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The Effect of *Alhagi maurorum* Medik on Mitochondrial Function

Juwairiya Zulfiqar Butt

A thesis submitted in partial fulfilment of the requirements of the University of Westminster for the degree of Doctor of Philosophy

September 2023

Abstract	1
List of Conference Posters	2
Acknowledgements	3
List of Abbreviations	5
List of Tables	8
List of Figures	9
Statement of Problem	
Aims and Objectives	11 12
Hypotnesis	12 12
Unapter 1	13 12
1 1 Mitochondria- Impact Life	13 13
1.2 Mitochondrial Morphology	
1.3 Mitochondrial Cellular Crosstalk	
1.4 Mitochondrial Disorders	23
1.5 Mitochondrial Functions Evolving Therapeutic Targets	24
1.6 Mitocans	28
1.7 Mitochondria and Phytotherapy	31
1.8 Scope of Complementary Medicine in Pakistan	34
1.9 Alhagi maurorum Medik: Multipurpose Herb	34
1.10 Phenethylamine Alkaloid: Hordenine	40
In vitro and in vivo studies of hordenine	43
Chapter 2	54
Materials and Methods	54
2.1 Plant Collection	54
2.2 Plant Identification	55
2.3 Plant Extraction	56
2.3.1 Pulverization	56
2.3.2 Solvent Extraction	56
2.4 High Performance Thin Layer Chromatography analysis of <i>Alhagi</i> rextract	<i>naurorum</i> 58
2.5 Liquid Chromatography-Mass Spectroscopic Analysis of Alhagi maurori	<i>ım</i> extract
2.6 Cell Culture	60
2.6.1 Cell Lines Used in Research	60
2.6.2 Thawing Cell Lines from Liquid Nitrogen	
2.6.3 Subculture of Cell Lines (Cell Passaging/Cell Splitting)	67
2.6.4 Cell Seeding	02 67

Table of Contents

2.6.5 Freezing Cells for Storage in Liquid Nitrogen	63
2.6.6 MTT Reduction Assay (Cell proliferation Assay)	63
2.6.7 Detection and Quantification of Cellular ROS	64
2.6.8 Detection and Quantification of Mitochondrial Membrane Potential	65
2.6.9 Detection and Quantification of Intracellular Ca ⁺²	66
2.6.10 Seahorse Cell Mito Stress Assay	67
2.6.11 Morphological Analysis Using Fluorescent Microscopy	69
Chapter 3	71
Results of <i>Alhagi maurorum</i> extract	71
3.2 Characterization of <i>Alhagi maurorum</i> with Liquid	
Chromatography-Mass Spectroscopic	
3.3 Measurement of Cell Viability	82
3.3.1 Effects of <i>Alhagi maurorum</i> Extract on Cell Viability	82
3.4 Measurement of Oxidative Stress	84
3.4.1 Effects of Alhagi maurorum Extract on ROS Levels	84
3.5 Measurement of Mitochondrial Membrane Potential (MMP)	85
3.5.1 Effects of Alhagi maurorum Extract on MMP	
3.6 Measurement of Intracellular Calcium Concentrations	
3.6.1 Effects of <i>Alhagi maurorum</i> Extract on Intracellular Ca ²⁺	
3.7 Measurement of Mitochondrial Respiration	
3.7.1 Effects of Alhagi maurorum Extract on Mitochondrial Respiration	
Chapter 4	92
Results of Hordenine	
4.1 1 Effects of Usedening on Coll Vichility	
4.1.1 Effects of Hordenine on Cell Viability	
4.2 Measurement of Oxidative stress	
4.2.1 Effects of Hordenine on ROS Levels	94
4.3 Measurement of Mitochondrial Membrane Potential (MMP)	95
4.3.1 Effects of Hordenine on MMP	
4.4 Measurement of Intracellular Ca ²⁺	97
4.4.1 Effects of Hordenine on Intracellular Ca ²⁺	97
4.5 Measurement of Mitochondrial Respiration	97
4.5.1 Effects of Hordenine on Mitochondrial Respiration	
4.6 Observation of Morphology Using Fluorescent Microscopy	101

Chapter 5	104
Discussion	104
5.1 Effect of AM extract and hordenine on Cell Viability	104
5.2 Effect of AM extract and hordenine on ROS Levels	106
5.3 Effect of AM extract and hordenine on Mitochondrial Membrane Potential	111
5.4 Effect of AM extract and hordenine on Intracellular Calcium Concentration	112
5.5 Effect of AM extract and hordenine on OXPHOS	114
5.6 Effect of AM extract and hordenine on mitochondrial morphology	116
Future Work	119
Conclusion	121
References	122

Abstract

Mitochondrial bioenergetics are crucial for the optimal activity of the cell and are emerging therapeutic targets for the treatment of several metabolic syndromes and cancer. *Alhagi maurorum* plant is traditionally used in several clinical conditions due to its potential anti-inflammatory, hepatoprotective and gastrointestinal effects.

In this study changes in cell viability, reactive oxygen species (ROS) levels, mitochondrial membrane potential (MMP), oxidative phosphorylation (OXPHOS), calcium homeostasis and mitochondrial morphology in MCF-7 and MCF-10-A (control) cells after treatment with Alhagi maurorum for 24hr and its main bioactive compound hordenine for 6hr, 24hr, and 48hr are observed. Results indicate that cells treated with Alhagi maurorum have significantly reduced cell viability, ROS levels, MMP, and intracellular calcium concentrations at higher concentrations that suggest cytotoxic effects, while moderately reduced effects at 100 μ g/ml, indicating possible stimulation of several other cellular processes such as cell death, antioxidant systems, and immune responses. Furthermore, Alhagi maurorum extract modulated various parameters of mitochondrial respiration too including, basal respiration, ATP production, proton leak, spare respiratory capacity, maximal respiration, and non-mitochondrial respiration. In the cells treated with hordenine, the key bioactive compound of Alhagi maurorum, results indicate that hordenine has no significant effect on MCF-7 and MCF-10-A after 24hr and 48hr of exposure, including cell viability, ROS levels, MMP, intracellular calcium concentrations and mitochondrial respiration. However, in MCF-10-A after 6hr of the treatment ROS levels are significantly reduced at concentrations 0.1 μ M, 10 μ M, 500 μ M, and 1000 μ M. Furthermore, no significant changes to mitochondrial morphology are observed at lower concentrations of Alhagi maurorum and hordenine tested in MCF-7 and MCF-10-A, though only small, rounded networks of mitochondria, increased in number in case of hordenine suggest possible occurrence of the process of fission.

In conclusion, my work shows that *Alhagi maurorum* extract modulates mitochondrial function, and effects are concentration-dependent while one of the main bioactive compounds of hordenine phenethylamine alkaloid, has more limited effects. Characterisation of the extract through HPTLC and LC-MS analysis showed presence of more than one compound in the extract.

List of Conference Posters

Hordenine Regulates Redox state and Mitochondrial Functions in Malignant and Nonmalignant Breast Cell Lines in Concentration Dependent Manner. The 21st International Congress of International Society for Ethnopharmacology (Ethnopharmacology 2022) taking place at China Medical University, Taichung, Taiwan, May 28-31, 2022.

Hordenine: Redox State and Mitochondrial Functions in Malignant and Non-Malignant Breast Cell Lines. Poster presented at 70th International Congress and Annual Meetings of the Society for Medicinal Plant and Natural Product Research (GA), Thessaloniki, Greece, August 29, 2022.

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Declaration of Contributors

I declare that all the material contained in this thesis is my own work. Any assistance is detailed below:

LC-MS Analysis

Francesca Scotti performed LC-MS experiments of plant *Alhagi maurorum* and data is analysed by the author.

List of Abbreviations

ATP	Adenosine triphosphate
AMPA	α-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid
ANOVA	Analysis of variation
ANT	Adenosine nucleotide translocator
ADP	Adenosine diphosphate
APAF1	Apoptotic protease activating factor
AIF	Apoptosis-inducing factor
Bad	Bcl-2 associated death promoter.
Bak	Bcl-2 homologous antagonist killer
Bax	Bcl-2 associated protein X
Bcl	Cell lymphoma
Bcl-2	B-cell lymphoma
BSA	Bovine serum albumin
Ca ⁺²	Calcium ion
СоА	Coenzyme A
Cyt c	Cytochrome c
CSCs	Cancer stem cells
DCF	2',7' –dichlorofluorescin
DCFDA	2',7' –dichlorofluorescin diacetate
DNA	Deoxyribonucleic acid
DAMPs	Damage-associated molecular patterns
DMEM	Dulbecco's modified Eagle's medium
DI	Deionized
DMEM	Dulbecco's modified Eagle's medium
DNA	Deoxyribonucleic acid
Drp1	Dynamin-related protein
ETC	Electron transport chain
FAD/FADH	Flavin adenine dinucleotide
FADD	Fas-associated protein with death domain

FasL	Fas ligand
FBS	Foetal bovine serum
FCCP	Carbonyl cyanide-p-trifluoromethoxyphenylhydrazone
GIT	Gastrointestinal tract
IMM	Inner mitochondrial membrane
HL60	Human acute myeloid leukemia cells
HPTLC	High Performance Thin Layer Chromatography
MAPK	Mitogen activated protein kinase
MAVS	Mitochondrial antiviral-signalling protein
MCU	Mitochondrial calcium uniporter
MEME	Modified Essential Media Eagle
miRNA	Micro ribonucleic acid
MOMP	Mitochondrial outer membrane permeabilization
M.W	Molecular weight
mPTP	Mitochondrial permeability transition pore
mtDNA	Mitochondrial DNA
MitoTAM	Mitochondrially targeted tamoxifen
MTT	3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide
MMP	Mitochondrial membrane potential
MS	Mass Spectrometry
NAD/NADH	Nicotinamide adenine dinucleotide
NADP/NADPH	Nicotinamide adenine dinucleotide phosphate
NF-ĸB	Nuclear factor kappa-light-chain-enhancer of activated B cells
NO	Nitric oxide
NOX	NADPH oxidase
Nrf2	Nuclear factor (erythroid-derived 2)-like 2
Na+	Sodium ion
NMR	Nuclear Magnetic Resonance
OCR	Oxygen consumption rate
OMM	Outer mitochondrial membrane
OXPHOS	Oxidative phosphorylation

PAMPs	Pathogen-associated molecular patterns
PBS	Phosphate buffered saline
PCD	Programmed cell death
PI3K	Phosphoinositide 3-kinase
P53	Upregulated modulator of apoptosis
PRL	Prolactin
PRRs	Pattern-recognition receptors
RIPA	Radioimmunoprecipitation assay
RNA	Ribonucleic acid
ROS	Reactive oxygen species
RPMI	Roswell Park memorial institute medium
RIPK3	Receptor-interacting serine/threonine protein kinase 3
Smac/DIABLO	Second mitochondria derived activator of caspases
TCA	Tricarboxylic acid cycle/Kreb's cycle/citric acid cycle
TMRE	Tetramethyl rhodamine ethyl ester
TNF-α	Tumour necrosis factor-a
TP53	Tumour protein 53, commonly referred to as p53
tRNA	Transfer RNA
UTI	Urinary tract infections
UV	Ultraviolet
VDAC	Voltage dependant anion channel

List of Tables

Table 1: Mitochondrial morphology and structural diversity	20
Table 2: Role of pro- and anti-apoptotic proteins in morphological remodelling	21
Table 3: Anti-cancer drugs and their mechanism of action	30
Table 4: Natural compounds affecting mitochondrial functions	33
Table 5: Alhagi maurorum: Pharmacological effects	39
Table 6: Biological activities of hordenine	53
Table 7: Working concentration of DCFDA	57
Table 8: Working concentration of TMRE	65
Table 9: Working concentration of Rhod-2	65
Table 10: Working concentration ETC Modulators	66
Table 11: Equations of key parameters of Seahorse mitostress assay	69
Table 12: List of extraction solvents	69
Table 13: List of developing solvents for HPTLC	72
Table 14: LC-MS analysis: Tentative identification of compounds	78

List of Figures

Figure 1: Mitochondrial research overview	16
Figure 2: Mitochondrial morphology	19
Figure 3: Schematic of process of fission and fusion	21
Figure 4: Mitochondrial crosstalk with other organelles	22
Figure 5: Illustration of ROS and MMP	27
Figure 6: Hallmarks of cancer and their inhibitors	31
Figure 7: Alhagi maurorum Medik: Traditional background	36
Figure 8: Commercially available dietary supplements of hordenine	41
Figure 9: Chemical structure of hordenine	42
Figure 10: Biosynthesis of hordenine	42
Figure 11: Hordenine modulation of MAPK pathway	44
Figure 12: Hordenine and synergistic effect	45
Figure 13: Hordenine and pyruvate dehydrogenase kinase 3 (PDK3)	46
Figure 14: Renal protective activity: hordenine and Nrf2	47
Figure 15: Hordenine and cAMP	48
Figure 16: Effect of hordenine on critical cellular pathway	52
Figure 17: Cell viability assay	54
Figure 18: Acute intracellular Ca ⁺² assay	55
Figure 19: ETC Targets: Site of action of ETC modulators	56
Figure 20: Sampling area of Alhagi maurorum	58
Figure 21: Desiccated sample of Alhagi maurorum	64
Figure 22: Mini Mill: Pulverization of Alhagi maurorum	67
Figure 23: Schematic of extract preparation of Alhagi maurorum	68
Figure 24: HPTLC analysis of Alhagi maurorum	72
Figure 25: Densitometric view of Alhagi maurorum	73
Figure 26: LC-MS analysis of Alhagi maurorum	77
Figure 27: Effect of solvent on Cell viability	82
Figure 28: Effect of Alhagi maurorum on Cell viability	83

Figure 29: Cell viability assessment of Alhagi maurorum	83
Figure 30: Effect of solvent on ROS levels	84
Figure 31: Effect of Alhagi maurorum on ROS levels	85
Figure 32: Effect of Alhagi maurorum on mitochondrial membrane potential	86
Figure 33: Effect of <i>Alhagi maurorum</i> on intracellular calcium levels (Ca ⁺²)	87
Figure 34: Effect of Alhagi maurorum on OXPHOS in MCF-7	89
Figure 35: MCF-7 OCR profile	89
Figure 36: Effect of Alhagi maurorum on OXPHOS in MCF-10-A	90
Figure 37: MCF-10-A OCR profile	90
Figure 38: Effect of hordenine on Cell viability	93
Figure 39: Effect of hordenine on ROS levels	94
Figure 40: Effect of hordenine on mitochondrial membrane potential	96
Figure 41: Effect of hordenine on intracellular calcium levels (Ca ⁺²)	97
Figure 42: Effect of hordenine on OXPHOS of MCF-7	98
Figure 43: OCR Profile of MCF-7	99
Figure 44: Effect of hordenine on OXPHOS of MCF-10-A	100
Figure 45: OCR Profile of MCF-10-A	101
Figure 46: Morphological analysis	103

Statement of Problem

Mitochondrial dysfunction has been reported in several acute and chronic diseases, for example fatigue, obesity, diabetes, hypertension, and cancer. Therefore, regulating mitochondrial functions may offer new therapeutic opportunities. *Alhagi maurorum* Medik plant has been reported to have potential anti-inflammatory and anti-cancer activities however, it has not been assessed in reference to possible impact on mitochondrial functions.

Aims and Objectives

Aim: To assess the impact of *Alhagi maurorum* Medik and its bioactive compounds on mitochondrial function.

Objectives:

- 1. To determine if cell viability is affected by *Alhagi maurorum* extract and its bioactive compound hordenine.
- To assess the effect of *Alhagi maurorum* extract and its main bioactive compound on mitochondrial functions, including reactive oxygen species (ROS) production, mitochondrial membrane potential (MMP), intracellular Ca⁺² concentrations, and mitochondrial respiration.
- **3.** To determine the effect of *Alhagi maurorum* extract and hordenine on mitochondrial morphology.
- **4.** To identify the principal bioactive compounds associated with the mitochondrial effects induced by *Alhagi maurorum* extract.

Hypothesis

I hypothesize that *Alhagi maurorum* extract and selective secondary metabolite hordenine can regulate mitochondrial functions, including mitochondrial membrane potential (MMP), adenosine triphosphate (ATP) production, reactive oxygen species (ROS) levels, intracellular Ca⁺² concentrations, oxidative phosphorylation (OXPHOS) and mitochondrial morphology.

Chapter 1

Introduction

1.1 Mitochondria- Impact Life

Mitochondria are critical semi-autonomous organelles essential for the optimal activity of the cell. Mitochondria meet the functional needs of the cell because their size, structure and number vary considerably, according to their physiological role (Liu et al., 2019; Calvo and Mootha, 2010).

The mitochondrion is commonly known for its classically appreciated role as the powerhouse of the cell, but now after intensive research it is also considered as master regulator of the cellular system. It serves as a biosynthetic hub for the synthesis of macromolecules essential for nucleotides, fatty acids, cholesterol, amino acids, glucose, heme, and iron-sulphur proteins by taking part in critical bioenergetic circuitries such as the tricarboxylic acid (TCA) cycle (Martinez-Reyes and Chandel, 2020) and oxidative phosphorylation (OXPHOS) (Ait-Aissa et al., 2019; Allen et al., 2011). It also serves as a waste management hub, degrading misfolded mitochondrial proteins intrinsically through the system of proteases in their compartments, for example the ubiquitin proteasome system (UPS) which is a part of mitochondrial quality control system and removes dysfunctional mitochondria involved in critical cellular processes of cell death most importantly mitophagy to maintain healthy cellular environment (Xiao and Loscalzo, 2019; Spinelli and Haigis, 2018; Wang and Youle, 2009). The mitochondria are also important to the cell signalling circuitry, either serving as a physical platform where protein-protein signalling interactions occur or regulating production and levels of different signalling molecules such as reactive oxygen species (ROS) and Calcium (Ca^{2+}), implicated in several other important cellular processes including initiation of immunogenic responses, growth, and differentiation. However, here focus will be mainly on the first three main aspects of mitochondrial functions because they are interrelated, crucial for mitochondrial health and significant therapeutic targets (Wang and Youle, 2009; Liu et al., 2019; Gibellini et al., 2015; Antico et al., 2012)

Mitochondria, being sensitive organelles, rapidly detect and respond to any changes in their cellular environment caused by endogenous or exogenous stimuli such as nutrients (selenium), co-factors, enzymes, toxins, alcohol and even frequently prescribed drugs as seen in the case of carnitine, which is an essential co-factor required for the transport of long chain fatty acids into the mitochondrion for β -oxidation and depletion of Co-enzyme Q10 (CoQ10), essential for the transport of high energy electrons across the electron transport chain (ETC). At present, mitochondrial damage is assessed indirectly through several clinical markers for example, altered blood levels of lactate, pyruvate, and organic acid in urine (Yun and Finkel, 2014).

Since discovery, mitochondria continued to be a subject of research in various disciplines of biological and medical sciences, now their emerging role as a pharmacological target is a subject of research in the field of modern medicine (Ernster, and Schatz, 1981; Pagliarini and Rutter, 2013) (For an overview of mitochondrial research see Figure 1).





Figure 1: Mitochondrial research overview: Mitochondria are interesting organelles for researchers of various discipline (Created with BioRender).

1.2 Mitochondrial Morphology

Mitochondria are compartmentalized intracellular organelles normally occupying an area between 0.75 and 3 μ m², round to oval in shape and range in size from 0.5 to 10 μ m, found in the cytoplasm of the eukaryotic cells (Veltri et al., 1990; Piomboni et al., 2012). The outer mitochondrial membrane (OMM) is porous for most of the molecules such as metabolites and products of cellular respiration and faces cytosol. Proteins residing on OMM play significant role in mitochondrial fusion proteins (MFN1, MFN2), anti-apoptotic proteins and mitochondrial voltage-dependent anion channels (VDACs) also termed as porins (De Stefani et al., 2012). The inner mitochondrial membrane (IMM) protrudes into the mitochondrial matrix and forms invaginations called cristae that hold the main protein complexes of the main energy-generating system, the ETC. In addition to this, mitochondrial fusion protein optic atrophy 1 (OPA1) also resides in the inner mitochondrial membrane (Rastogi et al., 2019). The organelle's central mass is a gel like matrix which contains mitochondrial DNA (mtDNA), also called mitogenome.

Common geometric features that describe morphology of mitochondrion includes shape, size, position, and dynamics measured at nanometre scale to the micron scale and all the above-mentioned geometric features are interrelated and collectively define mitochondrial morphology. Remodelling of mitochondrial architecture in response to the cell's metabolic requirements results in either punctuated mitochondria (<1 μ m in length) or interlocked connected networks of mitochondria (>3 μ m in length) (Collins et al, 2021; Dowling et al, 2021). Different mitochondrial morphological states have been linked to various physiological as well as pathophysiological states. Since, fragmented mitochondria are observed under stressed and different states of cellular death where, elongated morphology has been linked to the efficient production of ATP though, full impact of mitochondrial morphology on the mitochondrial matrix and OXPHOS still needs to be fully explored (Figure 2).



Figure 2: Mitochondrial morphological remodeling: Different mitochondrial shapes indicate dynamic nature of mitochondria under physiological and pathological conditions. Fibroblasts (Differentiated cells) possess elongated mitochondria with increased OXPHOS. Fragmented mitochondrial network with higher glycolytic rate and sensitive to death (Created with BioRender).

Although the dynamic nature of mitochondria was observed over 100 years ago, the main proteins involved in the maintenance of mitochondrial morphology have only been identified in the past few decades and lack full mechanistic information (Lewis and Lewis, 1914; Garcia et al., 2019; Lotz et al., 2014). Also, the mitochondrial proteome is found to be dynamically regulated as observed in the case of human cardiac cells that contain fewer mitochondria as compared to muscle cells, so relatively low number of proteins too. This indicates the impact of mitochondrial density on the mitochondrial functions depending upon the metabolic requirements of the cells even in two highly metabolically active cells (for more information about mitochondrial morphology and cell type, see Table 1) (Martinez-Reyes and Chandel, 2020; Ait-Aissa et al., 2019; Xiao and Loscalzo, 2019; Liu et al., 2019; Gibellini et al., 2015;). Similarly, higher concentration of mitochondria are also found in the active growth cones of the developing neurons further supporting the relationship of mitochondrial density and mitochondrial remodelling based on metabolic requirements of the organs (Zhao et al., 2012; Mils, 2015; Benard et al., 2007).

