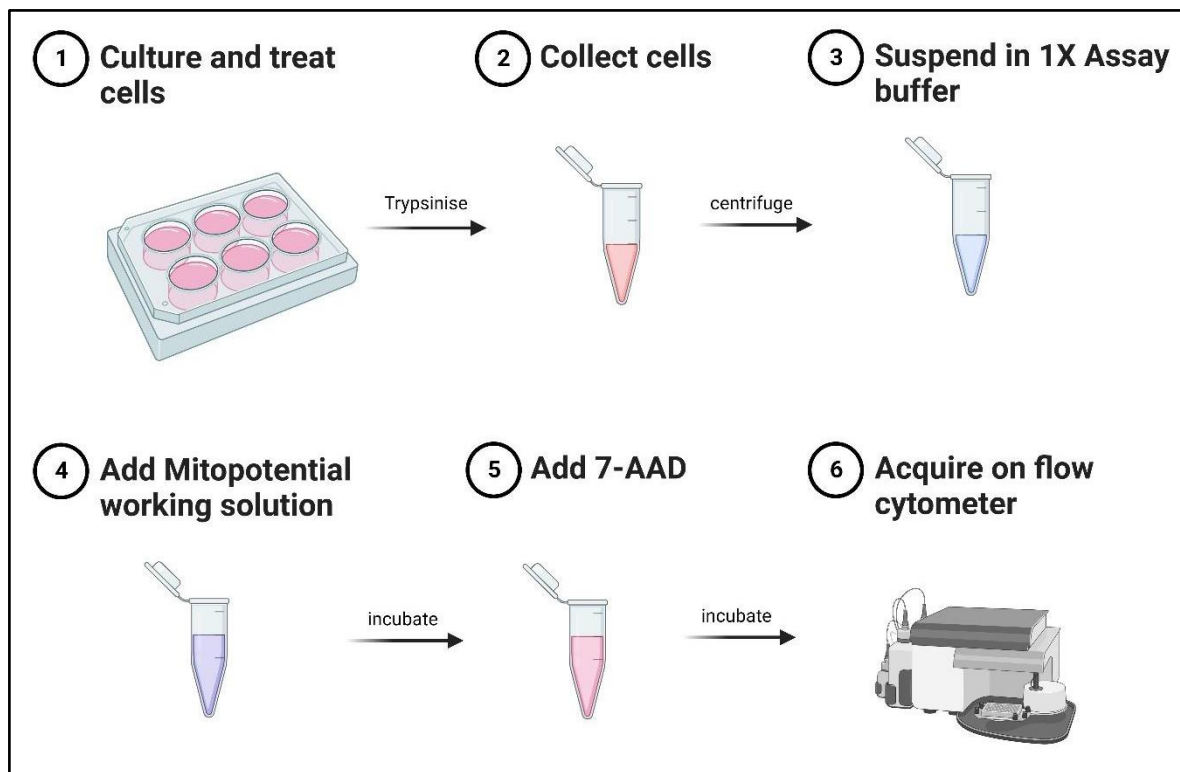


Figure 2.4 Overview of workflow for mitopotential experiments (Source: personal collection created with Biorender.com)

2.6.3 Methodology

As instructed by the Muse™ MitoPotential Kit User Guide⁶⁹, cells were seeded at a concentration of 1×10^6 cells/mL. Actinomycin D was used as a positive control. The experimental and control cells were trypsinised, collected in DMEM, centrifuged and re-suspended at 1×10^5 cells/mL in 1X Assay Buffer. A working solution was prepared by diluting the Muse™ MitoPotential Dye 1:1000 in 1X Assay buffer. Each sample required 95 μ L of the working solution and 100 μ L of cell suspension. The cells were incubated for 20 minutes in a 37°C incubator with 5% CO₂. Then 5 μ L of Muse™ MitoPotential 7-aminoactinomycin D (7-AAD) reagent was added to each sample and incubated for 5 minutes at room temperature. Cells were analysed using the Guava® Muse™ Cell Analyser. Negative and positive control samples were used for gating to exclude cellular debris. Dot plots of mitopotential vs viability were plotted by the Muse™ v1.8.0.3. software (Windows, Luminex Corporation, Austin, TX, USA).

2.7 Annexin V assay

2.7.1 Principle of assay

The primary goal of this assay is to identify the phase of apoptosis, specifically the early and late stages, by utilising Annexin V. This protein binds to externalised phosphatidylserines in the cell membrane. Additionally, including 7-AAD dye enables the differentiation of live cells from dead cells⁷⁰. The dot plots generated show the distribution of four cell populations, which are live, early apoptotic, late apoptotic and dead, and dead cells.

2.7.2 Materials and overview of assay

Muse™ Annexin V & Dead Cell Reagent (Part No. 4700-1485) was purchased from Merck (Modderfontein, South Africa). Micropipette tips and microcentrifuge tubes were purchased from Lasec (Cape Town, South Africa).

A four-step process was followed in annexin V experiments to analyse protein extracts from HeLa cell cultures (Figure 2.5).

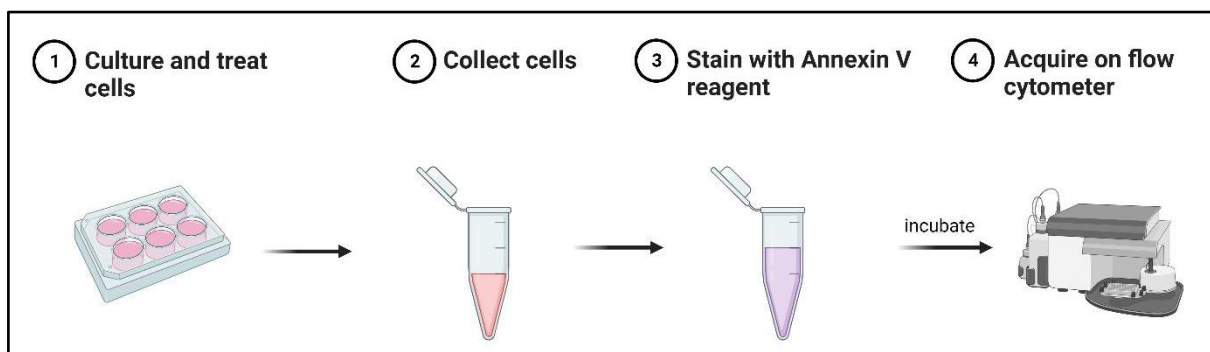


Figure 2.5 Overview of workflow for annexin V experiments (Source: personal collection created with Biorender.com)

2.7.3 Methodology

As instructed by the Muse™ Annexin V & Dead Cell Reagent Kit User' Guide⁷⁰, cells were seeded at a concentration of 1×10^6 cells/mL. Actinomycin D was used as a positive control. The experimental and control cells were treated with trypsin for 2 minutes to allow for detachment. The trypsin was discarded, and the cells were collected with DMEM. Each sample required 100 μ L of the cell suspension and 150 μ L of the Muse™ Annexin V & Dead Cell Reagent (warmed to room temperature) mixed by gentle vortexing. The samples were then

incubated at room temperature for 20 minutes, protected from light. Cells were analysed using the Guava® Muse™ Cell Analyser (Luminex; Austin, TX, USA). Negative and positive control samples were used for gating for method quality control, excluding cell debris and standardising plots. The Muse™ software v1.8.0.3(Windows, Luminex Corporation, Austin, TX, USA) plotted dot plots of annexin vs viability.

2.8 Caspase-3/7 assay

2.8.1 Principle of assay

This assay aimed to identify apoptosis by evaluating the activity of terminal caspases -3 and -7, which are detected using the DEVD peptide (amino acid sequence Asp-Glu-Val-Asp). Caspases-3 and -7 are executioner caspases activated in response to various apoptosis inducers. Once activated, the caspases cleave the initially colourless peptide, which results in the release of a fluorescent product⁷¹. The assay distinguishes between apoptotic and normal cells and live and dead cells⁷².

2.8.2 Materials and overview of assay

The Muse™ Caspase-3/7 Kit containing Muse™ Caspase-3/7 Reagent (Part No. 4700-1505,), Muse™ Caspase 7-AAD (Part No. 4700-1510), 1X Assay Buffer BA (Part No. 4700-1360) and 1X PBS (Part No. 4700-1515) was purchased from Merck (Modderfontein, South Africa). Micropipette tips and microcentrifuge tubes were purchased from Lasec (Cape Town, South Africa).

A six-step process was followed in caspase-3 and -7 experiments to analyse protein extracts from HeLa cell cultures (Figure 2.6).

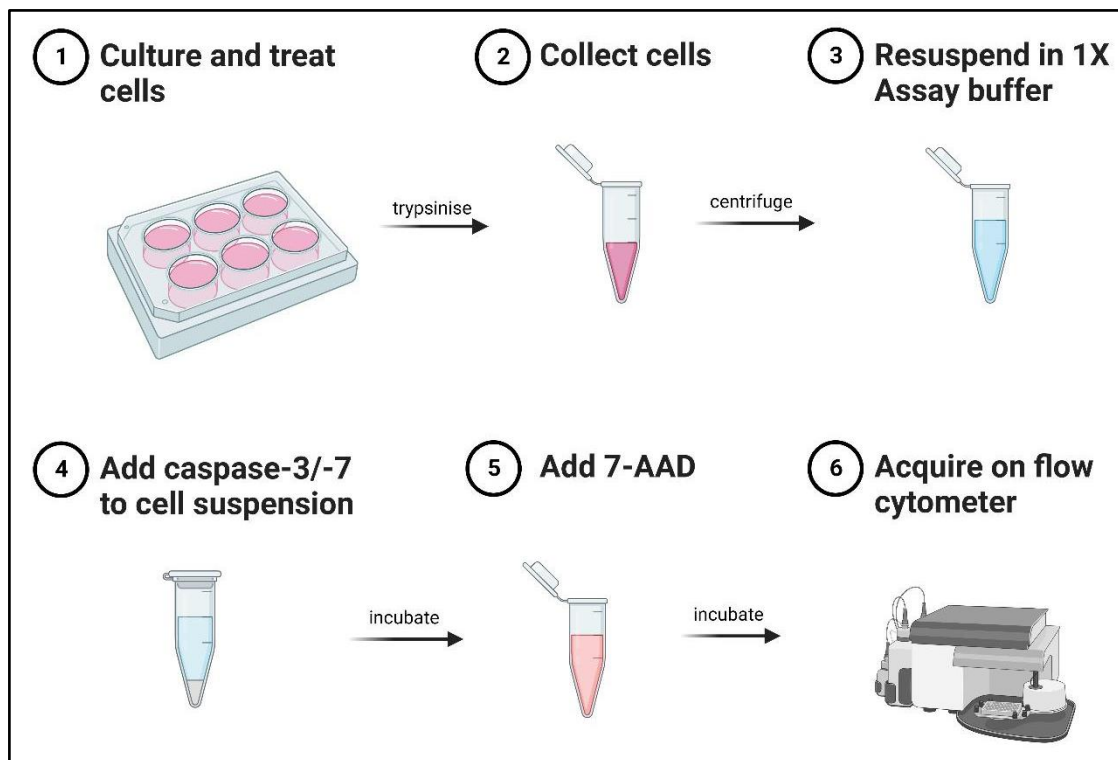


Figure 2.6 Overview of workflow for caspase-3/-7 experiments (Source: personal collection created with Biorender.com)

2.8.3 Methodology

As instructed by the manufacturer⁷², cells were seeded at a concentration of 1×10^6 cells/mL. Actinomycin D was used as a positive control. The Muse™ Caspase-3/7 Reagent, 1X Assay Buffer BA, 1X PBS (pH 7.4), and Muse™ Caspase 7-AAD were acclimatised to room temperature. A Muse™ Caspase-3/7 Reagent working solution was prepared with 3.1 μ L Muse™ Caspase-3/7 Reagent and 21.9 μ L 1X PBS. A Muse™ Caspase 7-AAD working solution was prepared with 10 μ L Muse™ Caspase 7-AAD and 740 μ L 1X Assay Buffer BA. The experimental and control cells were detached from the culture vessel using trypsin and suspended in 1X Assay Buffer BA. Each sample required a mixture of 50 μ L of the cell suspension and 5 μ L of Muse™ Caspase-3/7 Reagent working solution in microcentrifuge tubes. The caps of the tubes were loosened, and the tubes were incubated for 30 minutes at 37°C and 5% CO₂ in a humidified incubator. After incubation, 150 μ L of Muse™ Caspase 7-AAD working solution was added to each tube and mixed with a vortex at medium speed for 5 seconds before incubating at room temperature for 5 minutes, protected from light. Cells were analysed using the Guava® Muse™ Cell Analyser. Negative and positive control samples were used for gating to exclude cellular debris and standardise plots. The Muse™ software

v1.8.0.3 (Windows, Luminex Corporation, Austin, TX, USA) plotted dot plots of caspase-3/-7 vs. viability.

2.9 Autophagy LC3-II assay

2.9.1 Principle of assay

The purpose of this assay is to identify autophagic cells. The assay contains an anti-LC3 mouse monoclonal antibody conjugated to Fluor™ 555, which discriminates between cytosolic LC3-I and autophagosome-associated LC3-II by protecting LC3-II sequestered into the autophagosome during autophagy⁷³.

2.9.2 Materials and overview of assay

The Muse™ Autophagy LC3-antibody kit containing 20X Anti-LC3 Alexa Fluor™ 555, clone 4E12 (Part No. CS208164), Autophagy Detection Reagent Pack (Part No. CF200093), Autophagy reagent A (Part No. CS208212), Autophagy reagent B (Part No. CS208215), and 5X Assay Buffer (Part No. CS202124) was purchased from Merck (Modderfontein, South Africa). Microcentrifuge tubes were purchased from Lasec (Cape Town, South Africa).

A six-step process was followed in autophagy LC3-II experiments to analyse protein extracts from HeLa cell cultures (Figure 2.7).

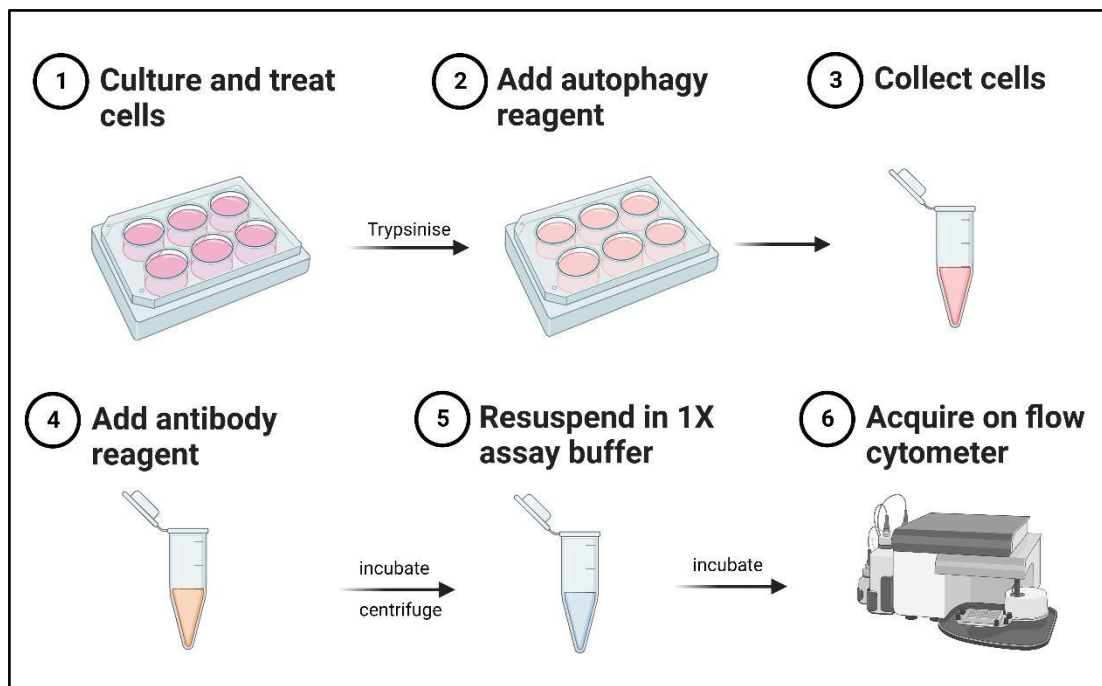


Figure 2.7 Overview of workflow for autophagy LC3-II experiments (Source: personal collection created with Biorender.com)

2.9.3 Methodology

Following the manufacturer's instructions⁷³, cells were seeded at a concentration of 1×10^6 cells/mL. Rapamycin was used as a positive control. Experimental and control cells were washed once with 1X Hank's Balanced Salt Solution (HBSS). The medium was then replaced with a solution of 200 μ L of EBSS and 0.2 μ L Autophagy Reagent A (1:1000 dilution) and was incubated at 37°C for 2-6 hours to induce autophagy under starvation conditions. After the 2-6 hours of incubation treatment, the wells were aspirated to remove culture supernatants and washed with 1X HBSS. To detach cells from the surface of the plate, 100 μ L of Accutase was added for 5 minutes at 37°C. The cells were transferred to Eppendorf tubes, and 100 μ L 1X HBSS was added to each sample tube. The sample tubes were spun down at 300 x g for 5 minutes at 4°C, and the supernatant was discarded. 5 μ L of Anti-LC3 Alexa Fluor™ 555 and 95 μ L of 1X Autophagy Reagent B were added to each sample tube. The sample tubes were incubated on ice for 30 minutes and protected from light. The sample tubes were spun down again at 300 x g for 5 minutes at 4°C. The supernatant was discarded, and the cells were washed with 1X Assay Buffer. The samples were re-suspended in 200 μ L 1X Assay Buffer, analysed immediately by Muse™ and evaluated with the Muse™ v1.8.0.3. software (Windows, Luminex Corporation, Austin, TX, USA) to ensure that event counts per μ L do not exceed 800.

