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# ROLE OF AUTOPHAGY IN ACRIDOCARPUS ORIENTALIS-INDUCED ANTI-BREAST CANCER ACTIVITY

Suhib Hisham Ahmed Saeed Altabbal

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United Arab Emirates University

<span id="page-1-0"></span>College of Science

Department of Biology

# ROLE OF AUTOPHAGY IN ACRIDOCARPUS ORIENTALIS-INDUCED ANTI-BREAST CANCER ACTIVITY

Suhib Hisham Ahmed Saeed Altabbal

This thesis is submitted in partial fulfilment of the requirements for the degree of Master of Science in Molecular Biology and Biotechnology

Under the Supervision of Dr. Yusra Al Dhaheri

November 2021

#### **Declaration of Original Work**

<span id="page-2-0"></span>I, Suhib Hisham Ahmed Saeed Altabbal, the undersigned, a graduate student at the United Arab Emirates University (UAEU), and the author of this thesis entitled "*Role of Autophagy in Acridocarpus Orientalis-Induced Anti-Breast Cancer Activity*", hereby, solemnly declare that this thesis is my own original research work that has been done and prepared by me under the supervision of Dr. Yusra Al-Dhaheri, in the College of Science at UAEU. This work has not previously formed the basis for the award of any academic degree, diploma or a similar title at this or any other university. Any materials borrowed from other sources (whether published or unpublished) and relied upon or included in my thesis have been properly cited and acknowledged in accordance with appropriate academic conventions. I further declare that there is no potential conflict of interest with respect to the research, data collection, authorship, presentation and/or publication of this thesis.

 $\overline{\mathcal{W}}$ 

Student's Signature: Date: \_11/11/2021

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#### **Abstract**

<span id="page-7-0"></span>Breast cancer is the most frequently diagnosed cancer in women worldwide. Triple Negative Breast Cancer (TNBC), which lacks the expression of the hormonal Estrogen Receptor (ER) and Progesterone Receptor (PR), and the amplification of Human Epidermal Growth Factor Receptor 2 (HER2), is not responsive to the hormonal therapy. Therefore, chemotherapy and radiotherapy, which cause severe side effects, are the current available choices to treat TNBC. Hence, there is an urgent need to find new therapeutic choices for TNBC. It is estimated that 50% of all drugs in clinical use during the 21st century are natural products and plants derived. *Acridocarpus orientalis* is a rare plant used in folk medicine to treat many health conditions. The aim of this study is to evaluate the anti-cancer activity of *Acriodocarpus orientalis* Ethanoic Extract (AOEE) against TNBC cell line MDA-MB-231, and to investigate the molecular mechanisms underlying its activity. The results revealed that AOEE inhibits cell proliferation in a concentration- and time-dependent manner. The antiproliferative effect of AOEE was found to be accompanied with the induction of cell cycle arrest at the G1/S phase. These changes were accomplished with upregulation of  $p21^{WAF1}$  and  $p27^{Kip1}$ , downregulation of PCNA, Cyclin D1, phospho-Rb. Moreover, AOEE induces autophagy through upregulation of autophagy related proteins LC3-II, Beclin-1, p62. Cellular senescence was induced in AOEE treated MDA-MB-231 cells confirmed by p16 upregulation and senescence-associated β-galactosidase (SA-β-gal) expression in the treated cells. AOEE induced activation of ERK and p38 pathways, which might be involved in autophagy induction and senescence. In conclusion, AOEE inhibits the proliferation of MDA-MD-231 breast cancer cells through induction of autophagy, cellular senescence and DNA double stranded breaks, suggesting that *Acridocarpus orientalis* could be a potential source for novel chemotherapeutic agents against TNBC.

**Keywords**: *Acridocarpus orientalis*, triple-negative breast cancer, autophagy, cell cycle arrest, cellular senescence, DNA damage.

### **Title and Abstract (in Arabic)**

# <span id="page-8-0"></span>**دور اإللتهام الذاتي في التأثير المضاد لسرطان الثدي لنبات القفاص** *Acridocarpus orientalis*

# **الملخص**

يعتبر سرطان الندي أكثر أنواع السرطانات تشخيصاً بين النساء في جميع أنحاء العالم. سرطان الثدي السلبي الثالثي ) TNBC)، الذي يفتقر إلى مستقبالت هرمون االستروجين ) ER )ومستقبل البروجسترون )PR)، وتضخيم عامل نمو البشرة البشري 2 ) 2HER)، ال يستجيب للعالج الهرموني، والعلاجات الكيميائية التي تسبب آثارًا جانبية شديدة هي الخيارات الحالية الوحيدة المتاحة لعالج سرطان الثدي السلبي الثالثي. لذلك، هناك حاجة ملحة إليجاد أهداف جديدة وخيارات عالجية جديدة لـ سرطان الثدي السلبي الثالثي. تشير االدراسات إلى أن %50 من جميع الأدوية المستخدمة في القرن الحادي و العشرين هي منتجات طبيعية أو مشتقة من النباتات. (نبات القفاص( *orientalis Acridocarpus* هو نبات نادر يستخدم في الطب الشعبي لعالج العديد من الحاالت الصحية. الهدف من هذه الدراسة هو تقييم التأثير المضاد للسرطان لـ مستخلص األوراق ل نبا القفاص ضد خاليا سرطان الثدي السلبي الثالثي -231MB-MDA، والكشف عن اآلليات الجزيئية الكامنة وراء التأثيرالمضاد للخاليا السرطانية. أظهرت نتائج هذه الدراسة أن مستخلص الأوراق ل نبات القفاص يمنع تكاثر الخلايا بطريقة تعتمد على التركيز والوقت. إضافة لذلك, أظهرت الدراسة أن التأثير المضاد لتكاثر خلايا السرطان الثدي السلبي الثلاثي مرتبطًا بإيقاف دورة الخلية في مرحلة S1/G. تم تحقيق هذ ا التأثير من خالل تحليل إنتاج البروتينات التي تنظم دورة انقسام الخلية مثل بروتين 21p 27,p الذي زاد انتاجهم مع استخدام المستخلص ، والذي أدى إلى تقليل تنظيم بروتينات PCNA و 1D Cyclin و Rb-phospho. باإلضافة إلى ذلك، أظهرت نتائج الدر اسة أن مستخلص الأوراق ل نبات القفاص يحفز الالتهام الذاتي الذاتي والشيخوخة الخلوية في خاليا -231MB-MDA. على المستوى الجزيئي, أظهرت النتائج أنه تم زيادة إنتاج بروتينات II3-LC و -1Beclin و 62p المرتبطة بااللتهام الذاتي، و زيادة إنتاج 16p و galactosidase-β( gal-β-SA )المرتبط بالشيخوخة في الخاليا المعالجة. كما أظهرت النتائج أن مستخلص نبات القفاص ينشط مسارات ) ERK )و 38p، التي قد تكون مرتبطة بتفعيل الالتهام الذاتي والشيخوخة الخلوية. في الختام، يمنع مستخلص الأوراق ل نبات القفاص تكاثر خلايا سرطان الثدي MDA-MD-231، ويرتبط التأثير بتحريض الالتهام الذاتي والشيخوخة الخلوية وضرر للحمض النووي، مما يشير إلى أن نبات القفاص *orientalis Acridocarpus* يمكن أن يكون مصدًرا غنيا لمركبات نشطة جديدة لعالج سرطان الثدي السلبي الثالثي TNBC.

**مفاهيم البحث الرئيسية** : سرطان الثدي الثالثي السلبي، االلتهام الذاتي، توقف الدورة الخلوية، الشيخوخة الخلوية، تلف الحمض النووي.

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<span id="page-11-0"></span>**Dedication**

*To my beloved parents and family*

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#### **Chapter 1: Introduction**

#### <span id="page-18-1"></span><span id="page-18-0"></span>**1.1 Overview of Cancer**

Cancer is a large group of diseases with more than 277 cancer types is known up to date. According to 2020 global cancer statistics, prostate, breast, and colorectum cancers for both sexes are among the highest prevalence cancer types (Sung et al., 2021). Cancer is the second leading cause of death worldwide and expected be the leading cause of death by 2060. According to the American cancer society, growing of the global burden to cancer is expected by 2040 to reach 27.5 million new cases and 16.3 million deaths. The growing in cancer mortality and morbidity rates is explained by the growing and aging of the global inhabitance (American Cancer Society, 2018). Among all human diseases, cancer causes the highest rates of clinical, social, and economic burden (Mattiuzzi & Lippi, 2019). Negative impact on the quality of life in cancer patients by its symptoms, treatment and duration is definite (Lewandowska et al., 2020; Nayak et al., 2017). Cancer cells are transformed cells characterized by the unlimited growth potential. Other characteristics of cancer cells distinguish them from normal cells include their ability to migrate and invade the surrounding or distant tissues, resist cell death, sustain proliferative signaling, induce angiogenesis, and avoid growth suppressors or immune destruction (Markopoulos et al., 2017; Senga & Grose, 2021).