Cell type	Mitochondrion size, shape, position	
Fibroblasts	Long filaments (1–10 μ m in length, diameter of ~ 700 nm)	
Hepatocytes	Uniform spheres or ovoids in shape	
Vascular smooth muscle	Ovoid or rod-shaped organelles	
Endothelium	Tubular mitochondrial network	
Skeletal muscle	Ovoid structures: Have two populations (one posi- tioned close to the sarcolemma that are round and smaller while other embedded among the myofibrils)	
Pancreatic acinar cells	 Three different regional groups, functionally unconnected. 1. Peripheral basolateral region near plasma membrane 2. Around the nucleus 3. In the periphery of the granular region separating the granules from the basolateral area 	
Cardiac myocytes	 Three distinct populations, 1. Perinuclear (more rounded & densely packed) 2. Subsarcolemmal 3. Interfibrillar 	
Cultured vascular smooth cells	Long filamentous entities, loops, and networks and re- curring structural classes such as small spheres, swol- len spheres, straight rods, twisted rods, branched rods, and loops	
Native vascular smooth cells	Solitary spheres and rods of various sizes	

Table 1: Mitochondrial morphology and structural diversity

Furthermore, a new role of mitochondrial morphology in the maintenance of cellular homeostasis has been shown to influence mitochondrial bioenergetic status (Navaratnarajah et al., 2021). Rapid but temporary morphological adaptations of mitochondria in response to metabolic flux are generally known to be mediated by the processes of fission and fusion [for detailed process of fission and fusion possibly involved in mitochondrial remodelling, see Figure 3 (Table 2)]. Fission and fusion processes are the most studied processes though the impact of some undefined processes that are also influencing mitochondrial morphology such as branching/de-branching and extension/retraction is still unknown. This is an emerging area of research and as mitochondrial morphological changes are observed in complex and rare diseases there is intense need for more studies to explore the connection between mitochondrial morphology and mitochondrial bioenergetics to design highly effective drugs to target mitochondrial morphology (Galloway, Lee and Yoon, 2012; Navaratnarajah et al.,

2021). Interestingly, mitochondrial morphological remodelling also has a significant impact on the initiation of immune cell metabolism in response to pathogens a process still needs understanding but critical for new emerging field of immunometabolism (O'Neill et al, 2016; Angajala et al, 2018; Rambold and Pearce, 2018).



Figure 3: Schematic of process of fission and fusion. Fusion (Increased oxidative capacity), Fission (Decreased oxidative capacity). MFN1 and MFN2 mediate fusion of the OMM which is followed by fusion of IMM mediated by OPA1 and results in elongated mitochondria with increased ATP production and OXPHOS rate. Various cycles of fission and fusion maintain mitochondrial health and help to eliminate dysfunctional mitochondria or repair any damaged mitochondria through a process called mitophagy controlled by Pink1 and Parkin molecules. Apoptotic cells further release messengers as signals for phagocytosis in the form of metabolites (inflammatory, ApoEVs (apoptotic extracellular vesicles), PS (Phosphatidylserine) molecules) (Rastogi, et al., 2019) (Created with BioRender).

Protein	Protein characteristics	References
Bcl-2 (Apoptosis regulator)	Increases mitochondrial size Increases mitochondrial complexity	Kowaltowski et al., 2002
Bax/Bak (Proapoptotic)	Increases fragmented mitochondria Increases apoptosis induction	Desagher and Martinou, 2000
Bid (Proapoptotic)	Increases apoptosis induction Increases mitochondrial cristae fusion Increases lipid translocase activity	Scorrano, Ashiya, et al., 2002 :

Table 2: Role of pro- and anti-apoptotic proteins in morphological remodelling

1.3 Mitochondrial Cellular Crosstalk

As mitochondria are considered critical organelles due to their role in cellular metabolism, they are also important in deciding the fate of the cell (Sun et al., 2015; Wiedemann and Pfanner, 2017). Recent studies about the functional roles of mitochondria suggest that mitochondrial activities and involvement of these organelles in multiple processes are tightly controlled by nuclear DNA through anterograde regulation of gene expression (Arnold, et al., 2006; Audano et al., 2020). More than 1000 mitochondrial proteins are encoded by nuclear genes and are synthesized as precursors in cytosol and are transported into mitochondria through five main transport pathways such as the carrier, β barrel and the pre-sequence pathways. Nuclear encoded mitochondrial proteins are transported into the mitochondrial matrix through the TOM/TIM complex located on OMM and IMM, respectively (Wiedemann and Pfanner, 2017; Suhm, et al., 2018; Picard, Wallace and Burelle, 2016). Mitochondrial protein import is a strictly controlled process and largely dependent upon mitochondrial energetics, though full molecular mechanism of transport of proteins is not known. Many scientific studies suggest mitochondria undergo intracellular stress and trigger specific nuclear gene transcription patterns through retrograde communication pathways (Chang, Shtessel and Lee, 2015; Braymer and Lill, 2017). This mito-cytosol-nuclear crosstalk is essential for the maintenance of cellular homeostasis (Xia, et al., 2019) (Figure 4).



Figure 4: Mitochondrial crosstalk with other organelles: Mitochondria endoplasmic reticulum (ER) crosstalk for Ca²⁺homeostasis. Mitochondria-lysosome crosstalk for lysosomal biogenesis. Mitochon-

drila peroxisomes crosstalk for fatty acid oxidation and ROS metabolism. Mito nuclear crosstalk, anterograde regulation and retrograde regulation for transcription proteins, mitochondrial biogenesis (Created with BioRender).

Mitochondria and lysosomes communicate for fine tuning of cell's functionality as both organelles take part in the metabolism and participate in the events of catabolism as well as anabolism (Scorrano et al., 2019). Lysosomes, which are membrane bound organelles in cells, are involved in cellular processes of breaking down of large molecules of carbohydrates, lipids, proteins and prevent infections by digesting pathogens due to the presence of hydrolytic enzymes. However, dysregulation of any of the mitochondrial activity affects the functions of lysosomes and results in many pathologies such as Parkinsonism disease and Huntington's disease (Demers-Lamarche et al., 2016; Baixauli et al., 2015). Mitochondria-lysosome crosstalk is thought to be mediated by AMPK and mTORC1 pathways (Mullins and Bonifacino, 2001).

Peroxisome, another dynamic organelle of the cell, coordinates with mitochondria in processes like redox regulation, inflammatory and immune signalling (Fransen, Lismont and Walton, 2017). Similarly, endoplasmic reticulum (ER) and mitochondria form microdomains, known as mitochondria-associated ER membranes, and are necessary for Ca^{2+} signaling, and mitochondrial dynamics. So, various studies on mitochondrial association with other organelles in the cell have shown these intracellular talks are essential to maintain cellular and mitochondrial homeostasis (Veeresh et al., 2019).

1.4 Mitochondrial Disorders

Mitochondrial disorders are multiple medical disorders produced due to defects in mitochondrial function (Garcia-Berumen et al., 2019). Symptoms due to mitochondrial defects range from mild to life threatening, such as fatigue, weakness, developmental abnormalities, diabetes, cognitive disabilities, liver failure, metabolic disorders, and cancer (Singh et al., 2015; Gorman et al., 2015; Wallace, 2010). Mitochondrial DNA contains 37 genes out of which 13 genes code for the components of respiratory chain complexes (Ruhoy and Saneto, 2014; McFarland et al., 2010; Wallace, 2010). Although, maternal inheritance of mtDNA mutations (deletions, duplications), is reported primarily, but most of the mitochondrial disorders are sporadic in nature and show different clinical presentations with similar phenotype that suggest this area needs to be further explored. However, lactic acidosis (metabolic acidosis) and Leigh's disease (neurological disorders) are some

examples of primary mitochondrial disorders (Lopez, 1998; Chinnery, 2021; Seifert et al., 2015; Marra et al., 2021; Taylor and Turnbull, 2005).

Secondary mitochondrial disorders occur during lifetime and are caused by alterations in some of the major steps of mitochondrial metabolism. Among several causes altering normal metabolism, the most commonly reported are the changes in mitochondrial bioenergetics or defects in substrate utilization, for example, PDH complex (Pyruvate dehydrogenase) which connects glycolysis to the TCA cycle and any alteration in the activity affects pyruvate metabolism and ultimately glycolysis, gluconeogenesis and the citric acid cycle as seen in diabetes, obesity, and even various types of cancers (Currie et al., 2013, Tennant et al., 2010; DeFronzo and Tripathy, 2009; Du, et al., 2013; DeFronzo and Tripathy, 2009; Du, et al., 2013; Kerr, 2013). Similarly, excess of fructose triggers apoptosis, uric acid and contribute to cardiovascular diseases (Longo et al., 2006; DiDonato, 1997; Stanley, 2004; Imbe et al., 2018; Ohara et al., 2018; Du, et al., 2013; Feinman and Fine, 2013; Calvo et al., 2014; Gnaiger, 2014) and fumarase deficiency causes mitochondrial encephalomyopathy (De Vivo, 1996; Luft et al., 1962; Liang, Ahmad and Sue, 2014). Furthermore, defects in OXPHOS coupling results in some of the abnormalities of the respiratory chain affecting ATP production and causes several neurodegenerative diseases and nonthyroidal hypermetabolism commonly known as Luft disease (Sivitz and Yorek, 2010). Neurodegenerative disorders like Parkinson and Huntington's disease are some examples of mitochondrial disorders due to defects in multi protein complexes of the ETC (Mimaki, et al., 2012; Fullerton et al., 2020; Fernndez-Vizarra and Zeviani, 2015). These defects are typically due to mitochondrial DNA mutations (Mayr et al., 2015; Vyas, Zaganjor and Haigis, 2016). Any defect in even one complex of electron transport chain ETC can alter the functions of other complexes for example, complex-III which is a master regulator of mitochondrial respiratory chain which can halt the assembly of complex I and biogenesis of Complex IV (cytochrome c oxidase) (Signes and Fernandez-Vizarra, 2018; Letts et al., 2019). Also, deficiency of Coenzyme Q_{10} (Co Q_{10}) an antioxidant and essential electron carrier of the mitochondrial respiratory chain has been linked to nephrotic conditions (Sacconi, et al., 2010; Shiau et al., 2022) and defects of Complex V are linked to fatal neonatal mitochondrial encephalopathy (Signes and Fernandez-Vizarra, 2018; Letts et al., 2019; Karolinska Institutet, 2021).

1.5 Mitochondrial Functions Evolving Therapeutic Targets

Cancer is a global health concern and one of the leading causes of death. It is a multifaceted disease caused by genetic, epigenetic, proteomic, transcriptomic, and metabolic changes (Hanahan and Weinberg, 2011; Alkhazraji et al., 2019; Cuyas et al., 2018). Nearly 10 million deaths are reported in 2020 according to the World Health Organization (WHO) report published in 2022, and among several benign and chronic types the most common types are, breast, lung, colon, rectum, and prostate cancers that are mainly caused by smoking, high body mass index (BMI), alcohol, poor lifestyle and certain infections in low-and middle-income countries such as hepatitis (Spivak et al., 2021; Schockel et al., 2015).

Breast cancer is the most common cancer particularly in women contributing 12.5% of the total number of newly diagnosed cancers in 2020 (Sung et al., 2021; Franze et al., 2020; Guo et al., 2019). The incredible inter- and intra- heterogeneity of cancer is attributed to mutated genes, their altered behaviors, and defective cellular pathways of respiration and metabolism (Rogalinska, 2016; Kozakiewicz, et al., 2021; Taanman, 1999; Vyas, Zaganjor and Haigis, 2016; Perciavalle et al., 2012; Gaude and Frezza, 2014; Stroud et al., 2016; Warburg, Wind and Negelein, 1927; Reichert and Neupert, 2004; Wallace, 2012). Based on various scientific studies published to date, it has become much clearer now that cancer cells show variable bioenergetics states, as some cancers rely on aerobic glycolysis while others on OXPHOS, and even metabolic coupling occurs in some aggressive cancers and cause chemoresistance. Otto Warburg has emphasized the occurrence of metabolic reprogramming in cancer cells since 1920 - the Warburg Effect. Though metabolic events are not always oncogenic drivers, it is more likely there is also some role in the activation of other associated cellular pathways that induce oncogenesis. For example, mitochondrial protein p32 is known to maintain high levels of OXPHOS, promote tumorigenesis in breast, thyroid, pancreas, skin, ovarian, prostate and colon cancers, by either acting as a metabolic switch between aerobic glycolysis and OXPHOS or activating oncogenic signalling pathways by stimulating Akt and mTORC proteins and promoting proliferation. The full impact of p32 on mitochondrial activity is still unknown, but it has also been shown to regulate Ca²⁺ concentrations in the mitochondrial matrix, which is coupled with the process of OXPHOS. P32 mitochondrial protein could be a potential therapeutic target considering the antiapoptotic role and as a positive modulator of Akt/mTORC signalling pathway. Based on current evidence, p32 is found to be primarily located in mitochondrial matrix, but it is also a multicompartment protein and found on plasma membrane and nucleus so, therefore its functional role needs further study as on nucleus p32 protein may interact with tumor suppressor protein p53, altering its function, and also increases nuclear localization in cancer cells and on plasma membrane, facilitating cell migration (Yenugonda et al., 2017; Sotgia et al., 2011; Egusquiza-Alvarez, et al., 2021). Similarly, high levels of mitochondrial STAT3 and mitochondrial transcription factor A are also significant to sustain mitochondrial respiration in the process of KRAS induced lung cancer (Sotgia et al., 2011; Pfanner, Warscheid and Wiedemann, 2019). The translocases of the inner membrane (TIM) complexes are among several other protein transport machineries also linked to electron transport transmembrane potential. Based on mitochondrial proteomic studies, multiple protein import pathways could be potential therapeutic targets as they relate to bioenergetics and mitochondrial morphology (Schmidt, Pfanner, and Meisinger, 2010).

Moreover, each cell experiences about 1.5×10^5 oxidative insults per day and mitochondria are very sensitive to internal and external stressors and release several signalling molecules to maintain cell homeostasis. ROS produced in the cell as byproducts of metabolism and mitochondria are the main site of ROS production. Several species of ROS are produced in the cell, however, the most reported are superoxide (O2-•), hydroxyl radical (OH^{*}), nitric oxide (NO), hydrogen peroxide (H_2O_2), and singlet oxygen (1O_2). ROS regulate various activities governed by kinase pathways such as such as mitogen activated protein kinases (MAPKs), Akt to promote cell growth (Yang, Anzo and Cohen, 2005; He et al., 2012; Latimer and Veal, 2016). Cell have naturally conserved mechanisms to balance ROS production, such as its ROS scavenging system through activation of the transcription factor FOXO3a. which induces expression of manganese superoxide dismutase (MnSOD), the chief ROS scavenging enzyme in mitochondria (Brand, 2016; Perillo et al., 2020; Sanz, 2016; Scialo, Fernandez-Ayala and Sanz, 2017b; Barcena, Mayoral and Quiros, 2018). It is very important to understand the role of levels of ROS, as a certain level of ROS is required to maintain normal cellular function. "Mitohormesis" is a cellular protective response where lower levels of ROS are induced to initiate diverse cytosolic and nuclear responses to improve life's expectancy through regulation of metabolism and immune system. Modulation of levels of ROS is very significant in designing ROS therapies used to either prevent or delay many chronic diseases (Ristow and Schmeisser, 2014; Palmeira et al., 2019; Lingappan, 2018; Holmstrom and Finkel, 2014).

The Mitochondrial membrane potential (MMP) is one of the most significant indicators of mitochondrial health and any prolonged fluctuations in MMP may cause bioenergetic stress. MMP, along with proton gradient (ΔpH), develops the transmembrane

potential of hydrogen ions which are used to synthesize ATP (Zorova et al., 2018). Modulation of OMM is not sufficient for the release of cytochrome c protein from the cristae, however, depolarization causes structural remodeling of the inner membrane which leads to the release of cytochrome c from cristae to the intermembrane space (Gottlieb et al., 2003; Sanz, 2016; Xie et al., 2020) (Figure 5).



Figure 5: Illustration of ROS and MMP. Normoxia: is considered normal physiological state where slight increase in ROS levels promotes differentiation and development whereas under conditions of Oxidative stress with increase in ROS levels above celluar threshold limit leads to cell death due to depolarisation of mitochondrial membrane (Created with BioRender).

Impaired OXPHOS is another signal of mitochondrial dysfunction which is critical to produce ATP inside mitochondria. Mitochondria are the major source of ATP in the cell due to the presence of ETC on the IMM. Nicotinamide adenine dinucleotide (NAD) coenzyme is another indicator of mitochondrial health. It is synthesized from the simple amino acid building blocks aspartic acid and tryptophan and exists in two forms; an oxidized (NAD⁺) and a reduced form (NADH) (Blacker et al., 2014; Yang et al., 2007; Srivastava, 2016; Williamson, Lund and Krebs, 1967; Wang et al., 2013). NAD⁺/NADH ratio has significant effect on the redox state of the cell by controlling key enzymes such as glyceraldehyde 3-phosphate dehydrogenase (GAPDH) and pyruvate dehydrogenase (PDH) (Zhang, Wang and Mo, 2018; Stein and Imai, 2012; Pittelli et al., 2011;Pumiputavon et al., 2017). The main role of NAD⁺ in redox regulation is the reduction to NADH, as part of beta oxidation, glycolysis, and the citric acid cycle (Hopp, Gruter and Hottiger, 2019;). Emerging

roles of NAD⁺ as an extracellular signalling molecule is also a significant discovery, although full impact on health and disease is still not clear, but it has been known to enhance glycolysis (Srivastava, 2016; Williamson, Lund and Kreb, 1967). Boosting NAD⁺ through dietary supplementation improves range of physiological and pathological conditions (Xie, et al., 2020; Kotnik et al., 2017; Kuriyama and Maeda, 2019).

Intracellular Ca^{2+} concentration is another indicator of mitochondrial health. Though the ER is one of the main storage sites of Ca^{2+} , other organelles such as mitochondria, Golgi apparatus, lysosomes and peroxisomes also participate in Ca^{2+} signalling. How much Ca^{2+} homeostasis is crucial for the cellular system is yet to be explored, however Ca^{2+} overload results in oxidative stress and several pathological conditions, for example cardimyopathies (Shimizu et al., 2015; Marchi and Pinton, 2014; Carafoli, 2003; Peitzsch et al., 2017; Arnold et al., 2020).

There is exceptional potential of mitochondria as a therapeutic target for several diseases and various types of cancers and a recent focus of research is to target these intriguing organelles, however due to some discouraging findings that indicates variability of tumours for example multiform of glioblastoma (Reviewed by Hanif) and inter and intra tumor heterogeneity there is a need for more intensive research (Hanif et al., 2017). Even the same tumors differ in type of mutations from individual patients indicating tumors are unlikely to be treated by targeting a single pathway, gene or signalling molecule (Fisher, Pusztai and Swanton, 2013). So, a current trend is to design anti-cancer drugs that can specifically destabilise mitochondria named as "Mitocans", which will be explained later in detail (Section 1.6).

Moreover, the body's immune system is trained enough to recognize cancer cells but not cancer stem cells (CSCs) that are reproduced as normal, sustain harmful effects, cause relapse after treatment and result in failure of many immunotherapies (The Ludwig Center, 2021; De Francesco, Sotgia, and Lisanti, 2018; Cui, Wen and Huang, 2017; Kalyana-raman et al., 2018; Bobrovnikova-Marjon and Hurov, 2014; Falkowska et al., 2015).

1.6 Mitocans

After several years of research, now there is unprecedented focus on the discovery of more efficient drug candidates and invariant, clinically relevant therapeutic targets, due to drug resistance, side effects of existing therapies, disease relapse and the differences in the patterns of even the same disease in different individuals. Developing mitocans is one such attempt in progress to develop anticancer agents that can specifically target certain genes, cellular pathways of complex diseases, including cancers, because aberrant mitochondrial metabolism in cancer also makes them more attractive therapeutic targets (Ralph et al., 2006; Neuzil et al., 2013; Spivak et al., 2021). Mitocans produce effects by destabilizing and altering the functions of mitochondria, such as ETC blockers (Figure 6) (Ralph et al., 2006; Rohlena et al., 2011; Neuzil et al., 2007; Dong et al., 2007; Chen et al., 2009; Guzzo et al., 2014). Scientists are expecting these mitocans will have fewer side effects on normal healthy cells. As the process of tumorigenesis and development are regulated by mitochondria, they could also offer a general approach for the effective treatment because of their functional role in most of the diseases like cancer, inflammation and aging (Baggetto and Testa-Parussini, 1990; Bernal et al., 1983; Ferrarini et al., 2021; Dong et al., 2011; Rohlenova et al., 2017; Neuzil et al., 2013; Ashkenazi et al., 2017; Ghoneum and Said, 2019; Du et al., 2019; Dong and Neuzil, 2019). Some of the main classes of mitocans are hexokinase inhibitors, ETC inhibitors, BcL-2 family of proteins inhibitors, inhibitors of VDAC/ANT channels, and lipophilic cations which target inner membrane, the citric acid cycle and mtDNA (Figure 6). Some examples of anticancer drugs showing effects through mitochondria are already in clinical trials and are listed in Table 3.

Mitochondrial	Mechanism of action/Tumor type	Clinical	References
function		trial	
ETC	Papuamine inhibits ATP production /lung	NA	Min et al., 2020
	Metformin inhibits Complex I/ Colon, Ovary,	СТ	Wheaton et al., 2014
	Breast,		
	Prostate		
	Tamoxifen inhibits Complex I/ Breast	FA	Daurio et al., 2016
	Mito-Tam inhibits Complex I/ Breast	СТ	Bryant et al., 2017
	α-TOS inhibits Complex II/ Breast	PC	Dong et al., 2008
	MitoVES inhibits Complex II/ Breast	PC	Dong et al., 2011
	VLX600 inhibits Complex IV/ Colon	PC	Zhang et al., 2014
	Tigecycline inhibits Complex I	FA	Skrtic, et al., 2011
	Inhibits Complex IV/ Leukemia		
	Gamitrinib inhibits ATPase activity/ Prostate	PC	Chae et al., 2012
TCA Cycle	AGI-5198 inhibits IDHs activity/	CT	Golub et al., 2019
	Glioblastoma		
	Dichloroacetate inhibits IDHs activity/ Brain	СТ	Dunbar et al., 2014
Glycolysis &	2-deoxyglucose (2-DG) competitor for	СТ	Xu, et al., 2018
OXPHOS	binding hexokinase /Lung,		
	Prostate,Ovary,Breast		
	Metformin/2DG inhibits ATP production/	СТ	Cheng et al., 2012
	Lung, pancreas		
	ABT737/2DG inhibits OXPHOS/ Ovary	NA	Xu et al., 2018
Signalling	Venetoclax Bcl-x _L inhibitor/ Leukemia,	FA	Aghvami et al., 2018
pathways	Lymphoma		
	Navitoclax Bcl-Xl/Bcl2 inhibitor/ Breast,	СТ	Cui, Wen, and
	Lung		Huang, 2017
	Prostate, Colon		
	ECPU-0001 Bcl2 inhibitor/ Lung	PC	Mongre et al., 2019
	Gossypol LDHA inhibitor,	СТ	Missiroli et al.,2020
	NADH competitor/ Breast, Brain, Prostate		

Table 3: Anti-cancer drugs and their mechanism of action on mitochondrial function.

Abbreviations: ETC (electron transport chain); TCA Cycle (tricarboxylic acid cycle); OXPHOS (Oxidative phosphorylation); IDH (isocitrate dehydrogenase); ATP (adenosine triphosphate); LDHA (lactate dehydrogenase); NADH (nicotinamide adenine dinucleotide) (Dong et al., 2020). Abbreviation: CT(Clinial trial), PC(Preclinical),FA (FDA Approved)



Figure 6: Hallmarks of cancer and their inhibitors: Original hallmarks (self-sufficiency in growth signals, insensitivity to anti-growth signals, tissue invasion and metastasis, limitless replicative potential, sustained angiogenesis (blood vessel growth) and evasion of apoptosis), enabling factors (Inflammation and instability genes), emerging hallmarks (Reprogramming metabolism and immune evasion) (Created with Biorender).