The software developed plots of LC3 intensity vs cell count that indicate the distribution of autophagy induction.

2.10 Lactate dehydrogenase assay

2.10.1 Principle of assay

The lactate dehydrogenase assay was used to determine the presence of necrotic cells. One key feature of necrotic cells is the permeabilization of the plasma membrane, releasing intracellular contents into the extracellular milieu. Lactate dehydrogenase (LDH) is a cytoplasmic enzyme that catalyses the oxidation of lactate to pyruvate while simultaneously reducing nicotinamide adenine dinucleotide (NAD⁺) to NADH. When NADH is in the presence of the electron acceptor, 2-p-iodophenyl-3-p-nitrophenyl-5-phenyl tetrazolium chloride (INT), a tetrazolium salt, is converted to INT formazan, a colourful compound^{66,74}. The colour change is detected with spectrophotometry.

2.10.2 Materials and overview of assay

Acetic acid, DL-Lactic Acid, 2-p-iodophenyl-3-p-nitrophenyl-5-phenyl tetrazolium chloride (INT), N-methylphenazonium methyl sulphate (PMS), nicotinamide adenine dinucleotide (NAD), Tris buffer, and Triton X-100 were purchased from Sigma-Aldrich (St. Louis, MO, USA). The 96-well culture plates and micropipette tips were purchased from Lasec (Capetown, South Africa).

A four-step process was followed in lactate dehydrogenase experiments to analyse protein extracts from HeLa cell cultures (Figure 2.8).

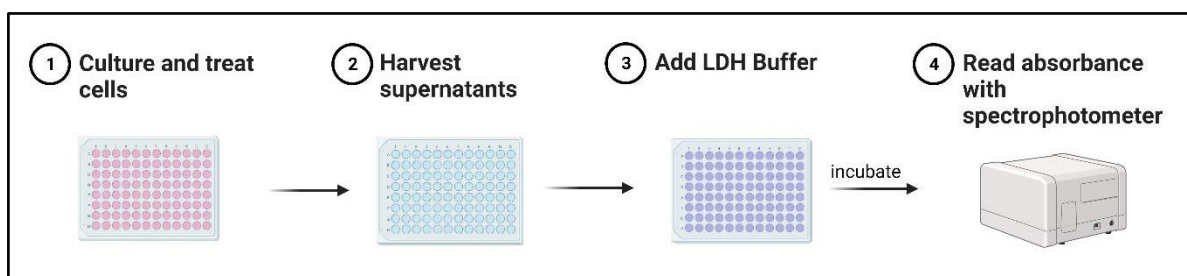


Figure 2.8 Overview of workflow for Lactate dehydrogenase experiments (Source: personal collection created with Biorender.com)

2.10.3 Methodology

Following the protocol by Chan *et al.*⁷⁴, 5000 cells in 100 µL of culture medium were seeded per well in a sterile 96-well cell culture plate, also considering experimental controls; maximum LDH release, Medium alone and volume correction control. Cells were incubated for 12 hours before treatment and then cultured in a 37°C incubator with 5% CO₂. To the maximum LDH release and volume correction control wells, 10 µL of 10X lysis solution (Appendix B) was added, and the plate was incubated for 45 minutes in a 37°C incubator with 5% CO₂. The 96-well plate was centrifuged for 5 minutes at 200 x g. A multi-channel pipette was used to transfer 50 µL of the culture supernatant to a new 96-well plate, taking care not to transfer any cell materials. The 2X LDH assay buffer was reconstituted, and 50 µL was added to the supernatants. The samples were mixed by gently shaking the plate for 30 seconds. The plate was then incubated at room temperature for 30 minutes, protected from light, before mixing 50 µL of 1M acetic acid into each well. The optical density (OD) was measured at 490 nm using the ELx800 Universal Microplate Reader (Bio-Tek Instruments Inc., Winooski, VT, USA). The percentage cytotoxicity was calculated using the following equation:

$$\% \text{ cytotoxicity} = 100 \times \frac{\text{corrected OD from Test well} - \text{corrected OD from untreated well}}{\text{corrected maximum LDH release control} - \text{corrected OD from untreated well}}$$

2.11 Haematoxylin and eosin stain

2.11.1 Principle of assay

Haematoxylin and eosin (H&E) staining assessed the gross morphological changes of 25(OH)D₃-treated HeLa cells. Haematoxylin dye is basic, and when conjugated with an aluminium salt, it stains the acidic structures of the cell, such as the nucleus, blue. Eosin dye is an acidic dye and is used as a counterstain. It stains the basic structures of the cell, such as the cytoplasm, pink⁷⁵.

2.11.2 Materials and overview of assay

Microscope slides and coverslips were purchased from Lasec (Capetown, South Africa). Bouin's fixative was purchased from Sigma-Aldrich (St. Louis, MO, USA). Mayer's

haematoxylin solution, Eosin, Xylol and Entellan mounting fluid were purchased from Merck (Modderfontein, South Africa).

A six-step process was followed in haematoxylin and eosin staining experiments to analyse protein extracts from HeLa cell cultures (Figure 2.9).

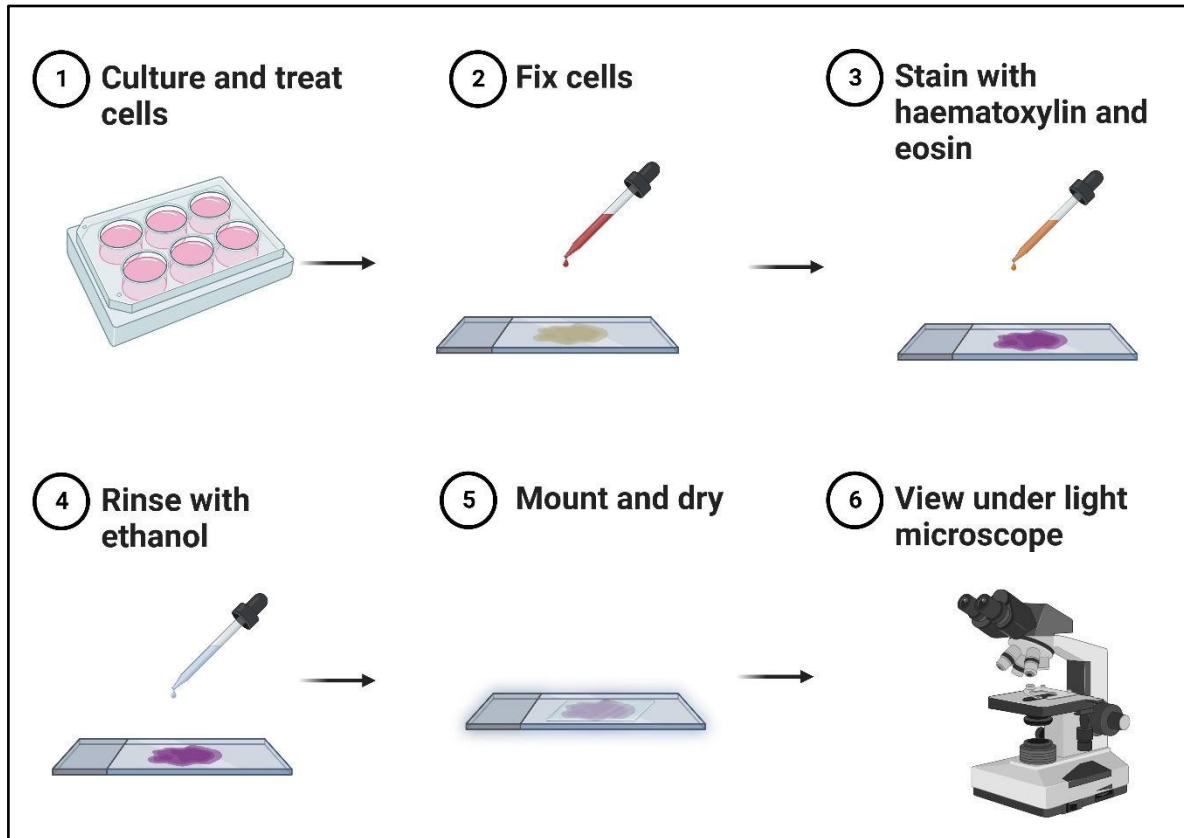


Figure 2.9 Overview of workflow for haematoxylin and eosin experiments (Source: personal collection created with Biorender.com)

2.11.3 Methodology

The protocol was adapted from Ausubel *et al.*⁷⁶ and Conn *et al.*⁷⁷. Cells were seeded onto glass coverslips in a 6-well plate at a seeding density of approximately 5×10^4 cells per well in 3 mL of growth medium. The coverslips were gently rinsed with 1X PBS and then fixed with 500 μ L Bouin's fixative for 30 minutes. The fixative was discarded and replaced with 500 μ L 70% ethanol for 20 minutes. The coverslips were rinsed with tap water and left in Mayer's Hemalum for 20 minutes. The coverslips were rinsed using running tap water for 2 minutes, followed by 70% ethanol. The cells were stained with 500 μ L 1% eosin for 7 minutes. The coverslips were rinsed with 70% ethanol, 95% ethanol, 100% ethanol and xylol, successively for 5 minutes each. This process was repeated once. The coverslips were mounted to microscope slides with

resin and left to dry. The samples were viewed, and images were captured using an Olympus DP74 camera. Cells were counted, measured, and analysed for gross morphological changes using ImageJ 1.53e (Java 1.8.0_172; National Institutes of Health, USA).

2.12 Transmission electron microscopy

2.12.1 Principle of assay

The purpose of this experiment is to assess the ultrastructural features of 25(OH)D₃-treated HeLa cells to identify features of cell death. Transmission electron microscopy allows the visualisation of specimen electrons to pass through the specimen, revealing ultrastructural features⁷⁸.

2.12.2 Materials and overview of assay

Agar resin (product number: AGR1045), Araldite resin (product number: AGR1042), HAZ Benzyl Dimethylamine (product number: AGR1062), Dodecyl Succinic Anhydride EM grade redistilled (product number: AGR1053), UA-Zero® EM Stain (product number: AGR1000), HAZ Lead Citrate (product number: AGR1210-25) and Agar Grids 300 Mesh Copper 3.05mm (product number: AGG2300C) were purchased from Wirsam Scientific and Precision Equipment (Pty) Ltd. (Johannesburg, South Africa).

An eight-step process was followed in transmission electron microscopy experiments to analyse protein extracts from HeLa cell cultures (Figure 2.10).

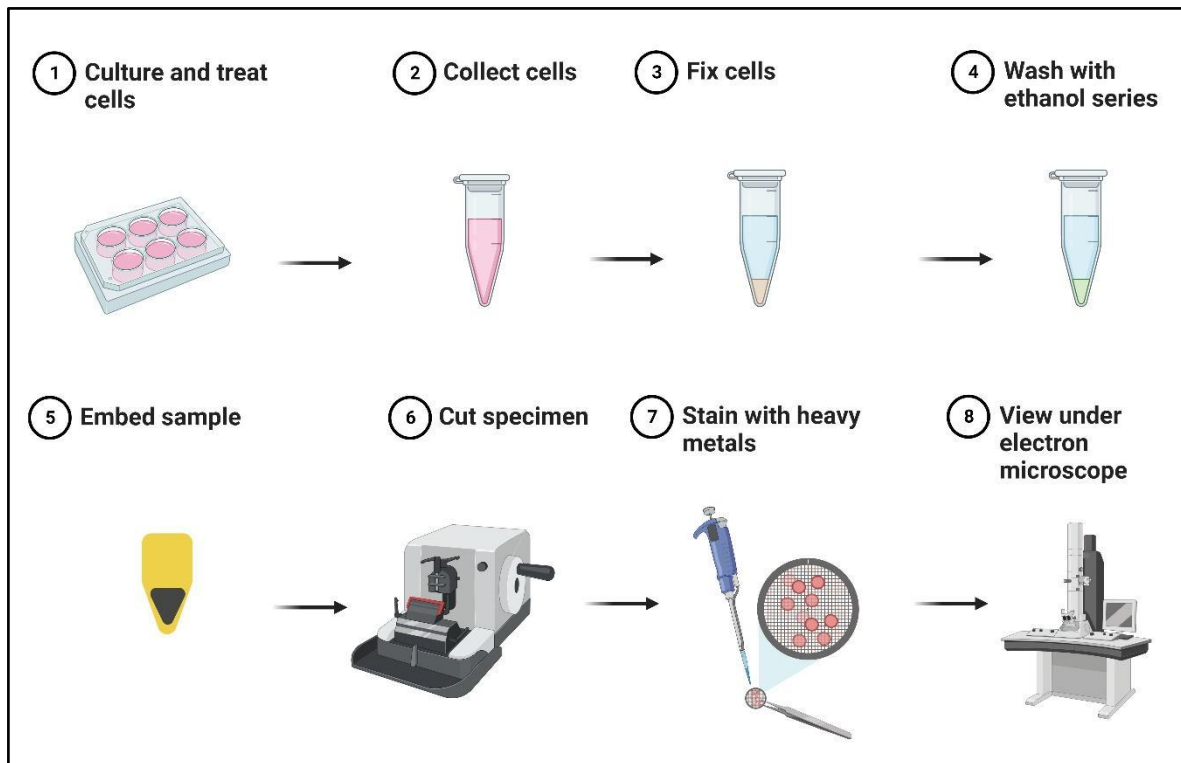


Figure 2.10 Overview of workflow for transmission electron microscopy experiments (Source: personal collection created with Biorender.com)

2.12.3 Methodology

The protocol for electron microscopy from the Anatomical Pathology department of the University of Pretoria was used, adapted from Weakley⁷⁹. Treated cells were washed with 1X PBS and collected in Eppendorf tubes, suspended DMEM, after trypsinisation. The cells were centrifuged for 3 minutes at 300 x g, the supernatant was discarded, and cells were washed with 1 ml of 1X PBS (pH 7.4) and centrifuged.

The supernatant was discarded and replaced with 10 drops of Palade's fixative (Appendix C) for 1 hour of incubation at room temperature. The fixative was removed and replaced with 10 drops of 50% alcohol for 15 minutes. This step was repeated using 70% alcohol. After removing the 70% alcohol, 96% alcohol was added for 30 minutes, followed by two 30-minute incubations with 100% alcohol. Propylene oxide was added for 30 minutes, after which a 50/50 mixture of propylene oxide and embedding medium (Appendix C) was added for 90 minutes. The cells were centrifuged for 3 minutes and then transferred to gelatine capsules. 100% embedding media was added to the capsules, and they were placed in a 65°C oven overnight to allow the media to harden.

Excess embedding media was cut from the edges of the capsule using a blade. Thin golden slices (50 nm) were cut with a Reichert-Jung Ultracut ultramicrotome. The slices were collected on Agar Grids 300 Mesh Copper discs (Wirsam Scientific) and stained with UA-Zero® EM Stain (Wirsam Scientific) for 1 hour, followed by lead citrate (Wirsam Scientific) for 3 minutes. The specimens were viewed using a camera from JEOL Ltd. (Tokyo, Japan) at the Veterinary campus of the University of Pretoria. The images were analysed qualitatively for ultrastructural changes.

2.13 Quantitative Reverse Transcription Polymerase Chain Reaction (RT-qPCR)

2.13.1 Principle of assay

The expression VDR, CYP27B1 and CYP24A1 genes were assessed using real-time reverse transcription polymerase chain reaction. The cells are lysed, and total RNA is extracted, synthesised to complementary deoxyribonucleic acid (cDNA) and amplified with PCR. Real-time reverse transcription polymerase chain reaction (RT-qPCR) uses fluorescent dyes to measure the amount of DNA produced during each cycle of the reaction⁸⁰.

2.13.2 Materials and overview of assay

The iScript™ cDNA Synthesis Kit (Catalogue number: 1708891), iTaq™ Universal SYBR Green (Catalogue number: 1725120) and TubeONE microcentrifuge tubes were purchased from Lasec (Capetown, South Africa). Sterile pipette tips, thin-walled cap tubes, and Lab mark PCR-grade water were purchased from Celtic Molecular Diagnostics (Capetown, South Africa). Qiazol lysis reagent and the Rotor gene were purchased from Whitehead Scientific (Lethabong, South Africa).

A four-step process was followed in PCR experiments to analyse protein extracts from HeLa cell cultures (Figure 2.11).