Cancer in solid organs forms tumor, which can be benign or malignant. Benign tumors are localized in the affected tissue without noticed spreading to other adjacent or distant tissues and they tend to grow slowly. On the other hand, malignant or cancerous tumors grow rapidly and can metastasize to the tissues and lymph nodes surrounding

them. Some types of benign tumors such as colon polyps can transform to malignant ones, making their diagnosis, early monitoring and treatment a priority (Patel, 2020). Metastasis is the spread of cancer cells from the primary site of tumor to the adjacent and surrounding tissues or to distant organs in the advance cases (Seyfried & Huysentruyt, 2013). The most common sites for the metastasis of most cancer types are liver, lungs, brain, and bones (Patel, 2020). During metastasis, cancer cells invade from their primary site to the surrounding tissue. Colonization in the distant organ after entering the circulation and survival from the immune cells must occur for cancer cells to metastasize (Elia et al., 2018). Mutations, which change the cell function and cause dysregulation in life cycle, are believed to have a major role in cancer pathogenesis and progression (Hassanpour & Dehghani, 2017). Cancer is multifactorial disease and several factors can interact to initiate and develop different types of cancer (Wu et al., 2016). Table 1 summarize the most common identified causes of cancer.

Viruses	Human T-cell lymphotrophic virus Human immunodeficiency virus Hepatitis B virus Hepatitis C virus Human papillomavirus Epstein-Barr virus
	Human herpes virus $\bullet$
<b>External factors</b>	Sunlight
	Tobacco
	Alcohol
	Salted fish
	Wood dust
	Radiation rays $\bullet$
Food	Beta carotene
	Red meat
	Processed meats
	Low fiber diets

<span id="page-19-0"></span>Table 1: Common causes of cancer (Blackadar, 2016; Hassanpour & Dehghani, 2017)

Lifestyle		Obesity			
	$\bullet$	Increased adult height			
		Non-breast feeding			
		Sedentary lifestyles			
Pharmaceuticals			Nitrogen mustards $(\beta$ -chloroethyl	sulphides	and
carcinogenic and		amines)			
chemicals		Melphalan			
		<b>Busulfan</b>			
		Cyclophosphamide			
		β-naphthylamine			

Table 1: Common causes of cancer (Blackadar, 2016; Hassanpour & Dehghani, 2017) (Continued)

#### <span id="page-20-0"></span>**1.2 Overview of Breast Cancer**

Breast cancer is the most frequently diagnosed cancer in women worldwide. According to the Global Cancer Observatory, breast cancer accounts for 11.7% of new cancer cases in 2020 worldwide (WHO, 2020). More than 248,000 new cases were estimated in the United States during 2021, with more than 2,600 cases of them in men (Siegel et al., 2021). Although the incidence of breast cancer increases with advanced aging, breast cancer incidence rates among women aged 20 to 29 years, increased by 2% every year in the last 10 years, whereas the annual increasing in the incidence of breast cancer among women in their 30s for the same period is 0.2%. In adolescents, no increase in the incidence rates was noticed in the last 10 years (Miller et al., 2020; Society, 2018). While the incidence rate of breast cancer is higher in developed countries, the mortality rate of breast cancer is higher in less developed countries (Momenimovahed & Salehiniya, 2019). The development of more invasive forms of breast cancer is affected by race. Black women are at higher risk to develop invasive cancer from pre-malignant breast lesions (Dania et al., 2019).

#### <span id="page-21-0"></span>**1.2.1 Types and Clinical Staging of Breast Cancer**

Breast cancer can be divided into several subtypes based on the expression of several molecular markers. Cellular expression status of Estrogen Receptor (ER) and/or Progesterone Receptor (PR) along with the amplification of Human Epidermal Growth Factor Receptor-2 (HER2) can divide the breast cancer into 3 types: HR+/HER2-, which express either estrogen or progesterone receptors; HER2+, which express HER2 receptor; and Triple-Negative Breast Cancer (TNBC), which lacks the expression of the two hormonal receptors and the amplification of HER2 (Waks & Winer, 2019).

Breast cancer clinical staging depends on many anatomical features including: the primary tumor extent and size, presence and extent of lymph node involvement, and presence or absence of metastases. TNM system created by the American Joint Committee on Cancer (AJCC), which depends on the anatomical spread of the disease, is the most used clinical staging system for breast cancer worldwide. In TNM system, (T) refers to tumor size and take value from 0-4, (N) refers to nodal status and can take value from 0-4 based on the lymph nodes affected by the cancer cells, and (M) refers to the presence of metastasis and can be 0 or 1. Other staging systems depend on both the TNM staging and the expression status of ER/PR and HER2 can be used to stage breast cancer, the American Joint Committee on Cancer (AJCC) staging system is one example of such staging systems (Boster et al., 2020; Giuliano et al., 2018).

Based on the TNM staging system, breast cancer can be simply divided into 4 stages as below (Boster et al., 2020).:

- Stage 0 represents cancer cells without involvement of basement membrane of breast tissue.
- Stage I represent small invasive tumor without lymph node involvement.
- Stage II represent small invasive tumor with involvement of regional lymph nodes.
- Stage III represent large invasive tumor with extensive nodal involvement is
- Stage IV or metastatic disease refers to breast cancer with metastases to distant organs.

#### <span id="page-22-0"></span>**1.2.2 Triple Negative Breast Cancer**

Triple negative breast cancer is the most aggressive subtype of breast cancer. Despite the low incidence rates of TNBC among breast cancer subtypes, mortality rates for TNBC are remain the highest and the prognosis is the poorest among the other subtypes. Furthermore, more complex pathogeneses are believed to control the overgrowth of TNBC cells compared to the other breast cancer subtypes (Lee  $\&$ Djamgoz, 2018; Waks & Winer, 2019). Since TNBC lack the expression of ER, PR and HER2 amplification, TNBC is not responsive to the hormonal therapy. The conventional chemotherapies, which cause severe side effect, are the only current available choices to treat TNBC. Therefore, new targets and therapeutic choices are needed to control TNBC (Waks & Winer, 2019). Additionally, based on the gene expression profile, TNBC can be subdivided into six subtypes, each subtype displays unique gene expression and ontologies. Surprisingly, each subtype has specific features, makes it susceptible to specific chemotherapies not the others. This can be explained by the specific signaling pathways controlling the overgrowth and excessive proliferation for each subtype (Diana et al., 2018; Lehmann et al., 2011). Table 1 summaries the main 6 subtypes of TNBC as divided based on the gene expression status.

Many germline mutations associated with TNBC were identified. *BRCA1/2* gene mutation, which is tumor suppressor gene related to the Homologous Recombination (HR) repair of double-strand DNA breaks, is highly prevalent and its frequency reaches up to 18.2% in TNBC. The current guidelines recommended *BRCA1/2* gene status screening for women with TNBC diagnosed at age below 60 years regardless the positive cancer family history. Other germline mutations such as *PALB2,* and *FANCM* genes mutations were identified among TNBC patients (Hahnen et al., 2017). Mutation in the tumor suppressor gene P53 in human cancer cells is also believed to be associated with the pathogenesis of TNBC. p53 has a key role in the regulation of cell cycle, senescence, and apoptosis in cancer cells (Zeng et al., 2019).

<b>TNBC Subtype</b>	<b>Characteristics</b>	
Basal-like 1 (BL1)	- Heavily enriched in cell cycle-related genes and pathways	
	involved in the repair of DNA damages	
	- High $ki-67$	
Basal-like 2 (BL2)	- Enriched in growth factor signaling, such as: epithelial	
	growth factor (EGF), MET and insulin growth factor	
	receptor (IGF1R) pathway	
	- Enriched in signaling of glycolysis and gluconeogenesis	
Immunomodulatory	- Enriched for gene involved in immune cell processes, such	
(IM)	as: B, natural killer cell and T cell signaling; cytokine	
	signaling; chemokine receptors and ligands; complement	
	cascade and antigen presentation	
	- High levels of infiltration of immune cells defined tumor-	
	infiltrating lymphocytes (TILs)	

<span id="page-23-0"></span>Table 2: Characteristics of different subtypes of TNBC based on the genes expression (Diana et al., 2018)

Table 2: Characteristics of different subtypes of TNBC based on the genes expression (Diana et al., 2018) (Continued)

<b>TNBC Subtype</b>	<b>Characteristics</b>		
Mesenchymal-like	- High expression of vimentin		
(M)	- Decreased of expression of E-cadherin		
	- Activation of:		
	c-MET		
	epithelial growth factor (EGF)		
	mammalian target of rapamycin (mTOR)		
	fibroblastic growth factor (FGF) $\bullet$		
	insulin growth factor (IGF) $\bullet$		
	transforming growth factor β (TGF-β)		
	$Wnt/\beta$ catenin pathways		
Mesenchymal	- Similar to M with enrichment in genes involved in		
stem-like (MSL)	angiogenesis, including VEGFR2 and some components of		
	immune signaling		
	- High expression of stem cells genes		
	- Low expression of proliferation genes and epithelial-		
	related genes involved in the maintenance of cellular		
	junction, such as claudin (claudin-low breast cancer)		
Luminal androgen	- Androgen receptor (AR) positivity		
receptor $(LAR)$	- Enrichment in hormonally regulated pathways, especially		
	steroid synthesis and metabolism		

# <span id="page-24-0"></span>**1.2.3 Diagnosis of Breast Cancer**

Imaging techniques including Mammography, Magnetic Resonance Imaging (MRI), Positron-Emission Tomography (PET), Computed Tomography (CT), and Single-Photon Emission Computed Tomography (SPECT) are the main techniques used to have definitive diagnosis for breast cancer (Jafari et al., 2018). Mammography is considered as the gold standard imaging modality for breast cancer diagnosis and proven to reduce mortality in breast cancer patients. MRI is considered more sensitive technique to detect smaller breast masses at earlier stage, duo to the high resolution of the results (Wellings et al., 2016). Biological markers including CA 27-29, CA 15-3, CA27.29, carcinoembryonic antigen, tissue polypeptide specific antigen, p53, cathepsin D, cyclin E, nestin, Estrogen Receptor (ER), Progesterone Receptor (PR) and HER-2 are usually expressed in breast cancer patients and can be used as tool for better diagnosis and monitoring of the disease (Kabel, 2017). Depending on the risk of having breast cancer, women with *BRCA1* or *BRCA2* mutations are encouraged to screen for breast mass by clinical breast examination and mammography starting at age of 30 years and additional screening with MRI is recommended for higher risk patients (Jafari et al., 2018). New investigational biomarkers including Epidermal Growth Factor Receptor (EGFR) and 8-Hydroxy-2'-deoxyGuanosine (8-OHdG) might be used in the future to detect early stages of noninvasive breast cancer with high sensitivity (Bayo et al., 2018; Boster et al., 2020; Eldin et al., 2019).