1.7 Mitochondria and Phytotherapy

The tradition and knowledge of drug development from natural sources, particularly plants based, is centuries old, for example morphine (*Papaver somniferum* L.) and digoxin (*Digitalis purpurea* L./ (Wachtel-Galor and Benzie, 2011). Plants, either used for medicinal purposes or as functional foods, have provided several classes of bioactive compounds (micro molecules and macro molecules), including phenolics, alkaloids, flavonoids, which are rich in valuable bioactivities, such as antioxidant, anti-inflammatory, anticancer, antibacterial, or antiviral activities (Koo, Song and Bae, 2018; Rahman et al., 2011; Shati et al., 2020). However, the exact pathways through which phyto-active compounds produce their effects are still not clear, due to the diverse nature of most of these bioactive compounds. How-
ever, some compounds show characteristic and distinctive mechanisms, such as the sulforaphane effect through cancer stem cell self-renewal pathway (Liu, Min and Bao, 2009), curcumin targeting a plethora of signalling pathways in pancreatic cancer (Glienke et al., 2009; Schutte et al., 1998), capsaicin homovanillic acid derivative targeting Trx-ASK1 signalling and mitochondrial transport chain during apoptosis (Pramanik and Srivastava, 2012), epigallocatechin-3-gallate stimulating mitochondrial membrane depolarization (Qanungo et al., 2005; Urra et al., 2013; Li et al., 2017), resveratrol inhibiting proliferation through hedgehog signalling pathway (Mo et al., 2011), polymethoxylated flavones inducing Ca⁺² mediated apoptosis, magnolol inhibiting EGFR/PI3K/Akt signalling pathway (Lee, Park and Roh, 2019), lectins targeting programmed cell death pathways (Fu et al., 2011), ascorbic acid affects bone cancer, chlorbenzoyl berbamine (natural derivative of berbamine) modulating NF-kB and JNK signalling pathways (Yun et al., 2016;Zhu et al., 2020; Wang et al., 2015; Plitzko, Kaweesa and Loesgen, 2017). Considering the above evidence and the association of mitochondria with different diseases and signalling pathways, mitochondria prove to be a promising model for testing of biological activities of new bioactive compounds (Table 4) (Yu et al., 2022; Liu et al., 2019; Sinha et al., 2019; Fernandes et al., 2017; Arauna et al., 2019; Cho et al., 2019).

Table 4:	Natural	Compounds	Affecting	Mitochondrial	Functions
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Compounds	Cell line	Mechanism	Reference
Lupeol (Bombax ceiba L.)	Renal cell carcinoma SK-RC-45 cells	Altered mitochondrial dynamics Increased fission Increased apoptosis	Sinha et al., 2019 Zhang et al., 2015 Zhang et al., 2017
Fucoidan (Marine brown algae)	MCF-7, MDA-MB-231, HeLa	Increased DNA fragmentation Decreased MMP Increased apoptosis	Zhang et al., 2011 Ayman, May, Mohammad, 2021
Ganoleuconin (<i>Ganoderma leucocontextum T.</i>)	Liver cancer cell line Huh7.5	Decreased ATP production Increased apoptosis Decreased MMP	Sinha et al., 2019 He et al., 2016
Betulinic acid (Betula pendula R.)	HeLa cells	Increased ROS, Decreased MMP	Xu, et al., 2017
Genipin (Gardenia jasminoides J.)	Lipid bilayer membranes	Decreased uncoupling protein 2 (UCP2)	Kreiter, et al., 2019
Alternol (Taxus baccata L.)	PC-3 prostate cancer cells	Decreased mitochondrial respiration, Decreased isocitric acid, fumaric acid, malic acid	Li et al., 2019 Liu et al., 2007 Zhang et al., 2018 Chan, et al., 2015
Dihydromyricetin (Ampelopsis grossedentata/ Nekemias grossed- entata (HandMazz.) J.Wen & Z.)	HepG2 cells	Increased Bax and Bad Decreased apoptosis	Zhang, et al., 2017
Cathachunine (Catharanthus roseus L.)	HL60 cells	Increased ROS mediated intrinsic apoptosis Decreased MMP	Wang et al., 2016
Dentatin (<i>Clausena excavate</i> B.)	HepG2 cells	Increased cytoplasmic cytochrome <i>c</i> Increased Bax Decreased Bcl-2	Gong et al., 2017
Plumbagin (Plumbago zeylanica L.)	MG63	Increased ROS Altered protein levels of Bcl-2, Bax, Bcl-xL, Bak	Chao et al., 2017
Aellinane (Euphorbia aellenii R.)	Ovarian cancer cell line	Increased apoptosis via mitochondrial pathway	Nabatchian et al., 2017

1.8 Scope of Complementary Medicine in Pakistan

Pakistan has a rich tradition of using medicinal plants for the treatment of various health related issues including cancer and rheumatism (Kanwal and Sherazi, 2017; Welz, Emberger-Klein and Menrad, 2018). Along with advanced medical treatments, traditional medicine is still widely practiced in Pakistan as complementary and alternative medicine (CAM), mainly due to limited health care facilities and poverty. Ayurvedic and homeopathic system of medicines have been accepted as a part of national health care system of Pakistan.

Pakistan holds a great diversity of flora due to its distinct geographical position and diverse climatic conditions. More than 6,000 vascular plant species and 5,600 species having 22 families and 150 genera are described in the flora of Pakistan (Stewart, 1972; Ali and Qaiser, 1986; Mallon, 1991). Over 1,000 species have been evaluated for possible phytochemicals and around 350-400 species are traded and used by the manufacturers of Unani and Homeopathic medicines. There is very high demand of exploring the medicinal plants of Pakistan as there are around 40,000-50,000 practitioners of Greco-Arabic medicine, Ayurvedic and folk medicine locally known as Tabibs and Vaids in addition to unregistered practitioners of rural areas utilize more than 200 plants in the preparation of their medicinal recipes. Though there are some challenges that need consideration, such as over exploitation of resources due to unsustainable use of medicinal plants by local people and herbal medicine manufacturers, scientists (pharmacologist/biochemist/botanist) and pharmaceutical companies, who process large quantities of medicinal plants to obtain small quantities of biologically active compounds (Shinwari and Shah, 2007; Shinwari and Shinwari, 2010; Ekor, 2014).

1.9 Alhagi maurorum Medik: Multipurpose Herb

Alhagi maurorum M. (Fabaceae) is a multipurpose herb commonly used in the Greco-Roman and the Unani system of medicines (Measer, 2019; Xu et al., 2021). Folklore claims about the use of Alhagi species are reported in various countries such as Egypt (gastrointestinal track - GIT, liver and UTI disorders), India (laxative, diuretic, asthma, rheumatism, fever), Iran (kidney stones, laxative, cardiac pains, diuretic), China (rheumatism, cancer), Palestine (urinary tract disorders), Turkey (tonic), Jordan (kidney stones), Afghanistan (diarrhea, jaundice, skin

wounds, appetizer), Pakistan (eyesight improvement, antirheumatic, stomachache), Qatar (migraine, rheumatism, jaundice), other South Asian countries (hemorrhoids, GIT), Saudi Arabia (analgesic, antioxidant, diuretic, cough, angina, antipyretic, anti-tussive, dysenteries, gastritis), Uzbekistan (eczema) (reviewed by Tavassoli et al., 2020). However, it is mainly explored due to its antioxidant potential and nutritive value (Sarkar et al., 2020; Davinelli et al., 2013). Alhagi *maurorum* is a rich source of biologically active compounds, including phenolics and alkaloids (Samejo et al., 2012; Loizzo et al., 2014; The Plant List, 2010; EOL, Encyclopedia of the life, 2014). Almost all parts of *Alhagi maurorum* plant are used in traditional system of medicine. Its phytochemistry and its use in Greco-Roman and Unani system of medicine make it a promising medicinal plant. It is marketed under the name of "Toranjabin" (Atta and Abo, 2004; Measer., 2019; Ara et al., 2020; Aslam et al., 2016). Ardekani reported in his study about temperament of this plant in Iranian traditional medicinal system as hot and dry and suggested it is due to the presence of its constituent alkaloidal amines (Parvinroo et al., 2014; Ardekani, Rahimi, and Mohammad, 2010; Atta et al., 2010). Furthermore, previously published research studies have ascribed therapeutic properties of anti-rheumatism and nutritive value to alkaloids present in plant extract, for example phenethylamine (tyramine, hordenine, β - phenethylamine (β-PEA), N-methyl mescaline (Irsfeld, Spadafore and Prub, 2013:Rafehi et al., 2021), isoquinoline alkaloid (salsolidine) (Tundis et al., 2012) and pyrrole (pyrrolizidine, pyrrolezanthine) alkaloids (Guan et al., 2020; Thu et al., 2020). Some pharmacological evaluation of different extracts prepared in different solvents have already been reported as cytotoxic effects by Sulaiman using leaves and flowers extract on the human leukaemia cell line (IC₅₀ 16.0-22.0 µg/ml) (Asghari et al., 2016; Sulaiman et al., 2013). Its gastroprotective effects in rabbit using aqueous ethanolic extract and aqueous extract, anti-inflammatory effect, antipyretic effect using ethanolic extract of root, hepatoprotective effect using ethanolic extract of aerial parts in albino rats, and antifungal effect using methanolic extract have already been investigated (Sheweita et al., 2016; Neamah, 2012; Alqasoumi et al., 2008; Ara et al., 2020; Aslam et al., 2016; Ahmad et al., 2010).

Alhagi maurorum M. (Camelthorn)



Figure 7: *Alhagi maurorum's* commercially available samples, folklore claims in different regions of the world and different form of traditional preparations used to achieve traditional medicinal effects in different traditional system of medicine mainly Unani and Greco- Roman (Alchetron, 2018; Asghari and Tewari, 2007) (Created with Biorender).

In addition to pharmacological evaluation, chemical investigation using fresh stems of *Alhagi maurorum* by Ghosal and Srivastava showed the presence of mixture of alkaloids such as tyramine derivative, isoquinoline and pyrrole alkaloids in addition to other chemical compounds (Ghosal and Srivastava, 1973; Asghari and Tewari, 2007) (Table 5). Most of the alkaloids are under investigation but very limited data is available about their pharmacological effects, pyrrolizidine alkaloids have been investigated for their anti-inflammatory effects (Wei, Ruan and Vrieling, 2021; Hoang et al., 2015; Kaltner, 2020; Kazemi, Fatemi, and Wink, 2018), salsolidine has a potential role in the treatment of Alzheimer's disease (Tundis et al., 2009), tyramine and it's derivatives acts as catecholamine releasing agent and shares pharmacological pharmacokinetics in humans (Rafehi et al., 2019), N-methyl mescaline has been investigated as a pharmacological chaperone design for reducing risk factor of Parkinson's disease (dysfunction of glucocerebrosidase enzyme that result in mitochondrial dysfunction) to treat misfolded protein (Huang, 2014) and β - phenethylamine (β -PEA) acts as a CNS stimulant in humans (Irsfeld et al., 2013) and hordenine consumed as supplement.

Alhagi maurorum is considered as a cheaper and ever-available source for lupeol (a bioactive triterpenoid) which is also found in root bark. *Alhagi* honey contain sugar exudates polysaccharides (galactose, uronic acid) (Gonchaov, 2001), epigeal parts of plants contain catechins, proanthocyanidins (Islambekov et al., 1982) whereas whole plant extract reported to contain, octadecane, eicosane; di-, tri- and tetra-cosane, triterpene and derivatives (Ghosal and Srivastava, 1973; Ghosal and Srivastava, 2016), alhacidin and alhacin (Muhammad et al., 2015). There are some studies which have reported alhagifoline A, pseudalhagin, alhacin, alhacidin (Al-Snafi, 2015; Li et al., 2010), lupeol (Laghari et al., 2011) in addition to steroids, alkaloids from the aerial parts of this plant (Sultan et al., 2011; Xiuwei et al., 1996; Guijie et al., 2010 and Laghari et al., 2012b) and showed presence of twelve different compounds of new iso flavonolignan and flavanol, alhagitin and alhagidin from the root bark. Furthermore, some studies have been published about the contents of the seed oils of plant *Alhagi maurorum* and found presence of saturated and unsaturated fatty acids and some metals such as Cu, Al, Mn (Hashem and Alfarhan, 1993; Singh et al., 1999) (Table 5).

So far, quantitative data of the most of the compounds of Alhagi spp. is not available, though some studies have reported presence of 4.22 mg/g of phenolic acid (200 g root powder), 8.36 ± 1.33 of alkaloids The n-butanol fraction exhibited the highest total phenolic content ($20.78 \pm 0.63 \mu g$ gallic acid eq./mg), while the ethyl acetate fraction exhibited the highest total flavonoids ($14.41 \pm 0.03 \mu g$ quercetin eq./mg) and tannins ($9.44 \pm 0.36 \mu g$) (Mostafa et al., 2019)

Biological System	Pharmacological Effects	Extract	References
Gastrointestinal	Curing stomach, intestinal complaints, liver ailments (Hepatoprotective)	Ethanolic	Alqasoumi, 2007
	Anti-inflammatory (reduces the thickness of paw edema)	Aqueous	Neamah, 2012
	Anti-diarrheal (Increased contraction duode- nal smooth muscles (low conc.), sedative ef- fect (high conc.)	Methanolic	Gutierrez et al., 2007
	Anti-ulcerogenic effect (Decreased acid out- put)	Ethanolic	Shaker, Mahmoud, Mnaa, 2010
Systemic	Antioxidant effect (Decreased malondialde- hyde levels) Antioxidant effect (scavenging free radicals)	Aqueous Alcoholic	Neamah, 2012 Awaad et al., 2011
	Anti-tumour (increase immune activity)	Extract	Laghari et al., 2012
Urinary tract	Enlargement of ureter to expel stones	Ethanolic	Marashdah, and Al- Hazimi, 2010.
	Diuretic, reduction of pH and crystalluria	Extract	Mirdeilami, et al., 2011
Antimicrobial	Antibacterial activity	Methanolic	Bonjar, 2004
	Antifungal activity	Ethanolic	Al-Askar, 2012
Blood	Decrease bilirubin levels, serum creatinine phosphate level and lipid per oxidation	Extract	Nabavizadeh and Nabavi, 2010 Khushbaktova et al., 1992
CNS	Antinociceptive (protection against writhing)	Ethanolic	Awaad et al., 2011
	Antinociceptive (traditional use)	Methanolic	Hudaib et al., 2008
Genome	Genotoxicity (causes DNA damage at a conc. Of 5µg/ml)	Extract	Etebari et al., 2012

Table 5: Alhagi maurorum: Pharmacological effects

1.10 Phenethylamine Alkaloid: Hordenine

There is now enough evidence available to support that alkaloids have had a significant role in medicine since ancient times and remained a source of medicines. For example, psilocin affecting perception, caffeine stimulating CNS, morphine and cocaine relieving pain (Avois et al., 2006), cephalosporin, and penicillin being used as antibiotics (Cushnie, and Lamb, 2014), quinine as antimalarial drug (Achan et al., 2011). Phenethylamine (PEA) is a natural monoamine alkaloid and trace amine marketed as dietary supplement mainly to increase cognitive functions and to reduce weight (Andersen and Fogh, 2001; Galitzky et al., 1988; Seneca, 2007; Kaltner et al., 2020). Hordenine is an organic compound synthesized from tyrosine (Figure 9) (for detailed steps of synthesis see Figure 10), also known as eremursine, N,N-dimethyltyramine, peyocactine, p-hydroxy-N,N-dimethylphenethylamine, 4-[2-(Dimethylamino) ethyl] phenol and occurs naturally in wide range of plants such as Ariocarpus fissuratus E., Hordeum vulgare L., (Ghosal and Srivastava et al., 1972; Schultes, 1937), Panicum miliaceum L. (Brady and Tyler, 1958), Lophophora williamsii L. (Rao, 1970), Schinus terebinthifolia R., Citus aurantium F. (Mahmoud et al., 2021; Smith, 1977; Weaton and Stewart, 1970; Kim, Lee et al., 2013; Reilly, 1981) and Anahalonium fissuratum E (Spath and Roder, 1922; Spath, 1919; Kapadia and Fayez, 1970; Lima et al., 2015). It has taken its name from Hordeum vulgare (Barley) member of grass family which is one of the major cereal grains cultivated 10,000 years ago (Guo et al., 2018; Roman, Betz and Hildreth, 2007; Xiong, Li, and Jin, 2014). *Tamarindus indica* L. is traditionally used as an antiasthmatic agent and is reported to be a promising anti-diabetic drug containing hordenine in its leaves, bark and even flowers (Saxena et al., 2022). So, possibly diuretic, psychostimulatory, and anti-obesity activities of Tamarindus indica are due to hordenine (Ghosh and Das, 2010). In addition to the above-mentioned plants, hordenine is also one of the interesting alkaloids reported from *Cannabis sativa* L. (Atakan, 2012). Hordenine is structurally related to ephedrine, a central nervous system stimulant, has adrenergic-like actions and releases neurotransmitters epinephrine (adrenaline) and norepinephrine (noradrenaline) to increase energy and focus. Hordenine is also a vital ingredient of nutritional supplements marketed with the claim as CNS stimulant, nootropic agents (increase energy level), it lifts the mood, promotes weight loss, and increases athletic performance. Some examples of these supplements are synephrine, higenamine and hordenine, which are marketed as the best pre-workout stimulants, improve ability to use fat for fuel and promote mental arousal. Some examples of supplements available in the market are shown in Figure 8 (Efimova, et al., 2020; Lovett, Hoult, and Christen, 1994; Li et al., 2020).



Figure 8: Commercially available dietary supplements containing hordenine.

Dietary supplements are easily available in pharmacies, food stores and groceries without any prescription, so, they are used as complementary therapy globally (Kotnik et al., 2017; Kuriyama and Maeda, 2019; Anders and Schroeter, 2017; Li et al., 2017; Liu, Ye and Guo, 2020). However, like all other medicines, dietary supplements can also cause allergic reactions, adverse effects and interact with prescription medicines (Dwyer, Coates and Smith, 2018; Omar et al., 2019; NCCIHT, 2021) therefore these should also be consumed based on scientific evidence (Brady and Tyler, 1958; Webmed, 2022).



Figure 9: Structure of hordenine



Figure 10: Biosynthesis of hordenine in three steps: Hordenine is produced in plants from tyramine in a stepwise methylation process (Leete and Marion, 1954; Stribney and Kirkwood, 1954; Thomas, 1963; Massicot and Marion, 1957; Brady and Tyler, 1958. In 1937 Raoul proposed based on the structural similarity of hordenine and tyrosine following series of three step reaction of biosynthesis of hordenine: (I) L-Tyrosine decarboxylated to tyramine (II) Methylation of tyramine to N-methyltyramine (III) Methylation of N-methyltyramine to hordenine (Raoul, 1937; Frank and Marion, 1956; Liu and Lovett, 1993a-1993b).

Additionally, not only the availability of hordenine from different sources but also the pharmacological features and physicochemical properties make it a potential candidate for research as it is a low molecular weight compound with amphoteric in nature, selectively deaminated by MAO-B, crosses the blood brain barrier, and has short half-life of 1-2hr (Barwell, 1989; Lebecque et al., 2018).

Pharmacokinetics studies of hordenine have shown bioavailability of 66.2% after oral administration in a rat model and in humans after one hour of beer consumption maximum plasma levels of 12.0-17.3 nM were achieved (Ma et al., 2015; Sommer et al., 2020). In 2020, Sobiech designed and utilized magnetic core-shell molecularly imprinted

sorbent to study pharmacokinetics of the hordenine present in human plasma after consuming dietary supplements and noted highest concentration of $16.4 \pm 7.8 \ \mu g/l \ (65 \pm 14 \ min)$, $t_{1/2} 54 \pm 19 \ min$ and V_d (volume of distribution) $6000 \pm 2600 \ (66 \pm 24 \ l/kg)$ (Sobiech et al., 2020).

In vitro and in vivo studies of hordenine

In vitro: Hordenine and Cellular Processes

1) Mitogen Activated Kinase (MAPK) Signalling Pathway

As the prevalence of pituitary tumors in general population is very high and prolactinomas are the most recurrent pituitary tumors (Daly et al., 2009; Ciccarelli et al., 2005). To restore pituitary functions therapeutic aim is to lower excessive serum prolactin (PRL) levels and the size of the tumor. Dopamine under physiological conditions is responsible for inhibiting prolactin production. Dopamine D2 receptors are expressed on the membranes of lactotroph cells (Ben-Jonathan and Hnasko, 2001; Wong et al., 2015). Wang et al., (2020) investigated the effect of hordenine extracted from traditional Chinese herb Hordeum vulgare (Wang et al., 2014) on prolactinomas. Wang et al., (2020) investigated the antiprolactinoma effect of hordenine in a rat model and showed decreased level of the p38, ERK1/2, and JNK protein expression and the production of inflammatory cytokines TNF- α , IL-1 β , and IL-6 possibly through MAPK signalling pathway. Hordenine significantly reduced the size of pituitary tumors in rats and restored serum prolactin levels, and he proposed hordenine inhibited the production of PRL by regulating MAPKs signalling (Lu, Wang and Zhan, 2019; Weichert et al., 2015; Hu et al., 2017). MAPK is a key signalling pathway involved in several physiological and pathological functions by activating three sequentially activated protein kinases (See Figure 11 for detailed process) (Zhan, 2019).



Figure 11: Hordenine modulation of MAPK pathway. Extracellular signal-regulate kinase (ERK1/2), Jun kinase (JNK), MAPK. Each MAPK pathway is initiated by diverse extracellular as well as intracellular stressors such as oxidative stress, ER stress, reductive stress, hormones, and growth factors. Stress response is a very broad term encompassing several molecular mechanisms that are initiated depending on the nature, strength, duration of the stress and cell types. Protein phosphorylation is the most rapid and effective response initiated against stress among many others. (Created with BioRender).

Wang et al., in 2014 further showed hordenine has inhibited PRL secretion through DRD2 agonist which further regulate cyclic adenosine monophosphate (cAMP)/PKA/CREB pathways in two cells; high expression of estrogen induced rat pituitary tumor MMQ cells and low expression of radiation induced rat pituitary tumor GH3 cell. However, this study also investigated total barley maiya alkaloids and proposed synergistic phenomena responsible for efficient way of reducing PRL levels (Figure 12) (Sussman et al., 2020; Gong, Tao, Wang, et al., 2021; Recouvreux et al., 2016; Yang et al., 2020; Gao et al., 2017).