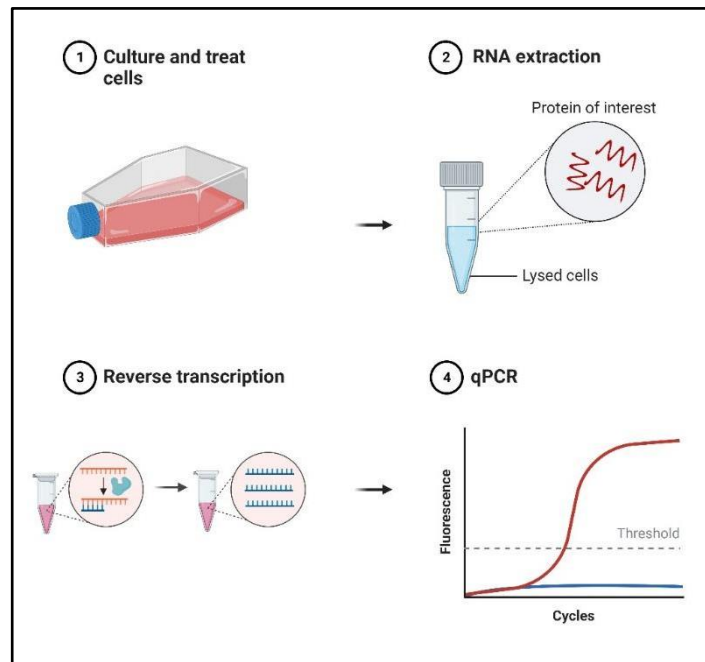


Figure 2.11 Overview of workflow for PCR experiments (Source: personal collection adapted from Biorender.com)

2.13.3 Methodology

Cells were cultured and treated on a 25cm² flask. Total RNA was extracted and quantified following a protocol by Chomczynski and Sacchi⁸¹. The experimental and control cell cultures were washed with cold PBS. About 500 µL of Qiazol lysis reagent was pipetted into each flask. The flasks were then incubated on ice for 5 minutes. Cells were then scraped off the flasks, and the residue was transferred to 1.5 mL tubes. The tubes were then frozen for 1 hour at -80 °C. After thawing the samples, 100 µL of chloroform was added, and the samples were vigorously shaken for 1 second to mix. Samples were incubated at room temperature for 3 minutes and then centrifuged at 4 °C for 15 minutes at 12 000 x g. Centrifuging separated samples into layers of RNA, DNA, and cellular debris. Then 250 µL of the RNA was transferred to a clean, sterile 1.5 mL tube and stored on ice. To each 1.5 mL tube, 250 µL of isopropanol was added, and the tubes were flicked to mix. The samples were incubated at -80 °C for 1 hour. The samples were then centrifuged at 12000 x g for 20 minutes at 4 °C, and the supernatant was discarded. The pellet was washed with 500 µL of cold 75 % ethanol and centrifuged at 7400 x g for 15 minutes at 4°C. The ethanol was discarded, and the pellet was allowed to air dry for 5 minutes before re-suspending in 15 µL of nuclease-free water. The samples were incubated at room temperature for 3 minutes and then placed on ice. The RNA was quantified using an Optizen

NanoQ™ (KLAB Keen Innovative Solutions, Daejeon, Republic of Korea). The total RNA was stored at -80°C.

The total RNA extracted from experimental and control cultures was used to synthesise cDNA following the optimised iScript™ Reverse Transcription Supermix for RT-qPCR kit guidelines (Bio-Rad)⁸². All total RNA samples were normalised to a concentration of 400 ng/μL in a total volume of 10 μL. This 10 μL of RNA was mixed with 4 μL of 5x iScript™ RT Supermix and 6 μL nuclease-free water in a PCR tube to make a 20 μL reaction mixture. The PCR tubes were incubated in a Rotor-Gene Q thermal cycler (QIAGEN, Hilden, Germany). Thermal cycling followed the protocol: priming for 5 minutes at 25°C, reverse transcription (RT) for 20 minutes at 26°C, RT inactivation for 1 minute at 95°C and a hold step at 4 °C. The synthesised cDNA was then diluted with 40 μL of nuclease-free water. The cDNA was then aliquoted and stored at -80°C.

The cDNA was amplified using the iTaq™ Universal SYBR® Green Supermix kit from BioRad⁸³. The iTaq™ Universal SYBR® Green Supermix and cDNA were thawed to room temperature and centrifuged before storing on ice, protected from light. Reaction mixtures containing 5 μL of iTaq™ Universal SYBR® Green Supermix, forward and reverse primers (design in Appendix D) and nuclease-free water were prepared in qPCR tubes, on ice, for all qPCR reactions based on previously determined reaction conditions (Appendix D). The reaction mixtures were mixed by vortexing ensure homogeneity before adding the cDNA of experimental and control samples. A no template control, no reverse transcriptase control and 18S housekeeping gene control were also prepared on ice. The qPCR tubes were then sealed and vortexed for 30 seconds to mix and remove all bubbles in the samples. Amplification was conducted using the Rotor-Gene Q thermal cycler (QIAGEN, Hilden, Germany) programmed according to the following protocol: Polymerase activation and DNA denaturation at 95°C for 30 seconds; Amplification: Denaturation at 95°C for 5 seconds, Annealing, extension and reading for 30 seconds, for 40 cycles; Melt curve analysis: 65-95°C in 0.5°C increments at 5 seconds per step. Annealing temperatures for each primer set can be found in Appendix D. Amplification and melt curves were obtained and used to assess the specificity of the primers. The fold change in gene expression was determined using the delta-delta Ct method.

2.14 Western blot

2.14.1 Principle of assay

Protein expressions of VDR, CYP27B1, and CYP24A1 were measured using Western blotting. Total protein was extracted from cells and separated by size using gel electrophoresis. The proteins are then transferred or blotted to a second matrix where they can be probed for proteins of interest using enzyme-labelled antibodies⁸⁴.

2.14.2 Materials and overview of assay

M-PER™ Mammalian Protein Extraction Reagent (Catalogue number: 78501), EDTA-free Halt™ Protease Inhibitor Cocktail (Catalogue number: 87785), Pierce™ Bicinchoninic acid (BCA) protein assay kit (Catalogue number: 23227), 4X NuPAGE™ Lithium dodecyl sulphate (LDS) Buffer (Catalogue number: NP0008), NuPAGE™ 4-12% Bis-Tris protein gels (Catalogue number: NP0321PK2) and Clarity™ Western ECL Substrate (Catalogue number: 1705061) were purchased from ThermoFisher (Waltham, MA, USA). Methanol (Catalogue number: 34860) was purchased from Merck (Modderfontein, South Africa). Tween 20, 2-Mercaptoethanol (Catalogue number: M6250), and β -actin antibody were purchased from Sigma-Aldrich (St. Louis, MO, USA). Bovine serum albumin (BSA) was purchased from Inqaba Biotech (Pretoria, South Africa). Filter paper, Precision Plus Protein™ Dual Colour Standards (Catalogue number: 1610374), Western blot chemiluminescent substrate, 20X MOPS running buffer (Catalogue number: 161-0788), 10X Tris/Glycine buffer (Catalogue number: 1610734) and Immun-Blot® polyvinylidene fluoride (PVDF) membranes (Catalogue number: 1620177) were purchased from Lasec (Cape Town, South Africa). Anti-rabbit IgG, monoclonal (CYP27B1, Catalogue number: ab206655), Anti-rabbit IgG, polyclonal (CYP24A1, Catalogue number: ab175976) and Goat anti-rabbit IgG, HRP Conjugated (Elabscience, Catalogue number: E-AB-1003) were purchased from Biocom Africa (Centurion, South Africa). Anti-rabbit IgG, HRP-linked antibody (VDR) was purchased from Anatech (Randburg, South Africa).

A six-step process was followed in Western blot experiments to analyse protein extracts from HeLa cell cultures (Figure 2.12).

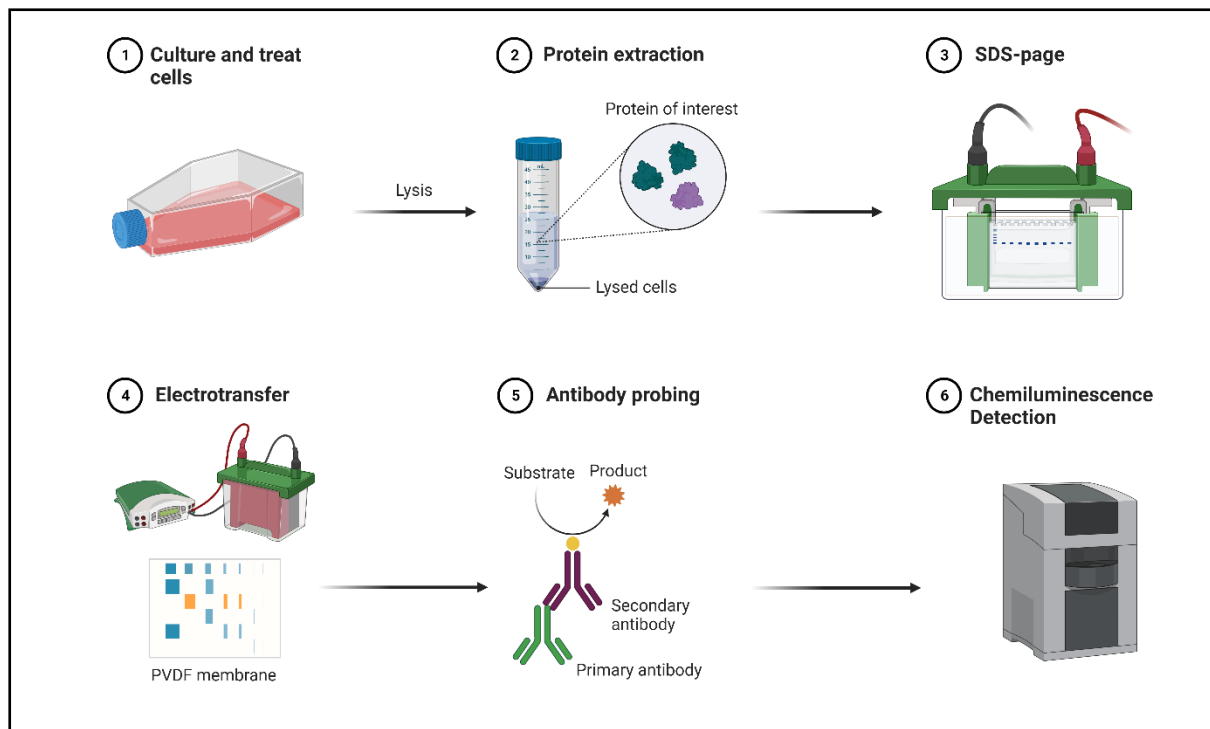


Figure 2.12 Overview of workflow for Western blot experiments (Source: personal collection adapted from Biorender.com)

2.14.3 Methodology

Western blot experiments were conducted according to the Abcam Western Blot protocol⁸⁵. Cells were seeded at a concentration of 100000 cells/mL in a 6-well plate. After 72 hours of incubation with the treatment, experimental and control cultures were washed twice with ice-cold 1X PBS (pH 7.4). After washing, cells were incubated with 500 μ L of M-PER™ Mammalian Protein Extraction Reagent (ThermoFisher) and 100x EDTA-free Halt™ Protease Inhibitor Cocktail (ThermoFisher) at a ratio of 100:1 on ice for 5 minutes. The cells were then scrapped off the plates with plastic cell scrapers and were transferred to pre-cooled 1.5 mL tubes. Cells were then centrifuged at 4°C for 20 minutes at 14000 x g. The lysate was removed, placed in clean 1.5 mL tubes, and kept on ice.

The total protein extracted in the lysate was quantified using the Pierce™ BCA protein assay kit (Thermo Fisher, Catalogue number: 23227) as described by the manufacturer⁸⁶. A BCA working reagent (WR) solution was prepared and divided into 2 mL aliquots. 100 μ L of the experimental and control lysates were added to the BCA WR. The samples were mixed and then incubated in a 60°C water bath for 30 minutes. Samples were then cooled to room temperature, and 200 μ L of each sample was transferred to a 96-well plate. The optical density

was measured at 562 nm using an ELx800 Absorbance Reader (BioTek Instruments Inc., Vermont, United States of America). Bovine serum albumin (BSA) reference standards, ranging from 5-250 µg/mL, were used to generate a standard curve for determining the protein concentration of each lysate. The remaining lysates were aliquoted and stored at -80°C.

The protein concentrations were standardised to 20 µg using M-PER Mammalian Protein Extraction Buffer. To each sample, 6.3 µL of 4X NuPAGE™ LDS Buffer (ThermoFisher) and 0.626 µL 2-Mercaptoethanol (Sigma-Aldrich) were added. The samples were then denatured on a 95°C hot plate for 7 minutes. The samples were allowed to cool to room temperature before loading 20 µL into the wells of a NuPAGE™ 4-12% Bis-Tris protein gel, starting at the second well. The first well was loaded with 5 µL of the Precision Plus Protein™ Dual Colour Standards (Bio-Rad). The gel was loaded into an Invitrogen™ XCell SureLock™ Mini-Cell and XCell II™ Blot Module, which was then filled with 1X XT MOPS running buffer (Bio-Rad). A current of 200 volts was applied to the system for 80 minutes.

After running the gel, the proteins were transferred to an Immun-Blot® PVDF Membrane (Bio-Rad) using the wet transfer sandwich method. Before transfer, the PVDF membrane was activated in methanol (Merck), whilst the sponges and filter paper were soaked in 1X Tris/Glycine buffer (Bio-Rad). The transfer sandwich was assembled (Appendix E) and placed in a Mini Trans-Blot® Electrophoretic Transfer Cell. The system was covered in ice and placed in a walk-in fridge. A current of 110 volts was applied to the system for 90 minutes.

Once the transfer was complete, the PVDF membrane was gently washed with deionised water and then blocked with 5 ml of 0.2% PBS Tween and 2.5% BSA for 1 hour at room temperature. The membrane was then washed three times with 0.2% PBS Tween. The membrane was incubated with primary antibody (anti-rabbit IgG VDR, anti-rabbit IgG CYP27B1 and anti-rabbit IgG CYP24A1) overnight at 4°C (Appendix F). The membrane was then washed three times with 0.2% PBS Tween before incubating with the secondary antibody, HRP Conjugated Goat anti-rabbit IgG (Appendix F), for 1 hour at room temperature. The membrane was washed thrice with 0.2% PBS Tween before viewing with the ChemiDoc™ MP System (Bio-Rad, Catalogue number 1708280).

To enhance viewing, an 800 µL 1:1 dilution of the Clarity™ Western ECL Substrate (Bio-Rad) was added to the membrane before viewing. The membrane blot was analysed semi-quantitatively using the Image Lab™ v6.1 software (Windows, Bio-Rad, Hercules, CA, USA).

The membrane was then washed with 0.2% PBS Tween and stained with the loading control, anti- β -actin (Appendix F), to check the specificity and effectiveness of the transfer.

The membrane was stripped and re-probed with a different antibody (VDR, CYP27B1 and CYP24A1). The membrane was incubated twice with 5 mL of stripping buffer (Appendix G), with agitation, at room temperature for 10 minutes. The membrane was then washed twice with 1X PBS (pH 7.4) for 10 minutes each, followed by two washes with 0.2% PBS Tween for 5 minutes each. The membrane was blocked, and the antibody staining process was repeated.

2.15 Statistics

Raw data was captured using Microsoft Excel (Microsoft), and statistical analysis was performed with GraphPad Prism v9.4.1 (GraphPad Software, San Diego, California, USA). At least three biological and three technical repeats were conducted for each experiment. All data are expressed as mean \pm standard error of the mean (SEM). Statistical variations in means were tested using the one-way analysis of variance (ANOVA) test and the *post hoc* Bonferroni test. $p \leq 0.05$ was considered significant.

Qualitative data for microscopy were analysed using ImageJ (Java 1.8.0_172, Windows, National Institutes of Health, USA) and Western blots using Image Lab™ v6.1 software (Windows, Bio-Rad, Hercules, CA, USA).

Chapter 3

Results

3.1 Determining growth inhibition concentration of 25(OH)D₃ in HeLa cells

The GI₅₀ is defined as the concentration where cells are reduced by 50% from their initial value, and this can be determined through colorimetric assays⁶⁴. A growth inhibition curve was constructed using the crystal violet assay in order to determine the treatment concentrations for assessing the anti-cancer actions of 25(OH)D₃ on HeLa cells.

The cells were subjected to incubation for 72 hours after treatment with 10-fold dilutions of 25(OH)D₃ in the range of 5.0x10⁻⁶ to 5.0x10⁻¹³ M. The resulting cell numbers were used to calculate the percentage growth inhibition. A non-linear best fit line revealed that the highest treatment dose of 5.0x10⁻⁶ M (5000 nM) 25(OH)D₃ inhibited HeLa cell growth by 20% (Figure 3.1). The lowest treatment dose of 5.0x10⁻¹³ M inhibited the growth of HeLa cells by approximately 4%. The supraphysiological concentration of 5000 nM was selected for treatment and used in all subsequent assays. A second treatment dosage, 260 nM, in the physiological range of 25(OH)D₃, was also used for comparison in all subsequent assays. This physiological concentration resulted in approximately 12% growth inhibition.

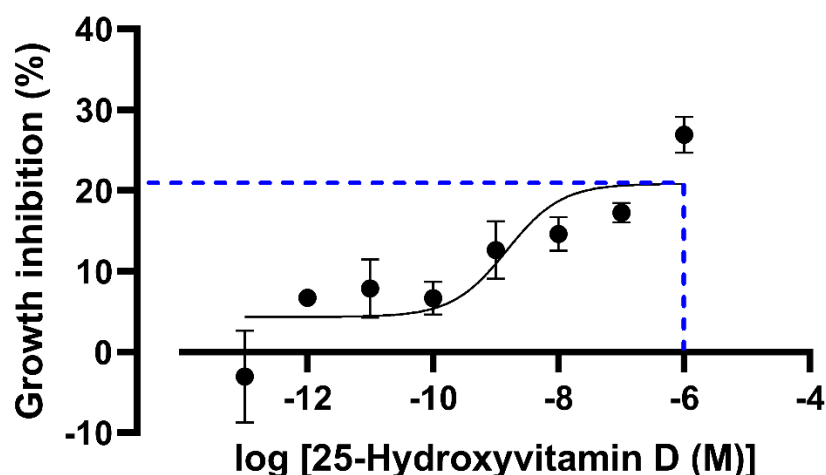


Figure 3.1 Growth inhibition of HeLa cells with various concentrations of 25(OH)D₃. HeLa cells were treated with a series of concentrations of 25(OH)D₃, and a growth inhibition curve was constructed from crystal violet assay results. Data are presented as mean±SEM. The blue dashed line indicates 20% growth inhibition.