#### <span id="page-25-0"></span>**1.2.4 Treatment of Breast Cancer**

Treatment choices for breast cancer are highly variable. The most important factors determine the choice of treatment are the disease subtype and the disease stage. Additionally, the treatment goal differs among the disease stage at the time of diagnosis. Total cure from the disease is the treatment goal in stage I to III of the disease. Choices with curative intent include breast surgery, radiotherapy, and adjuvant/neoadjuvant systemic treatment (cytotoxic chemotherapy, endocrine treatment, and targeted agents). Furthermore, supportive care of cancer-related pain and chemotherapy related cytotoxicity and side effects are essential (Mutebi et al., 2020). In contrast, palliative care strategies including pain management, as well as psychosocial and spiritual support are the treatment choices used in the metastatic breast cancer. The treatment choices used in stage I to III can also be used in the metastatic breast cancer with specific goals. For example, surgery can be used to remove breast masses with very few metastatic deposits; and hepatic surgery can be done if the disease presenting with liver metastases to prolong overall survival. In contrast, removal of primary tumor in metastatic disease is controversial and usually not recommended for cure goal. Radiation can be used if the primary breast mass is concurrent with few malignant cells. In case of bone metastasis, radiotherapy is used to palliate the symptoms associated with it. Interestingly, radiotherapy have curative role in case of Central Nervous System (CNS) metastasis. Hormonal therapy and chemotherapy may be used in metastatic breast cancer to prolong the overall survival and quality of life; depending on the subtype of the cancer cells (Sambi et al., 2019).

# <span id="page-26-0"></span>**1.3 Phytomedicine**

#### <span id="page-26-1"></span>**1.3.1 Overview of Phytomedicine**

Phytomedicine is the traditional practice that involves the use of plants and herbs as medications. Before the scientific revolution, ancients' cultures relied on the error and trial to investigate the medicinal effects of the plants. In the ongoing era, several chemical, biological techniques and methods can be used to explore the active compounds responsible for the potential activities of the medicinal plants for the treatment of the diseases, making the plants rich sources for finding novel plant-based medications (Nigam, 2021). Many widely used medications such as Aspirin, Atropine, Codeine, Ephedrine and Digoxin are either plant derived or discovered after the study and analysis of the traditionally used plants (Saad et al., 2017). Its estimated that 50% of all drugs in clinical use during the  $21<sup>st</sup>$  century are natural products and plants derived (Shakya, 2016).

#### <span id="page-27-0"></span>**1.3.2 Biological Activities of Medicinal Plants**

Based on the active compounds synthesized by the plant, medicinal plants can have wide biological activities including antispasmodic, antimalarial, analgesic, diuretic, antiviral, anthelmintic, antibacterial, anticancer, antimalarial and anti-inflammatory properties (Shakya, 2016).

Acute diseases such as myocardial infarction and chronic diseases such as osteoarthritis, rheumatoid arthritis, inflammatory bowel disease, atherosclerosis, chronic heart failure and cancers are associated with chronic inflammation and autoimmune responses. *Zingiber officinale Roscoe* (known as Ginger), *Curcuma longa Linn* and *Camellia sinensis* are few examples of medicinal plants containing potential anti-inflammatory active compounds such as Curcumin and epigallocatechin-3-gallate with varied molecular mechanisms (Tasneem et al., 2019). Some medicinal plants are characterized by their antipyretic and analgesic activities in addition to the antiinflammatory properties they have. *Asparagus officinalis*, *Avena sativa*, *Brassica rapa*, *Capsicum annuum* and *Capsicum frutescens* are few examples of them (Al-Snafi, 2016).

More than 80 medicinal plants as reviewed by Jacob and Narendhirakannan (2019) found to have several effects on diabetes mellitus. Anti-diabetic, anti-hyperglycemic, hypoglycemic and insulin mimetic properties were screened from such plants. Additionally, such plants gained by the anti-lipidemic effects of them can reduce diabetes mellitus mediated hyperlipidemia. Medicinal plants contain active compounds effective for diabetes mellitus include *Coriandrum sativum L.* (Coriander), *Zingiber officinale Roscoe* (Ginger) and *Syzygium cumini (L.) Skeels* (Black plum) (Jacob & Narendhirakannan, 2019).

Other diseases causing high mortality rates such as atherosclerosis can be prevented by the active compounds in some medicinal plants. Phenols, flavonoids, and antioxidants are examples of plant-derived compounds proven to be effective in preventing atherosclerosis via reducing the levels of cholesterol and excessive free radicals' production, which are responsible for the development of vascular plaque, and by decrease the vascular resistance responsible for the development of the disease. Such plants include *Gynostemma pentaphyllum, Triticum aestivum* and *Panax* ginseng (Qadir et al., 2018; Sedighi et al., 2017).

# <span id="page-28-0"></span>**1.3.3 Medicinal Plants for Cancer Prevention and Treatment**

Numerous numbers of plants were studied and screened for potential compounds to prevent and treat cancer. Initial phase of carcinogenesis can be suppressed or reversed using natural plants-derived biological agents leading to prevention of tumorigenesis. Capsaicin derived from *Capsicum* (chili pepper), catechines derived from green tea and lycopene derived from tomatoes are examples for natural plant derived active compounds effective both for cancer prevention and treatment (Ranjan et al., 2019). Furthermore, many currently used chemotherapeutic agents used for many types of cancer are either natural products isolated from various plants or semisynthetic products based on the modification of the original compounds isolated from the plants. Vincristine and Vinblastine extracted from *Vinca rosea* are vinca alkaloids chemotherapeutic agents, both used for the treatment of various types of cancer such as acute lymphoblastic leukemia, lymphomas, breast cancer and others. Similarly, Paclitaxel is classified as taxane chemotherapeutic agent extracted from *Taxus brevifolia* and current guidelines include it within the treatment choices for several solid tumors such as lung, ovarian and breast cancers (Dragoi & Alexandru, 2020).

#### <span id="page-29-0"></span>**1.3.4 Molecular Mechanisms of Anti-Cancer Activity of Medicinal Plants**

Due to the high diversity of the active compounds found in the medicinal plants, various molecular mechanisms and pathways are involved in their anti-cancer pharmacological effects. In addition to the induction of cell death pathways such as apoptosis and necrosis, other pathways involved in the inhibition of cancer cells proliferation, the active compounds in the medicinal plants can inhibit invasion and metastasis.

#### <span id="page-29-1"></span>**1.3.4.1 Apoptosis**

Apoptosis is the programmed cellular death (type I cell death), which is activated by cellular stresses such as DNA damage or Endoplasmic Reticulum (ER) stress (Green & Llambi, 2015). Apoptosis induction can be stimulated by external (receptor mediated) or internal mechanisms. In the external pathway, apoptosis is activated by the interaction of the proapoptotic signaling molecules with specific receptors on the surface of the cells, whereas proapoptotic proteins are released from the internal organelles or expressed by specific genes in the internal pathway. Both internal and external mechanisms of apoptosis activate specific cytosolic caspase cascade to induce intracellular organelles degradation or to prepare the cell for phagocytosis (Savitskaya & Onishchenko, 2015). Figure 1 summarizes the major steps in the external and internal mechanisms of apoptosis (D'Arcy, 2019). *Garcinia quaesita* (Fruits Hexane) extract, a commonly used plant to flavor food in Sri Lanka., is one example of phytomedicinal plants shown to inhibit the growth of TNBC MDA-MB-231 cells via induction of apoptosis (Colamba Pathiranage et al., 2020).



<span id="page-30-0"></span>Figure 1: Major steps in the external and internal pathways of apoptosis (D'Arcy, 2019)

#### <span id="page-31-0"></span>**1.3.4.2 Autophagy**

Autophagy is a process of recycling cellular components. The non-functional macroproteins or whole organelles are degraded inside lysosomes after sequestering them by the formation of the autophagosomes. Fusion of autophagosomes with the lysosomes to form the functional autolysosomes degrade the sequestered proteins. Despite the fact that autophagy is highly conserved catabolic process in eukaryotes, extensive autophagy can lead to cellular destruction in both cancer and aged cells (D'Arcy, 2019; Green & Llambi, 2015). Autophagic cell death (type II cell death) is another major cellular death pathway in mammalian cells targeted by the phytochemicals (Green & Llambi, 2015). Figure 2 summarizes the steps and signaling pathways involved in autophagy (Green & Llambi, 2015).

While autophagy is usually termed as macroautophagy, other forms of autophagy are also existed. Microautophagy is a more specific form of autophagy than macroautophagy. Microautophagy is triggered for recycling specific damaged organelles, for example, mitophagy is one type of microautophagy specific for mitochondria (D'Arcy, 2019).