Figure 12: Hordenine and synergistic effect. cAMP Response Element Binding Protein (CREB), Dopamine D2 Receptor (DRD2), Mesenger RNA (mRNA), PKA (Protein Kinase Pathway), PRL (Prolactin), Protein Kinase A (PKA), Total Barley Maiya Alkaloid (TBMA) (Wang et al., 2014; Lu et al., 2019)

2) Pyruvate Dehydrogenase Kinase (PDK)

Protein kinases are important drug targets, governing many cellular pathways in various diseases, including cancers and metabolic disorders. Nuclear-encoded mitochondrial multienzyme complex the pyruvate dehydrogenase (PDH) complex provides primary link between glycolysis and TCA cycle. The enzymatic activity of PDH involves phosphorylation and dephosphorylation cycles, regulated by products and substrates of the reaction inside mitochondria (Gudi, et al., 1995; Xu et al., 2019). The phosphorylation of PDH results in inactivation of its enzymatic activity completely due to highly specific pyruvate PDK. Pyruvate Dehydrogenase Kinase 3 (PDK3) plays a pivotal role in metabolic switching during progression of cancer. Anwar, et al. (2020) investigated effects of hordenine in lung cancer cell lines and suggested, hordenine is a potential inhibitor of PDK3 gene that in chemoresistance drives glycolysis by forming a positive feedback loop with HSF1. So, targeting this mitonuclear communication may represent a novel approach to overcome chemoresistance (Figure 13) (Anwar et al., 2021; Xu et al., 2021).



Figure 13: Hordenine binds and inhibits PDK3 gene and eventually activity of Pyruvate Dehydrogenase (PDH) is altered that offers metabolic switch between glycolysis and TCA cycle. Heat Shock Transcription Factor 1(HSF protein coding gene highly conserved and regulate transcriptional responses mainly non stress related such as metabolism and development), Glycogen Synthase Kinase-3 beta (GSK3b), Ubiquitin (Ub), Tricarboxylic Acid (TCA) (Anwar et al., 2021; Xu et al., 2021) (created with Biorender).

3) Nuclear Factor Erythroid 2–related Factor 2 (Nrf2)

The prevalence of diabetes and its associated complications is growing globally (Harding, et al., 2019). Diabetes is a disorder of glucose metabolism and may result in multiple organ failure. There is need for novel agents which either enhance or complement the actions of insulin (Wang et al., 2017). Shuhao et al., (2018) investigated the effect of hordenine alone and in combination with insulin. They observed significant effects in reduction of fasting and postprandial blood glucose levels. In addition to this, hordenine has shown significant renal protective activity (diabetic nephropathy) in combination with insulin. Authors suggested based on evidence the hordenine/insulin combination produced effects through inhibition of Nrf2 expression (Zhao et al., 2018; Su et al., 2018). Under unstressed conditions low levels of Nrf2 offer basal expression of its targeted genes while under stressed conditions cysteine modified KEAP1 (Kelch-like ECH-associated protein 1 (KEAP1) is unable to degrade Nrf2 and higher levels of antioxidant and pro-oxidant genes are produced (Figure 14).

Renal Protective Activity of Hordenine



Figure 14: Renal protective activity of hordenine: Hordenine protects kidney through Nrf2 essential transcription factor and regulate multiple homeostatic functions such as inflammation, redox metabolism, and proteostasis. Nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB) (Created with BioRender).

4) Cyclic Adenosine Monophosphate (cAMP)

Melanin protects human skin by absorbing free radicals and shielding from ultraviolet light. However, abnormal levels of melanin result in skin disorders. Many studies have shown cAMP-mediated pathways are the major signaling pathways involved in this process (Bos, 2006). cAMP is a second messenger used for intracellular signal transduction and is associated with the function of kinases in many biochemical processes such as sugar regulation and lipid metabolism. Activation of associated genes is linked to growth of cancer (Ali et al., 2016). Hordenine is reported to be produced as secondary metabolite during germination stage in barley so when germinated barley extract and ungerminated barley extracts were used in study, germinated barley extract exhibited potent inhibitory effect. Further studies confirmed hordenine involved in pigmenting activity through cAMP pathway and altered CREB (cAMP response element binding protein) phosphorylation as well as microphthalmia-associated transcription factor (MITF) associated genes expression (Bertolotto et al., 1998; Busca and Ballotti, 2000; Kim et al., 2013) (Detailed pathway see Figure 15).



Figure 15: Hordenine and cAMP. Microphthalmia-associated transcription factor (MITF), CREB (cAMP response element binding protein) (created with Biorender).

5) Akt Signaling and Nuclear Factor κB (NF-κB)

Clinical syndromes of acute respiratory failure increase morbidity as well as mortality. If a patient survives an acute lung injury their quality of life remains affected. Prevalence of acute lung injury and acute respiratory distress is so common that there is need to find new drugs (Johnson and Matthay, 2010). In a recent study, the effect of hordenine is investigated in acute lung injury induced by lipopolysaccharide. Protein kinase B (Akt) involves in various important cellular processes such as cell proliferation, apoptosis, metabolism of glucose and a complex protein NF-κB which leads to cytokine production, survival of cell and even DNA transcription. Hordenine has shown effect on the activation of both inflammatory factors. It also has been shown to affect the MAPK signalling, which is associated with heat shock, pro-inflammatory cytokines. Zhang et al., (2021) recently reported the effectiveness of hordenine treatment of acute lung injury both *in vitro* and *in vivo*.

6) Dopaminergic Pathway

Dopaminergic pathways are involved in hedonic food intake and can lead to overconsumption. Dopamine D2 receptor is particularly involved in food reward and ultimately increased consumption (Hoch, Kreitz, et al., 2015; Baik, J.H., 2013). Sommer, Hubner, Kerdawy, et al., (2017) observed hordenine caused D2R-promoted G-protein activation (Shonberg, Lopez et al., 2014). It also has shown functional selectivity by antagonized D2-

mediated β -arrestin recruitment. So, this study has associated mood elevating effects of beer to hordenine a component of beer though its concentration in beer depends on malting process (Brauers, Steiner and Daldrup, 2013). Hordenine is identified as food-derived dopamine receptor agonist (Sommer et al., 2020).

7) Hordenine Activating Serotonin Receptor

Serotonin (5-hydroxytryptamine) neurotransmitter is involved in complex, multifaceted biological functions. In humans it has diverse effects on mood, anxiety, sleep, appetite, GIT motility, eating, temperature (Beattie and Smith, 2008; Rosen, 2009; Collet et al., 2008). In humans 95% of 5-HT receptors are present in GIT which promote transmittance of signals from the gut lumen to nerve cells and smooth muscles. So, many studies have reported it regulates digestive functions, bone mass and organ development too. Hordenine (antibiofilm) activates 5 HTR 4b in human colon epithelial cells which is associated with wound healing and increased cell motility (Yasi et al., 2019).

8) Hordenine Selectivity for Monoamine Oxidase (MAO)

Monoamine oxidases (MAO) catalyze oxidation of amines and are present in the outer membrane of mitochondria in various cell types. Neurotransmitters are inactivated by MAOs. Normal levels of MAOs have clinical significance as imbalance is linked to many diseases such as depression, attention deficit disorder, schizophrenia, migraines (Domino and Khanna, 1976; Meyer et al., 2006; Filic et al., 2005). In a study deamination of hordenine was conducted in rats which showed its high selectivity for MAO-B in liver (Barwell et al., 1989; Fagervall and Ross, 1986) and is not deaminated by intestinal MAO-A. So, dietary hordenine is unlikely to be deaminated by intestinal MAO-A more likely will be absorbed and affect the sympathetic nervous system.

9) Hordenine Antibiofilm Agent

Hordenine extracted from sprouting barley has shown quorum sensing inhibitory activity against *Pseudomonas aeruginosa* (food borne pathogen) at lower concentrations. It also has inhibited acyl-homoserine lactones levels which is an important intercellular signaling molecule through which bacteria monitor density of their population. It is synthesized by members of Luxl family of proteins. Treatment with hordenine decreased the quorum sensing-related extracellular factors and genes (lasI, lasR, rhlI) (Zhou, et al., 2018; Zohary and Hop, 2000).

In vivo studies

Hordenine remained the focus of research over a century ago, Hefter in 1894, found hordenine induced CNS paralysis in frogs without affecting blood circulation at lower doses however, with increased doses caused hypertension and eventually induced death. Rietschel, (1937), proposed the cause of through stimulation of the heart muscle (Rietschel, 1937; Jensen et al., 2021). Schweitzer and Wright, in 1938, studied derivatives of hordenine and reported dimethylcarbamic ester of hordenine hydrochloride, (D.-C.-Hord. -HCI) and dimethylcarbamic ester of hordenine methiodide, (D.-C.-Hord. -MCI) have similar anticholinesterase activity, group of drugs used to neurodegenerative diseases for example, Alzhemier disease and Parkinson diseases (Kapadia and Fayez, 1970). Frank et al., observed respiratory distress, increased heart rate and flehman response in horses after administration of hordenine (Frank et al., 1990; Hart, 1983). Overall, these studies highlighted the significance of hordenine for the treatment of CNS disorders and called for further study.

Hordenine Supplements

Thermogenic supplements are used by adults to boost energy and control body fat without scientific evidence (Hoffman et al., 2009). Many natural compounds are added to these supplements, for example, caffeine in combination with ephedra produces thermogenic effects targeting lipolytic metabolic pathway in fat adipocytes and reduces depression with addition of phenethylamine (Bell et al., 2004; Pittler and Ernst, 2004; Hoffman et al., 2006; Dulloo et al., 1999).

In a recent study on Meltdown® (SUP), which contains caffeine (increases resting metabolic rate), yerba mate extract (a stored fat mobilizer and appetite suppressant), tetradecylthioacetic acid (boosts mitochondrial fatty acid oxidation), methyl synephrine (acts as a mild stimulant), phenylethylamine (promotes mood elevation and lipolysis), yohimbine (facilitates lipolysis), and hordenine, researchers measured parameters including oxygen consumption (VO2) and respiratory quotient (RQ), energy expenditure, heart rate, fat oxidation rate, and blood pressure, the investigators concluded that the supplement effectively increased

acute energy expenditure and had a strong inotropic effect on heart rate and systolic blood pressure. However, the study also suggested supplements possess cardiovascular risks therefore should be used with caution (Roberts et al., 2005; Heck and De Mejia, 2007; Ro et al., 2015).

In 2017 a recipe of nutraceutical capsule containing hordenine, caffeine and β -phenylethylamine was patented with the claims that it enhances mental clarity and improves stamina, and energy (Allison et al., 2018; Jason, Guillermo and John, 2018).

In the most recent work on hordenine, Xu et al., showed effects of hordenine through shingilipid pathway for the treatment of ulcerative colitis which is also involved in several signal transduction pathways of cell proliferation, cell migration as well as inflammation. Overview of hordenine biological activities and cellular pathways reported in recent publications, see Table 6 and Figure 16. So, based on available evidence, hordenine has some potential to be researched as a new drug lead.



Figure 16: Hordenine and cellular pathways: Hordenine produce anti-prolactinoma effect in rat model through MAPK pathway supported by decline in Extracellular signal-regulated kinase (ERK1/2), Jun kinase (JNK), P38 protein expression level and the production of inflammatory cytokines TNF- α , IL-1 β , and IL-6 and inhibited PRL secretion through DRD2 agonist which is regulated through cAMP /PKA/CREB pathways in two cell lines MMQ cells (high expression)- oestrogen-induced rat pituitary tumour cells and GH3 cells (low expression)- radiation-induced rat pituitary tumour cells: Hordenine is also potential inhibitor of PDK3 gene, an important drug target to block PDH complex a switch between glycolysis and citric acid cycle targeting this mitonuclear communication may represent a novel approach to overcome chemoresistance. Hordenine enhance or complement actions of insulin acitivating Nuclear Factor Erythroid 2–related Factor 2 (Nrf2), regulate melanin through cyclic Adenosine Monophosphate (cAMP):cAMP-mediated pathways that is CREB (cAMP response element binding protein) phosphorylation, treat acute lung injury through Protein kinase B (Akt) Signalling and Nuclear Factor κ B (NF- κ B) and heals wound through serotonin (5-hydroxytryptamine) (Created with Biorender).

Table 6: Biological Activities of Hordenine

Biological Activities	Concentration	References
In mouse colonic epithelial cells (MCECs).	Dextran sodium sulphate	Xu et al., 2023
Effects on ulcerative colitis: reduced	(DSS) DSS+hordenine 50	
disease activity index (DAI), relieved colon	mg/kg, DSS+hordenine 25	
oedema, lesions, inflammatory cells	mg/kg, DSS+hordenine 12.5	
infiltration through sphingolipid pathway.	mg/kg	
PBCEC, CCK8 kit contain (lymphocytes (T	0.1 μM to 100 μM	Hahn et al., 2022
cells, B cells, and NK cells), monocytes,		
and dendritic dendritic cells), human colon		
adenocarcinoma (Caco-2)		
Hordenine protects against	12.5, 25, 50, 75–100 µg/ml	Zhang et al., 2021
lipopolysaccharide-induced acute lung		
injury by inhibiting inflammation		
Hordenine a potent inhibitor of PDK3 in	5.4 µM	Anwar et al., 2020
human lung cancer cells (A549 and H1299)		
Antiprolactinoma effect of hordenine by	152.8 (p < 0.01), 76.4	Wang et al., 2020
inhibiting MAPK signaling pathway activation in rats	(p< 0.01), 38.2 mg/kg (p < 0.05)	
Quorum sensing (QS) inhibitory activity	0.5 to 1.0 mg ml ⁻¹	Zhou et al., 2018
Binds to and activates the dopamine D2	13 µM	Sommer et al.,
		2017
Anti-melanogenic effects of hordenine and its mechanism in human melanocytes	500 μM (Treated duration 5 days)	Kim et al., 2013
Hordenine exerts anti-melanogenic effects	0-100 μM,	Kim et al., 2013
by inhibiting the cAMP signaling pathway	(Treatment duration up to 5 days)	
No measurable contraction of vasa	150 μM	Ghosh and Das,
deterentia in rat.		2010
Significantly potentiated the tonic response	25 µM	Barwell, 1989
inhibited those to Tyramine		

Chapter 2

Materials and Methods

2.1 Plant Collection

Whole plant of *Alhagi maurorum* was collected from Pattoki, Pakistan. *Alhagi maurorum* is not very hardy in nature and grows in sunny season and alkaline soil. Around 10 kg of the plant was collected by farmers when it was in its flowering stage and were transported under the supervision of Assistant professor Shumaila Arshad, (The University of Lahore), Lahore, Pakistan, to the Botany department of the Government College University, Lahore, Pakistan (Figure 17). All samples used in this research were collected in a single lot from August – September 2020. *Alhagi maurorum* was washed three times with water to remove soil and any contaminant and then left to dry under shade which is a traditional method to preserve maximum phyto-constituents of the plants (Zhang et al., 2021; Gasecka et al., 2020; Roshanak, Rahimmalek and Goli, 2016).



Figure 17: Sampling area Alhagi maurorum (Map of Pakistan, 2022)

2.2 Plant Identification

Alhagi maurorum was authenticated for this research by Professor Zaheer-ud-Din (Botany department of Government College University (GCU), Lahore, Pakistan).



Figure 18: Sample of desiccated Alhagi maurorum used in this study.

Following organoleptic evaluation *Alhagi maurorum* was identified, briefly as thorny shrub, intricately branched, aerial parts were not very hard and greenish in colour. It generally grows 1 m tall, but roots penetrate up to 6 m (deep root system) in soil and were very hard along with rhizome. Greenish stems bear axillary spreading very sharp spines. Leaves are alternate, simple, ovate, entire, and elliptic. Petioles are short up to 2 mm long while stipules are 1 mm in length. Leaves are hairy and light green in colour. Flowers are pinkish to maroon having short pedicel. Flowers arise from axil and are not too many in number. Fruits are pod shaped and are reddish brown in colour. Authenticated voucher specimens were deposited in the Herbarium of GC University (**Herbarium code**: GC. Herb. Bot. 3332). Samples of the desiccated plant were transported to the University of Westminster, London, UK fully sealed contained in cardboard boxes by air using DHL service (Figure 18).

2.3 Plant Extraction

2.3.1 Pulverization

Dried soft stem parts were separated and pulverized to fine powder using a laboratory scale mini mill/commercial blender. Then the powder obtained after milling was passed through sieve of stainless steel of mesh number 70 (Aperture 125 µm) (Figure 19).



Figure 19: Mini Mill: Pulverization of Alhagi maurorum

2.3.2 Solvent Extraction

Different solvents were used in the extraction procedures based on some published studies (Chemat, Vian, Cravotto, 2012; Zhang, Lin & Ye, 2018; Gallo, 2022), including methanol, chloroform, ethanol, and water. 0.5gm of powdered sample was mixed with 5ml of each solvent in clean glass test tubes. All tubes were shaken and were allowed to stand for 30 minutes. The solutions obtained were then observed under visible light as well as UV light for the detection of their colour. Based on the colour results and published literature ethanol was selected as an appropriate solvent for the extraction. After selecting solvent different concentrations of ethanol (50%, 70%, 99.8%) were also tested for extraction. Plant material was macerated for 72hr and 70% ethanol was found to be more appropriate solvent for extraction because 99.8% ethanol extracted all types of organic compounds this includes polar and non-

polar compounds and even chlorophyl of the plant. So, based on these results 70% ethanol was chosen as the most effective solvent for extraction (Sultana, Anwar, and Ashraf, 2009; Cao et al., 2022; Urabee et al., 2021) (For the details of extracting solvent see Table 7).

Solvent	Percentage
Methanol (CH ₃ OH)	99.8% (Analytical grade) (Sigma, UK)
Chloroform (CHCl ₃)	Anhydrous, \geq 99%, (0.5-1.0% ethanol) as stabilizer (Sigma, UK). Analytical grade
Water	Distilled water
Ethanol (CH ₃ CH ₂ OH)	 a. 99.8 % (Absolute ethanol) (Sigma, UK) b. 70% (70:30) c. 50% (50:50)

Table 7: List of extraction solvents

Fine powder of *Alhagi maurorum* was dissolved in 70% ethanol in a 1:10 ratio and solution was agitated three times a day. After 72hr of maceration, the macerant was removed and filtered through Whatmann filter paper #1 (medium fast) to remove residue/debri.

Then filtered extract was concentrated using rotary evaporator (Buchi Rotavapor R-215) under reduced pressure 100 mbar and at temperature not exceeding 40 °C. Concentrated extract was dried in incubator at 37 °C in petri dishes to increase the surface area. Dried extract was homogenized using pestle mortar and kept at -20 °C for experiments in tightly closed bottles. Extract was found to be soluble in 70% ethanol and thereafter extract was reconstituted in 70% ethanol for pharmacological testing.



Figure 20: Schematics of extract preparation of *Alhagi maurorum* with 70% Ethanol Solvent A .Fluorescence analysis for the selection of suitable solvent B. Maceration of pulverized powder in 70% Ethanol for 72hr C. Filtration of macerated extract through Whatmann filter paper # 1 D. Filtered extract E. Solvent evaporation using rotary evaporator F. Concentrated extract G. Dried extract at 37°C H. Homogenization of extract using pestle and mortar I. Extract completely soluble in 70% ethanol which is used as solvent for cell culture experiments.

2.4 High Performance Thin Layer Chromatography analysis of *Alhagi* maurorum extract

High Performance Thin Layer Chromatography (HPTLC) (CAMAG, Muttenz, Switzerland) provided an excellent separation of components of the extracts (Varghese, 2013; Reich and Schibli, 2006). For HPTLC analysis chapters on HPTLC about herbal materials were used as a reference, published by The United States Pharmacopoeia (USP-NF 2015) and the European Pharmacopoeia (Ph. Eur. 2017).

After extracting the *Alhagi maurorum* (for details of extraction see Section 2.3.2) with ethanol 70%, characterization was performed with HPTLC. For quantitative analysis of constituents of dried extract of *Alhagi maurorum* dried extract 100 mg was reconstituted in 70% ethanol (1ml). The reconstituted sample was centrifuged at 3000 rpm for 5 minutes.

Reconstituted sample was filtered using 0.45 μ m Nylon syringe filter. Various solvent combinations (mobile phase) were used to develop plates (for details of solvent see Table 13) and 2 μ l sample was applied to HPTLC plate (Precoated Silica gel plates 60 F₂₅₄ (Merck or equivalent), 10 x 10 cm) by Automatic TLC Sampler (ATS4) (CAMAG, Muttenz, Switzerland) using proper needle type on Silica coated plate band wise (8 mm) at a flow rate of 1.0 ml/min and column temperature was kept at 25°C. Plates were then dried for 20 minutes at 100°C. For the preparation of reference 100 mg hordenine powder was dissolved in 1ml of ethanol which was later used as a reference compound.

After successful sample application plate was moved to the Automatic Development Chamber (ADC2) (CAMAG, Muttenz, Switzerland) for the development of chromatogram saturated with preconditioning solvent and developing solvent under controlled conditions of humidity to final solvent migration distance of 70mm.

2.5 Liquid Chromatography-Mass Spectroscopic Analysis of *Alhagi* maurorum extract

To determine the exact molecular weight of the compounds in *Alhagi maurorum* extract Liquid Chromatography-Mass Spectroscopic (LC-MS) analysis was performed on Shimadzu LC-MS system in Chemistry Mass Spectrometry Facility, School of Pharmacy, University College London, and London, United Kingdom (Heyman and Meyer, 2012; Kamboj and Saluja, 2011). LC-MS spectrometer technique used combination of liquid chromatography (LC) and mass spectroscopy (MS) where LC allowed physical separation of analytes. Ions having strongest interaction with stationary phase left the column in the end and the time an analyte spends in the column is the characteristics of that analyte and is called retention time (RT) and response was captured in the form of peaks. It is one of the common methods used to identify compounds. Later separated ions in LC were driven into the mass spectrometer. A mass spectrometer (MS) an analytical instrument produces a beam of gas phase ions from analytes (sample) as a result sorting of mixture of ions based on m/z ratio when electrical or magnetic fields or both were applied. As a result, digital output signals in the form of peaks appeared and the m/z ratio and abundance or intensity of each ionic species detected was determined. 10 μ l of Alhagi maurorum extract sample was introduced through direct infusion through syringe connected to the pump. Separation was achieved using mobile phase containing 0.2% formic acid in water (A) and a mixture of ethanol and acetonitrile (1:1, v/v) used at the flow rate 0.3 ml/min. Shimadzu was used for data processing and reporting (Kowalczyk and Kwiatek, 2018; Hoffmann et al., 2014; Vassiliadis et al., 2019).

2.6 Cell Culture

All cell culture work was carried out aseptically to ensure that all cell lines were not infected with fungi, bacteria, or cross contaminated with other cell lines. Following are the processes, materials and equipment used in this research.