3.2 Cell count determination using the crystal violet assay

Cell count was assessed using the crystal violet assay. Crystal violet is an organic chloride salt that binds to the protein and DNA of cells⁸⁷. During cell death, adherent cells detach from the surface of cell culture plates and are washed away, leaving live cells attached to the culture plates⁶⁵. The live cells are stained with the crystal violet dye, and the absorbance values are read with a spectrophotometer. The absorbance is directly proportional to the cell number.

Significant decreases in cell number were observed with 5000 nM 25(OH)D₃ treatment (82.05%±1.758%) relative to the solvent control (101.0%±1.691%, p<0.0001) and to the medium control (100.7%±2.048%, p<0.0001). Significant reductions in cell number were also observed at the 260 nM treatment dose (90.64%±1.448%) compared to the medium (p=0.0023) and solvent (p=0.0016) controls and to the 5000 nM treatment (p=0.0257) (Figure 3.2).

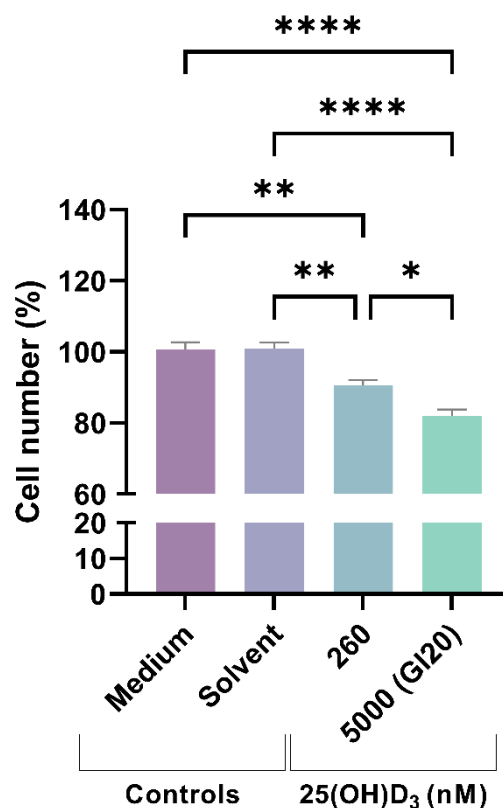


Figure 3.2 The effect of 25(OH)D₃ on cell count in HeLa cells. The crystal violet assay observed significant changes at the supraphysiological 25(OH)D₃ treatment dose. Data are expressed as mean±SEM (* p<0.05; ** p<0.01; **** p<0.0001).

3.3 Assessment of cell viability with alamarBlue®

Cell viability was assessed with the AlamarBlue® assay. AlamarBlue® incorporates resazurin, a colorimetric oxidation-reduction (REDOX) indicator which detects the cellular metabolic activity of the mitochondria⁶⁶. Metabolically active cells reduce Resazurin (blue) to resorufin (red). The colorimetric endpoint is measured by absorbance spectrophotometry. The absorbance is directly proportional to cell viability⁶⁶⁻⁶⁷.

Significant decreases in cell viability were observed with the 5000 nM 25(OH)D₃ treatment dose (87.79%±1.834%) relative to the solvent control (99.27%±1.357%, p=0.0005) and the medium control (98.51%±1.525%, p=0.0012). Treatment at the physiological dose (91.43%±2.082%) significantly decreased cell viability relative to the solvent (p=0.0156) and medium controls (p=0.0360). There was no significant variation noted between the two 25(OH)D₃ treatments (Figure 3.3).

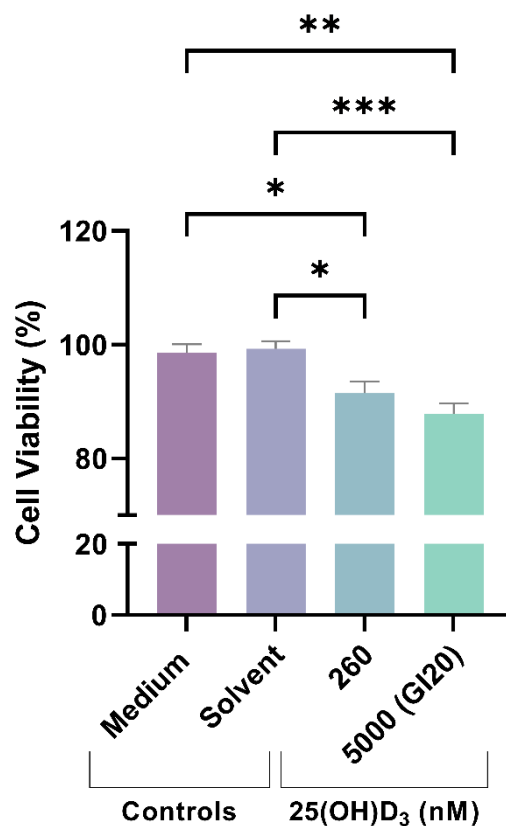


Figure 3.3 The effect of 25(OH)D₃ on cell viability in HeLa cells. Histogram plot of cell viability using the alamarBlue® assay. Data are expressed as mean±SEM (* p<0.05; ** p<0.01; *** p<0.001).

3.4 Cell cycle analysis using flow cytometry

The cell cycle regulates cell growth and division. This flow cytometric assay discriminates cells in the distinct phases of the cell cycle using a formulation of propidium iodide (PI) and RNase A⁶⁸. PI stains nucleic acids and is used to identify RNA and DNA. RNase A in the proprietary formation degrades RNA, and therefore cells containing DNA are identified.

Treatment with the 5000 nM dose of 25(OH)D₃ (8.30%±0.7506%) significantly increased the sub-G₁ cell population relative to the solvent control (5.57%±0.1202%, p=0.0305), medium control (5.63%±0.2404%, p=0.0348) and 260 nM treatment dose (5.37%±0.6227%, p=0.0206). (Figure 3.4)

There were no significant changes in the G₀/G₁ cell population observed with 5000 nM 25(OH)D₃ treatment (68.83%±2.341%) relative to the solvent control (65.95 %±3.384%) and medium control (66.48%±2.308%). Treatment at 260 nM 25(OH)D₃ (67.05%±2.775%) resulted in no significant changes relative to the controls and the supraphysiological treatment dose.

Treatment with the 5000 nM dose of 25(OH)D₃ (12.17%±0.8212%) showed no significant alteration in the number of cells in the S phase of the cell cycle relative to the solvent control (13.80%±0.4583%), medium control (14.13%±0.4333%) and 260 nM treatment (14.07%±0.1202%).

Treatment with the 5000 nM dose of 25(OH)D₃ (12.30%±1.457%) did not significantly increase the G₂/M cell population relative to the solvent control (16.57%±3.180%), medium control (15.70%±1.888%) and 260 nM treatment (15.47%±1.634%).

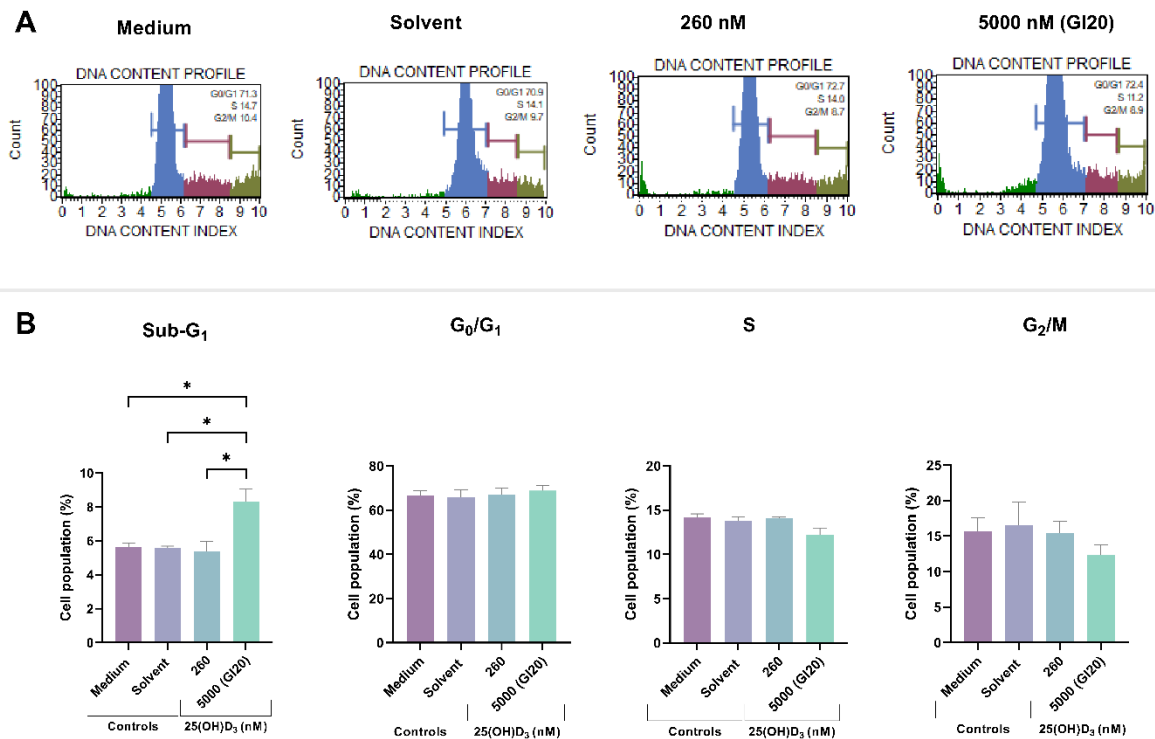


Figure 3.4 Cell cycle analysis with flow cytometry. (A) Histogram plots and (B) bar charts displaying the percentage of total cells in each phase of the cell cycle (sub-G₁, G₀/G₁, S and G₂/M) for experimental and control cultures. Data are expressed as mean±SEM (* p<0.05).

3.5 Assessing mitochondrial membrane depolarisation using flow cytometry

Changes to the mitochondrial membrane potential during apoptotic cell death were analysed using the Luminex Muse Mitopotential Assay. Changes in mitochondrial membrane potential can disrupt the function of the mitochondria and result in apoptotic cell death⁴⁰. The flow cytometric assay used discriminated cells with damaged mitochondria and cells with normal mitochondrial, as well as live and dead cells⁶⁹.

Treatment with 5000 nM of 25(OH)D₃ significantly decreased the number of live (viable) cells (84.25%±1.156%) and relative to the solvent (97.28%±0.9052%, p=0.0006) and medium controls (96.22%±0.2689%, p=0.0012). A significant decrease in the cell population was observed in the 260 nM treatment (87.64%±1.9052%) relative to the solvent (p=0.0050) and medium (p=0.0105) controls. No significant differences in cell population size were observed between the two 25(OH)D₃ treatment doses. (Figure 3.6).

For live cells with depolarised mitochondrial membranes, there was a significant decrease in the cell population with 5000 nM treatment ($9.158\% \pm 1.490\%$) relative to the solvent ($0.4167\% \pm 0.2619\%$, $p=0.0028$) and medium ($0.7500\% \pm 0.1504\%$, $p=0.0037$) controls. No differences were observed in the cell population sizes between the 5000 nM and 260 nM ($5.407\% \pm 1.190\%$) treatments. No statistical significance was observed in cell population with the 260 nM treatment dose compared to the solvent and medium controls.

The number of dead cells with depolarised mitochondrial membranes and dead cells did not differ significantly across the experimental and control treatments.

The total number of depolarised cells was significantly higher with 5000 nM treatment ($13.86\% \pm 0.6120\%$) compared to the 260 nM treatment ($9.537\% \pm 1.302\%$, $p=0.0362$), the solvent control ($2.177\% \pm 0.4359\%$, $p<0.0001$) and the medium control ($2.147\% \pm 0.05364\%$, $p<0.0001$). The total number of depolarised cells in 260 nM treatment culture were significantly higher than in the solvent ($p=0.0012$) and medium ($p=0.0012$) controls.

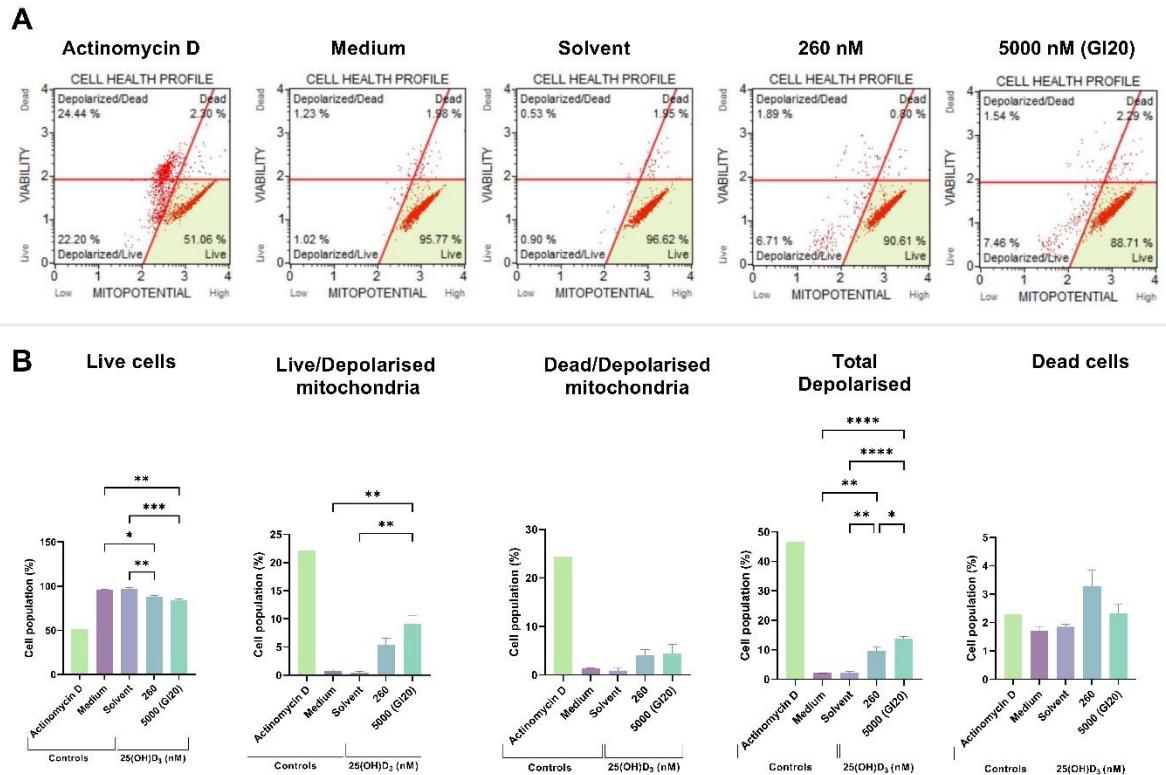


Figure 3.5 Detection of mitochondrial membrane depolarisation with flow cytometry. Two-dimensional dot-plot diagrams (A) and bar graphs (B) displaying live (viable), live cells with depolarised mitochondrial membranes, dead with depolarised mitochondrial membranes, total depolarised and dead cells for experimental and control cultures (* $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$; **** $p < 0.0001$).

3.6 Assessment of phosphatidylserine externalisation using flow cytometry

Stimuli that induce apoptosis cause the translocation of phosphatidylserine residues in the inner leaflet of the cell membrane to the outer leaflet³⁷. Flow cytometric analysis was conducted using Annexin V and 7-AAD. Annexin V binds to externalised phosphatidylserine residues, and 7-AAD binds to DNA⁷⁰.

Treatment at 5000 nM significantly decreased the number of healthy live (viable) cells ($83.33\% \pm 1.274\%$) relative to the solvent control ($96.10\% \pm 1.1815\%$, $p < 0.0001$), the medium control ($96.48\% \pm 0.2258\%$, $p < 0.0001$) and the 260 nM treatment ($91.18\% \pm 1.275\%$, $p = 0.0024$). A significant decrease in the cell population was observed with the 260 nM treatment compared to the solvent ($p = 0.0430$) and medium ($p = 0.0283$) controls. (Figure 3.5).

25(OH)D₃ treatment at 5000 nM (7.787%±1.477%) significantly increased the live, apoptotic cell population compared to the solvent (2.550%±0.1914%, p=0.0216) and medium (2.397%±0.3263%, p=0.0182) controls. No statistical differences were observed with the 260 nM treatment dose (3.460%±0.6751%) and the controls.

For cells that were apoptotic and dead, there was a significant increase in the cell population with 5000 nM treatment (8.193%±0.5980%) relative to the solvent control (0.3733%±0.1910%, p<0.0001), medium control (0.1233%±0.09939%, p<0.0001) and the 260 nM treatment (4.683%±0.08239%, p=0.0141). Treatment at 260 nM significantly increased the cell population compared to the controls of the solvent (p=0.0038) and medium (p=0.0026).

Significant increases in the total apoptotic cell population were observed with 25(OH)D₃ treatments at 260 nM (8.147%±1.487%) and 5000 nM (15.98%±1.195%) relative to the medium control (2.917%±0.2490%, p=0.0495 and p=0.0001, respectively). Treatment at 5000 nM significantly increased the total apoptotic cell population compared to the solvent control (p=0.0001) and 260 nM treatment (p=0.0043).

There were no significant changes observed in the number of dead cells across the experimental treatments and controls.