Selective autophagy (chaperone-mediated autophagy) involves the degradation of misfolded proteins with the specific peptide motifs (such as KFERQ peptide motif) by selective interaction of specific receptor on the surface of phagophors and proteins motifs to guide them for degradation (Hosaka et al., 2020; Johansen & Lamark, 2020). Chebulinic acid is a polyphenolic compound naturally found in many medicinal traditional plants such as *Phyllanthus emblica* and *Terminalia arborea*. Chebulinic acid was found to inhibit the growth and the metastatic potential of the TNBC cells MDA-MB-231 cells through Autophagy induction (Sharma et al., 2020). Interestingly,

some medicinal plants such as St. John's Wort can inhibit the growth of TNBC by targeting both autophagy and apoptosis (You et al., 2020).

#### <span id="page-32-0"></span>**1.3.4.3 Necrosis**

Necrosis (type III cell death) is the non-programmed cell death in response to severe environmental or pathological changes inside the cells. Cell swelling, distension of various cellular organelles, clumping and random degradation of nuclear DNA, and extensive injury and/or rapture of the plasma membrane are the distinctive cellular changes accompanied with necrosis. Various regulators are responsible for specific subtypes of necrosis such as necroptosis and oncosis (D'Arcy, 2019).



<span id="page-33-0"></span>Figure 2: Steps involved in autophagy and it's signaling pathways (Green & Llambi, 2015)

#### <span id="page-34-0"></span>**1.3.4.4 DNA Damage**

Normally, both healthy and cancer cells can be exposed to various endogenous and exogenous DNA damage stimuli such as Reactive Oxygen Species (ROS) and the exposure to the Ultraviolet (UV) radiation. If DNA damage occurs in normal cell, DNA repair is activated through specific network of proteins called the DNA Damage Response (DDR). DDR can also regulate the inhibition of the proliferation in DNA damaged cells. However, the DNA repair capacity in the cell is limited and excessive DNA damage can activate several growth inhibitory pathways such as apoptosis and senescence (Reuvers et al., 2020; Srinivas et al., 2019). DNA damage has emerged as effective target to inhibit the proliferation of cancer cells. Many anti-cancer treatment options such as radiotherapy and cytotoxic chemotherapy exert their effect by inducing several forms of DNA damage such as Single-Strand Breaks (SSBs) and Double-Strand Breaks (DSBs). For example, the antimetabolite 5-Fluorouacil (5-FU) can lead to chain termination, by inhibiting the synthesis of the nucleoside thymidine and by being incorporated in the DNA after metabolism, and therefore induce SSB and DSB (Reuvers et al., 2020). Kaempferol, a natural derived flavonoid, was reported to inhibit the proliferation of MDA-MB-231 TNBC cells by inducing DNA damage (Zhu & Xue, 2019).

#### <span id="page-34-1"></span>**1.3.4.5 Cell Cycle Arrest**

For the normal cell to divide and duplicate, it must pass through sequence of events called "cell cycle" (van den Heuvel, 2005; Wenzel & Singh, 2018):

• G1 phase: during the G1 phase of the cell cycle, the cells grow in preparation for DNA replication. Depending on the environmental and developmental signals, cells in G1 may temporarily or permanently leave the cell cycle and enter arrested phase known as G0 phase. For example, the removal of the growth factors signals during early G1 will enter the cells into G0 phase, while the removal of such signals in the late G1 phase will enter the cells into the S phase.

- S phase: during the S phase of the cell cycle, DNA replication occurs, and each chromosome duplicates to become two sister chromatids.
- G2 phase (separates the S and M phases): during the G2 phase of cell cycle, synthesis of the materials needed for mitosis, such as RNA and proteins, occurs.
- M phase (mitosis): during the M phase of the cell cycle, the duplicated sets of chromosomes separate to the two formed daughter cells. Figure 3 illustrates the cell cycle phases.



<span id="page-35-0"></span>Figure 3: Cell cycle phases (Alberts et al., 2009)
Specific regulatory proteins such as Cyclins, Cyclin-Dependent Kinases (CDKs), oncogenes and tumor suppressor genes regulate the cell cycle progression through specific "checkpoints", in which they either allow the cell cycle to proceed or not. Those checkpoints ensure that each phase is completed before the progression to the next phase. Each checkpoint in the cell cycle is regulated by various regulatory proteins and signals, which lead to either the progression of cell cycle or cell cycle arrest. For example, a regulatory sensory signal to error in the cell cycle phases (e.g., DNA damage) can induce cell cycle arrest at one or more checkpoint(s). During the G1 phase for example, specific Cyclins/CDKs complexes (Cdk4/cyclin D and Cdk2/cyclin E) must be formed to phosphorylate the Retinoblastoma protein (Rb), which activates the transcription of further proteins responsible for cell cycle transition through G1/S phases (Lim & Kaldis, 2013). Figure 4 shows the checkpoints involved in the cell cycle regulation.



Figure 4: Cell cycle checkpoints (Ma, 2017)

Many medicinal plants with anti-cancer activities act by inducing cell cycle arrest at different checkpoints. For example, bacopaside II, derived from the medicinal plant *Bacopa monnieri*, was found to inhibit the growth and the proliferation of colon cancer cells by inducing G2/M arrest and apoptosis (Smith et al., 2018). Similarly, Ethyl Acetate Fractions of Ajwa Dates (EAFAD) induced S phase cell cycle arrest and decreased the percentage of cells in G2 phase in prostate cancer cell line PC3 (Mirza et al., 2018).

# **1.3.4.6 Cellular Senescence**

Cellular senescence is irreversible cell cycle arrest in response to different intrinsic or external stresses (oxidative damage, ultraviolet and chemotherapeutic drugs) that cause persistent DNA damage. Interestingly, senescent cells, unlike quiescent cells, resist mitogenic or growth factor stimuli, which make them unable to enter the cell cycle again in response to such stimuli (Herranz & Gil, 2018). Figure 5 represents the molecular pathways controlling the growth arrest during senescence.



Figure 5: Molecular pathways controlling growth arrest during senescence (Herranz & Gil, 2018)

Senescent cells are characterized by the change in their morphological features such as cell flattening, and the enlargement caused by the rearrangement of the cytoskeleton. Other features of senescence include mitochondrial enlargement and dysfunction and the upregulation of the lysosomal enzyme Senescence-Associated βgalactosidase (SA-β-gal). Thus, SA-β-gal is considered as biological marker for senescent cells (Lee & Lee, 2019). Wogonin, a natural derived compound, was found to inhibit the proliferation of MDA-MB-231 cells by Reactive Oxygen Species (ROS) accumulation that lead to senescence induction (Yang et al., 2020).

### **1.4** *Acridocarpus orientalis*

### **1.4.1 Overview of** *Acridocarpus orientalis*

*Acridocarpus orientalis* (Arabic name *Qafas*) is a rare plant that belongs to the family Malpighiaceae. It is a small perennial shrub with highly branched, hairy stems and yellow flowers (Rehman, Hussain, et al., 2019). Figure 6 shows *A. orientalis* growing in Oman*.* Even though *A. orientalis* is prone to extinction, 30 species of this genus are known to date and distributed in Africa, Asia, New Caledonia, and Arab countries including UAE and Oman (Kisksi et al., 2012; Rehman, Hussain, et al., 2019). In 2012, *A. orientalis* distribution in UAE was restricted to specific regions such as Jebel Hafit in Al-Ain, further information for the distribution and availability of *A. orientalis* are not available (Kisksi et al., 2012).

#### **1.4.2 Chemical Constitutions of** *Acridocarpus orientalis*

Phytochemical investigations of the chemical composition of *A. orientalis* collected from Oman revealed several active compounds (Table 3). Similarly, phytochemical screening *A. orientalis* fresh young leaves collected from Jabal Hafeet, Al-Ain revealed similar constituents including flavonoids, phenolics, and tannins (Lotfy, Al-Hammadi, et al., 2020). Phenolic content of *A. orientalis* stem and the leaves showed  $128.5 \pm 0.17$  mg/100 g and  $46.6 \pm 0.14$  mg/100 g, respectively (Ali et al., 2018). For *A. orientalis* extracted from Jabal Hafeet, quantification of total phenolic and flavonoid contents of the dry leaves extract revealed 154.2 mg/g total phenolic content and 79.9 mg/g total flavonoid content (Lotfy, Al-Hammadi, et al., 2020).



Figure 6: *Acridocarpus orientalis* growing in Al-Hamra, in AdDakhiliyah region of the Sultanate of Oman (Rehman, Mabood, et al., 2019)

Table 3: Chemical compositions of *A. orientalis* collected from Oman (Rehman, Hussain, et al., 2019)



### **1.4.3 Biological Activities of** *Acridocarpus orientalis*

Traditionally, *A. orientalis* has been used in folk medicine for many health conditions. In UAE and Oman, paste is made by crushing the seeds or seeds oil and applied on the forehead to relief headache (Divakar et al., 2016; Sakkir et al., 2012). Furthermore, *A. orientalis* leaves have been used to relieve swellings, muscle pains, and to treat arthritis (Rehman, Mabood, et al., 2019), dermatological and topical disorders, nerve disorders (hysteria and epilepsy) and eye and urinary disorders (Ghazanfar & Al-Al-Sabahi, 1993; Hinai et al., 2020). Additionally, villagers in Oman have used *A. orientalis* plant as a source of yellow dye (Divakar et al., 2016).