Chemicals

- Hordenine, M.W (165.23). Analytical grade (HPLC (Assay > 97.5%); Impurities <3.0% water)
- 2. Foetal Bovine Serum -heat inactivated (Gibco®, Thermo Fisher Scientific UK)
- Dulbecco's modified Eagle medium (DMEM)- liquid, high glucose with L-glutamine, D-glucose, sodium pyruvate (Gibco®, Thermo Fisher Scientific UK)
- 4. Phosphate Buffer Saline (PBS), pH 7.4 with potassium phosphate, sodium chloride and dibasic sodium phosphate (Gibco®, Fisher Scientific UK)
- 5. 0.25% TrypLE-EDTA liquid (Gibco®, Thermo Fisher Scientific UK)
- 6. Penicillin-Streptomycin (100X solution) (Gibco®, Thermo Fisher Scientific UK)
- 7. Dimethyl sulphoxide: DMSO (AnalaR BDH, UK)
- 8. Horse serum (Sigma, UK)
- 9. Glutamine (Sigma, UK)
- 10. Cholera toxin, 100 ng/ml of final volume (1 mg/ml stock) (Sigma, UK)
- 11. Insulin, 10 µg/ml of final volume (10 mg/ml stock) (Sigma, UK)
- 12. Extracellular growth factor, 20 ng/ml of final volume (100 µg/ml stock) (Sigma, UK)
- 13. Hydrocortisone, 0.5 mg/ml final volume, (1mg/ml stock) (Sigma, UK)

2.6.1 Cell Lines Used in Research

Two cell lines were used in this research MCF-7 and MCF-10-A. After defrosting, cells were seeded and incubated in humidified atmosphere, with in appropriate size T75-flasks, at 37°C, at 5% CO₂. Cells reaching 70% confluency were detached by using TrypLE-EDTA and were used for further analysis. Following a standard protocol, cells were seeded in 96 well plate for cell viability, ROS, MMP, intracellular Ca⁺² assays and 24 well plate for Seahorse mitostress assay throughout the experiments.

1. Breast Cancer Cell Line (MCF-7)

MCF-7 is an adherent, human epithelial cancer cell line derived from a metastatic mammary adenocarcinoma cultured in Minimum Essential Media (MEM) (Sigma, UK) (Soule et al., 1990). MCF-7 cells cultured in Minimum Essential Media (MEM) (Sigma, UK) was also supplemented with 10% Fetal Bovine Serum (FBS), 1% l-glutamine and 1% penicillin/streptomycin.

MCF-7 is estrogen receptor-positive (ER+); progesterone receptor positive (PR+) and human epidermal growth factor receptor-2 negative (HER2-) breast cancer cell line. Low metastatic potential and presence of steroid receptors make MCF-7 cell line suitable *in vitro* model for several studies such as endocrine therapy, drug sensitivity (Horwitz, Costlow and McGuire, 1975). In addition to the above-mentioned features MCF-7 cells form domes, capable to grow in monolayers and are sensitive to cytokeratin (CK) structural proteins expressed by epithelial cells which help them to withstand mechanical stress (Camarillo et al., 2014).

2. Non-Cancerous Breast Cell Line (MCF-10-A)

MCF-10-A is an adherent, human epithelial non-cancerous breast cell line and found to exhibit immortality in low Ca²⁺ media (Soule et al., 1990). These were cultured in a mixture of Dulbecco's Modified Eagle's Media and Ham's F12 nutrient mix (DMEM: F12) (Life Sciences, UK), supplemented with 5% horse serum (Sigma, UK), 1% penicillin/streptomycin, 20 ng/ml epidermal growth factor (Sigma, UK), 0.5 mg/ml hydrocortisone (Sigma, UK), 100 ng/ml cholera toxin (Sigma, UK) and 10 μ g/ml insulin (Sigma, UK).

MCF-10-A (human breast epithelial cell line) is a widely used *in vitro* non-cancerous breast epithelial cell model. MCF-10-A is estrogen receptor negative (ER-) and forms acinus-like

spheroids (3D model) a good feature to study the microenvironment of the cell. However, unlike MCF-7 cell line MCF-10-A lack anchorage-independent growth (Qu et al., 2015). Cells between passages 5 to 15 were used in this research and tested periodically for mycoplasma infection using the Plasmo Test Mycoplasma Detection Kit (Invivo Gen, UK).

2.6.2 Thawing Cell Lines from Liquid Nitrogen

Cell lines bought commercially and received in frozen form. So, cells must be thawed and cultured before use. Cells must be thawed rapidly and diluted slowly using warm growth medium to eliminate dimethyl sulfoxide, an organosulfur compound (DMSO) which is used as a cryoprotectant. DMSO also prevents water crystallization that would cause cell lysis during cryopreservation. Cells kept in liquid nitrogen were defrosted, transferred to T75 flask, and kept in incubator at appropriate temperature 37° C with 5% CO₂.

2.6.3 Subculture of Cell Lines (Cell Passaging/Cell Splitting)

Cells should be observed often to check their confluency as adherent cell lines continue to grow until the medium is depleted of nutrients. Once cell lines covered the surface area of the flask, they were sub-cultured after three days to prevent the culture dying as well as to increase the number of cells for different assays. Following aseptic procedure, media was aspirated off and discarded from flask. Cells were washed with PBS solution to remove dead cells from the flask and later detached following five to seven minutes incubation using TrypLE (10% of the total volume of media) which hydrolyses proteins that attach the cells to the bottom of the flask in case of adherent cells. After incubation detached cells transferred to 15ml falcon tube after adding fresh media and centrifuged at 1000G for about 5 minutes. Then supernatant was aspirated off and pellet is dispersed using fresh media. Then cells were transferred to new flasks labeled containing fresh media and kept in an incubator at 37°C, 5% CO₂.

2.6.4 Cell Seeding

Cells were seeded in consistent numbers for experiments. Old media from cells was aspirated and cells were washed using PBS buffer and were left in incubator after adding TrypLE in incubator at 37°C, 5% CO₂. Once cells were detached after diluting with fresh media were transferred to falcon for centrifugation at 1000 G for 5 minutes. After centrifugation old media was aspirated off and pellet was resuspended in fresh media. For cell counting, cells (20 μ l) were diluted with Trypan blue. Then cells were counted using Countess automated cell counter. A consistent number of cell concentrations were added to wells with the appropriate volume of media. Two different plates were used,

- 1) 100 µl total well volume for 96 well plates (25,000 cells per well)
- 2) 500 µl total well volume for Seahorse assay 24 well plate (30,000 cells per well)
 Following equation is used to calculate the number of cells / well required for the experiments,

Seeding Density = Desired number of cells / Total number of cells X 1000

2.6.5 Freezing Cells for Storage in Liquid Nitrogen

Cells are frozen to be stored for future use, but the procedure should be carried out slowly to avoid osmotic shock which is caused by extracellular and intracellular ice formation and to minimize the effect of heat produced by DMSO when it is mixed to the water. Cells should be viable up to 90% at the time of freezing. Trypsinize cells using TrypLE followed by centrifugation step as explained previously. After the removal of supernatant from pellet add 90% of FBS and 10% DMSO and resuspend that are later stored into cryogenic vials and these vials are then stored into Liquid Nitrogen containers.

2.6.6 MTT Reduction Assay (Cell proliferation Assay)

The MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-tetrazolium bromide) assay is a colourimetric assay used to determine cell proliferation.). Principle of this assay is based on MTT (yellow tetrazole) reduction by mitochondrial reductase enzyme which is NADPH-dependent to Formazan (1-(4,5-Dimethylthiazol-2-yl)-3,5-diphenylformazan, Thiazolyl blue formazan) (purple) (Mosmann, 1983) (Figure 21).

For MTT Assay cultured cells were seeded in a 96-well plate at a density of 2.5×10^4 cells per well and left overnight to adhere. Cells were treated with different concentrations of AM extract/hordenine prepared in fresh media. Post treatment (AM extract for 24hr and hordenine for 6, 24, 48hr) cells were washed with PBS. Then fresh phenol red-free media (100 µL) was added in each well containing treated cells followed by addition of MTT solution (10 µl/well). Plates were covered in aluminum tinfoil to protect from light and incubated at 37°C, 5% CO₂, for 3hr. After incubation, media was aspirated, and resultant crystals were dissolved in 100 µl MTT solvent (DMSO) (Lysis buffer). Viability is quantified by ultraviolet (UV)-Vis spectrometry (SPECTROstar Nano, BMG Labtech, and Germany) with a microplate reader.

Equation,

Absorbance normalized = Absorbance (570nm) – Absorbance (690nm) (Background) % Cell viability = Absorbance (Treated) / Absorbance (untreated control) X 100



Figure 21: Cell Viability Assay: MTT assay for determination of Cell viability (Created with BioRender).

2.6.7 Detection and Quantification of Cellular ROS

Cellular ROS levels were investigated in this study using a 2',7'–dichlorofluorescein diacetate (DCFDA) assay (Abcam, UK). The fluorescence produced is due to presence of ROS which converts DCFDA into the highly fluorescent compound 2',7'–dichlorofluorescein (DCF) is detected and quantified using fluorescence spectroscopy (FLUOstar Optima, BMG Labtech, UK) (Excitation 485 nm, Emission 520 nm) (Henley et al., 2017; Keston and Brandt, 1965; LeBel et al., 1992). Cultured cells were seeded in a 96-well plate at a density of 2.5 x 10^4 cells per well and left overnight to adhere. Cells were treated with AM extract and hordenine prepared in fresh media. Post treatment, media was aspirated off and cells were washed with PBS (100 µl). Later the buffer was replaced with DCFDA stain prepared in 1X buffer. Treated plates were

incubated at 37°C for 45 minutes. DCFDA stain was removed and replaced with PBS (100 μ l). Plates were covered in aluminum tin foil to avoid photobleaching effect. Fluorescence was measured using fluorescence plate reader using excitation/emission filters (Table 8).

Fable 8:	Typical	working	concentrations	of DCFDA
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Preparation of reagents		
1 x Buffer	Sterilize ddH ₂ O using syringe filter.	
	Mix 9 ml of sterile ddH ₂ O with 1ml of 10X buffer	
DCFDA Stain	2.1 ml of 1X buffer and 2.1 µl of DCFDA stain (20	
	μΜ)	

Equation,

% of $ROS = As-Cb/C-Cb \times 100$

Where,

C= Fluorscence of control (Average of at least 3-5 wells) Cb= Fluorscence of blank (Average of at least 3-5 wells) As= Fluorscence of sample

2.6.8 Detection and Quantification of Mitochondrial Membrane Potential

TMRE (tetramethylrhodamine, ethyl ester) (Abcam, UK) is a cell based qualitative assay to assess the changes in membrane potential of active mitochondria. It is a red-orange dye, due to positive charge readily accumulates in mitochondria having relatively negative charge (Xu et al., 2021; Jaune et al., 2021). Cultured cells were seeded in a 96-well plate at a density of 2.5 x 10^4 cells per well and left overnight to adhere. Post treatment, cells were stained with TMRE prepared in fresh media (100 µl). Stained plates covered in aluminium foil were incubated at 37°C, 5% CO₂ environment for 30-35 minutes. The stain was aspirated off and cells were washed twice with PBS to remove background fluorescence caused by cell culture media and replace with PBS. Fluorescence was measured using fluorescence spectroscopy (FLUOstar Optima BMG Labtech, UK) and selecting appropriate filter set (excitation/emission (544 nm / 590 nm) (Table 9).

Table 9: Typical working concentrations of TMRE

Working Concentrations

Serial Dilution	1 μl of TMRE in 99 μl media (10 μM)
	50 μl of 10 μM solution in 2.5ml (200 nm)

Equation,

% of MMP = $As-Cb/C-Cb \times 100$

Where,

C= Fluorscence of control (Average of at least 3-5 wells) Cb= Fluorscence of blank (Average of at least 3-5 wells) As= Fluorscence of sample

2.6.9 Detection and Quantification of Intracellular Ca⁺²

The acute intracellular Ca²⁺ levels are measured with the red fluorescent dye Rhod-2 (Abcam, UK). Basic principle involves the hydrolysis of the ester to Rhod-2 by cellular endogenous esterase which will bind to Ca²⁺ and once compound is in cells, localizes to mitochondria because ester itself is non-fluorescent and does not bind to Ca²⁺ directly. Chemical name of Rhod-2 is 1-[2-Amino-5-(3-dimethylamino-6-dimethylammonio-9-xanthenyl) phenoxy]-2-(2-amino-5-methylphenoxy) ethane-N,N,N',N'-tetraacetic acid, chloride (Dajindo, 2022)(Figure 22). Cultured cells were seeded in a 96-well plate at a density of 2.5 x10⁴ cells per well and left overnight to adhere. Post treatment cells were stained with Rhod-2 prepared in fresh media (100 μ l/well). Stained plates covered in aluminum foil and incubated at 37°C, 5% CO₂ environment for 40minutes. Then stain was aspirated from wells and cells were washed with PBS twice and replaced with PBS (100 μ l/well). Fluorescence was measured using fluorescence spectroscopy (FLUOstar Optima, BMG Labtech, UK) and selecting appropriate filter set (excitation/emission (544 nm/590 nm) (Table 10).

 Table 10: Typical working concentrations of Rhod-2

Working Concentration			
Dilute stock in media (1 μ M-5 μ M)	2.1 μ l stock in 2.1 ml media (1 μ M)		

Equation,

% Of Intracellular $Ca^{2+} = As-Cb/C-Cb X 100$

Where,

C= Fluorscence of control (Average of at least 3-5 wells) Cb= Fluorscence of blank (Average of at least 3-5 wells) As= Fluorscence of sample



Figure 22: Acute Intracellular Ca²⁺ Assay. (Created with BioRender).

2.6.10 Seahorse Cell Mito Stress Assay

The mitochondrial respiration was measured using the Seahorse cell mitostress assay, the standard assay often used for assessing the mitochondrial respiration. The Agilent Seahorse XF delivers real time and direct measurements of mitochondrial metabolism with high specificity. Oxygen consumption rate (OCR) of plated cells was measured in real time by modulating ETC in cell well by adding modulators, oligomycin (Cheng et al., 2012; Kim et al., 2020), carbonyl cyanide-4 (trifluoromethoxy) phenylhydrazone (FCCP) (Reda et al., 2019), antimycin and rotenone (Sanchez-Alvarez et al., 2020) through built in injection ports on XF sensor cartridges (Figure 23) (Table 11).


Figure 23: Mitochondria: Site of action of ETC modulators on large mitochondrial protein respiratory complexes located in inner mitochondrial membrane. Proton pumping complexes of ETC are mainly targeted in Mitostress assay. Oligomycin (Complex V/ATP Synthase inhibitor), FCCP (ETC uncoupling agent), Antimycin (Complex III/ ubiquinone-cytochrome c oxidoreductase inhibitor), Rotenone (Complex I/ NADHubiquinone oxidoreductase inhibitor) [(Agilent, 2021) (Image created with BioRender)].

On the first day cultured cells were seeded using XF24 cell culture microplates following two step seeding process. First, cells seeded to the desired final concentration in 100 μ l of the growth medium without seeding in background correction wells (A1, B4, C3, D6). The seeded plate was incubated for 3hr to allow cells to adhere completely then 150 μ l of growth medium was added to make the total volume in well 250 μ l. Seeded plates were incubated overnight at 37°C, 5% CO₂. On the second day of treatment various concentrations of AM extract and hordenine were prepared in fresh media. Old media from wells was aspirated and fresh media containing treatments was added while keeping cell volume at this stage around 500 μ l. Post treatment seeded plates were incubated at 37°C, 5% CO₂. 1 ml of XF Calibrant was added into each well of the Utility Plate and incubated overnight at 37°C in a non-CO₂ incubator. On the third day, the assay media was prepared by warming 49 ml of XF Base at 37°C and adding Sodium pyruvate (500 μ l) and Glucose 200g/ml solution (500 μ l). Treated cells were washed with assay media (500 μ l) twice. After addition of assay media (500 μ l) in each well including blanks, the plate was placed at 37°C in non-CO₂ incubator for 45 minutes. ETC modulators were prepared in the assay media as follows,

Modulators	Stock	Final Conc.	Function	Effects on OCR
A. Oligomycin	5 mg/ml	1 μg/ml	ATP synthase inhibitor	Decrease
B. FCCP	10 mM	10 µM	Uncoupling agent	Increase
C. Antimycin + Rotenone	10 mM	10 μM	Complex III and I inhibitors	Decrease

Table 11: Typical working concentrations of ETC modulators

Prepared modulators as described in Table 10 were loaded to ports A (55 μ l), B (60 μ l) and C (65 μ l) of the utility plate. This calibration plate was loaded to Agilent Seahorse XF Analyzer and replaced after calibration step (15 minutes) with cell culture plate from non-CO₂ incubator (Kosgodage et al., 2018)

 Table 12: Equations of key parameters of seahorse mito-stress assay (Agilent, 2021)

Parameter	Rate measurement equation					
Non-Mitochondrial Respiration	Minimum rate measurement after rotenone/antimycin injection					
Basal respiration	(Last rate measurement before injection of oligomycin) - (Non-mitochondrial respiration rate)					
Maximal respiration	(Maximum rate measurement after FCCP injection) - (Non- mitochondrial respiration)					
Proton (H+) leak	(Minimum rate measurement after oligomycin injection) – (Non-mitochondrial respiration)					
ATP production (Last rate measurement before injection of oligomycii (Minimum rate measurement after oligomycin injection)						
Spare respiratory capacity	(Maximal Respiration) - (Basal respiration)					

2.6.11 Morphological Analysis Using Fluorescent Microscopy

MCF-7 and MCF-10-A cells were seeded in 96 well plates and left overnight to adhere. After treating cells with *Alhagi maurorum* and hordenine they were observed under microscope. The Mito tracker deep red stain was used to observe the changes in mitochondrial morphology. MCF-7 and MCF-10-A cells were washed with PBS and replaced with phenol red free media and stained with 100nM Mito tracker deep red dye for 35 minutes and were kept in incubator at 37 °C, 5% CO₂. Cells were replaced with PBS and observed under microscope.

Statistical analysis

All statistical analysis was performed on Prism 5.04 (GraphPad) (San Diego, USA). For multiple comparisons and to find specific pair wise differences Ordinary One-way ANOVA test was applied, where mean of each group is compared to the mean of the control group was corrected by Dunnett's Multiple Comparison Test. Outlier test was applied to the data using Rout (robust regression and outlier removal) method based on false discovery rate which is recommended as it can find any number of outliers (definitive/likely) in the data. For some analysis one sample t test was also applied. Values are presented as mean \pm standard error and p value =/<0.05 is considered statistically significant. Significant differences are indicated with asterisks (* if p< 0.05, **if p<0.01, *** if p<0.001, **** if p<0.001).

Chapter 3

Results of *Alhagi maurorum* extract 3.1 Characterization of *Alhagi maurorum* with HPTLC

The high-performance thin layer chromatography (HPTLC) fingerprints of ethanolic extract of plant Alhagi maurorum showed the presence of a several phytochemicals as illustrated by different bands in (Figure 24). Various developing solvents were tried to develop chromatograms to find out the most suitable solvent. First, mobile phase used was composed of toluene, ethyl acetate and ethanol in the ratio of 70:30:5 reported for AM extract in literature (Chen et al., 2021; Hashim et al., 2016a-2017b) which separated the components of AM extract which is indicated by different bands in the chromatograms D (Figure 24). Then a mobile phase is composed of chloroform, methanol and con. Ammonia having ratios of 80:20:10 reported for hordenine was used for the AM extract and hordenine as a reference compound was also added (Ghosal and Srivastava, 1973; Varghese and Alka, 2013). Reference compound hordenine used was also dissolved in ethanol (100 mg/ml). A band appeared relative to the reference compound (hordenine) which indicated possible presence of hordenine in AM extract as shown in chromatogram B (Figure 24). Furthermore, one mobile phase composed of toluene, ethyl acetate, ethanol, and ammonia 25% was used which yielded less bands and indicated incomplete separation of constituents of AM extract as shown in chromatogram C (Figure 24). Mobile phase composed of ethyl acetate: methanol: ammonia 35% separated bioactive compounds of AM extract (Figure A) and then these developed plates were then dried. After drying plates were analysed at 254nm and 366nm using TLC Visualiser which were later derivatized and heated (CAMAG TLC Plate Heater) at 70 °C for 5 min. Finally, the cooled plates were analysed at different wavelengths of light (254nm, 366nm, white light (TLC Visualiser, CAMAG, Muttenz, Switzerland). A summary of the developing solvents is in Table 13 for reference. The peak tables were also generated and the RF values, peak height of unknown compounds were recorded (Hashim et al., 2016; Bian et al., 2022; Thammarat et al., 2021).



Figure 24: HPTLC Analysis: results view at wavelength 366nm. Separation of *Alhagi maurorum* extract in HPTLC analysis. Chromatograms for *Alhagi maurorum* obtained with different mobile phase. A. Ethyl acetate: Methanol: Ammonia 35% (Phenethylamine), B. Chloroform: Methanol: Conc. Ammonia (hordenine), C. Toluene: Methanol: Ammonia 25% (*Alhagi* extract) D. Toluene: Ethyl acetate: Ethanol (*Alhagi* extract).

Sample	Mobile Phase /	References
Alhagi maurorum extract	Toluene: Ethyl acetate: Ethanol (70:30:5)	(Chen et al., 2021)
Alkaloids	Toluene: Ethyl acetate: Methanol: Ammonia 25% (30:30:15:1)	HPTLC Book
Phenethylamine	Ethyl acetate: Methanol: Ammonia 35% (85:10:5)	(<u>www.camag.com</u> , 2017)
Hordenine	Two solvent system: Solvent 1: Chloroform: Methanol: Conc. Ammonia (80:20:10) for hor- denine detection Solvent 2: Chloroform: Methanol: Acetic Acid (75:15:10) beta phene- thylamine	(Ghosal and Srivastava, 1973)

 Table 13: List of developing solvents for HPTLC analysis.

These sets of fingerprints of sample at 254 nm and 366 nm, and at white light are to derive values for migration distances (Rf) and corresponding intensities (AU). So, these

chromatograms hold two-dimensional information (Rf vs. AU). Fingerprints generated by HPTLC are richer source of information as various bands of different colours indicate different compounds (Islam et al., 2021). Previous literature has shown phenolic compounds exhibit different spot colouration when exposed to UV light specifically, flavonoids, show orange or yellow spots, flavones are dull brown in colour. Purple-red-orange indicates the presence of anthocyanins that are coloured water-soluble pigments. They appear as red, blue, and purple as different shades of blue colour were also detected which possibly indicate presence of different classes of phenolics. Simple phenolics (salicylic acids, gallic acids) dot fluoresces (UV light) and so, some dark absorbing blue spots were also observed. Dark and light blue indicate presence of coumaric, ferulic acid (Locher et al., 2017; Milojkovic et al., 2016). Violet blue indicates saponins and blue terpenoids. As different numbers of blue bands appeared when different developing solvents were used, therefore, more repeats are required to confirm these results. Bright reddish orange colour is also observed which indicates the presence of alkaloids. As it is very common to detect alkaloids because there are more than 10,000 alkaloids distributed in 300 different species of plants. Our results indicate presence of higher contents of flavonoids and alkaloids in 70% ethanol extract of Alhagi maurorum. Previously, reported literature indicated alkaloids are detected at Rf values 0.18-0.92 and AUC 2800-above 10,000 (Karthika and Paulsamy, 2015; Kulkarni et al., 2014) (Figure 25).



Figure 25: Detection of tracks of *Alhagi maurorum extract* and reference (hordenine) at different wavelengths (254 nm, RT White, 366 nm) in HPTLC analysis.