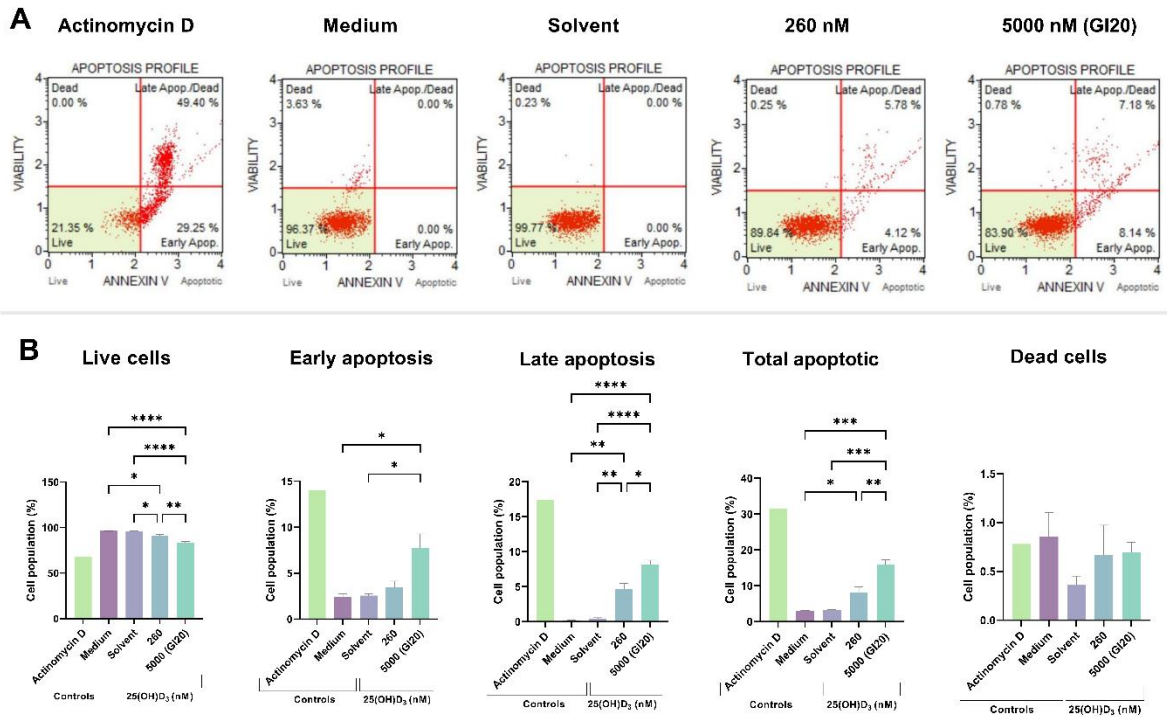


Figure 3.6 Annexin V detection with flow cytometry. Two-dimensional dot-plot diagrams (A) and bar graphs (B) displaying live, early apoptotic, late apoptotic, total apoptotic, and dead cell populations for experimental and control cultures (* $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$; **** $p < 0.0001$).

3.7 Activation of caspase-3 and -7 activity in 25(OH)D₃ treated HeLa cells

Caspase-3 and -7 activity was measured using the Luminex Muse Caspase-3/7 assay. Caspases are proteases that play a role in a cascade of reactions leading to apoptotic cell death⁴⁰. This flow cytometric assay discriminated between apoptotic and normal cells and live and dead cells⁷².

Treatment with 5000 nM of 25(OH)D₃ significantly decreased the number of healthy live (viable) cells (90.08%±1.919%) and relative to the solvent (98.08%±0.2589%, $p=0.0095$) and medium controls (97.26%±0.3044%, $p=0.0184$). No significant differences were observed in the 260 nM treatment (88.38%±1.055%) relative to the 5000 nM treatment. The 260 nM treatment significantly decreased the number of healthy live cells relative to the solvent ($p=0.0027$) and medium controls ($p=0.0048$). (Figure 3.7)

For live cells with activated caspases -3 and -7, there were no significant differences between the 5000 nM treatment (1.077%±0.5235%) relative to the solvent control (0.3800%±0.1943%),

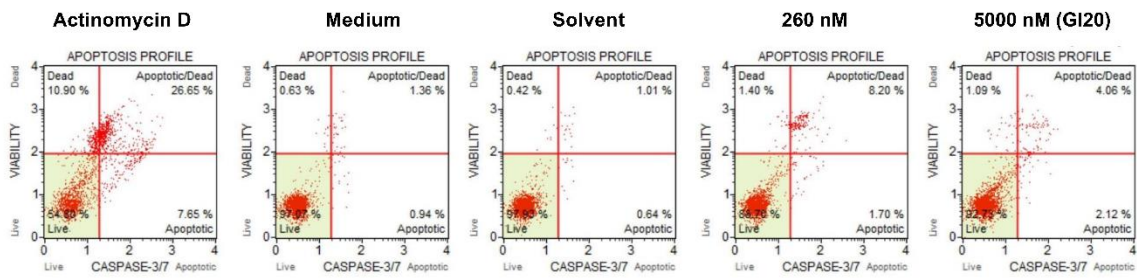
medium control ($0.3333\% \pm 0.3038\%$) and 260 nM $25(\text{OH})\text{D}_3$ treatment ($1.270\% \pm 0.2150\%$) treatments. Both treatments of $25(\text{OH})\text{D}_3$ had a higher cell population of live cells with activated caspases than the controls.

Treatment at 5000 nM treatment ($5.123\% \pm 0.8802\%$) significantly increased the number of dead cells with activated caspases -3 and -7 relative to the solvent ($0.3933\% \pm 0.3122\%$, $p=0.0345$) and medium ($0.5500\% \pm 0.4136\%$, $p=0.0416$) controls. Treatment at 260 nM treatment ($8.027\% \pm 1.273\%$) significantly increased the cell population relative to the solvent ($p=0.0017$) and medium ($p=0.0019$) controls.

Significant increases in the total apoptotic cell population were observed with $25(\text{OH})\text{D}_3$ treatments at 260 nM ($9.297\% \pm 1.308\%$) and 5000 nM ($6.207\% \pm 0.7477\%$) relative to the solvent control ($0.7700\% \pm 0.4761\%$, $p=0.0012$ and $p=0.0219$, respectively) and medium control ($0.8833\% \pm 0.7155\%$, $p=0.0013$ and $p=0.0248$, respectively).

Treatment with $25(\text{OH})\text{D}_3$ at both doses increased the percentage of dead cells, but no significant differences were observed relative to the controls.

A



B

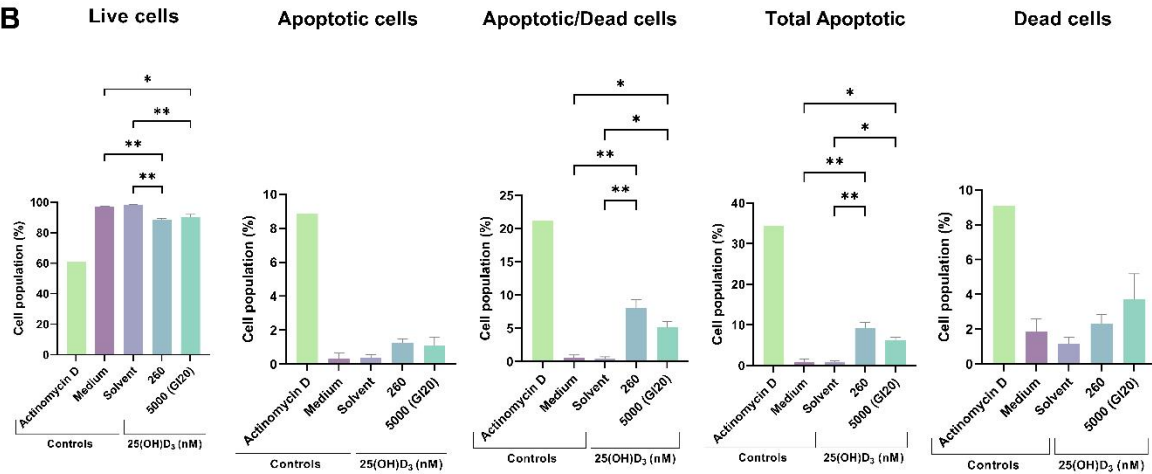


Figure 3.7 Detection of caspases -3 and -7 activity with flow cytometry. Two-dimensional dot-plot diagrams (A) and bar graphs (B) displaying live, apoptotic, apoptotic and dead, total apoptotic, and dead cell populations for experimental and control cultures (* $p < 0.05$; ** $p < 0.01$).

3.8 Induction of LC3-II in 25(OH)D₃ treated HeLa cells

The presence of autophagic cell death was assessed using the Luminex Muse Autophagy LC3-II assay. LC3 is a protein essential for the formation of autophagosomes. This assay discriminated between cells with cytosolic LC3-I and autophagosome-associated LC3-II^{73,88}.

Treatment with 25(OH)D₃ at 260 nM (4.600 ± 0.4619) and 5000 nM (8.167 ± 0.4667) did not significantly induce autophagic cell death when compared to the medium (5.733 ± 0.4842) and solvent (5.300 ± 0.2082) controls (Figure 3.8).

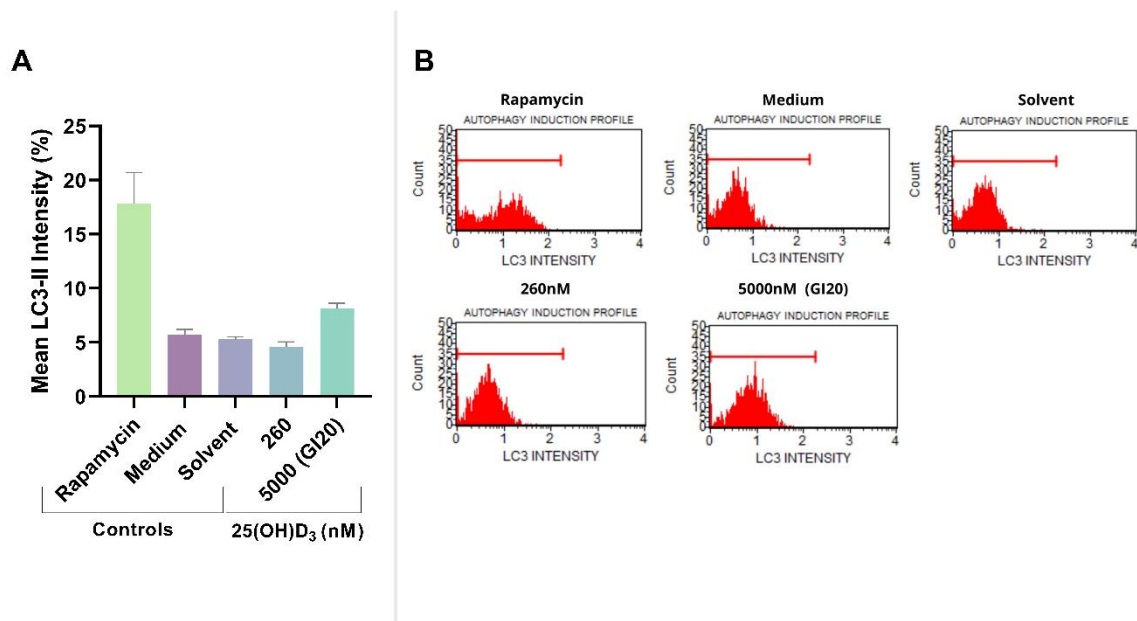


Figure 3.8 Detection of LC3-II activity with flow cytometry. (A) Bar graphs and (B) histogram plots of LC3-II intensity for experimental and control cultures detected through flow cytometry.

3.9 Lactate dehydrogenase release in 25(OH)D₃ treated HeLa cells

Necrotic cell death (cell cytotoxicity) was assessed using the lactate dehydrogenase (LDH) assay. The assay measures cytosolic LDH released from damaged cells.

Treatment with 5000 nM 25(OH)D₃ (6.173%±2.705%) increased the percentage of cell cytotoxicity; however, the increase was not significant relative to the solvent control (0.4325 ±2.156%) and medium control (1.514%±0.9826%) controls and the 260 nM treatment dose (0.3309%±1.755%). (Figure 3.9)

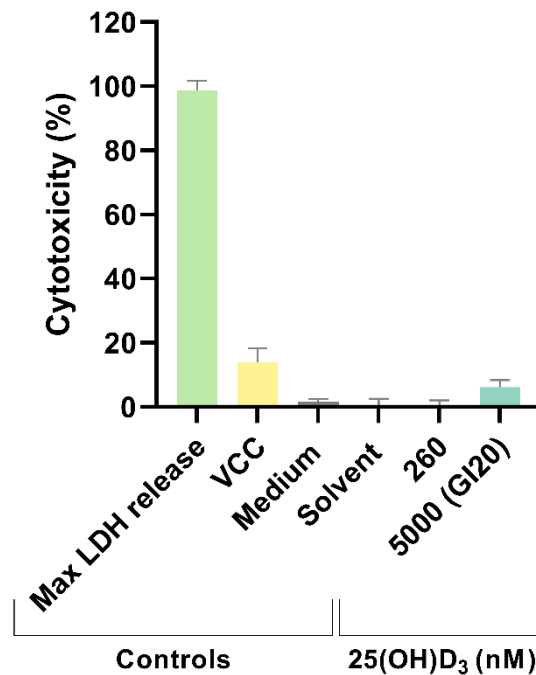


Figure 3.9 Detection of lactate dehydrogenase. Lactate dehydrogenase (LDH) release was measured in experimental and control cultures. A maximum LDH release and a volume correction control (VCC) were also included in the experiments (* $p < 0.05$; ** $p < 0.01$).

3.10 Assessment of gross morphological changes in 25(OH)D₃ treated HeLa cells

Gross morphological changes were analysed using light microscopy. Experimental and control samples were stained using haematoxylin and eosin, and samples were viewed with a brightfield microscope. Haematoxylin stains acidic structures blue and eosin stains basic structures pink⁷⁵.

Solvent and medium controls exhibited minimal cell death features, with most cells appearing healthy. Cells in experimental samples showed morphological features of cell death also present in the actinomycin treatment (positive control). Notable observed features include cell membrane blebbing, apoptotic bodies, hyper-condensed chromatin (pyknosis) and cytoplasmic shrinkage. Positive control and 25(OH)D₃ samples showed decreased cell density compared to medium and solvent controls (Figure 3.10A-I).

The samples were semi-quantitatively analysed. Approximately 1200 cells were counted for each sample, and the percentage of cells with features of apoptotic cell death was calculated.

There was a significantly higher percentage of cells with morphological features of apoptosis in the 5000 nM 25(OH)D₃ treatment (7.715%±1.443%, p=0.0159) relative to the medium control (3.064%±0.6881%). No significance was observed relative to the solvent control (4.228%±0.9296%) and 260 nM treatment (7.350%±1.215%) (Figure 3.10J).

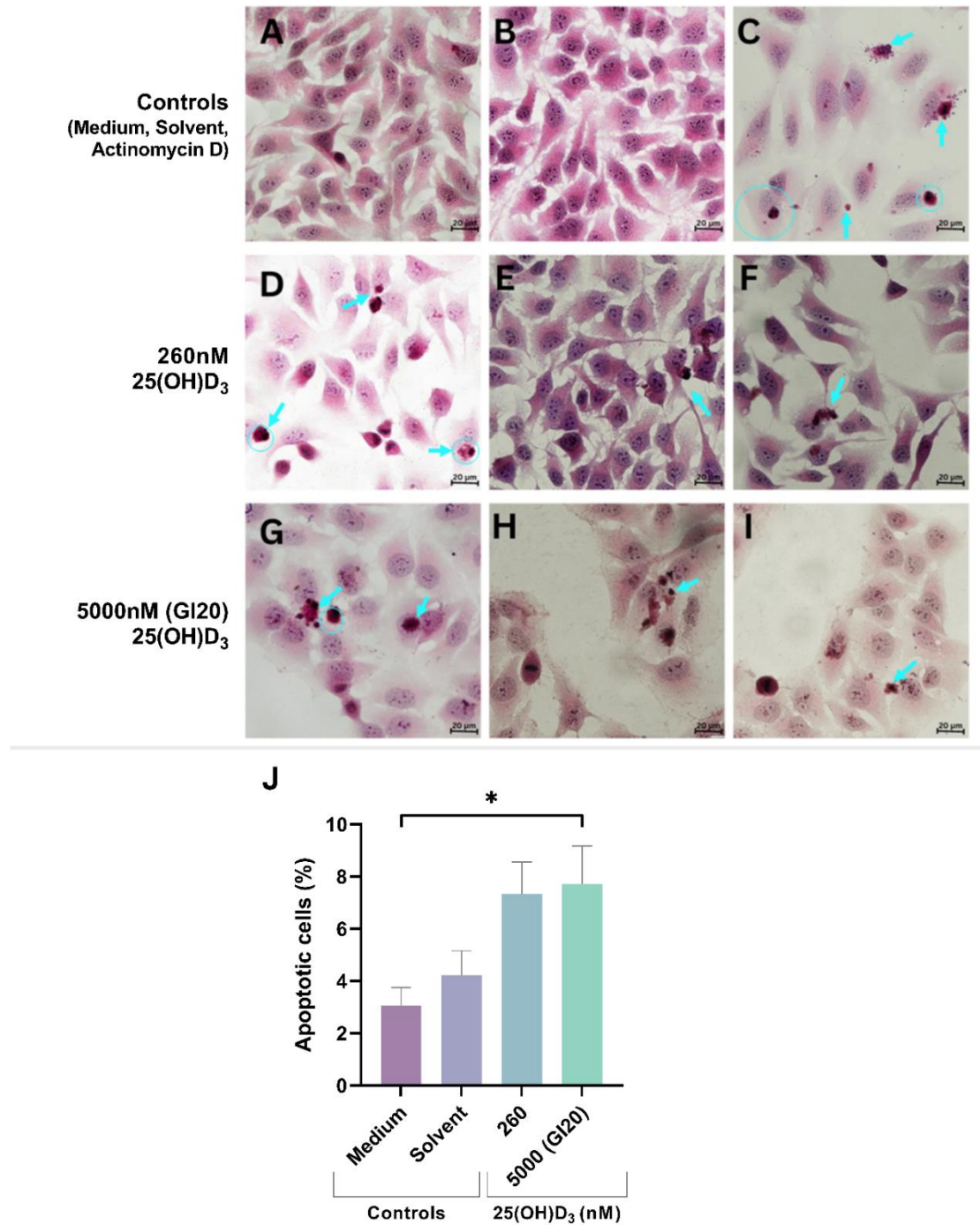


Figure 3.10 Analysis of morphological changes using brightfield microscopy. Growth medium (A) and solvent (B) treated cells showed a dense population with minimal signs of cell death. The positive control (actinomycin D) (C) and vitamin D treatments of 260 nM (D-E) and 5000 nM (G-I) revealed apoptotic bodies, blebbing, cytoplasmic shrinkage, hyper-condensed chromatin, and compromised cell density. Approximately 1200 cells were counted from each treatment and analysed for features of apoptosis and cell death. The percentage of apoptotic cells was calculated (J). The 5000 nM treatment significantly increased the percentage of apoptotic cells. Scale bar = 20µm. * $p < 0.05$.