Despite the limited studies on *A. orientalis*, two flavonoids morin and morin-3-O-β-D-glucopyranoside isolated from the crude methanolic extract of *A. orientalis* collected from Oman showed various biological activities, including Antifungal, allelopathic and antioxidants effects (Hussain et al., 2014). Interestingly, ethanolic crude extract collected from Al Ain and Oman showed stronger antioxidant, antilipoxygenase and anti-inflammatory effects of *A. orientalis* compared to the activity of *A. orientalis* samples collected from Oman (Ksiksi & Hamza, 2012; Ksiksi et al., 2017). Furthermore, *A. orientalis* fresh young leaves ethanolic extract found to increase the levels of reduced Glutathione (GSH) in male albino mice model, indicating the hepatoprotective potential of this extract (Lotfy, Al-Hammadi, et al., 2020).

A study by Lotfy, Ksiksi et al. (2020) revealed significant antidiabetics effects of *A. orientalis* on sterptozocin induced diabetes mellitus rat model. Blood glucose levels and glucagon-immunoreactive cells were reduced after treatment with *A. orientalis* ethanolic extract. Additionally, the number of insulin-positive cells and the serum level of superoxide dismutase were increased, indicating the promising effects of *A. orientalis* active compounds on diabetes (Lotfy, Ksiksi, et al., 2020). Moreover, αglucosidase and urease enzymes inhibition by various *A. orientalis* fractions support the antidiabetic effects of the this plant (Rehman, Mabood, et al., 2019).

## **1.5 Statement of the Problem**

Breast cancer cases among females are continuously increasing. Finding new novel therapeutic options for breast cancer with less side effects is urgently needed. Traditional medicinal plants are rich source of new compounds with novel mechanisms of action. *A. orientalis* is a rare medicinal plant in UAE with promising anticancer effects against various types of cancer cells. However, the anticancer effect of *A. orientalis* leaves extract against TNBC and its underlying molecular mechanism is not investigated.

## **1.6 Hypothesis**

This study hypothesized that *A. orientalis* Ethanolic Extract (AOEE) can inhibit the proliferation of TNBC through inducing autophagy process and cell cycle arrest.

## **1.7 Research Objectives**

The objectives of the current study are:

- 1. To determine cell growth inhibition activity of AOEE in three different subtypes of human breast cancer cell lines, MDA-MB-231, MCF-7 and Hs578t.
- 2. To assess whether AOEE induces autophagy in breast cancer using MDA-MB-231 cell line as *in vitro* model.
- 3. To evaluate expression levels of cell cycle control and autophagy related proteins markers, which are associated with the cell cycle and autophagy induction.
- 4. To investigate whether other oncogenic signaling pathways such as NF-kB, mTOR, and PI3K are implicated in AOEE mediated growth inhibition and autophagy.

## **Chapter 2: Materials and Methods**

## **2.1 Preparation of** *A. orientalis* **Ethanolic Extract**

An air-dried and ground aerial part of *A. orientalis* (10 g) was extracted with 70% (v/v) ethanol (200 ml). The mixture was macerated for 72 h at room temperature. The resulting extract was then filtered dried under reduced pressure in a rotary evaporator at 40°C and an aqueous ethanol crude extract was generated. This crude extract was weighed, dissolved in 50% ethanol (typically 50 mg/ml) and kept at −20°C for further analysis.

# **2.2 Cell Culture**

The three human breast cancer cell lines used in this study (MDA-MB-231, MCF-7 and Hs578T) were maintained in Dulbecco Minimal Essential Medium (DMEM) at 37°C under a humidified atmosphere containing 5% CO2. The DMEM (Gibco/Invitrogen, UK) was supplemented with 10% Fetal Bovine Serum (FBS) (Gibco/Invitrogen, UK) and antibiotics (100 U/ml penicillin/streptomycin) (Hyclone, Cramlington, UK). Daily monitoring of the cells was done using EVOS™ XL Core Imaging System (Invitrogen, UK), and media was changed when necessary. Cells were passed as required as follows: media was removed, Phosphate Buffer Saline (PBS) solution was used for washing the cells and then 0.25% Trypsin/EDTA (Gibco/Invitrogen) for 3-5 minutes at 37°C to allow cell to detach, then neutralized with 1:1 ratio of media and adjust the final volume with the growth medium. For each experiment, cells were counted and then seeded at desired density according to the cell/assay type.

### **2.3 Cellular Viability Measurement**

 $5 * 10<sup>3</sup>$  MDA-MB-231 cells and  $7 * 10<sup>3</sup>$  Hs578t and MCF-7 cells were seeded in 100  $\mu$ L of culture medium per well in triplicate in 96-well plates. The cells were left for 24 hours to attach after seeding, then the media was replaced with treatment-added media with increasing concentration of AOEE or an equivalent volume of vehicle (50% EtOH) for control cells and incubated for another 6, 24, 48, and 72 hrs. EVOS™ XL Core Imaging System (Invitrogen, UK) was used to observe the morphological changes upon treatment at several time points. CellTiter-Glo Luminescent cell viability assay (Promega Corporation, Madison, USA), which measure the amount of ATP signaling as indicative of the metabolic activity of the cells, was used to measure the viability at 6 (for MDA-MB-231 cells), 24, 48 and 72 hours post treatment. For each well, 1:1 ratio of new media and CellTiter-Glo reagent were added after removing the media. With gentle shaking, the plate was incubated after covering it with aluminum foil for 15 minutes at room tempreture. GloMax® Explorer Multimode Detection System (Promega Corporation, Madison, USA) was used to detect the luminescence of each well. Data was provided as proportional viability (%) by equating the data for treated cells to untreated cells, which is assumed to be 100%. All experiments were performed in triplicate. The results are representative of an average of at least three independent experiments. Cell viability was calculated as follow:

% of cell viability = (average luminescence values for treatment wells / average luminescence value for control) \* 100.

The results were plotted against the range of AOEE concentrations in Excel.

### **2.4 Senescence Associated-β-galactosidase (SA-β-gal) staining**

 $2.5 * 10<sup>5</sup>$  MDA-MB-231 cells were seeded and cultured in each well of 6-wells plate. The cells were left overnight to attach then treated with AOEE (200 and 400  $\mu$ g/mL or 50% EtOH for 96 hours). Cells were washed with PBS, then fixed with 2% formaldehyde/0.2% glutaraldehyde for 5 min at room temperature (Al Dhaheri et al., 2013). The senescent cells were counted manually using microscope at 48, 72, and 96hours post-treatment.

## **2.5 Cell Lysate Preparation**

1.8 \* 10<sup>6</sup> of MDA-MB-231 cells were seeded in 10 cm tissue culture dish and treated with AOEE (200, 400 and 600 μg/ml) or vehicle (50% EtOH) for 48 hrs. After that, cells were washed twice with ice-cold PBS, scraped, pelleted, lysed with sonication in Radioimmunoprecipitation Assay (RIPA) lysis buffer (Pierce) supplemented with protease inhibitor cocktail (Roche) and phosphatase inhibitor (Roche). Cell lysates were centrifuged at 14,000 rpm at 4°C for 30 min after incubation for 30 min on ice. BCA protein assay kit (Thermo Scientific) was used to quantify protein concentrations of the supernatants, and the lysates were adjusted with lysis buffer. The supernatants were aliquoted and stored at - 80°C.

#### **2.6 Western Blotting Analysis**

Depending on the molecular weight of the protein of interest, 6 - 15% Sodium Dodecyl Sulfate-Polyacrylamide Gel Electrophoresis (SDS-PAGE) gels were prepared and loaded with equal amounts (15 μg) of cell lysates along with PageRuler plus prestained protein ladder (Thermo Scientific) for electrophoresis. The proteins separated in the gel were then transferred onto Polyvnylidene Fluoride (PVDF) membranes (Millipore). The membranes were then blocked in 5% non-fat skim milk prepared in TBST (Tris-buffered saline with 0.05% Tween 20) for one hour at room temperature, then were washed 3 times with TBST to remove the excess blocking buffer. After that, the membranes were incubated with the specific diluted primary antibodies in blocking buffer overnight at 4°C. The membranes were then washed 3 times with TBST and anti- mouse, -rabbit or -goat Horseradish peroxidase-conjugated anti-IgG were used as secondary antibodies depending on the primary antibody used. Immunoreactive bands were detected by enhanced chemiluminescence ECL/ SuperSignal West Femto chemiluminescent substrate (Thermo Scientific) using C-DiGit® Blot Scanner (LI-COR). Restore western blot stripping buffer (Thermo Scientific) was used to strip the membranes when needed according to the manufacturer's instructions. Image Studio Digits Ver 5.2 software **(**LI-COR) was used to obtain the bands and measure their intensity.

Antibodies to Cyclin D1 (04-1151),  $p21^{WAF1}$  (05-655), Phospho-Rb (Ser807/Ser811) (07-899) and PCNA were obtained from Millipore (Millipore, Hayward, CA, USA). Antibodies to β-actin-HRP (sc-47778), goat anti-mouse IgG- HRP (sc-2005), and goat anti-rabbit IgG-HRP (sc-23575), were obtained from Santa Cruz Biotechnology, Inc (USA). Antibodies to SQSTM1/p62 (ab91526) was obtained from Abcam (Abcam, Cambridge, UK). Antibodies to Cyclin E1 (#4129), Phospho-Histone H2A.X (Ser139) (#9718), LC3B (#2775), Beclin-1 (#3495), p27<sup>Kip1</sup> (#3686), Phospho-p44/42 MAPK (ERK1/2) (Thr202/Tyr204) (#9101) and Phospho-p38 MAPK (Thr180/Tyr182) (#9211) were obtained from Cell Signaling. Antibody to p16 (551154) was obtained from BD Biosciences.