3.2 Characterization of *Alhagi maurorum* with Liquid Chromatography-Mass Spectroscopic

Liquid chromatography and mass spectroscopic analysis was performed. Data was acquired in the positive ionization mode (ESI+) and negative ionization mode (ESI-). Several compounds have been identified including phenolics and derivatives. Detection at 254 nm (Figure 26).

Screening of chemically diverse Alhagi maurorum for medicinal compound.



Peak 1[ES+/ES-]

2.000e6-

1.500e6

1.000e6

5.000e5

311.00

200.0

366.85

400 0

446.95 599.15

457.10

600 0

633.10

640.85

742.85 774.75

800 0

1000.0

1200.0

1400.0

1600.0

1800.0

m/7



Peak 3



Peak 4

MS Spectrum Group#1 - LC Peak: 4, RT Int 1.500e6-]3(12: 4.47 to 4.60 min 18: 15 18: 15 14: 15 14: 10 14: 10 15: 10 15: 10 15: 10 10 10 10 10 10 10 10 10 10	33.05 563.10		1000.85	5				
1.000e6 5.000e5 0 156.70 156.70 200.0	441.05 440.0	625.35 648 600.0	95 729.35 ⁸³⁸ 800.0	95 1002.70 960.85 110 1000.0	5.55 1200.0	1400.0	1600.0	1800.0	m/z
MS Spectrum Group#2 - LC Peak: 4, RT	: 4.47 to 4.60 min								
Int	001 70								
6.000e6 4.000e6 2.000e6 158.65 306.80 2.000e6 262.75	20.80 366.35 482.90	593.10 671.10							
200.0	400.0	600.0	800.0	1000.0	1200.0	1400.0	1600.0	1800.0	m/z

Peak 5



Peak 6



Peak 7



Peak 8





Figure 26: (A) LC-MS Chromatograms of *Alhagi maurorum* extract. Detection is done at -254, 2+20. 155-2000mz.Sample infused through autosampler and the mass-to-charge ratio (m/z) 155-2000mz. (B) Chemical structures of some important compounds.

LC-ESI-MS method was used in this study, due to high efficiency in separation as well as sensitivity of alkaloids in the positive mode to identify potential bioactive compounds. Tentatively mainly alkaloids were identified at this stage by aligning the molecular weight based on proposed charge to mass ratio with PubChem, google scholar, dictionary of natural products and MassBank databases. Spectra was acquired in the mass range of 155-2,000. A group of alkaloids were detected, possibly phenethylamine hordenine (m/z 158.65) though tyramine (m/z 138)/ N-methyltyramine (151.2) was not detected on selected spectra range as reported previously but N-methyltyramine (151.2) is proposed based on peak observed (156.70). Furthermore, lysergic acid ergot derivatives, protopine, stylopine were also detected and possibly have fragmentation pattern as of hordenine (m/z 158.65). Isoquinoline alkaloids identified for example protopine, cryptopine, stylopine which have shown to have hepatoprotective activity and antiproliferative activity based on observed peaks in positive ionization mode. Five peaks reported in PubChem for a group of heterohexacyclic alkaloids and considering molecular weight closest compounds identified were protopine (m/z 353.4), hydroprotopine (m/z 354.38), stylopine (m/z 323.3) and cryptopine (m/z 368.85) based on reported peaks (m/z 354.275:171,149:288), and (m/z 370,369.8,369.9,369.7,370.1) respectively. (S)-N-Methylcanadine (m/z 354.4) benzylisoquinoline alkaloid, a quaternary ammonium ion that was S-canadine (m/z 339.4) bearing an N-methyl substituent. Canadine was reported to be immediate precursor of berberine and act as a calcium channel blocker. Aurantiamide (m/z 402.5), an isoindole alkaloid previously being reported from *Alhagi sparsifolia* was potentially detected which previously being reported to have an anti-neuroinflammatory activity and block activation of NF-kB, phosphorylation of JNK and p38 of MAPK pathway identified based on reported peaks (m/z 385,400, 402). Salsolidine (m/z 207.27), a tetrahydroisoquinoline alkaloid, known as a stereoselective competitive MAO A inhibitor identified. Caffeoyl-hexose-deoxyhexoside potentially also found based on the following $[M - H]^-$ at m/z 487.0 and fragment ion peak detected at m/z 308 (Li et al., 2018; Jianshe et al., 2015) as reported in literature (Table 14).

Some other classes of organic compounds of pharmacological significance were also tentatively identified as well such as diclofenac (NSAID), posaconazole (Antifungal), catechol etc.

Table	14: LC-MS	Analysis:	Tentative	identification	of alkaloids	and	other	compounds	in Alhagi	maurorum
				ех	tract					

No.	RT (min)	Peak m/z	Tentative Identification	Chemical Formula	[M–H] ⁻ /[M–H]+ (m/z) literature	Reference
1	3.94-4.20	366.85	13-Hydroxyoxyberberine	C20H17NO6	367.4	PubChem
2	6.47-6.65	355.15	Hydroprotopine	$C_{20}H_{20}NO_5^+$	354.4	PubChem
3	4.20-4.47	158.65	Ephedrine/hordenine	C ₁₀ H ₁₅ NO	166.123/165.	(Hahn et al.,
		(-/+)	_		23	2022);
						Mass bank
4	4.47-4.60	156.70	N-methyltyramine		152	NIST
		(+)				
5	6.47-6.65	355.15	Protopine	$C_{20}H_{19}NO_5$		Lehner et al.,
					353.13	2004
6	3.94-4.20	323.25	Stylopine	$C_{19}H_{17}NO_4$	323.3 (-)	PubChem
7	3.94-4.20	368.85	Cryptopine	$C_{21}H_{23}NO_5$	369.4 (-)	PubChem
8	4.47-4.60	320.80	7-Methoxy-17,19-dioxa- 11-azapentacy- clo[12.7.0.03,11.04,9.016, 20]henicosa- 1(21),2,4,6,9,14,16(20)- heptaen-8-on	C ₁₉ H ₁₅ NO ₄	321.3	PubChem
9	6.84-7.18	408.75	alpha-Fagarine hydrochlo- ride	C ₂₁ H ₂₄ ClNO ₅	405.9	PubChem
10	6.84-7.18	408.75	16,17-Dimethoxy-14-(2- oxopropyl)-5,7-dioxa-13- azapentacy- clo[11.8.0.02,10.04,8.015,	C ₂₃ H ₂₃ NO ₆	409.4	PubChem

			20]henicosa-			
			2,4(8),9,15(20),16,18-hex-			
			aen-21-one			
11	4.47-4.60	364.70	Corycavine	$C_{21}H_{21}NO_5$	367.4 (-)	PubChem
12	6.47-6.65	355.15/3	trans-Canadine-N-oxide/	$C_{20}H_{21}NO_{5}/$	355.4	PubChem
		44.95	D-Tetrahydropalmatine	$C_{21}H_{25}NO_4$		
13	11.42-	520.65-	Dimethyl 14,16,17-trimet-	$C_{27}H_{25}NO_{10}$	523.5	PubChem
	11.97/7.1	529.30	hoxy-21-oxo-5,7-dioxa-			
	8-7.38	(-)	13-azahexa-			
			cyclo[12.7.2.01,13.02,10.0			
			4,8.015,20 tricosa-			
			2,4(8),9,13(20),10,18,22- henteene 22,23 dicar			
			hoxylate			
14	7 18-7 38	363.85	Leptopidinine/Hypeco-	C20H17NOc	367.4	PubChem
17	7.10 7.50	505.05	rinine (possible isomers)	C2011/1106	507.4	T ubenem
15	3.94-4.20	366.85	16.17-Dimethoxy-13.21-	$C_{22}H_{24}NO_4^+$	366.4	PubChem
			dimethyl-5,7-dioxa-13-	- 2224- + - 4		
			azoniapen-			
			tacyclo[11.8.0.02,10.04,8.			
			015,20]henicosa-			
			1(21),2,4(8),9,15(20),16,1			
			8-heptaene			
16	7.18-	355.15/3	(S)-N-Methylcanadine	$C_{21}H_{24}NO_4^+$	354.4	PubChem
	7.38/11.4	52.90				
17	2-11.97	266.95	Destaurtieren	C II O	266.2	Dath Charm
17	3.94-4.20	300.83	Pachyrrnizone	$C_{20}H_{14}O_{7}$	300.3	PubChem
18	11.42-	352.90	8-Oxocanadine		353.4	PubChem
10	11.97	552.90	0 Oxoculturite	0201191(03	555.1	rubenem
19	3.94-4.20	368.85	1,2-Dimethoxy-6-methyl-	$C_{21}H_{23}NO_5$	369.4	PubChem
			5,7,8,15-tetrahydro-6H-			
			benzo[c][1,3]diox-			
			olo[4',5':4,5]benzo[1,2-			
			g]azecin-14-one			
20	6.84-7.18	358.50	1-(4-fluorophenyl)-2-	$C_{20}H_{20}FNO_4$	357.4	PubChem
			[(5R)-4-methoxy-6-me-			
			thyl-7,8-dihydro-5H-			
			[1,3]d10x010[4,5-g]180-			
21	1 17	366 35()	1 Mathoxybarbarinium	CarHasNO ⁺	366 /	PubChom
21	4.47-	3// 95	Mundoserone	$C_{21}T_{20}TO_5$	342.3	PubChem
22	3 94-4 20	368.85	2-(4-Methoxy-6-methyl-	$C_{19}H_{18}O_6$	369.4	PubChem
23	5.91 1.20	500.05	5.6.7.8-tetrahydro[1.3]di-	021112311003	507.1	rubenem
			oxolo[4,5-g]isoquinolin-5-			
			yl)-1-(3-methoxy-			
			phenyl)79xoazeti/			
			(16,17-Dimethoxy-5,7-			
			dioxa-1-azapen-			
			tacyclo[11.8.0.03,11.04,8.			
			014,19]henicosa-			
			3(11),4(8),9,14,16,18-he-			
			xaen-12-yl)methanol			

24	3.94-4.20	340.80	16,17-Dimethoxy-5,7-	$C_{20}H_{22}NO_4^+$	340.4	PubChem
			dioxa-13-azoniapen-			
			tacyclo[11.8.0.02,10.04,8.			
			015,20]henicosa-			
			2,4(8),9,15(20),16,18-he-			
			xaene			
25	3.94-4.20	368.85	2-(4-Methoxy-6-methyl-	$C_{21}H_{23}NO_5$	369.4	PubChem
			5,6,7,8-tetrahydro[1,3] di-			
			oxolo[4,5-g]isoquinolin-5-			
			yl)-1-(2-methoxy-			
			phenyl)ethanone			
26	3.58-3.94	398.85	1-(2,4-Dimethoxyphenyl)-	$C_{22}H_{25}NO_6$	399.4	PubChem
			2-(4-methoxy-6-methyl-			
			5,6,7,8-tetrahydro[1,3] di-			
			oxolo[4,5-g]isoquinolin-5-			
27		294 (0()	yl)ethanone	C U NO ±	204.4	Dath Charm
27		384.00(-)	4-Methoxy-5-[2-(2-meth-	$C_{22}H_{26}NO_5^{+}$	384.4	PubChem
			6.6 dimethyl 5.6.7.8 tetra			
			hvdro[1,3] dioxolo[4,5-			
			glisoquinolin-6-jum			
28		446.95	4-Methoxy-6 6-dimethyl-	$C_{24}H_{20}NO_7^+$	444.5	PubChem
20		110.95	5-[2-oxo-2-(3.4.5-tri-	02411301107	111.5	rubenem
			methoxyphenyl)ethyl]-			
			5,6,7,8-tetrahydro[1,3]di-			
			oxolo[4,5-g]isoquinolin-6-			
			ium			
29	3.94-4.20	340.80(-)	2-(4-Methoxy-6-methyl-	$C_{20}H_{21}NO_4$	339.4	PubChem
			5,6,7,8-tetrahydro[1,3]di-			
			oxolo[4,5-g]isoquinolin-5-			
			yl)-1-phenylethanone			
30	4.47-	487.05/	caffeoyl-hexose-deoxy-	C15H18O9	487.05/308.7	Chen et al.,
	4.60/7.18	308.70/	hexoside		0/308.15	2012
21	-7.38	308.15	NT N (41 - 1 1	C12U10NO2	225 295	CLEDI
31	6.47-6.65	222.95/2	N-Methylmescaline	CI2HI9NO3	225.285	Chebi
20	11.42	21.85	Calcolidina	C II NO	207.07	
32	11.42-	65	Saisondine	$C_{12}H_{17}NO_2$	207.07	
33	6.84-7.18	401.80(-)	Aurantiamide	CarHacNaOa	402.5	
55	0.04-7.10	401.00(-)	Aurannannae	$C_{2511261} V_{2} O_{3}$	402.5	
		+01.10(- /+)				
34	4.47-4.60	441.05	dia-aurantiamide acetate	C27H28N2O4	444.522	ChemSpider
35	3,58-3.94	245.85	Pyrrolezanthine	$C_{14}H_{15}NO_3$	245.85	ChemSpider
36	3.58-3.94	291.85	Fusarentin 6-Methyl Ether	C ₁₅ H ₂₀ O ₆	296.31	ChemSpider
						L
- 1	LC-MS	Analysis: Te	entative identification of othe	er compounds in	Alhagi mauroru	<i>m</i> extract
1	3.58-3.98	475	4-((3-chloro-2-(2-hydrox-	C26H22CIN3	4/5.13	10.1002/jnet.2
			ypneny1)-4-	04		549
			80xoazetidine-1-v1)(2-			
			hydroxyphenvl)-methvl)-			
			1,2-dihydro-5-methyl2-			
			phenyl-1h-pyrazol-3(2h)-			
			one (antifungal)			

2	3.58-3.98	291.85	Diclofenac (NSAID)	C14H11Cl2N O2	292.00	PMID:- 20470236
3	3.58-3.98	250.85	Bromocatechol (phenol)	C6H5BrO2	251.05	CN- 113480534-A
4	3.58-3.98	627.10	5-(2-Ethoxy-5-((4-methyl- 1,4-diazepan-1- yl)sulfonyl)phenyl)-1- methyl-3-propyl-1,6- dihydro7H-pyrazolo[4,3- d]pyrimidin-7-one (Phosphodiesterase inhibitor)	C34H39N6O 4S	627.30	WO- 2017060874- A1
5	3.58-3.98	459.05	4-((4-chlorophenyl)(4- hydroxy-2-oxo-2H- chromen-3-yl)methyl)-5- methyl-2-phenyl- 1Hpyrazol-3(2H)-one (coumarin derivative)	C26H19CIN2 O4	460.10	PMID:- 9635556
6	3.58-3.98	773.15	(1,3-Dibenzyl-4,5- diphenylimidazol-2- ylidene)gold(I) p- Mercaptobenzoic Acid (Gold compound)	C36H29N2O 2SAu	773.25	PMID:- 30110951
7	3.58-3.98	795.20	(S)-Azelastine N-Oxide (Antiallergy)	C22H24CIN3 O2	794.48	PubChem
8	3.58-3.98	773.15	Posaconazole (Antifungal)	C37H42F2N8 O4	687.00	71313564
9	3.58-3.98	308.25	Bisdemethoxycurcumin (curcuminoid)	C19H16O4	308.04	Patent CN- 113980063A
10	3.58-3.98	563.95	4-[(3-bromophenyl) methoxy] -N-[(2-hydroxy- 5-methoxyphenyl) methylideneamino]benza mide (palladacycle inhibitor)	C22H19BrN2 O4 Pd	564.27	Compound CID: 1378937
11	3.58-3.94	727.30	N,N-bis(4-carbazol-9- ylphenyl)-4-(4- phenylphenyl)aniline (Aniline derivative)	C54H37N3	728.01	Compound CID: 158394886
12	3.58-3.94	624.95	Lyngbyabellin M (non- ribosomal peptide synthetase)	C25H34N2O 8Cl2S2	624.91	10.3390/mole- cules25173986
13	3.48-3.96	727.20	Phosphatidylserine (Phospholipid)	C45H84NO9 P	726.95	10.7150/thno.4 5125
14	3.48-3.96	647.05	DL-Thioctamide (lipoamide)	C8H15NOS2	647.50	PubChem CID 863

3.3 Measurement of Cell Viability

Alhagi maurorum 70% ethanolic extract was investigated using cancer breast cell line (MCF-7) and non-cancer breast cell line (MCF-10-A). The first step before assessing the cell viability of extract on cell line, various concentrations of ethanol on MCF-7 were tested. According to the published literature ethanol 0.3% is physiologically considered safe (Figure 27).



Figure 27: Breast cancer (MCF-7) cells treated with increasing concentrations of ethanol for 24hr. Data is represented as the average of n=4 independent experiments \pm SEM. Statistical analysis: One sample t test, * indicates significance p<0.05. (ET: Ethanol solvent control)

3.3.1 Effects of Alhagi maurorum Extract on Cell Viability

Alhagi maurorum extract prepared in 70% ethanol at different concentrations ranging from 10 μ g/ml to 1000 μ g/ml were studied using standard MTT assay. The stock of Alhagi maurorum extract was prepared in 70% ethanol (stock 1 g /5 ml), then diluted with culture medium and different concentrations for the treatment were prepared by serially diluting stock. The concentration of vehicle control (ethanol 70%) was kept at or below 0.5%. Untreated cells receiving the same volume of culture medium served as control and viability was measured by comparing absorbance of treated and untreated cells. Effects of AM on MCF-7 and MCF-10-A

cells were measured after 24hr of exposure (Figure 28). The formation of purple colour of formazan was the indicator of the assay formed by the activity of mitochondrial dehydrogenase in viable cells.



Figure 28: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with increasing concentrations of *Alhagi maurorum* extract for 24hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001, ****indicates p<0.001. (ET:Ethanol solvent control)

AM was observed to be toxic at higher concentrations of 500 μ g/ml and 1000 μ g/ml to MCF-7 cells (up to 60-70%, p < 0.0001) and 250 μ g/ml, 500 μ g/ml and 1000 μ g/ml to MCF-10-A cells (up to 60-80%, p< 0.0001) relative to control. It was also observed that AM has altered the morphology of cells at higher concentrations when observed under microscope



Figure 29: Cell viability Assay of *Alhagi maourorum:* MTT assay used to assess cytotoxic potential. Purple colour observed is proportional to the number of viable cells. Side images are microscopic view where cell death is visible at higher concentrations. (ET:Ethanol solvent control)

3.4 Measurement of Oxidative Stress

ROS signalling molecules are vital to maintain cellular homeostasis and an imbalance in ROS levels results in oxidative stress. In this study the effects of AM prepared in 70% Ethanol were analysed on breast cancer (MCF-7) and normal (MCF-10-A) cell lines. Intracellular ROS levels were measured in this study using a cell-based assay DCFDA used as an indicator of ROS levels (Figure 30).



Figure 30: Effect of solvent on ROS levels. Breast cancer (MCF-7) cells treated with increasing concentrations of ethanol for 24hr. Data is represented as the average of n=3 independent experiments \pm SEM. Statistical analysis: One sample t test, * indicates significance p<0.05. (ET:Ethanol solvent control)

3.4.1 Effects of Alhagi maurorum Extract on ROS Levels

AM Extract significantly reduced ROS levels at higher concentrations (100 μ g/ml, 250 μ g/ml, 500 μ g/ml, and 1000 μ g/ml) (up to 50%, p<0.0001) and at lower concentration (up to

20%, p<0.0053) in case of MCF-7 cell line when compared to untreated control. However, in MCF-10-A, it significantly reduced the ROS levels up to 10-20% at higher concentration of 250 μ g/ml, 500 μ g/ml, and 1000 μ g/ml (up to 20%, p<0.048) (Figure 31).



Figure 31: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with *Alhagi maurorum* ethanolic extract for 24hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001), ****indicates p<0.0001. (ET:Ethanol solvent control)

3.5 Measurement of Mitochondrial Membrane Potential (MMP)

MMP was measured using a cell-based assay where a fluorescent dye TMRE accumulates in polarized mitochondria due to their relative negative charge of the membrane and high intensity of fluorescence was generated. However, depolarized, or defective mitochondria will not retain dye and generate fluorescence of low intensity.

3.5.1 Effects of Alhagi maurorum Extract on MMP

AM extract significantly reduced MMP of MCF-7 and MCF-10-A cells after 24hr of treatment compared to untreated control at high concentrations 250 μ g/ml, 500 μ g/ml, 1000 μ g/ml (up to 50 to 80%, p < 0.0001) (Figure 32).



Figure 32: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with *Alhagi maurorum* ethanolic extract for 24hr. Data is represented as the average of n=5 independent experiments ± SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001), ****indicates p<0.0001. (ET:Ethanol solvent control)

3.6 Measurement of Intracellular Calcium Concentrations

 Ca^{2+} is an abundant intracellular messenger participating in diverse cellular functions. Concentration of Ca^{2+} is maintained by different compartments within a cell. Any change in the level of Ca^{2+} was measured in this study through fluorescence signals generated by labeled molecules used as indicators of Ca^{2+} in the cells. The Rhod-2 AM ester forms are cationic in nature and are selective indicator for mitochondrial Ca^{2+} via membrane potential driven uptake. Exact loading concentration, time and temperature were determined empirically for both cell lines (MCF-7 and MCF-10-A). Incubation time is a critical factor to allow complete deesterification of intracellular Rhod-2 AM esters. However, prolonged incubation eliminates cytosolic staining but mitochondrial retained. Further, results could be improved by reducing rhod-2 AM to dihydrorhod-2 AM (non-fluorescent) to distinguish between cytosolic and mitochondrial dye location. Main process involves oxidation of Rhod-2 AM ester of dihydrorhod-2 that led to the release of rhod-2 indicator a process occurs in mitochondrial environment.

3.6.1 Effects of Alhagi maurorum Extract on Intracellular Ca²⁺

Cultured and treated cells with *Alhagi maurorum* extract for 24hr were loaded with the Rhod-2 AM ester forms of the Ca²⁺ indicators. Then fluorescence signals generated from these stained cells were measured. In MCF-7 cells the levels of Ca²⁺ were reduced at higher concentrations 250 µg/ml (up to 30%, p<0.01) and 500 µg/ml, 1000µg/ml (up to 50-60%, p<0.0001) while no significant change in Ca²⁺ level was observed at lower concentrations compared to untreated control (Figure 33). However, in the case of MCF-10-A the opposite trend was observed that is higher levels of Ca²⁺ at higher concentrations after exposure of 24hr treatment with *Alhagi maurorum* though more repeats are required to draw conclusion.



Figure 33: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with *Alhagi maurorum* extract for 24hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One

way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001. (ET:Ethanol solvent control)

3.7 Measurement of Mitochondrial Respiration

To get a deep insight into the effects of *Alhagi maurorum* on mitochondrial function, Seahorse mitostress assay was performed in which six parameters were measured in one assay: basal respiration, ATP production, proton leak, spare respiratory capacity, maximal respiration, and non-mitochondrial respiration [(For detailed formula see Section 2.2.10: Seahorse cell mito stress assay (Table 11)]. These parameters serve as an indicator of how cells respond to stress induced by mitochondrial ETC modulators/ inhibitors, Oligomycin (Complex V), FCCP (Uncoupler), antimycin A (Complex III) and rotenone (Complex I).