In addition to characteristics of cell death, mitotic cells were also visible in the different treatments. Cells were identified in the prophase, metaphase, anaphase, and telophase phases of mitosis (Figures 3.11A and 3.11B).

Approximately 1200 cells were counted in each experimental and control culture. The percentage of mitotic cells in each culture was calculated. The 5000 nM treatment has 5.56% of cells (Figure 3.11C) undergoing mitosis, which compared to the percentage of cells in the solvent control (4.63%), medium control (4.96%) and 260 nM 25(OH)D₃ treatment (3.95%).

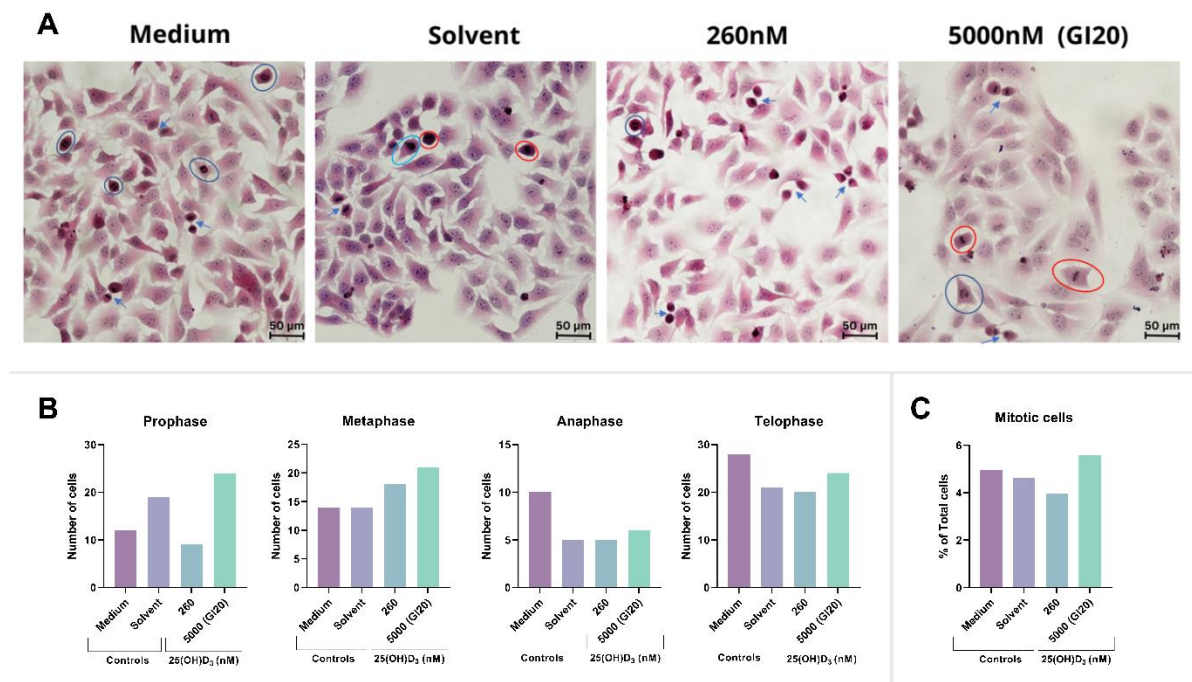


Figure 3.11 Brightfield microscopy images and semi-qualitative analysis of mitotic HeLa cells. Experimental and control cultures contained cells that were visibly undergoing mitosis. The cells were visualised using H&E staining (A). Cells in prophase (light blue circle), telophase (red circle), anaphase (dark blue circle) and telophase (blue arrow) were observed across treatments and controls. Cells in each phase of mitosis did not differ statistically across the cultures (B). The percentage of mitotic cells present did not differ across cultures (C).

Interestingly, multinucleated (polyploid) cells (MNCs) were observed in all cultures (Figure 3.12A). The percentage of MNCs cells varied across all cultures, with 2.884%±0.7720% of cells in the 5000 nM treatment, 1.775%±0.2171% in the solvent control, 1.552%±0.2702% in the medium control and 1.552%±0.2702% in the 260 nM treatment (Figure 3.12B). The size of the MNCs in the 5000 nM treatment averaged 2.413±0.2308 times larger relative to the size of normal cells in the same culture (Figure 3.12C). This was significantly larger than the MNCs

in the solvent control and medium control cultures, which averaged 1.364 ± 0.1056 ($p=0.0374$) and 1.555 ± 0.09751 ($p=0.0346$), respectively. The MNCs in the 260 nM treatment culture were 1.675 ± 0.4886 times larger than normal cells. In addition, the number of nuclei in each MNC was counted. MNCs in 5000 nM had an average of 3.200 ± 0.4078 nuclei, the solvent control 2.000 ± 0.000 , the medium control 2.917 ± 0.5833 and the 260 nM treatment 2.500 ± 0.5000 (Figure 3.14D).

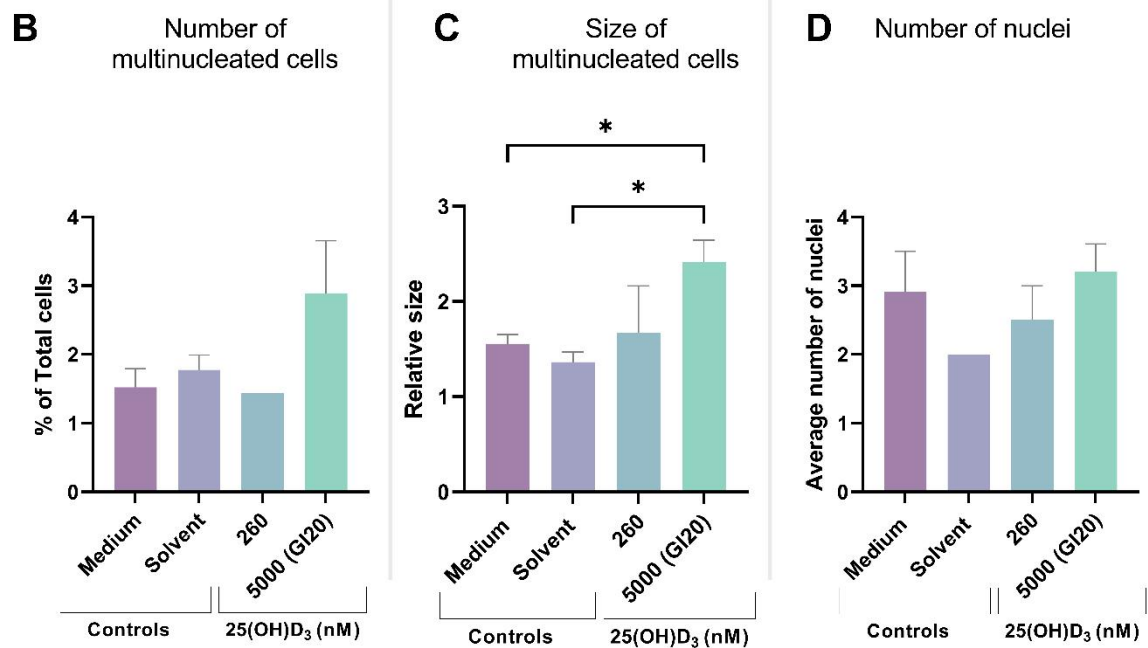
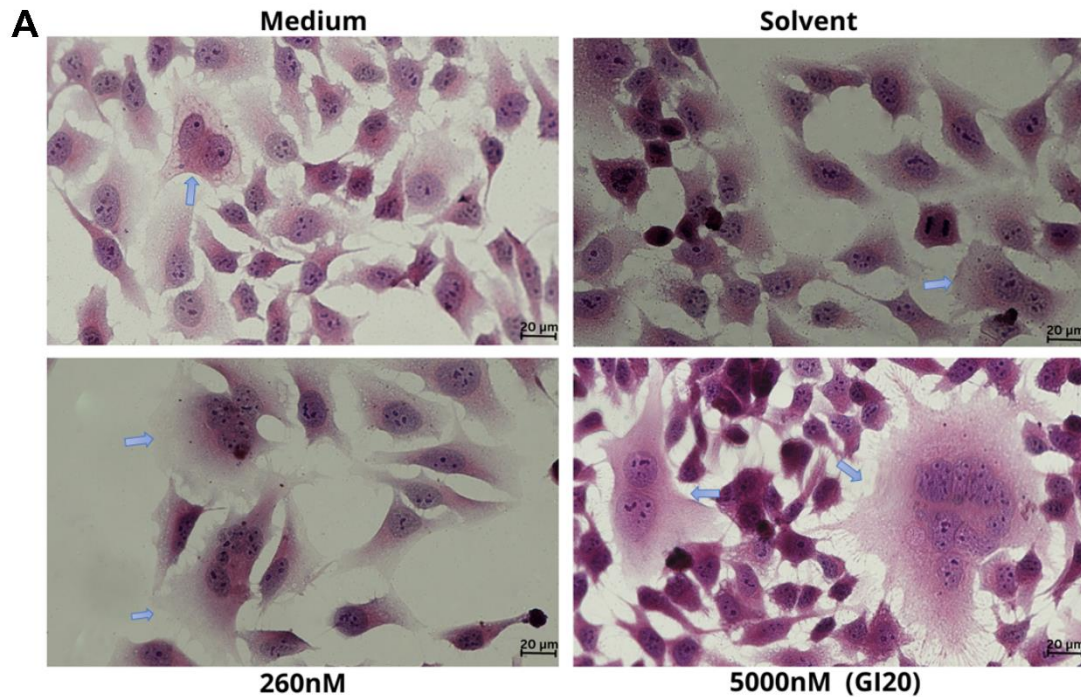


Figure 3.12 Multinucleated HeLa cells visualised with H&E stain. Experimental and control cultures were visualised with brightfield microscopy using H&E staining (A). Multinucleated cells (blue arrows) were present in all cultures. Bar graphs of the number of multinucleated cells (B), the size of multinucleated cells relative to the average size of normal cells (C) and the number of nuclei in multinucleated cells (D) show that 5000 nM treatment increased the number and size of multinucleated cells.

3.11 Ultrastructural changes in 25(OH)D₃ treated HeLa cells

Ultrastructural changes were observed using transmission electron microscopy (TEM). In TEM, an image is mirrored by scattered electron waves from heavy metal-stained samples⁷⁸. Cells were fixed, dehydrated, embedded, sectioned, and stained to allow for viewing of cellular organelles.

Cells in the solvent control appeared normal, indicating no signs of cellular damage or cell death. Cells treated with a supraphysiological dose (5000nM) displayed features of apoptosis such as nuclear membrane damage (Figure 3.13), apoptotic bodies (Figure 3.14) and membrane blebbing. The chromatin was not visible in images due to poor staining.

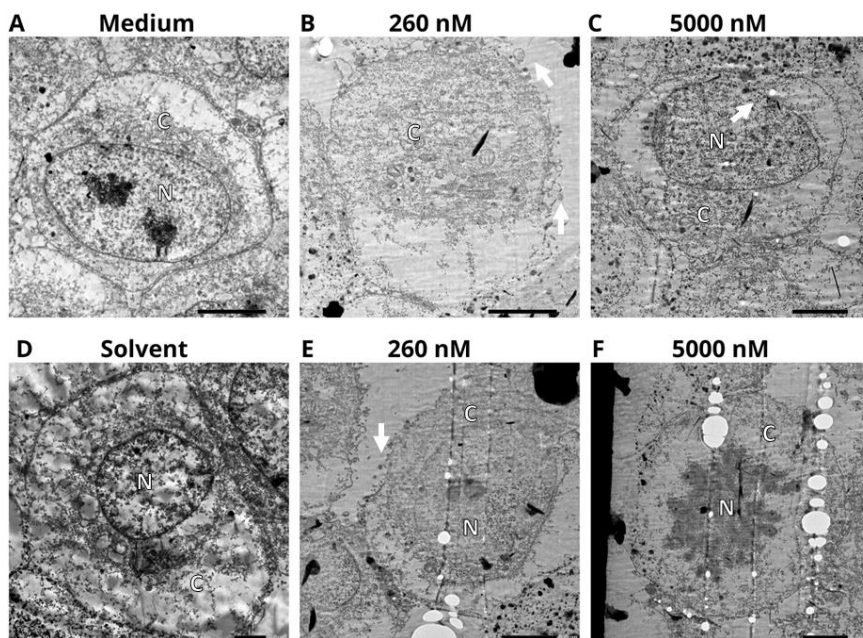


Figure 3.13 Electron micrographs of experimental and control cultures. Medium (A) and solvent (D) cultures show normal cell morphology with no signs of apoptosis. The 260nM (B and E) and 5000nM (C and F) treatments show features of apoptosis, including membrane blebbing, disrupted nuclear membrane, and nuclear damage. C, cytoplasm; N, nucleus.

At higher magnification, changes in mitochondria structure were visible for cells treated with 5000nM 25(OH)D₃. Some mitochondria appeared normal, whilst others were rounded with a swollen matrix (Figure 3.15). Nuclear membrane damage, apoptotic bodies, and membrane blebbing were also observed.

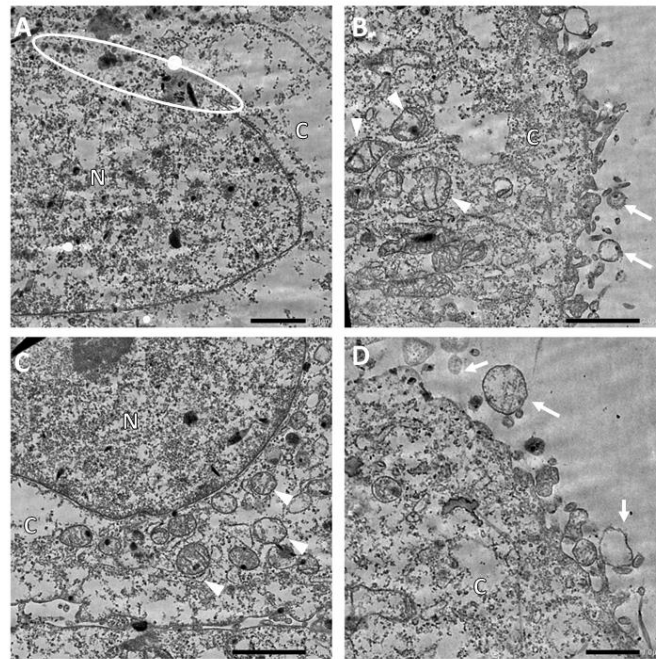


Figure 3.14 Transmission electron micrograph of HeLa treated with 5000nM 25(OH)D₃. Cells displaying features of apoptosis such as (A) nuclear membrane damage (oval), (B-C) round mitochondria with swollen matrix (arrowhead), (D) apoptotic bodies (arrow) and cytoplasmic blebbing. C, cytoplasm; N, nucleus.

3.12 Gene and enzyme expression of vitamin D metabolising system in 25(OH)D₃ treated HeLa cells

Total RNA extracts from cell cultures were synthesised into cDNA. The cDNA was amplified with real-time PCR. The mRNA expression for CYP27B1 was significantly upregulated at a 5000 nM treatment (1.335 ± 0.01878) in comparison to the solvent control (1.000 ± 0.06477 , $p=0.0151$), medium control (0.8674 ± 0.04684 , $p=0.0019$) and 260 nM treatment (0.8532 ± 0.07261 , $p=0.0015$). There were no significant differences observed between the 260 nM 25(OH)D₃ treatment and the control cultures (Figure 3.15A).

Total protein was extracted from experimental and control cultures. Proteins were separated with gel electrophoresis and identified with immunoblotting. The CYP27B1 protein is approximately 47.5 kDa, and β -actin is approximately 42 kDa in size. There was a significant increase in CYP27B1 protein expression observed in the 5000 nM treatment (2.409 ± 0.6266) relative to the solvent control (1.000 ± 0.000 , $p=0.0210$) and medium control (0.9861 ± 0.09133 , $p=0.0358$). There were no significant differences observed between the 260 nM 25(OH)D₃ treatment (1.502 ± 0.1152) and the control cultures (Figures 3.15B and 3.15C).