# **2.7 Statistical Analysis**

The data are expressed as the mean  $\pm$  Standard Error (SEM), and are derived from at least three independent experiments, unless specified otherwise. The difference between experimental and control values was assessed by ANOVA followed by Least Significant Difference (LSD) post-hoc multiple comparison test. Student t-test was used to calculate the significance between the two independent groups. The statistically significant difference was set at p values of  $* 0.05, ** < 0.005, ** < 0.001$ between control and treated groups.

## **Chapter 3: Results**

## **3.1 AOEE Decreases Cell Viability of Human Breast Cancer Cell Lines**

To investigate the anti-breast cancer activity of AOEE, three breast cancer cell lines (MDA-MB-231, MCF-7 and Hs578T) were treated with increasing concentration of AOEE and cell viability was measured at three time points (24, 48, and 72 hours) using CellTiter-Glo® assay. As shown in Figure 7 (a-c), AOEE decreases cell viability of the three human breast cancer cell lines in time- and concentration- dependent manner compared to control group (50% ethanol). Notably, the three human breast cancer cell lines showed relatively similar response to the AOEE treatment except for the highest concentration of AOEE of 600 μg/mL, where more cell viability reduction was detected in MDA-MB-231 and Hs578T cells, which implies a higher sensitivity of both cell lines to AOEE when used at high concentrations. The estimated  $IC_{50}$  (the concentration that leads to 50% inhibition) value for MDA-MB-231, Hs578T, and MCF-7 cells were approximately ~300, 350, and 325 μg/mL respectively at 72 hours with less than 20% cell viability of MDA-MB-231 observed at higher concentration (600 μg/mL) (Table 4). Since MCF-7 and MDA-MB-231 exhibited comparable sensitivity, this implies that estrogen receptor expression does not affect the AOEEinduced growth inhibition in human breast cancer cells. Collectively, these data indicate that AOEE exerts a negative action on the viability of the breast cancer cells independently of estrogen receptor and p53 status.





Figure 7: Inhibition of cell viability of human breast cancer cells by AOEE after 24, 48, and 72 hrs. (a) MDA-MB-231, (b) MCF-7, (c) Hs578T. Data represent the mean of three independent experiments carried out in triplicate. Statistical analysis for cell viability data was performed using one-way ANOVA followed by LSD Post-Hoc test (\* p < 0.05, \*\* p < 0.005, \*\*\* p < 0.001).



Figure 7: Inhibition of cell viability of breast cancer cells by AOEE after 24, 48, and 72 hrs. (a) MDA-MB-231, (b) MCF-7, (c) Hs578T. Data represent the mean of three independent experiments carried out in triplicate. Statistical analysis for cell viability data was performed using one-way ANOVA followed by LSD Post-Hoc test (\* p< 0.05, \*\* p< 0.005, \*\*\* p< 0.001). (Continued)

<b>Cell line</b>	<b>Estrogen Receptor Status</b>	Estimated IC50 ( $\mu$ g/mL) at 72 hours
$MDA-MB-231$	Negative	$\sim$ 311 $\pm$ 23
MCF-7	Positive	$-390 \pm 36$
Hs578t	Negative	$\sim$ 325 ± 12

Table 4: The estimated IC50 ( $\mu$ g/mL)  $\pm$  SEM of AOEE in three breast cancer cell line, MDA-MB-231, MCF-7 and Hs578t at 72 hrs.

In this study, the mechanism(s) by which AOEE exerts its anti-cancer activity on the highly proliferative and invasive Estrogen Receptor (ER)-negative, mutant p53 breast cancer cell line MDA-MB-231, was investigated.

# **3.2 AOEE Induces Morphological Changes in MDA-MB-231 Cells**

To investigate the molecular mechanism(s) responsible for the anti-cancer effect of AOEE in MDA-MB-231 cells, the morphological changes under the microscope at 48 hours post-treatment were monitored. As seen in Figure 8, the control group showed the morphological features of healthy and normally proliferated MDA-MB-231 cells. On the other hand, at low concentration of AOEE (200 μg/mL), the cells started to lose the cell-cell contact and they started to look flattened and more elongated, and cells showed a kind of membrane extensions as they lost cell to cell contact. Interestingly, the cytoplasmic vacuolation (black line arrow) was clear at 200 μg/mL of AOEE and the percentage of cells showed such vacuolation increased in concentration- dependent manner. The cytoplasmic vacuolation was reported to be induced by the induction of autophagy (Chen et al., 2005). At high concentrations of AOEE (400 and 600 μg/mL), the cells were enlarged with flattened shape (dashed black line arrow) compared to the untreated cells. The enlarged and flattened cell morphology is one of the characteristics of cells undergoing cellular senescence (Wang & Dreesen, 2018). The minimal appearance of morphological features associated with cell death, such as cellular shrinkage and floating cells, might indicates that the anticancer effect of AOEE is caused by non-apoptotic pathway(s). The molecular mechanisms for the anti-cancer effect of AOEE in MDA-MB-231 cells were further investigated in the next sections of the study.



Figure 8: Representative micrographs of the cellular morphology of MDA-MB-231 treated with increasing concentrations of AOEE after 48 hrs. Cells were viewed using EVOS XL Core Cell Imaging System (Life Technologies). Magnification: 40X. The black arrows show cytoplasmic vacuolation, the black dashed-arrows show flattened enlarged cells. The white arrows show the cell extensions and elongation.

### **3.3 AOEE Inhibits the Proliferation of MDA-MB-231 Cells**

The minimal appearance of cellular death morphological features under the microscope prompting us to count the number of viable cells in the control group and the treated group at different AOEE concentrations (100, 200, 400 and 600 μg/ml) and time points (24, 48 and 72 hours) after treatment. As shown in Figure 9 and Table 5, the number of cells after 24 hours of treatment did not change significantly, which indicates minimal effect of AOEE on the proliferation at this time point. Similar pattern was observed After 48 and 72 hours of treatment with 100 μg/ml indicating a minimal effect of AOEE on cell proliferation when treated with 100 μg/ml. However, number of cells treated with 200 and 400 μg/ml of AOEE decreased significantly compared to the control group, which indicates that cell growth inhibition started at 200 μg/mL after 48 hours. Interestingly, at highest concentration of AOEE (600 μg/ml), a significant reduction in the number of viable cells after 48 and 72 hours of treatment was observed. The results obtained confirm that AOEE inhibit the proliferation of MDA-MB-231.



Figure 9: AOEE inhibits the proliferation of MDA-MB-231 breast cancer cells. MDA-MB-231 cells were treated with increasing concentration of AOEE for 24, 48, and 72 hrs, then, cell number was measured using Muse<sup>®</sup> Count & Viability Kit (Millipore). Data represent the mean of three independent experiments carried out in triplicate  $\pm$  SEM.

	Total Viable Cells $(x 104)$				
	Day 0 (treatment)	Day 1	Day 2	Day 3	
control	$12.019 \pm 0.51$	$21.51 \pm 0.8$	$30.77 \pm 3.8$	$42.13 \pm 5.6$	
100		$21.04 \pm 1.5$	$ 29.28 \pm 3.8$	$42.97 \pm 5.5$	
200		$22.09 \pm 1.3$	$26.58 \pm 2.9$	$130.97 \pm 6.8$	
400		$20.51 \pm 1.2$	$19.49 \pm 1.9$	$ 20.38 \pm 3.0$	
600		$20.03 \pm 1.6$	$14.43 \pm 3.5$	$10.43 \pm 3.1$	

Table 5: The number of viable cells in AOEE treated MDA-MB-231 cells. Data represent the mean of three independent experiments carried out in triplicate  $\pm$  SEM.

Using Western blotting, the antiproliferative effect of AOEE on MDA-MB-231 cells was confirmed by analyzing the expression levels of Proliferating Cell Nuclear Antigen (PCNA) protein in the control and AOEE treated cells. PCNA plays a major

role in the proliferation of the cells through its role in nucleic acid metabolism during DNA replication and repair (Kelman, 1997). As shown in Figure 10, PCNA protein expression levels decreased in the treated MDA-MB-231 cells starting at 200 μg/mL of AOEE treatment.



Figure 10: AOEE decreases the expression of PCNA in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of PCNA were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

# **3.4 AOEE Induces Autophagy in MDA-MB-231 Cells**

Three major types of cell death are known to occur in cancer cells: caspase-dependent apoptosis, necroptosis, and autophagic cell death. For this, the type(s) of cell death induced by AOEE in breast cancer cells MDA-MB-231 was determined. To confirm that apoptosis is not involved in the AOEE-induced growth inhibitory effect on MDA-MB-231, cells were the pre-treated with Z-VAD-FMK - Pan-Caspase inhibitor, potent apoptosis inhibitor, and the viability of the cells were measured after 48 hours of AOEE treatment (400 and 600 μg/ml). As seen in Figure 11, Z-VAD-FMK pretreatment showed no significant effect on cell viability after AOEE treatment compared to the group treated without Z-VAD-FMK. Altogether, the results suggest that the inhibitory effect of AOEE on MDA-MB-231 cells is independent of apoptosis.



Figure 11: Cell viability of MDA-MB-231 cells treated with 400 and 600 µg/mL of AOEE with or without 50 µM of Z-VAD-FMK - Pan-Caspase inhibitor for 48 hrs. CellTiter-Glo Luminescent cell viability assay was used to measure the cell viability.