3.7.1 Effects of Alhagi maurorum Extract on Mitochondrial Respiration

Alhagi maurorum extract after treatment for 24hr has shown no significant effect on MCF-7 cell line only a significant decrease in proton leaks up to 60% was observed at 250 μ g/ml. There is no significant change is observed at lower concentrations of 10 μ g/ml in MCF-7 cell line (Figure 34). Also, AM extract reduced the basal respiration, ATP production and maximal respiration at 100 μ g/ml (up to 70-80%, p< 0.019) and at 250 μ g/ml (up to 80%, p<0.008) in MCF-10-A. In the same way, significant decrease in proton leak at concentration 250 μ g/ml (up to 80-90%, p<0.007) was also observed whereas, at 100 μ g/ml decline was up to 70% though it didn't reach significance. Furthermore, at 10 μ g/ml concentration there a significant increased effect of proton leak was observed (up to 10-20%, p<0.0198) in MCF-10-A (Figure 36). Additionally, significantly reduced ATP production was observed at 250 μ g/ml (up to 80-90%, p<0.0327) and at 100 μ g/ml (up to 80%, p< 0.0491). There is significant reduction in levels of spare respiratory capacity and non-mitochondrial respiration observed at higher concentrations though none reached significance after five repeats. Maximal respiration was also significantly reduced at concentrations 100 μ g/ml (p<0.0126) and 250 μ g/ml up to 70-80% (p<0.007) [(For OCR profile of MCF-7 and MCF-10-A see (Figure 35) (Figure 37)].



Figure 34: Effect of *Alhagi maurorum* on OXPHOS: Breast cancer (MCF-7) cells treated with *Alhagi maurorum* extract for 24hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001), ****indicates p<0.0001. (ET:Ethanol solvent control)



MCF-7 treated with A. maurorum

Figure 35: MCF-7 OCR Profile. Seahorse XF Cell Mito Stress test profile of MCF-7 Cell Line treated with *Alhagi maurorum* extract for 24 hr (Agilent, 2021). Schematic, showing interpretation of OCR levels in response to the injection of Oligomycin, FCCP, Rotenone and Antimycin.



Figure 36 : Effect of *Alhagi maurorum* on OXPHOS. Breast cells (MCF-10-A) treated with *Alhagi maurorum* extract for 24hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001, ****indicates p<0.0001. (ET:Ethanol solvent control).



Figure 37: MCF-10-A OCR profile. Seahorse XF Cell Mito Stress test profile of MCF-10-A Cell Line treated with *Alhagi maurorum*e extract for 24 hr (Agilent, 2021). Schematic, showing interpretation of OCR levels in response to the injection of Oligomycin, FCCP, Rotenone and Antimycin.

Chapter 4

Results of Hordenine

4.1 Measurement of Cell Viability

The bioactive compound hordenine was investigated using cancer breast cell line (MCF-7) and non-cancer breast cell line (MCF-10-A).

4.1.1 Effects of Hordenine on Cell Viability

Standard colorimetric MTT assay was used after treating MCF-7 and MCF-10-A cells with a range of concentrations of hordenine (0.1 μ M-10,000 μ M) for 24hr. Stock solution of hordenine was prepared in DMSO, then diluted serially in culture medium to prepare range of concentrations. However, concentration of vehicle DMSO was kept at and below 1% and untreated cells served as control after receiving the same volume of cell culture medium.

The results suggest that hordenine had no significant effect on cell growth on both cell lines though a trend of slightly lower cell viability compared to normal breast cell line was observed. A significant decrease in cell viability at much higher concentrations (5,000 μ M and 10,000 μ M) was also observed but that could be the consequence of higher solvent concentration. Furthermore, effects of hordenine by exposing cells (MCF-7 and MCF-10-A) for 6hr and 48hr were also studied. No significant effects observed at most of the concentrations after 6hr and 48hr after five independent experiments in both cell lines (Figure 38).



Figure 38: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with hordenine for 6 hr, 24hr and 48hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnett's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001, **** indicates p<0.001.

4.2 Measurement of Oxidative stress

The bioactive compound hordenine was analysed on breast cancer (MCF-7) and normal (MCF-10-A) cell lines.

4.2.1 Effects of Hordenine on ROS Levels

The effects of hordenine on ROS levels were also assessed fluorometrically using DCFDA assay. Hordenine stimulated ROS levels after 6hr of exposure in MCF-7 cell line. There were no significant effects observed after 24hr and 48hr of exposure of cells with hordenine at most of the concentrations but only at very high concentration of 1000 μ M after 24hr in MCF-7. However, ROS levels significantly reduced after 6hr in MCF-10-A at concentrations 0.1 μ M, 10 μ M, 500 μ M and 1000 μ M (up to 20-30%, p=0.005) while no significant effect after 24hr and 48hr of exposure of cells with hordenine was observed as compared to untreated control (Figure 39) at almost all concentrations.





Figure 39: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with hordenine for 6 hr, 24hr and 48hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001, **** indicates p<0.001.

4.3 Measurement of Mitochondrial Membrane Potential (MMP)

MMP was measured after treating MCF-7 and MCF-10-A cells with hordenine.

4.3.1 Effects of Hordenine on MMP

Effects of hordenine were assessed on MMP, a key indicator of mitochondrial activity using cell-based assay in which a fluorescent dye was used to measure the intensity of fluorescence generated by polarized and depolarized mitochondria in both cell lines (MCF-7 and MCF-10-A). Cells were exposed to hordenine prepared in DMSO for 6hr, 24hr and 48hr. No significant change in MMP of MCF-7 and MCF-10-A cells was observed compared to untreated control at most of the concentrations used after at least five independent experiments across three timepoints (Figure 40).



Figure 40: Breast cancer (MCF-7) and normal (MCF-10-A) cells treated with hordenine for 6 hr, 24hr and 48hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001, ****indicates p<0.001.

4.4 Measurement of Intracellular Ca²⁺

The effect of hordenine on intracellular Ca^{2+} was analyzed on breast cancer (MCF-7) and normal (MCF-10-A) cell lines.

4.4.1 Effects of Hordenine on Intracellular Ca²⁺

Cells cultured and treated with hordenine were stained to measure intracellular Ca^{2+} level at three time points, 6hr, 24hr and 48 hr. There is no significant change was observed in the levels of Ca^{2+} at most of the concentrations across three timepoints on MCF-7 cell line (Figure 41).



Figure 41: Breast cancer (MCF-7) cells treated with hordenine for 6 hr, 24hr and 48hr. Data is represented as the average of n=5 independent experiments \pm SEM. Statistical analysis: One way ANOVA followed by Dunnetts's multiple comparison tests, * indicates significance p<0.05; ** indicates p<0.01, *** indicates p<0.001.

4.5 Measurement of Mitochondrial Respiration

Effects of hordenine on mitochondrial respiration Seahorse analyzer was used to perform Seahorse mitostress assay. To get a deep insight into the effect of hordenine on mitochondrial functions. The following six parameters were measured in one assay, basal respiration, ATP production, proton leak, spare respiratory capacity, maximal respiration, and non-mitochondrial respiration. These parameters are an indicator of how cells respond to stress caused by mitochondrial inhibitors, Oligomycin (Complex V), FCCP (Uncoupler), antimycin A (Complex III) and rotenone (Complex I) after exposure of hordenine for 24hr.

4.5.1 Effects of Hordenine on Mitochondrial Respiration

No significant effect is observed after treatment with hordenine at all concentrations in MCF-7 and MCF-10-A on all parameters of OCR (Figure 42).



Figure 42: Seahorse mitostress assay. Key parameters of mitochondrial respiration measured after treatment of MCF-7 cell line with hordenine (0.05 μ M-1000 μ M) for 24hr. Data is presented as percentage of the control ± SEM of n=4 independent experiments. Statistical Analysis: One way ANOVA for pairwise comparison. * Indicates significance (p<0.05).



MCF-7 treated with Hordenine

Figure 43: OCR profile of MCF-7 after sequential injection of ETC modulators. Schematic, showing interpretation of OCR levels in response to the injection of Oligomycin, FCCP, Rotenone and Antimycin.



Figure 44: Seahorse Mitostress assay. Key parameters of mitochondrial respiration measured after treatment of MCF-10-A cell line with hordenine (0.05 μ M-1000 μ M) for 24hr. Data is presented as percentage of the control ± SEM of 3 independent experiments. Statistical analysis: One way ANOVA for pairwise comparison. * Indicates significance (p <0.05).



MCF-10-A treated with Hordenine

Figure 45: OCR profile of MCF-10-A. Seahorse mitostress assay MCF-10-A OCR profile after sequential injection of ETC modulators. Schematic, showing interpretation of OCR levels in response to the injection of Oligomycin, FCCP, Rotenone and Antimycin.

4.6 Observation of Morphology Using Fluorescent Microscopy

Mitochondrial imaging of MCF-7 and MCF-10-A after treating cells for 24hr with hordenine was done to see any effect on morphology of mitochondria (Figure 47). No significant changes in morphology were observed confirming the abovementioned results. Furthermore, low concentrations of *Alhagi maurorum* extract were also tested and morphology was analysed which confirmed the above findings that it has not affected the morphology at lower concentrations.







Figure 46: A. Morphological analysis of MCF-7 and MCF-10-A after treatment with hordenine at low and high concentrations for 24hr (top two figures). B. Morphological analysis of MCF-7 and MCF-10-A after treatment with *Alhagi maurorum* for 24hr. Mito Tracker Deep Red Images. Morphology quantified by observing changes in the intensity of Mito Tracker Deep Red (Created with BioRender).
Chapter 5

Discussion

Advance understanding of mitochondrial biology has thrown light on the mitochondrial impact on health and disease beyond ATP and macromolecule production. Mitochondrial involvement is now noticeable in several cellular processes such as redox and Ca^{2+} homeostasis, cell death, critical biosynthetic pathways but mitochondria undergo dramatic changes in morphology as cells grow, divide, and die, and the relationship between mitochondrial structure and functions is still unclear. Although, mitochondrial role in cancer progression has been identified in the last century but it is still not fully understood mechanism how changes in mitochondrial shape and organization can lead to cancer and how altered mitochondrial functions could be manipulated for therapeutic benefit.

Natural compounds and their structural analogues have historically made major contribution to new drug leads and Alhagi species are under study due to their antiinflammatory and anti-cancer activities (Sarkar et al., 2020; Davinelli et al., 2013; Parvinroo et al., 2014; Ardekani, Rahimi, and Mohammad, 2010). In this study the effects of *Alhagi maurorum* extract (AM extract) on mitochondrial functions of MCF-7 (breast cancer) and MCF-10-A (control) have been evaluated (Irsfeld, Spadafore and Prub, 2013; Rafehi et al., 2021; Tundis et al., 2012; Guan et al., 2020; Thu et al., 2020; Wang et al., 2020; Ghosh and Das, 2010; Zhou et al., 2018; Zhang et al., 2021).

5.1 Effect of AM extract and hordenine on Cell Viability

In the first part of this study the effect of AM extract on the cell viability of cancerous and non-cancerous breast cell lines is assessed. As the main objective is to investigate the impact of different concentrations on cell homeostasis, cell death initiation and induction so range of concentrations from low 10 μ g/ml to high up to 1000 μ g/ml are tested. Cell viability has not changed at lower concentrations tested and decreased consistently with increase in the concentration of AM extract. AM extract after 24hr of exposure has significantly reduced the cell viability of MCF-7 and MCF-10-A at high concentrations (250 μ g/ml, 500

 μ g/ml, and 1000 μ g/ml) up to 70%, though cell viability remained unchanged at low concentrations tested. Increased cell death at high concentrations in MCF-7 suggest cytotoxic effects of the AM extract. The highest cytotoxic activity is observed at 1000 μ g/ml concentration (See results 3.3.1 of Chapter 3) and indicate the level of cytotoxic activity is concentration dependent. Mitochondria are the main sites where most of the MTT is reduced, it passes through the mitochondrial inner membrane due to lipophilic nature and presence of positive charge (Ghasemi et al., 2021; Mueller et al., 2004; Riss et al., 2013). Formazan, a water insoluble molecule, is formed in viable cells due to the disruption of the core tetrazole ring by the activity of mitochondrial dehydrogenases, mainly succinate dehydrogenase. Bahamin et al., in his study of *Alhagi maurorum* extract on 4T1 breast cancer cells growth found significant reduction in cell growth up to 70-80% at 130 μ g/ml though he used DMSO as a solvent which in our study of hordenine has shown more growth inhibitory effects on MCF-7 and MCF-10-A (Bahamin et al., 2021).

Additionally, the bioactive compound hordenine was also tested on cell viability and it has shown no significant effects on cell viability after exposure for 6hr, 24hr and 48hr on MCF-7 and MCF-10-A cells (see Section 4.1.1). Only at very high physiological concentrations it reduced the cell viability in MCF-7 cell line but, it could be due to high concentrations of DMSO solvent used so these concentrations were excluded in later experiments. Cell viability results of the hordenine are consistent with the work of Anwar et al., 2020 who has also reported no effect of hordenine on the cell viability in lung cancer cell lines. However, unlike this, a study conducted by Zhang et al in 2021, investigated the effects of hordenine on RAW264.7 macrophages cells and showed decrease in cell viability at 100 µg/ml (Wang et al., 2010; Zhao et al., 2018; Lu, Wang, and Zhan, 2019; Zhang et al., 2021). In the most recent study of hordenine by Wang et al., it was shown that hordenine promoted cell proliferation and elevated the activity of primary mouse dermal papilla cells (to treat alopecia) through the Wnt signalling pathway, that promote angiogenesis. (Wang et al., 2023). As, hordenine has not altered the cell viability of different cell models as previously reported at lower concentrations while in this study even higher concentrations were tested in breast cell models and proved to be safe. However, several studies previously have shown hordenine regulate the MAPK signalling which is associated with several physiological processes of development, cellular proliferation, differentiation and apoptosis (Yasi et al., 2019; Zhao et al., 2018; Bertolotto et al., 1998; Busca and Ballotti, 2000; Kim,

et al., 2013 ; Brauers, Steiner and Daldrup, 2013) but due to the nature of the assay, it was not possible to identify any specific effect in this study (Van, Joubert and Cromarty, 2015).

MTT assay is primarily used to assess cellular metabolic activity and increased metabolic activity suggest proliferation and indirectly could be related to increase in mitochondrial mass succinate dehydrogenase complex (SDH-A), primary reducing enzyme of MTT. Succinate dehydrogenase (SDH-A protein) is encoded by the SDH-A gene in humans which has been linked to mitochondrial respiratory chain, citric acid cycle and is also considered as tumor suppressor gene (Rai, et al., 2018; Ghasemi et al., 2021; Alkahtani et al., 2022). Results obtained also suggest the effect of AM extract on glycolytic pathways too which are more active in cancer cells because cancer cells rely more on glycolysis to fulfill their ATP requirements. Interestingly, AM extract has initiated cell death at 250 μ g/ml concentration while induced apoptosis at higher concentrations so, different concentrations of AM extract can be used to manipulate cell death processes for example, autophagy, mitophagy and apoptosis and AM is a valuable source of novel anticancer compounds.

There are some limitations that need to be considered to draw full conclusion from the data because of cytotoxicity caused by the MTT assay. Moreover, the full mechanism of action of MTT reduction is still unknown because of the role of other organelles involved, enzymes and several biomolecules (glutathione, cysteine) in this process and need to be explored. So, more supportive experiments are required to confirm the findings, particularly the effect of AM extract on apoptosis or apoptotic proteins. Moreover, the principle behind MTT assay is, every viable cell in given cell line should have same metabolism so, quantification of total metabolism is directly proportional to number of viable cells. However, advanced research in this field also suggests cellular metabolic remodeling under stress meaning chemicals can modify metabolic potential of the cells. So, more relevant experiments must be considered to assess cytotoxic potential, growth inhibition and mitochondrial biogenesis because results simply indicate the metabolic state and not the cell number (Rai,et al., 2018; Mueller et al., 2004).

5.2 Effect of AM extract and hordenine on ROS Levels

AM extract has significantly reduced the ROS levels in MCF-7 at concentrations from 100 μ g/ml to 1000 μ g/ml in both cell lines. ROS signalling pathway is linked to the phosphorylation of JNK, p38 and ERK1/2. So, reduction in ROS levels in MCF-7 could be

possibly through modulation of JNK, p38 and ERK1/2 proteins. Reduction in ROS levels also indicates AM extract might have affected ETC which is the main source of ROS in mitochondria. Further, reduction in ATP levels while evaluating its effects on mitochondrial respiration confirmed the findings at high concentrations. However, unlike MCF-7 it is noted that AM extract has not affected the ROS levels of MCF-10-A at most of the concentrations tested, only up to 20-30% at higher concentration. Based on previously published studies, ROS species are considered as common and important signalling molecules and lower levels of ROS promote metabolic adaptation, moderate levels are significant to stimulate danger signals and immune responses while high levels activate apoptosis and autophagic cell death pathways so, interestingly AM extract has manipulated ROS levels in MCF-7 differently at different concentrations.

Further, the effect of various concentrations of hordenine on the redox state of both breast cell lines are also studied in this research and changes in ROS levels are quantified in the cells. After treating MCF-7 and MCF-10-A cell lines with hordenine for 24hr and 48hr no significant change in ROS levels is observed as compared to untreated control. Whereas, after 6hr of exposure with hordenine different response is observed in MCF-10-A where, ROS levels significantly reduced at 0.1 μ M, 10 μ M, 500 μ M and 1000 μ M and moderately, increased in MCF-7 at 0.05 μ M, 0.1 μ M and 10 μ M suggesting effects are concentration dependent.

ROS induces chronic inflammation through induction of inflammatory cytokines. Activation of NF-kB and chemokine receptors contribute to invasion, and metastasis of various tumor types. Therefore, slight increase in ROS levels activate antioxidant systems such as SOD and GSH. ROS levels can be increased as a therapeutic approach as higher levels of ROS above cellular tolerability threshold initiate cell death response pathways and this feature is very useful to induce apoptosis and even mitophagy. However, normal cell lines have higher capacity to cope with increased ROS levels therefore, not many changes in ROS levels observed in MCF-10-A after treatment with AM extract, only slight reduction in ROS levels at high concentrations (Choi et al., 2020). Anticancer agents could be developed based on dual role of ROS both as pro-oxidant and antioxidant though most of the concentrations have shown some effects though not reached significance, but modulation of ROS levels is not considered ideal as it increases adaptation of cancer cells to ROS and supports cancer stemness. Cancer cells tend to counterbalance ROS levels and steady state of ROS levels in cancer cells are determined by rate of ROS production and rate of ROS scavenge. High rate of ROS production is counterbalanced by equally high rate of antioxidant activity to maintain redox balance.

Research is ongoing to design ROS therapies where focus is to lower ROS levels to prevent signalling and induce to a certain level to selectively kill cancer cells for example, use of antioxidants, vitamin C in this regard. But the challenge here is if the ROS levels are not sufficiently raised within cancer cells, then therapy would simply further activate NFkB, P13K, HIFs and MAPK to promote tumorigenesis so, there is need to find compounds who selectively target antioxidant systems. There is also a need to identify which ROS species need to be targeted as ROS is a collective term used to describe various types of molecules produced as byproduct for example superoxide radical (O^{-2}) , hydrogen peroxide (H₂O₂) and hydroxyl radical (OH•) and some other and these are studied but still mechanism is not fully explored. Hydroxyl radical (OH•) have short life span generated from other ROS species and indiscriminately oxidise lipids, proteins and DNA and cause instability whereas, superoxide radical (O^{-2}) do not indiscriminately oxidises lipids, proteins, and DNA. Some information about mechanism of hydrogen peroxide (H₂O₂) mediated oxidation of cysteine residue within protein is known which exist as thiolate anion at physiological pH and is protonated by hydrogen peroxide (H_2O_2) and eventually alter functions of proteins of MAPK pathways (Zorov, Juhaszova and Sollott, 2014).

Furthermore, the site of ROS generation also plays a critical role as ETC process is not perfect and leakage of electrons more likely occur at complex I and Complex III though there are various redox centers on ETC. In a study targeting ROS for the treatment of inflammatory disease and cancer conducted by Li and his coworkers explains complex I leak superoxide radical $(O^{\bullet^{-2}})$ into mitochondrial matrix and Complex III leaks into the intermembrane space and mitochondrial matrix. Superoxide radical $(O^{\bullet^{-2}})$ dismutated to hydrogen peroxide by two dismutases located at different sites, superoxide dismutase 2 (SOD2) in mitochondrial matrix and superoxide dismutase 1 (SOD1) in mitochondrial intermembrane space. So, though both ROS are generated but superoxide radical $(O^{\bullet-2})$ cannot pass easily through mitochondrial outer membrane due to short half-life and electrophilic nature so very unlikely become the candidate of signalling molecule in the cell however within mitochondria may impact mitochondrial quality control system whereas, hydrogen peroxide (H₂O₂) is more stable and electrophobic in nature makes it an ideal signalling molecule (Li et al., 2013; Han et al., 2003). One more aspect needs consideration in designing ROS therapies, mitochondrial antioxidant enzymes are nuclear encoded and transported into the mitochondria so this mitochondria-nucleus crosstalk may also have some impact. There are some studies in this field who also reported three SOD isoforms where, Cu, Zn-dependent isoform SOD1 (Cu-Zn-SOD) found in mitochondrial intermembrane space and cytosol, manganese-dependent isoform SOD2 (Mn-SOD, SOD2) only in mitochondrial matrix and Cu, Zn SOD dependent isoform SOD3 in the extracellular space Cu, Zn SOD (EC-SOD). Moreover, impact of other mtROS scavenging enzymes and proteins such as catalase, GPx, peroxiredoxin and thioredoxins proteins need to be studied.

In addition to above mentioned factors, there is also the need to consider the role of metabolic state of mitochondria (that means rest and work state), a concept proposed by Chance and William in 1955. He explained the role of ADP and ATP in regulation of mitochondrial size by metabolism and found ADP as a key regulator of mitochondrial size, later this intramitochondrial regulation of respiratory rate was also confirmed by other scientists (Korzeniewski, 2015) though the impact of the optimal metabolic state of mitochondria is still not completely explored and hypermetabolism is reported in several mitochondrial diseases such as, Luft disease and complex diseases like cancer are some examples. Mitochondrial ROS production is largely dependent on O_2 concentration there is conflicting information available as several studies proposed linear relationship between mtROS production and oxygen concentration though some studies suggest increase in mtROS production even under hypoxic conditions therefore, to address this role of MMP cannot be ignored. It is known based on previous studies, physiological ROS signalling require optimised MMP and any fluctuation will cause oxidative stress the reason is there is an association between redox couples involved in oxidation of NADH and those involved in antioxidant defence systems (NADPH). NADP⁺ is necessary for mitochondrial antioxidant enzyme systems which is compromised with an increase in mitochondrial uncoupling of the ETC and increased ROS production but still molecular basis of this process needs more understanding as limited data is available to draw any conclusion because under hypoxic conditions increased mtROS production is linked to hypoxia-inducible factor 1α (HIF- 1α), high MMP, low electron flow and overexpression of mitochondrial uncoupling proteins (UCPs)(Busiello, Savarese S, Lombardi, 2015).