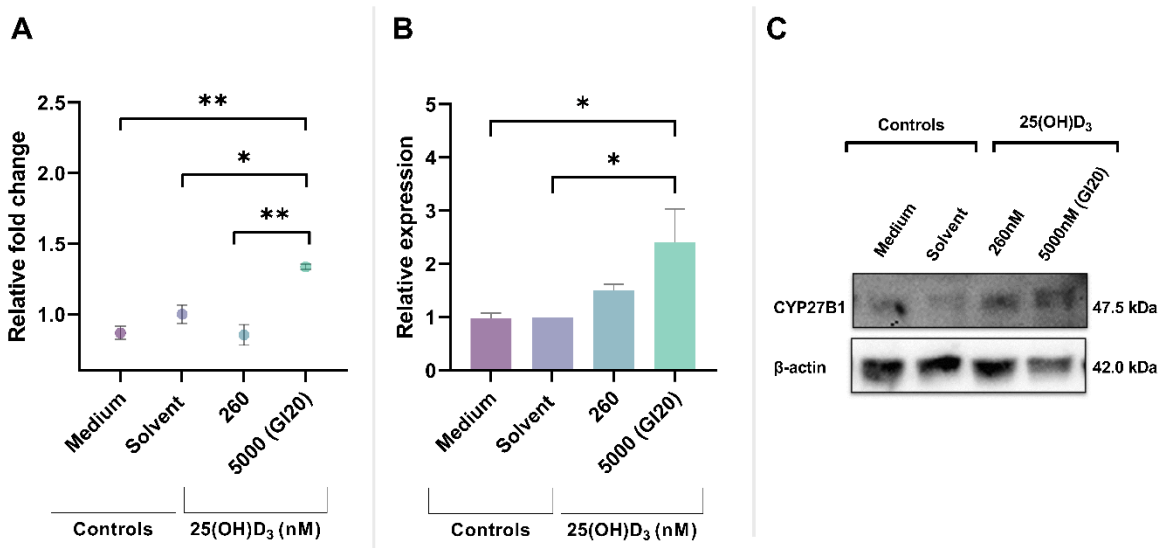


Figure 3.15 CYP27B1 gene and protein expression. Gene (A) and protein (B and C) were analysed with real-time qPCR and western blot analysis. Treatment with 5000nM of 25(OH)D₃ significantly increased CYP27B1 mRNA and protein expression. There was no statistically significant difference in protein expression (* $p < 0.05$; ** $p < 0.01$).

The mRNA expression for CYP24A1 was significantly upregulated at a 5000 nM treatment (1641 ± 444.4) in comparison to the solvent control (1.000 ± 0.05114 , $p=0.0107$), medium control (4.257 ± 0.2878 , $p=0.0108$) and 260 nM treatment (13.07 ± 4.195 , $p=0.0205$). There were no significant differences observed between the 260 nM 25(OH)D₃ treatment and the control cultures (Figure 3.16A).

The CYP24A1 protein is approximately 56.0 kDa in relative molecular weight (M_r). 5000 nM treatment (1.871 ± 0.3771) significantly increased protein expression relative to the solvent control (1.000 ± 0.000 , $p=0.0482$). No significant differences were observed between the 5000 nM 25(OH)D₃ treatment and the medium control (1.143 ± 0.07208) and 260 nM treatment (1.240 ± 0.05764). Additionally, no significant differences were observed between the 260 nM 25(OH)D₃ treatment and the control cultures (Figures 3.16B and 3.16C).

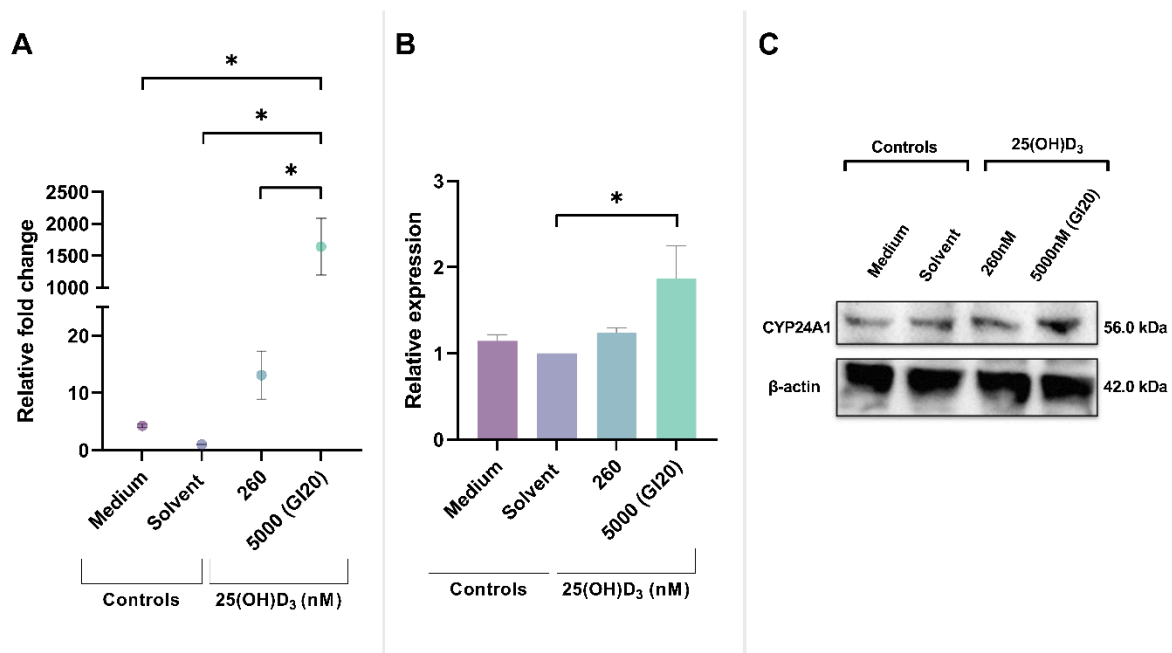


Figure 3.16 CYP24A1 gene and protein expression. Gene (A) and protein (B and C) were analysed with real-time qPCR and western blot analysis. Treatment with 5000nM of 25(OH)D₃ significantly upregulated CYP24A1 mRNA and protein expression (* p < 0.05).

3.13 Gene and protein expression of the vitamin D receptor in 25(OH)D₃ treated HeLa cells

The VDR mRNA was significantly upregulated with 5000 nM treatment (3.431 ± 0.1673) in comparison to the solvent control (1.000 ± 0.02935 , $p < 0.0001$), medium control (1.125 ± 0.1212 , $p < 0.0001$) and 260 nM treatment (1.433 ± 0.06940 , $p < 0.0001$). There was no significant difference between the 260 nM 25(OH)D₃ treatment and the control cultures (Figure 3.17A).

The VDR protein is approximately 46.7 kDa. Treatment with 5000 nM of 25(OH)D₃ (1.509 ± 0.1823) significantly increased protein expression compared to the solvent control (1.000 ± 0.000 , $p = 0.0344$). Treatment at 260 nM (1.678 ± 0.06220) protein expression compared to the solvent ($p = 0.0091$) and medium (1.020 ± 0.01598 , $p = 0.0173$) control cultures (Figures 3.17B and 3.17C).

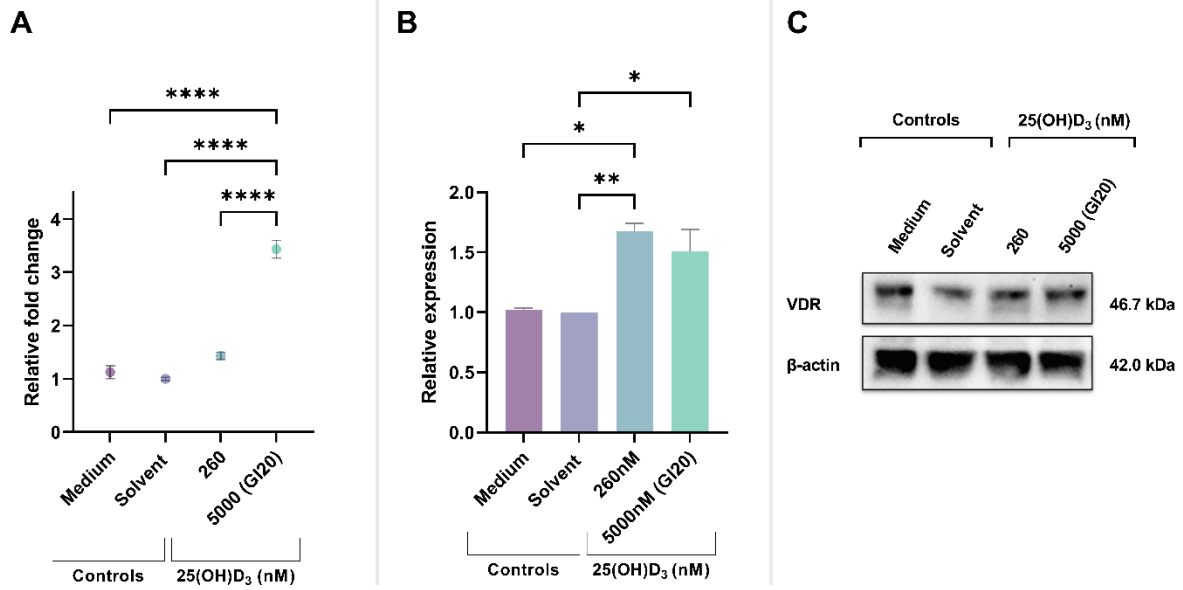


Figure 3.17 VDR gene and protein expression. Gene (A) and protein (B and C) were analysed with real-time qPCR and western blot analysis. Treatment with 5000 nM of 25(OH)D₃ significantly increased VDR mRNA expression. Both 260 nM and 5000 nM treatments statistically increased protein expression (* p<0.05; ** p<0.01; **** p < 0.0001).

Chapter 4

Discussion

Vitamin D and its metabolites are secosteroid hormones widely recognized for their ability to facilitate the absorption of calcium and phosphorous^{5,22}. In addition, *in vitro* and *in vivo* studies examining the anti-cancer effects of vitamin D metabolites in breast⁸⁹⁻⁹⁰, colon⁹¹⁻⁹², prostate⁹³, lung⁹⁴, and skin⁹⁵ cancers have established that vitamin D metabolites can effectively inhibit cancer cell growth and replication via a variety of pathways. However, the anti-tumour actions of vitamin D in cervical cancer remain insufficiently explored. The findings of this study demonstrate that 25(OH)D₃ reduced cell count and viability, prompted apoptotic cell death and upregulated gene and protein expression of CYP27B1, CYP24A1, and VDR.

25(OH)D₃ decreases cell number and cell viability in HeLa cells

The administration of 25(OH)D₃ caused a significant reduction in the cell number and viability of HeLa cells. Similar decreases in both parameters have been observed in other cervical cancer cell lines. CaSki cervical cancer cells, upon treatment with high concentrations of cholecalciferol (100 ng/mL and 1000 ng/mL), exhibited decreases in cell number to approximately 56.25% ± 7.08% (p<0.05) and 50.59% ± 5.14% (p<0.001), respectively³⁰. The treatments also resulted in a decline in cell viability to 73.94% ± 3.16% (p<0.01) and 62.41% ± 1.64% (p<0.0001). In another study, Shruthi *et al.* demonstrated a 50% inhibition in the growth of HeLa cells with 1250 μM cholecalciferol treatment⁹⁶. Together, these studies demonstrate that vitamin D metabolites negatively affect cell growth in cervical cancer cell lines.

25(OH)D₃ triggered accumulation in the sub-G₁ phase without inducing cell cycle arrest

Accumulation of sub-G₁ population was observed with 5000 nM 25(OH)D₃ treatment in cell cycle analysis. The sub-G₁ population is associated with nuclear fragments, micronuclei, nuclei

with different chromatin structures, clumps or chromosomes and apoptotic bodies⁹⁷. This suggests that high-dose treatment of 25(OH)D₃ potentially induced DNA damage. Similar observations were seen in the CaSki cell line with high-dose cholecalciferol treatments of 100 ng/ml and 1000 ng/ml, which increased the sub-G₁ populations to 7.30%±1.04% and 9.75%±1.65%, respectively³⁰. The sub-G₁ population is associated with cell death; however, supporting data is needed to confirm that cell death has I^{25,98}.

Vitamin D metabolites can induce cell cycle arrest in various cancer cell lines⁹⁸. No significant changes in G₀/G₁, S, and G₂/M cell populations were observed in this study. In addition, brightfield microscopy revealed features of mitosis, with insignificant differences in the number of cells in all stages of mitosis, confirming the cell population numbers observed by flow cytometry. This differs from observations made in HeLa S3 cells treated with 200 nM and 500 nM calcitriol treatments, which increased the G₁ population⁹⁹. The mechanism of G₁ phase arrest involves the upregulation of RB, p21 and p27, and the downregulation of c-Myc^{98,100}. 25(OH)D₃, therefore, does not negatively promote alterations in intracellular tumorigenic pathways that promote cell cycle arrest in the HeLa cell line. This also suggests that a cell-specific response between HeLa and HeLa S3 may be operational. Furthermore, vitamin D metabolites calcitriol and 25(OH)D₃ may exert differential responses in adenocarcinoma cervical cell lines, like the HeLa and HeLa S3.

25(OH)D₃ induced apoptotic cell death in HeLa cells

In the present study, apoptosis was demonstrated by the increased depolarisation of mitochondrial membrane potential, increased phosphatidylserine externalisation and activation of terminal executioner caspases -3 and -7 with both 25(OH)D₃ treatments. Furthermore, morphological features of apoptosis described by the NCCD were discernible with brightfield microscopy images for both treatments of vitamin D, including hyper-condensed chromatin (pyknosis), apoptotic bodies, plasma membrane blebbing and cytoplasmic shrinkage^{33,40}. Electron micrographs also showed signs of nuclear membrane damage, cytoplasmic blebbing, formation of apoptotic bodies and swollen, rounded mitochondria. The mitochondrial damage observed with flow cytometry and microscopy is supported by the cell viability assay results, which are associated with abnormal metabolic activity of the mitochondria. These results

collectively suggest irregular mitochondrial function and caspase activation associated with intrinsic apoptosis^{40,42}.

Phosphatidylserine externalisation is a marker for both apoptosis and necrosis. However, the LDH cytotoxicity assay, 7-AAD dye staining in flow cytometry and morphological assessment showed insignificant necrotic cell death. Morphological analysis and LC3-II induction findings collectively did not suggest autophagic dependant cell death. Therefore, this suggests that 25(OH)D₃ did not induce necrosis and autophagic-dependent cell death in HeLa cells. Similar outcomes were observed in HeLa cells treated with calcitriol in a dose-dependent and time manner¹⁰¹.

25(OH)D₃ enhances expression of CYP27B1, VDR and CYP24A1

Healthy and tumorigenic cervical tissue express VDMS⁵³. This study indicates an autocrine response to 25(OH)D₃ in HeLa cells. Treatment with 5000 nM 25(OH)D₃ significantly increased CYP27B1, CYP24A1 and VDR gene and protein expression in HeLa cells. In contrast, treatment with 260 nM significantly increased VDR protein expression with no change at the mRNA transcript level. Physiological treatment also did not change CYP27B1 and CYP24A1 gene and protein expression, implying that treatment did not affect activation and catabolism of 25(OH)D₃.

The upregulation of the CYP27B1 gene and protein expression with 5000 nM treatment substantiates the autocrine conversion of 25(OH)D₃ to 1,25(OH)₂D₃. Similar findings were observed in SiHa cervical cells treated with 2500 nM 25(OH)D₃, which caused a 5-fold increase in CYP27B1 gene expression¹⁰². Treatment with a physiological dose did not statistically affect CYP27B1 mRNA expression in HeLa cells. Similar findings were demonstrated by Kloss *et al.* with 100 nM of 25(OH)D₃⁶². This suggests that supraphysiological doses induce the negative feedback regulation of 25(OH)D₃ levels by enhancing CYP27B1 expression. However, given the high concentration of 25(OH)D₃ at supraphysiological treatment, inactivated 25(OH)D₃ may directly inhibit cell growth and induce cell death without activation through 1 α OHase, CYP27B1. This was demonstrated in LNCaP prostate adenocarcinoma cells when treated with 100 nM 25(OH)D₃ in combination with the CYP27B1 inhibitor genistein. The authors found that 25(OH)D₃ directly exerted an inhibitory effect on LNCaP cell growth and that the effect was enhanced by genistein⁹³. Therefore, precursors of calcitriol may operate

as a therapeutic agent in cervical cancer either directly or by means of autocrine activation of CYP27B1.

24-Hydroxylase is crucial in the catabolism of 25(OH)D₃ and 1,25(OH)₂D₃. Several studies show that its expression is markedly induced by 1,25(OH)₂D₃.⁸ Interestingly, inhibiting or knocking down CYP24A1 has been shown to impede tumour growth significantly. Conversely, overexpressing CYP24A1 has been found to promote tumour progression and metastasis¹⁰³⁻¹⁰⁵. In this study, treatment at 5000 nM resulted in the overexpression of the CYP24A1 gene and significantly increased protein expression. Although cell growth was inhibited, the maximum inhibition was about 20%. Approximately 15% growth inhibition has been achieved with only 100 nM 25(OH)D₃ in LNCaP cells⁹³ and 50% inhibition with 800-900 nM 25(OH)D₃ in BxPC-3 and AsPC-1 pancreatic cell lines¹⁰⁶. These concentrations are about 10-fold less than the concentration used in this study, suggesting that CYP24A1 may have exerted oncogenic effects on the HeLa cells with high doses of 25(OH)D₃.