Necroptosis, a programmed form of necrosis, is shown to be specifically inhibited by necrostatin-1. Next, necroststin-1 was used to test if necroptosis contributes to AOEEinduced cell death. The results show that necrostatin-1 does not inhibit AOE-induced cell death (Figure 12) indicating that AOEE-induced cell death does not involve necroptosis.



Figure 12: Cell viability of MDA-MB-231 cells treated with 400 and 600 µg/mL of AOEE with or without 50  $\mu$ M of Necrostatin-1 (Nec-1) for 48 hrs. #NS: Nonsignificant. CellTiter-Glo Luminescent cell viability assay was used to measure the cell viability. Data represent the mean  $\pm$  SEM of three independent experiments carried out in triplicate. Student t-test was used to evaluate the significance between the two groups.

To study whether autophagy is involved in AOEE-induced cell death, thr autophagy inhibitor 3-methyladenine (3-MA) was used confirm the role of autophagy as a key molecular mechanism in the anti-cancer effect of AOEE on MDA-MB-231 cells, the effect of 3-Methyladenine (3-MA), a widely used autophagy inhibitor acting by inhibiting class III Phosphatidylinositol 3-Kinases (PI3K) which block the formation of autophagosomes (Wu et al., 2010). The results showed that 3-MA decreases AOEEinduced proliferation inhibitory effect MDA-MB-231 cells (Figure 13). These findings suggest that autophagy contributes to anti-proliferative effect shown by AOEE treatment on MDA-MB-231.



Figure 13: Cell viability of MDA-MB-231 cells treated with AOEE (200, 400 and 600  $\mu$ g/mL) with or without 5 mM of 3-MA for 48 hrs. Data represent the mean  $\pm$  SEM of three independent experiments carried out in triplicate. Student t-test was used to calculate the significance between the two groups (cells treated with AOEE without 3- MA and cells treated with AOEE and 3-MA) at each concentration ( $* p < 0.05$ ). Oneway ANOVA followed by LSD Post-Hoc test was used to calculate the significance between the treated and control groups (\*  $p < 0.05$ , \*\* $p < 0.005$ , \*\*\*  $p < 0.001$ ).

Next, to confirm autophagy induction in breast cancer cells MDA-MB-231, the alteration in the cellular levels of different autophagy related proteins and markers was analyzed using Western blotting. The formation of the autophagosomes at the early step of autophagy is followed by the fusion with the lysosomes to form autolysosomes. Autophagosomes formation requires the cleavage of microtubules-associated light chain-I (LC3-I) to form LC3-II, which is recruited and conjugated to phospholipids and incorporated into the autophagosomes (Tanida et al., 2008). Therefore, LC3-II is used as indicator for the induction of autophagy. As shown in Figure 14, the levels of the conjugated form (LC3-II) clearly increased starting at cells treated with 200 μg/mL of AOEE, with average ~3-fold increase in cells treated with 600 μg/mL.



Figure 14: AOEE alters the levels of the conjugated form of LC3-I MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of LC3-I and LC3-II were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

The alteration in the level of Beclin-1, a protein with autophagy-promoting activity, was also determined using Western blotting. As shown in Figure 15, the protein levels of Beclin-1 increased starting at 400 μg/mL of AOEE treatment. In addition to its role in autophagy induction, Beclin-1 is considered as mammalian tumor suppressor protein, plays a role in inhibiting the anti-apoptotic protein Bcl-2, therefore inhibit the proliferation of the cancer cells. Notably, several cancer cell lines are reported for Beclin-1 deficient (Liang et al., 1999).



Figure 15: AOEE alters the levels of the Beclin-1 in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the proteins levels of Beclin-1were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

In similar pattern to Beclin-1, the levels of p62, another important autophagy related protein, was increased starting at 400 μg/mL of AOEE treatment (Figure 16). The p62 or [Sequestosome 1](https://www.sciencedirect.com/topics/biochemistry-genetics-and-molecular-biology/sequestosome-1) (SQSTM1) protein helps in linking and localizing the ubiquitinated non-functional or misfolded proteins to be degraded inside the autolysosomes, by binding to ubiquitin (Bjørkøy et al., 2009). Even though p62 is degraded during autophagy and its accumulation may indicates lack of autophagy , p62 protein accumulation can indicate also an abortive (prolonged) autophagy (Benhalilou et al., 2019). Altogether, the results confirm that autophagy could be one mechanism by which AOEE exerts its anti-cancer activity against MDA-MB-231 cells.



Figure 16: AOEE alters the levels of the p62/SQSTM1 in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of p62/SQSTM1 were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change

## **3.5 AOEE Induces Senescence in MDA-MB-231 Cells**

To investigate the role of cellular senescence in the anti-cancer effect of the AOEE on MDA-MB-231 cells, the Senescence-Associated β-galactosidase (SA-βgal) activity, a widely used marker for senescent cells, was measured using SA-β-gal staining at AOEE concentrations of 200 and 400 μg/mL after 48, 72 and 96 hours of treatment. Figure 17 shows the expression of SA-β-gal (black arrows) seen under the microscope, where the blue stained cells represent cells positive for SA-β-gal, which indicates the induction of senescence in AOEE treated MDA-MB-231 cells in concentration- and time- dependent manner. As seen in Figure 18, the percentage of senescent cells (blue stained cells counted) significantly increased in both AOEE concentrations used starting from 48 hours. To further confirm the role of cellular senescence in the anticancer effect of the AOEE, the variation in the level of the tumor suppressor gene p16, an indicator for cellular senescence (Rayess et al., 2012), was assessed using Western blotting. The results shown in Figure 19 demonstrate an increase in the levels of p16 protein at AOEE concentrations of 400 and 600 μg/mL. Overall, these results confirm that AOEE induces cellular senescence, suggesting its role in the AOEE-induced anticancer activity against MDA-MB-231 cells.





Figure 17: The expression of SA-β-gal in AOEE treated MDA-MB-231 cells after 96 hours of treatment. MDA-MB 231 cells were incubated with 400 μg/mL for 96 hrs and stained for SA-β-galactosidase activity to detect senescence after treatment.



Figure 18: AOEE induces senescence in MDA-MB-231 breast cancer cells. MDA-MB 231 cells were treated with AOEE (200 and 400 μg/mL) for 48, 72 and 96 hrs and stained for SA-β-Galactosidase activity to detect senescence. Data are representative of two independent experiments. Statistical analysis to determine the significance was performed using student t-test  $(*p < 0.05, **p < 0.005, **p < 0.001)$ .



Figure 19: AOEE alters the levels of p16 in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of p16 were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

### **3.6 AOEE Induces G1/S Cell cycle arrest**

To confirm further whether AOEE inhibited MDA-MB-231 cell proliferation by inducing cell cycle arrest, the alteration in the expression of the G1/S transition cell cycle regulatory proteins, Cyclin D1, Cyclin E1 and CDK2 were investigated using Western blotting. Cyclin D1 is one of the Cyclins proteins which accumulates in response to mitogenic growth factors to assemble with its cognate CDK4/6 subunit. The conjugation between Cyclin D1 and CDK4/6 is required for cells to progress through the restriction point at G1 phase. Interestingly, Cyclin D1 is overexpressed in many types of cancer due to several genetic alteration such as chromosomal translocation (Diehl, 2002). As seen in Figure 20, the level of Cyclin D1 decreased in the treated MDA-MB-231 cells in concentration- dependent manner, which further confirm the G1/S cell cycle arrest obtained by the cell cycle analysis.



Figure 20: AOEE alters the levels of the Cyclin D1 in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of Cyclin D1 were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

Like Cyclin D1, Cyclin E1 is another cell cycle regulatory protein which conjugate with CDK2 for the cell cycle transition through G1/S phases (Milioli et al., 2020). Surprisingly, the levels of Cyclin E1 and CKD2 proteins didn't change significantly upon treating MDA-MB-231 cells with AOEE, only slight increase in the protein level of Cyclin E1 when cells treated with 400 μg/mL of AOEE and slight decrease at 600 μg/mL concentration (Figure 21).



Figure 21: AOEE alters the levels of the Cyclin E1 and CDK2 in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the proteins levels of Cyclin E1 and CDK2 were examined by Western blotting. β-actin was used as loading control. Two independent experiments were performed to calculate the average fold change.

The alteration in the phosphorylated form of Retinoblastoma protein (p-Rb) was also investigated using Western blotting. For cells to enter the S phase of the cell cycle, both CDK4–Cyclin D and CDK2–Cyclin E must phosphorylate Rb protein. p-Rb next regulate the expression of many genes by controlling the activity of the transcription factor E2F (Rubin, 2013). As demonstrated in Figure 22, p-Rb levels decreased clearly in the treated MDA-MB-231 cells starting at 200 μg/mL of AOEE. In addition to its role in the metabolism of nucleic acid, PCNA has a role in the cell cycle progression in the S phase through the interaction with Cyclin A/CDK2 complex (Jurikova et al., 2016). As demonstrated here earlier in Figure 10, PCNA was downregulated in response to AOEE treatment in MDA-MB-231 cells, which further confirm the cell cycle arrest at G1/S. Overall, the results confirm that AOEE induces cell cycle arrest at G1/S phase in MDA-MB-231 cells, which contributes to the inhibitory effect on the proliferation and growth of MDA-MB-231 cells induced by AOEE.