Mitochondrial quantity is largely dependent on the energy demands of the cell so, if mitochondrial ROS generation increases, there is also an increase in the expression of antioxidant enzymes system GPx and SOD₂, so the role of mitochondrial biogenesis factor peroxisome-proliferator-activated receptor- γ coactivator 1 α (PGC1 α) needs understanding in addressing levels of ROS (Li et al., 2013; Zorov, Juhaszova and Sollott, 2014; Chance and William, 1955). Role of intracellular Ca²⁺ in monitoring the levels of ROS cannot be ignored and it regulates the metabolic state of mitochondria by moderately increasing mitochondrial Ca²⁺ concentration, that stimulates flow of electrons in ETC and decrease mtROS production (Peng and Jou, 2010; Li et al., 2013). MtROS are also involved with processes of fusion and fission; under stressed conditions fragmented mitochondrial networks are found but still limited information is available how mtROS affect process of mitochondrial fusions. Both processes of fission and fusion influence several other mitochondrial functions too because fresh mitochondria keep on adding in the system. Several nuclear transcription factors (mitoTFs). For example, p53 and NF- κ B influence ROS levels through different mechanisms, p53 induces apoptosis and inhibits SOD2 activity as well. Some of the epigenetic regulatory enzyme proteins like sirtuins, important for suppressing inflammation, are believed to play a role in mtROS regulation, especially mitochondrial SIRT-3, SIRT-4, and SIRT-5.

Also, combination of ROS-generating drugs with pharmaceuticals that can break redox adaptation could also be a better strategy to design new therapies and overcome issues of drug resistance and side effects. Some scientists suggest lack of ROS also contributes to the progression of cancer because some levels of ROS are needed to correctly fold proteins in ER that could be significant for other critical cellular processes (Zorov, Juhaszova and Sollott, 2014). DCFDA is used as a probe for overall oxidative stress measurements in this study and is not ideal to study specific radical species due to indiscriminate nature of the assay. More experiments are required to find which ROS species is mainly reduced and effects on antioxidant systems to interpret accurately the effect of AM extract and hordenine. Study in the field of mtROS is essential before designing ROS therapies because of impact it has on different organs for example hypertension is associated with increased mtROS in organs mainly kidney and central nervous system, more likely caused by angiotensin II hormone, artherosclerosis is caused by excessive mtROS as increased mtDNA damage reported in these conditions which directly encode genes for the protein complexes of ETC and also several inflammatory diseases like rheumatoid arthritis, thyroiditis and type 1 diabetes are also known to be impacted by the levels of mtROS (Togliatto, Lombardo, Brizzi, 1988;Wen, Gwathmey, Xie, 2012).

5.3 Effect of AM extract and hordenine on Mitochondrial Membrane Potential

Optimal MMP is essential for the normal metabolic activity of the mitochondria. It relies on the activity of citric acid cycle and is an intermediate form of energy which is used by ATP synthase (Haddad and Mohiuddin, 2023). The MMP also has significant influence on the process of mitophagy and facilitates the removal of dysfunctional mitochondria. Alternatively, a decline in MMP affects ATP synthesis and eventually arrest growth of the cells, as seen at higher concentrations tested in this study, when cells of both cell lines were treated with AM extract. Adenine nucleotide transporter (ANT) protein located in mitochondria plays an important role in the development of MMP which is the primary site of exchange of ATP and ADP from matrix to the intermembrane space. Another source of MMP is reverse operation of ATP synthase where the ATPase inhibitory factor 1(IF1) is involved and intracellular ATP is utilized to maintain MMP i.e., rotational direction of ATP synthase and reverse operation of ANT importing cytosolic ATP into mitochondria. ATP compound is essential for buffering stable MMP and this is very important feature in clinical conditions associated with decreased OXPHOS for example hypoxia, stroke, lactic acidosis and mitochondrial myopathies (Zorova et al., 2018; Walker, 2013

The results obtained from this study indicate AM extract modulated MMP and reduced it at high concentrations. Mitochondrial depolarization at high concentrations may induce opening of mitochondrial permeability transition pore formation and released cytochrome c, AIF (apoptosis inducing factor) and Smac into the cytosol from the inter membrane space of the mitochondria. Migration of cytochrome c is a significant event in the process of apoptosis. AIF release leads to chromatin condensation as well as DNA fragmentation. However, experiments are necessary to confirm role of AIF and MAPK pathway, though the results indicated intriguing role of AM extract in provoking mitochondrial damage at high doses, modulation of apoptotic proteins in MCF-7 and, in other words affecting chemiosmotic coupling within mitochondria in MCF-7 (Yuan et al., 2012; Xu et al., 2011). Furthermore, MMP after treating cells with hordenine showed no significant change across three time points at all concentrations in both cell lines used in this study. Interaction between mtROS, MMP has already been explained in Section 5.2.

Although some studies have reported that the MMP typically falls between 150-200 mV, which is higher than the cell membrane potential which values of 50-100 mV, determining the optimal values of MMP remains challenging despite intensive research in this field. Several factors contribute to this challenge, for example, it's understood that a higher MMP corresponds to a greater energy capacity of the IMM due to its role as an electrical insulator. However, the role of various other membrane transport proteins also needs to be investigated as they are part of the membrane mass; processes like ion leakage on the IMM rely on MMP, and a high MMP contributes to oxidative stress by increasing ATP production and ROS levels. Conversely, a low MMP results in reductive stress due to decreased ATP production and ROS levels, thus, there's a need to identify specific uncouplers that can lower MMP to a certain level. This is crucial to produce the required amount of ATP while maintaining lower levels of ROS production.

This feature could be a potential therapeutic target for disease associated with aging, obesity, heart attack, etc (Zorova et al., 2018; Gottlieb et al., 2003; Izyumov et al., 2004). Duration of depolarization and hyperpolarization is also very critical in design of therapies and needs more understanding and less data is available to draw any conclusion. So, data obtained in this study suggest the effect of AM extract on IMM. However, there is a need for more supportive experiments, because even cellular mitochondria have different magnitude of MMP, although the mechanism behind this is still not clear. Additionally, while fluorescent probes might non-specifically bind to phospholipid membranes of mitochondria, the probes themselves can impact the MMP for example, could actively pump other positively charged substances, and undergo intracellular modifications, though TMRE the most suitable probe used in this study based on literature (Zorova et al., 2018; Scaduto and Grotyohann, 1999).

5.4 Effect of AM extract and hordenine on Intracellular Calcium Concentration

Besides, the effect of AM extract and hordenine were also investigated on intracellular Ca^{2+} mobilization process. Ionized Ca^{2+} is a physiologically active form and is a ubiquitous signalling messenger involved in higher mitochondrial activity regions, for

example, skeletal muscle contraction and excitation-contraction in cardiac and smooth muscles.

Intracellular free Ca^{2+} concentration varies within cells, in cytoplasm, nuclear matrix and in mitochondrial matrix and ER serves as the primary Ca^{2+} store. Mitochondrial Ca^{2+} sensors are recently identified and are found in intermembrane space that control mitochondrial Ca^{2+} uptake and mitochondrial surface targeted Ca^{2+} sensors control mitochondrial motility though the distribution mechanism of mitochondrial transport of Ca^{2+} has not been fully explored yet. Mitochondrial Ca^{2+} sensing mechanism is still not fully explored, although the role of mtCU for Ca^{2+} uptake in the inner mitochondrial membrane (pore forming component) and intermembrane space and two proteins anchor mitochondria, Miro 1 target microtubular motor protein kinesin for anterograde movement and Miro 2 dynein for retrograde movement on OMM, has been identified (Bagur and Hajnoczky, 2017).

This signalling modulates vascular smooth muscle tone, stabilise plasma membrane by binding to phospholipids in the lipid bilayer, regulating plasma membranes to sodium ions affecting excitability of tissues in addition to mitochondrial morphology. AM extract significantly affected the intracellular Ca^{2+} levels at high concentrations in MCF-7 though opposite trend is observed in the case of MCF-10-A where there is a trend of an increase in intracellular Ca²⁺ levels at higher concentration. Pottle et al., in his study compared the intracellular basal concentrations of MCF-7 and MCF-10-A and found higher expression of IGF-1 in MCF-7 which increases Ca²⁺ permeability and reduced capacity to release Ca²⁺ suggesting higher concentrations of basal intracellular Ca^{2+} in MCF-7 (Pottle et al., 2013). He further suggested the functional role of T-type Ca²⁺ channels in proliferation of MCF-7 breast cancer cells (Melgari, et al., 2022; Pottle et al., 2013). Our results possibly produced effects through T-type Ca^{2+} channel in MCF-7 which can be a possible therapeutic drug target as it is also found in various other cancers for example, gastric, esophageal, and ovarian cancers are some examples. So, AM extract has potential to design T-type Ca²⁺ channel blockers like mibefradil T-type Ca²⁺ channel blocker. Differences in effects observed in both cell lines used in this study might be due to differences in expression of IGF-I signalling.

The effect of hordenine was also evaluated across three time points and no effect is observed on the intracellular concentration in MCF-7. Mitochondrial Ca^{2+} uniporter located on the IMM is the main site for the transport of Ca^{2+} into the matrix and process takes place

either via sodium- Ca^{2+} exchange protein or via Ca^{2+} -induced Ca^{2+} release pathways. These changes may affect MMP as it initiates Ca^{2+} waves which stimulate second messenger system proteins that performs several functions such as proliferation, mitosis, differentiation and possibly co-ordinate the release of hormones in endocrine cells (Neil et al., 1997; Pratt, Hernandez-Ochoa, and Martin, 2020). Influx of Ca^{2+} into mitochondrial matrix also impact mitochondrial bioenergetics and citric acid cycle (activate isocitrate dehydrogenase). In neurons, an increase in Ca^{2+} levels both in cytosol and mitochondria coordinate neuronal activity although mobilization decelerates with age. As AM extract mobilized Ca^{2+} levels though differently in both cell lines suggest it's impact on downstream signaling pathways including protein kinase (PKA) MAPK and nuclear translocation of NF-KB that are associated to Ca^{2+} mobilization. Under hypoxic conditions intracellular Ca^{2+} concentration increases immediately which is dangerous because it causes the activation of ATPase enzymes and uncontrolled release of neurotransmitters affecting normal synaptic transmission and affecting other cellular functions such as metabolism (Supnet and Bezprozvanny, 2010; Missiaen et al.,2000).

Now, interest is growing to target mitochondrial Ca^{2+} influx and efflux for the treatment of several diseases. Here are few things need to be considered that Ca^{2+} exist in different forms that is, protein bound, ionized Ca^{2+} and complexed form also extracellular and intracellular Ca^{2+} concentrations are closely regulated by different organelles such as, ER, mitochondria and sarcoplasmic reticulum. Mitochondria offer buffering Ca^{2+} activity, most likely to the organelles near OMM. However, targeting this mitochondrial communication is a very interesting area to design new drugs for the treatment of hypocalcaemia caused by hypothyroidism, renal diseases and hypercalcemia caused by bone resorption caused by hyperthyroidism and cancer, but this still needs intensive research (Slater and Cleland, 1953; Boyman, Greiser and Lederer, 2021; Lewis, 2023; Pallagi et al., 2020).

5.5 Effect of AM extract and hordenine on OXPHOS

In addition, the mitochondrial respiration rate of MCF-7 and MCF-10-A cells was also measured after treatment of cells with AM extract and hordenine for 24hr. As the mitochondrion engages in homeostatic processes because cellular homeostasis depends on the sustained ATP generation, OCR is also recognized as a traditional measure of mitochondrial functions in pathophysiology (Zhang et al., 1993).

Oligomycin inhibitor of ATP-synthase helps to indicate O₂ consumption directly coupled to ATP generation. Carbonyl cyanide-4-(trifluoromethoxy) phenylhydrazone (FCCP), a mitochondrial uncoupler, measures maximal OCR and antimycin A (complex III inhibitor) and rotenone (complex I inhibitor) together affects the ETC. In our study, hordenine has shown no significant effect on OCR in MCF-7 and MCF-10-A cell lines after exposure of 24hr at all concentrations tested (Lawrence et al., 2017; Clark, 1956). The Seahorse analyzer measures and quantifies the rate of ATP production from mitochondria and quantifies mitochondrial dysfunction in complex diseases, and is widely implemented to study the effects of natural compounds, including cytotoxicity study of bio-active components of licorice in HepG2 cells (Qian, et al., 2019), the marine sponge sesquiterpene on lung cancer cell apoptosis (Cheng et al., 2019; Mahmoud, May, and Mohammad, 2021; Rohlena et al., 2011), anti-proliferative effects of T-3-BP-AuNPs (mitochondria-targeted gold nanoparticle T-3-BP-AuNP) (Marrache and Dhar, 2015) and selective cytotoxicity of the natural compound acteoside.

AM extract was also tested on mitochondrial respiration of MCF-7 and MCF-10-A. AM extract has affected mitochondrial respiration which reduced from low to high concentrations on parameters tested such as basal respiration, non-mitochondrial respiration, ATP production, proton leak, though none of the effect reached significance in MCF-7. In the case of MCF-10-A it has significantly affected various parameters of mitochondrial respiration. This assay further shed light on the overall bioenergetics of the MCF-7 and MCF-10-A cell lines after sequential induction of the OXPHOS inhibitors. Drop in ATP linked OCR after addition of oligomycin (ATP synthase inhibitor) suggest inhibition of proton flux and this increases proton gradient across the inner mitochondrial membrane that prevents electron transport across through complexes I-IV eventually reducing oxygen consumption. As high concentrations of AM extract have severely affected ETC so reduction in other parameters indicate mitochondrial integrity is compromised which is previously observed in MMP assays. So, defects in ETC resulted in lower bioenergetics of both cell lines. Similarly, hordenine has not significantly affected most of the parameters after 24hr of exposure some effects observed could be random change as in case of spare reserve capacity. Based on other reported studies high basal respiration indicates high ATP turnover in cell, increased proton leak, increased non- mtROS and high proton leak indicates increased UCP activity that also damages inner membrane. Furthermore, high maximal capacity suggests high substrate availability, increased mitochondrial mass and good ETC integrity and finally high non-mitochondrial respiration suggests increased cytosolic ROS levels. In this study no significant changes at lower concentrations of AM extract were observed on mitochondrial respiration and differences in parameters observed are due to several factors and need further testing to shed light on specific cellular pathways affected.

According to Warburg theory proposed around 1930, tumour cells were believed to rely on glycolysis even under oxygen rich conditions. So, it was thought that mitochondria might be dysfunctional in cancers, and this was supported by certain studies of renal, melanoma and non-small cell lung cancer, where reduced OXPHOS was observed. However, this hypothesis was later challenged because upregulated OXPHOS, as in case of pancreatic cancer, was also observed in certain cases, whereas some cancer cells use mitochondrial fatty acid oxidation as a source of energy as seen in triple negative breast cancer. So, targeting OXPHOS to deprive cells of energy may be a questionable strategy because mitochondria may reprogram, and cancer cells use glycolysis as a source of energy under stress induced by ETC modulators in this study. There is need to find drug molecules that can specifically target OXPHOS sites as in case of metformin which is used to target complex I. So far, idea of targeting glycolytic pathway along with ETC is proposed and under study (Warburg, 1956; Ward and Thompson, 2012; Evans et al., 2005; Wheaton et al., 2014; Greene, Segaran and Lord, 2022).

5.6 Effect of AM extract and hordenine on mitochondrial morphology

Also mitochondrial morphological analysis is performed to observe any changes, after hordenine treatment mitochondrial morphology is not altered only small, interconnected mitochondrial networks in MCF-7 and MCF-10-A are formed that points to the process of fission rather fusion. Further AM extract at low concentration tested caused no change in the morphology of mitochondria of both cell lines (MCF-7 and MCF-10-A) used in this experiment (Rafelski, 2013; Rohlena et al., 2011).

Findings of this study greatly contributed to the understanding of the effect of AM extract on MCF-7 for any anticancer effect and MCF-10-A for nutritive value though mechanism through which these effects are produced are still not clear, and more work is considered necessary (Perini et al., 2018; Freitas et al., 2017). Alqasoumi reported in 2008 in his study that AM extract administration of 660 mg/kg had significant hepatoprotective effects, and significantly decreased the levels of serum glutamate oxaloacetate transaminase and serum glutamate pyruvate transaminase in mice. In another study, AM extract at a concentration of 4 μ g/ml produced skeletal muscle relaxant effect in frogs and a dose of 1.6 g/kg produced sedative effects, while 1 g/kg induced bradycardia and petroleum, while other extract has shown cytotoxic effects in dose and time dependent in human acute myeloid leukemia cell line (HL-60) (16.0 and 22.0 μ g/ml) (Abdellatif et 1., 2014; Gargoum et al., 2013; Sulaiman, 2013). AM is also a main ingredient of a Uyghur herbal preparation Abnormal Savda Munziq (ASMq) used in Uyghur medicine. Aikemu in 2012 investigated its antitumor activity and suggested affects are possibly through modulation of immune system (Aikemu et al., 2012; Wang et al., 2015).

Insulin-like growth factor type 1 receptor (IGF-1R) is found in human breast cancers however its expression in triple negative breast cancer is relatively low. IGF-1 is mostly secreted by stromal cells and acts through paracrine signaling to adjacent epithelial tumor cells and once the disease is well established within the primary breast tumor microenvironment, IGF-1 autocrine (within epithelium) and endocrine (via the systemic circulation) activity facilitates disease progression and metastasis respectively. Within the new tumor location, the interplay between metastasized breast cancer cells and host tissue cells drives the last to a more cancerous phenotype via IGF-1 paracrine signaling. PI3K/AKT and RAS-MAPK axes are two well-established downstream pathways of IGF/insulin signalling (Mauro et al., 2000; Peruzzi et al., 1999; Pan et al., 2016; Paradies et al., 2019). Therefore, modulation of AKT phosphorylation is potentially a therapeutic strategy for several diseases, especially inflammation of various organs. Although the exact mechanism has not yet been reported but some molecular docking studies have shown strong binding of hordenine with AKT which suggest hordenine can be a potential inhibitor of AKT signalling pathways which is linked to glucose metabolism, protein synthesis, apoptosis and cell survival and can serve as therapeutic target in breast cancer. IGF signaling is also critical for the maintenance of cellular stress homeostasis particularly ER stress which is caused by misfolded proteins and lead to ROS production so it can also be used to activate endoplasmic

stress response pathways to prevent apoptosis by inhibiting IGF signalling in breast cancer cells furthermore increasing adaptability and cell survival (Gao, et al., 2016).

AM extract has shown different effects compared to hordenine. As extracts are mixtures of several compounds and it is very difficult to find out which constituent is responsible for the purported biological activity. Urabee in 2021 reported in his chemical investigation of 98% ethanolic extract of AM the presence of more than 30 compounds (antioxidant compounds, essential oils, sterols, coumarins, vitamins and phenolic constituents) (Urabee et al., 2021). So, mass spectroscopic analysis was also carried out to find out about the bioactive compounds of the AM extract. HPTLC and LC-MS analysis has shown the presence of more than one compound in the AM extract and suggests further testing of the other compounds to interpret observed effects more appropriately.

Future Work

To address the limitations of this study some future work is required to explore pathways through which *Alhagi maurorum* and hordenine produced effects on mitochondria.

At higher concentrations, *Alhagi maurorum* induced cell death, while reducing ROS levels and altering OXPHOS. Thus, there is a need to examine changes in processes such as apoptosis, mitophagy, glycolysis, citric acid cycle of these cells. A complete footprint of mitochondrial morphology is required to complete morphological analysis in addition to quantification of mitochondrial markers of apoptosis for example mitofilin, which is the most abundant protein and maintains morphology of cristae, AIF, P53 (up-regulator of apoptosis), cytochrome c, Kras that require ROS for proliferation, VDAC that allows entry and exit between cytosol and mitochondria and PDK4 enzyme (metabolic switch between glycolysis and citric acid cycle) through Real Time Polymerase Chain Reaction (RT-PCR). Investigating the effects of *Alhagi maurorum* on mitophagy processes could be beneficial to remove dysfunctional mitochondria. Similarly, based on given claims of anti-inflammatory properties of this plant, a complete *in vitro* and *in vivo* analysis of key inflammatory markers may be of interest. ER being the most significant site of Ca^{2+} storage and *Alhagi maurorum* altered mitochondrial Ca^{2+} concentrations in both cell lines so, there is need to study interplay between the mitochondrion and ER with respect to Ca^{2+} .

Hordenine, an important constituent of *Alhagi maurorum* was also tested in this study and showed no prominent effects on mitochondrial functions, but with some specific differences such as effect on ROS levels after 6hr of exposure in both cell lines used in this study. These differences may arise from what is known as the additive, synergistic or antagonistic effects as seen in case of entourage effect observed in cannabis where mixtures of compounds are associated with modulation of the principal bioactive compound and work by Pan, showed combining baicalein and taxol induce mitochondria mediated apoptosis (Pan et al., 2016).

Mass spectroscopic analysis of *Alhagi maurorum* clearly indicates the presence of more than one compound in the mixture. Plant extracts with ability to regulate mitochondrial functions can offer new opportunities to treat various complex diseases that require more than one bioactive compound and to address issues of drug resistance and side effects by choosing combinations of different bioactive compounds (Caesar and Cech, 2019; Ferber et

al., 2020). Furthermore, there is also needed to carry out LC-MS analysis of hordenine to compare with the fragmentation pattern of hordenine observed in the LC-MS analysis of *Alhagi maurorum* extract.

Data gathered from this study must be supported by the above-mentioned experiments to get further in-depth insight into the observed effects and potential cellular pathways involved. This will open several therapeutic targets for the treatment of pathologies mainly due to altered metabolism, for example, diabetes, cancer, obesity, and neurodegenerative disorders.

Lastly, this study has utilized only breast cell models which are very valuable for the preliminary investigation due to the ease of handling, replication and consistency which makes them good model for comparison of results between experiments. However, it is well known there are several disadvantages such as possible genetic and phenotypic drift over extended passage numbers of cell lines, possible contamination, and lack of key characteristics of cancers such as complex microenvironment and cancer stem cells so, it is worth looking at more complex models, such as 3D cellular models.

Conclusion

In conclusion, the results from my PhD work indicate *Alhagi maurorum* has potential to regulate mitochondrial functions and is a potential source of compounds targeting mitochondrial functions. At this stage my work focused on the metabolic aspect of mitochondrial functions only and it would be interesting to undertake in depth study of proteomics and transcriptomics to get better understanding of the molecular mechanism underpinning the effects of *Alhagi maurorum* and its other constituents on associated mitochondrial and cellular functions. This work provided new insight into the effects of *Alhagi maurorum* on mitochondrial functions and will attract more in-depth study and may contribute to the design of more effective and targeted drug candidates considering metabolic flexibility of the cellular system.

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