The VDR protein is primarily known for regulating the expression of the target genes of vitamin D hormones. While precursor 25(OH)D₃ and active hormone 1,25(OH)₂D₃ can bind to VDR, they exhibit different potencies¹⁵. Notably, VDR protein expression increased with physiological and supraphysiological doses of 25(OH)D₃ treatment, indicating that VDR mediates cell growth inhibition and cell death mechanisms. VDR protein expression was increased with both physiological and supraphysiological dose treatments of 25(OH)D₃, suggesting that HeLa cell growth inhibition and cell death mechanisms were mediated by VDR. Interestingly, treatment at a physiological dose (250 nM) did not significantly change VDR expression in SiHa cervical cells and supraphysiological treatment dose significantly down-regulated VDR gene and protein expression ($p=0.0380$)¹⁰². This suggests that intracellular signalling by VDR is cell line and dose-dependent.

As previously mentioned, 25(OH)D₃ may exert direct effects in the supraphysiological treatment of HeLa cells. Whether this is through VDR or other mechanisms needs to be further investigated.

In summary, 25(OH)D₃ inhibits cell growth and directly induces cell death via CYP27B1 activation to calcitriol and VDR binding in a dose-dependent manner. Higher doses induce the overexpression of CYP24A1, which limits the anti-cancer actions of the 25(OH)D₃ through catabolic activity.

Reports have shown that splice variants may limit the enzymatic activity of CYP24A1¹⁰⁷ and CYP27B1¹⁰⁸. This raises the possibility that polymorphisms of VDMS enzymes may exist in the HeLa cell line, and they may also be affecting the growth inhibitory actions of vitamin D. Genetic sequencing and mRNA analysis may clarify understanding of the effects of 25(OH)D₃ on the VDMS and mechanisms by which anti-cancer actions are exerted in cervical cancer. In addition, 25(OH)D₃ is highly concentrated in blood serum and further studies are needed to assess the effects of high doses of 25(OH)D₃ in patients with cervical cancer.

The HeLa cell line contains multinucleated cells potentially resistant to 25(OH)D₃ treatment

The identification of isolated multinucleated cells (MNCs) in the experimental and control cultures was an unexpected observation. MNCs, also known as polyploid giant cancer cells (PGCCs), are believed to arise from chemotherapy-induced stress¹⁰⁹⁻¹¹². They can survive *in vitro* and produce daughter cells through budding, splitting and bursting¹¹³. Their function is not fully understood, but they have been identified in tumours of different cancers and have distinct morphological features such as enlarged cell size and multiple nuclei¹¹⁴.

Semi-quantitative analysis of microscopy images revealed that 25(OH)D₃ treatment significantly increased the size of MNCs in the HeLa cell line. The number of MNCs and number of nuclei in the 5000 nM dose treatment were higher than in the other cultures; however, there were no statistical differences observed. This suggests that the MNCs in the HeLa cultures may resist high doses of 25(OH)D₃ treatment. An increase in the size of MNCs supports the hypothesis that MNCs are senescent cells undergoing G₁ cell cycle arrest^{109,114}, however, cell cycle analysis on isolated MNCs is required to inform a conclusion.

Studies show that MNCs are common in tumours triggered by oncoviruses¹¹⁵. This supports claims that MNCs are resistant to the anti-cancer actions of chemotherapeutic drugs. Oncoviruses such as HPV can affect tumour heterogeneity, thereby hindering the anti-cancer actions of chemotherapeutic drugs. A noteworthy correlation has been identified between high-risk HPV oncogenic subtypes infection, the occurrence of high-grade squamous intraepithelial lesions, and the identification of polyploidy¹¹⁵. Studies have also shown that both E6 and E7 oncoproteins deregulate kinases essential for mitosis and cytokinesis, resulting in polyploid cells¹¹⁶, and E6 and E7 are implicated in HPV-mediated cervical tumorigenesis.

It is not clear if and how 25(OH)D₃ affected MNCs in the HeLa culture. Investigations with isolated cells are required to assess how the MNCs arise and how they are affected by vitamin D treatment. Also, investigations on the presence of MNCs *in vivo* with vitamin D treatment could give insight into the use of vitamin D as a therapeutic agent for cervical cancer treatment.

Chapter 5

Conclusions

Currently, most cervical cancer cases are treated with surgery and chemo-radiotherapy. Vitamin D treatment may prove to be a less invasive treatment option. It is well documented that $1,25(\text{OH})_2\text{D}_3$ is the most active and potent vitamin D metabolite; however, its administration is associated with dose-dependent hypercalcaemia. Therefore, there is immense potential in using $25(\text{OH})\text{D}_3$ as an alternative therapeutic due to its low calcaemic effects. This study revealed that $25(\text{OH})\text{D}_3$ inhibits cell growth and induces apoptotic cell death in the HeLa cell line via VDMS in an autocrine manner. Therefore, the cells can regulate their survival and growth with $25(\text{OH})\text{D}_3$ treatment through activation by CYP27B1 and VDR. A possibility exists that $25(\text{OH})\text{D}_3$ may exert its effects directly without activation by CYP27B1; however, further investigations are needed to conclude the mechanism of action. In addition, the catabolic CYP24A1 action that degrades early and activated vitamin D hormone may mitigate the anti-tumour action of $25(\text{OH})\text{D}_3$, especially at very high-dose of $25(\text{OH})\text{D}_3$ treatment via a negative autocrine feedback system.

Further studies are needed to determine how $25(\text{OH})\text{D}_3$ treatment affects serum $25(\text{OH})\text{D}_3$ levels and whether the anti-cancer actions of $25(\text{OH})\text{D}_3$ are similar *in vivo*. In addition, it is well known that there is a correlation between HIV infection and the development of cervical cancer. Investigations on the therapeutic effects of vitamin D in HIV-positive cervical cancer patients may broaden the understanding of the mechanisms of action of vitamin D in cervical cancer.

The collective results of this study suggest that the autocrine VDMS mediates the effects of $25(\text{OH})\text{D}_3$ on the growth and death of HeLa cells in a dose-dependent manner. These findings highlight the therapeutic potential of $25(\text{OH})\text{D}_3$ for treating cervical cancer and indicate the need for further clinical investigations to explore this potential.

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Appendices

Appendix A

Table 6.1: Preparation of 10X PBS Buffer stock solution (pH 7.4).

Sodium Chloride	80g
Potassium Chloride	2g
Sodium dihydrogen phosphate	10.5g
Potassium dihydrogen phosphate	2g
Distilled water	800ml

Once pH has been adjusted using hydrochloric acid (HCl) and sodium hydroxide (NaOH), the solution will be made up to 1 L with distilled water.

Appendix B

Table 6.2 Preparation LDH assay buffer.

Buffer	Concentration and pH	Components	Notes
Tris Buffer	200mM, pH 8.0	24.4 g Tris base 1 L milli-Q water	Sterilize by autoclave
2X LDH assay Buffer		223 mg INT 57 mg PMS 575 mg NAD 3.2 g lactic acid 480 ml Tris	Protect aliquots from light Store in small aliquots at -20°C
10X Lysis solution	9% Triton X-100	9 ml Triton X-100 91 ml milli-Q water	

Appendix C

Table 6.3: Palade's fixative preparation.

Palade's solution A	Barbital sodium	14.7 g
	Sodium acetate	9.7 g
	Distilled water	500 ml
Palade's solution B	(2%) Osmium tetroxide	1 g (capsule)
	Distilled water	50 ml
Palade's solution C	(0:1N HCC) HCL	1.96 ml
	Distilled water	200 ml
Palade's fixative	Solution A	5 ml
	Solution C	5 ml
	Distilled water	2.5 ml
	Solution B	27.5 ml

Table 6.4: Resin solution preparation for 5g mixture.

	5g
Quetol	1.94
MNA	2.23
DDSA	0.83
RD2	0.10

Table 6.5: Embedding solution preparation for 5g mixture.

	5g
EmBed812	2.617
MNA	1.767
DDSA	0.683
S1	0.05

Appendix D

Table 6.6: RT-qPCR primer sequences. Primer pairs¹¹⁷⁻¹¹⁸ for genes to be investigated in this study.

Amplicon	Forward (5' → 3')	Reverse (5' → 3')	Primer Concentration	Denaturation	Annealing	Extension
VDR	GGA GAA AAC ACT TGT AAG TTG CT	TGG TCA GGT TGG TCT CGA ACT	700 nM	95°C 15s	56°C 15s	72°C 10s
CYP27B1	CCT GGC AGA GCT TGA ATT GCA	GGG GAA GAT GTA TAC CTT GGT	700 nM	95°C 15s	58.5°C 15s	72°C 10s
CYP24A1	CGG ACT CTT GAC AAG GCA ACA	TGA GGC GTA TTA TCG CTG GCA	600 nM	95°C 15s	64.5°C 15s	72°C 10s
18s rRNA	ACA CGG ACA GGA TTG ACA GA	CAA ATC GCT CCA ACT AA	500 nM	-	-	-

Melt curve conditions:

- 65-95 °C
- 0.5°C each step
- 90 sec hold at start
- 5 sec after each step

Appendix E

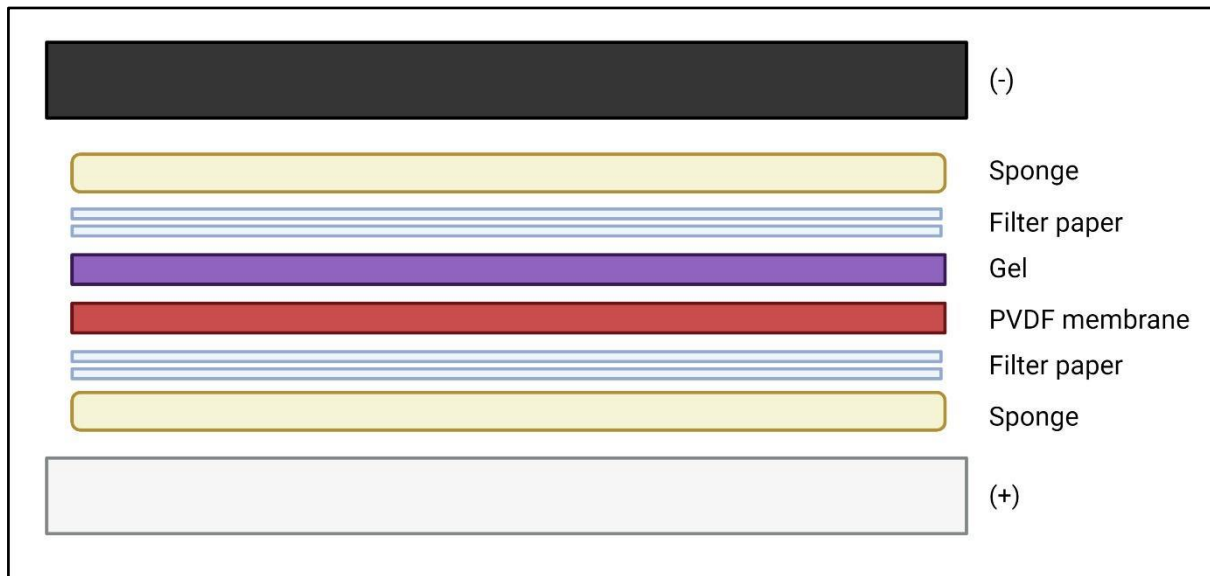


Figure 6.1: Western blot transfer sandwich orientation. Proteins contain a negative charge, so the gel will be placed near the cathode (negatively charged, in black). The polyvinylidene fluoride (PVDF) membrane will be placed on the side nearer to the anode (positively charged, in grey). The sandwich will be supported by filter paper (in dark blue) and sponges (in yellow).

Appendix F

Table 6.7: Western blot antibody concentrations and conditions.

Protein name	Antibody (catalogue number)	Calculated molecular weight (kDa)	Concentration	Incubation time and temperature	Blocking solution	Secondary antibody
CYP27B1	Anti-rabbit IgG, monoclonal CYP27B1 antibody (#ab206655)	56	1:500	4°C Overnight	0.125g BSA in 0.2% PBS-Tween	Goat Anit-Rabbit HRP-conjugated
CYP24A1	Polyclonal CYP241 antibody (#ab203308)	59	1:1000			
VDR	Rabbit IgG, monoclonal VDR antibody (#12550)	48	1:500			
β -Actin	Monoclonal anti- β -Actin peroxidase antibody (#A3854)	42	1:5000	1 hour at room temperature		-
2° Antibody	Goat Anit-Rabbit HRP-conjugated antibody (#E-AB-1003)	-	1:5000			-

Appendix G

Table 6.8: Stripping buffer recipe (pH 2.2)

Glycine	15 g
Sodium dodecyl sulphate (SDS)	1 g
Tween-20	10 mL
Distilled water	800 mL

Once pH has been adjusted using hydrochloric acid (HCl) and sodium hydroxide (NaOH), the solution will be made up to 1 L with distilled water.

Appendix H

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25-HYDROXYVITAMIN D₃ INDUCES BIOCHEMICAL AND MORPHOLOGICAL APOPTOSIS IN THE HELA CELL LINE

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Vitamin D and its metabolites have exhibited numerous anti-cancer actions, such as apoptotic cell death, in various pre-clinical cancer studies. Apoptosis is most commonly known as programmed cell death (PCD) type I and is essential during development, after injury and in disease prevention. Biochemical features include cell membrane asymmetry, caspase activation, changes in mitochondrial membrane potential ($\Delta\Psi_m$), and DNA damage. Morphological features of apoptosis include reduction of cellular volume, chromatin condensation (pyknosis), nuclear fragmentation (karyorrhexis), nuclear breakdown (karyolysis), cytoplasmic blebbing and apoptotic bodies.^{1,2}

Cervical cancer is one of the leading causes of cancer related deaths among South African women, affecting approximately 1 in 40 women. Few studies have explored the anti-cancer effects of vitamin D metabolites in cervical cancer.³ In this study, cervical adenocarcinoma cell line, HeLa was treated with 25-hydroxyvitamin D₃ (25(OH)D₃), and biochemical and morphological features of apoptosis were assessed.

HeLa cells were seeded at a concentration of 30 000 cells/ml and incubated for 16 hours. Cells were then treated with 25(OH)D₃ at a physiological concentration (260 nM) and a supraphysiological concentration (5000 nM). Medium and solvent (0.5 % ethanol v/v controls) were also used. After a 72 hours incubation, cells were trypsinised and resuspended in fresh culture media. The cells were then stained with either annexin V, tetramethyl rhodamine ethyl ester (TMRE) or DEVD peptide-linked DNA dye. All samples were then stained with 7-aminoactinomycin D (7-AAD). Samples were then acquired using the Luminex® Guava® Muse® Cell Analyzer. Biological and technical repeats were triplicated (expressed as mean \pm SEM) and analysed by one-way ANOVA with post-hoc Bonferroni testing at significance $p < 0.05$.

To assess morphological features of apoptosis, HeLa cells were seeded on glass coverslips at a concentration of 50 000 cells/ml and treated as previously described. After 72 hours incubation with treatment, cells were fixed and stained with Mayer's Hemalum and eosin. Cells were viewed with brightfield microscopy.

Annexin V staining and DEVD peptide-linked dye showed that 25(OH)D₃ treatment at 5000 nM significantly increased the percentage of apoptotic cells when compared to the solvent control ($p=0.0323$ and $p=0.0219$, respectively). In addition, 5000 nM treatment significantly induced mitochondrial membrane depolarisation when compared to the solvent control ($p=0.0061$).

Brightfield microscopy showed that treatment with 25(OH)D₃ at both the physiological and supraphysiological concentrations induced apoptosis. Vitamin D treated cells exhibited apoptotic bodies, cytoplasmic blebbing, hyper-condensed chromatin, shrunken nuclei and a reduction in cell volume (Fig. 1).

In the intrinsic pathway of apoptosis, the exposure of cells to external stimuli initiates a cascade of molecular events.^{1,2} Our results showed that annexin V staining was increased with 25(OH)D₃ treatment, indicating an increased externalisation of phosphatidylserine on the plasma membrane. Similar results were seen in osteosarcoma cell lines (SAOS-2 and 143B)⁴ treated with 25(OH)D₃ at 1000 nM. Depolarisation of the mitochondrial membrane releases proteins that activate caspases.¹ Our results show that 25(OH)D₃ treatment resulted in the depolarisation of the mitochondria and induction of caspase-3 and -7 activity², implying that 25(OH)D₃ induces the intrinsic pathway of apoptosis. Active caspase-3 and -7 are involved in the fragmentation of DNA during apoptosis. Brightfield microscopy revealed cells with hyper-condensed chromatin, indicating potential DNA damage. The presence of apoptotic bodies in 25(OH)D₃ further confirms the induction of apoptosis. In conclusion, 25(OH)D₃ induces apoptosis at a supraphysiological concentration, potentially via the intrinsic pathway. Further investigation of the cell cycle and DNA content need to be conducted to corroborate findings.

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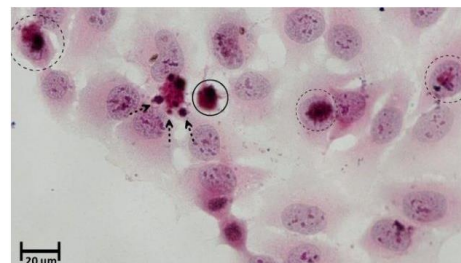


Figure 1. Brightfield microscopy image of HeLa cells treated with 5000 nM 25(OH)D₃. Cells with apoptotic bodies (arrow), shrunken cells (circle) and hyper-condensed chromatin (dashed circle) were visible.

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