Figure 22: AOEE alters the levels of the p-Rb in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of p-Rb were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

# **3.7 AOEE Upregulates the Cyclin-Dependent Kinases Inhibitors, p21WAF1 and p27kip1, in MDA-MB-231 cells**

Both  $p21^{WAF1}$  and  $p27^{Kip1}$  are cyclin-dependent kinases inhibitors and cell cycle regulators in which both proteins play a role in blocking the cell cycle at different stages in response to various stimuli (Zhang et al., 2021). To investigate the involvement of  $p21^{WAF1}$  and  $p27^{Kip1}$  in cell cycle arrest induced by AOEE in MDA-MB-231 cells, the alteration of their levels was investigated using Western blotting. As demonstrated in Figure 23 (a-b), both  $p21^{WAF1}$  and  $p27^{Kip1}$  were upregulated at AOEE concentrations of 200 and 400 μg/mL. Interestingly, the level of both proteins decreased at concentration of 600 μg/mL. In addition to its role in the induction of the cell cycle arrest, the upregulation of  $p21^{WAF1}$  was reported to induce senescence regardless to the p53 status (Fang et al., 1999). Furthermore,  $p21^{WAF1}$  dependent senescence pathway was also reported, independently to pRB/p16 and p53 (Jia et al., 2011).

**(a)** 48 h AOEE (µg/mL)  $\mathbf 0$ 200 400 600  $p21^{WAF1}$ **B-actin Average Fold Change**  $\mathbf{1}$  $2.0$ 2.18 1.19 **(b)**48h AOEE ( $\mu$ g/mL) 400  $\mathbf{0}$ 200 600  $p27^{kip1}$ **B-actin Average Fold Change**  $\mathbf{1}$ 1.16 1.92 0.66

Figure 23: AOEE Upregulates the Cyclin-Dependent Kinases Inhibitors,  $p21^{WAF1}$  and p27<sub>kip1</sub>, in MDA-MB-231 cells. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the proteins levels of p21<sup>WAF1</sup> (a) and p27<sup>Kip1</sup> (b) were examined by Western blotting. βactin was used as loading control. Three independent experiments were performed to calculate the average fold change.

# **3.8 AOEE Induces DNA Damage in MDA-MB-231 Cells**

To investigate the possible inducers responsible for the cell cycle arrest/senescence induced by AOEE in MDA-MB-231 cells, the variation in the level of the phosphorylated form of Histon-H2AX, which helps in DNA double-stranded breaks repair, was estimated using Western blotting. p-Histone-H2AX, also known as  $\gamma$ -H2AX, is considered as a novel biomarker for detecting DNA double-stranded breaks (Kuo & Yang, 2008). As seen in Figure 24, indeed p-Histone-H2AX level was dramatically increased starting at 400 μg/mL of AOEE compared to the untreated control cells. Therefore, the results suggest that DNA damage could be one key molecular initiator for the cell cycle arrest/senescence observed in the AOEE treated MDA-MB-231 cells.



Figure 24: AOEE induces a dose-dependent activation of p-Histone-H2AX, a marker of DNA double-strands breaks. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the protein levels of p-Histone-H2AX were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.

# **3.9 AOEE Activates ERK and p38 Signaling Pathways in MDA-MB-231 Cells**

The Extracellular Signal-Regulated Kinase (ERK1/2) and p38 pathways are both members of the Mammalian Family of Mitogen-Activated Protein Kinases (MAPKs) pathways, which regulate several cellular activities such as proliferation, differentiation, survival, and death in response to extracellular and intracellular stimuli (Kim & Choi, 2010). Many studies suggested the role of ERK 1/2 and p38 pathways in regulating and induction of autophagy (Huang et al., 2015; Sun et al., 2018; Wang et al., 2017). To investigate the effect of AOEE on ERK1/2 and p38 pathways, the alteration in the phosphorylated form of both proteins was analyzed using Western blotting. As demonstrated in Figure 25 (a-b), p-ERK1/2 (a) and p-p38 (b) levels started to increase at AOEE concentration of 200 μg/mL. Altogether, the results suggest that ERK1/2 and p38 pathways might be involved in the anti-breast cancer activity of AOEE.



Figure 25: AOEE induces phosphorylation of p38 and ERK1/2 proteins. MDA-MB-231 cells were treated with vehicle or increasing concentrations of AOEE (200, 400, and 600 µg/mL) for 48 hrs, then the proteins levels of p-p38 and p-ERK1/2 were examined by Western blotting. β-actin was used as loading control. Three independent experiments were performed to calculate the average fold change.
## **Chapter 4: Discussion**

Despite the extensive cancer research, important challenges are still going in cancer treatment. For example, conventional treatment choices of cancer such as chemotherapy causes severe side effects, which limit their use in high doses and the overall effect on metastatic cancers (Oun et al., 2018). Additionally, cancer is characterized by their plasticity, which leads to the development of chemotherapy resistance, and therefore treatment failure by various cellular mechanisms (Ramos & Bentires-Alj, 2015). TNBC is highly invasive in its nature and the absence of targeted therapy to defeat it makes the treatment challenging and highly subjected to relapse. Recently, several signaling pathways targeted by TNBC therapeutics were discovered, but they were hindered by either drug resistance or replace (Medina et al., 2020). Therefore, finding new treatment choices with novel targets to treat TNBC is necessary.

For decades, medicinal plants were used traditionally to treat many diseases. Despite the use of trial-and-error approach by the ancient people to discover the potential pharmaceutical effects of the medicinal plants, the current available biological techniques made it easier to screen for the biological activities of the plants and their derived compounds (Barkat et al., 2020). Interestingly, more than 50% of the currently used medications are plant derived. Most of the plants found to have anti-cancer potentials act by disrupting the abnormal cellular mechanisms which contributes to cancer cells hyperproliferation capacity. Large body of evidence revealed that medicinal plants often hit multiple molecular targets of multiple process associated with cancer development and progression such as proliferation, metastasis,

angiogenesis and cellular death (Karikas, 2010). Therefore, medicinal plants, indeed is a rich source for novel anti-cancer agents with novel molecular and cellular targets. *Acridocarpus orientalis* is a rare medicinal plant that is used in folk medicine to treat several illnesses. Furthermore, recent studies revealed anti-diabetic, antioxidant and hepatoprotective effects of *Acridocarpus orientalis* (Ksiksi & Hamza, 2012; Lotfy, Al-Hammadi, et al., 2020; Lotfy, Ksiksi, et al., 2020; Rehman, Mabood, et al., 2019).

*Acridocarpus orientalis* possesses strong inhibiting effects against a variety of cancer cells, including colorectal adenocarcinoma (HT29) (Hussain et al., 2014; Rehman, Hussain, et al., 2019), colorectal adenocarcinoma (HCT116) (Hussain et al., 2014; Rehman, Mabood, et al., 2019), human hepatoma derived cell line (HepG2) (Hussain et al., 2014; Rehman, Mabood, et al., 2019), hormone responsive breast cancer (MCF-7) (Balhamar et al., 2019), triple hormone receptor-negative breast cancer (MDA-MB-231) (Balhamar et al., 2019), cervical cancer (HeLa) (Balhamar et al., 2019), and mouse mammary carcinoma cell line (4T1 cells) (Jamshidi-Adegani et al., 2020), however, the underlying anticancer target(s)/mechanism(s) remains unknown. Therefore, the aim of the current study is to investigate the effect of *A. orientalis* on TNBC and to elucidate the underlying anti-breast cancer molecular target(s)/mechanism(s).

Different molecular mechanisms and death pathways are responsible for the anticancer effects of *A. orientalis*. For example, induction of apoptosis and activation of autophagy are the key mechanisms of various fractions of *A. orientalis* extract against HeLa cell line (Balhamar et al., 2019). In 4T1 cells, apoptosis and cell cycle arrest in G0/G1-phase are the key mechanisms of proliferation inhibition upon *A. orientalis* leaves and stem extracts treatment (Jamshidi-Adegani et al., 2020). Novel molecular mechanisms to induce cellular death were also reported by Ksiksi and Hamza (2012)*.* The ethanolic extract of *A. orientalis* was reported by to inhibit the Histone Deacetylases (HDACs) enzymes, which therefore can induce cell death and inhibit the angiogenesis in cancer cells (Ksiksi & Hamza, 2012).

The findings of this study revealed significant proliferation and growth inhibitory effect of AOEE on TNBC cell lines (MDA-MB-231 and Hs578T) and ER positive breast cancer cell line (MCF-7). The growth inhibitory effect of AOEE on the three studied human breast cancer cell lines is independent of the ER-status and p53 status. However, each may act through different molecular mechanisms. The current study, focused on studying the molecular mechanism(s) and identifying molecular target(s) associated with the anti-breast cancer activity of AOEE on MDA-MB-231.

Autophagy is an evolutionarily conserved catabolic mechanism, by which cells recycle or degrade internal proteins or organelles. In autophagy, cytoplasmic materials are directed to the lysosomes for degradation. In addition to its role in recycling of misfolded and degraded proteins, the induction of autophagy was found to have antiproliferative effects, mainly by regulating the cell cycle progression or cell death induction (Kocaturk et al., 2019). Indeed, AOEE induces autophagy in MDA-MB-231 cells confirmed by the upregulation of LC3-II protein, a marker for the formation of the autophagosomes. Beclin-1, which has a role in the initial steps of the autophagosomes formation (Vega-Rubín-de-Celis, 2020), was also upregulated upon AOEE treatment, which suggests that the autophagy induced by AOEE is Beclin-1 dependent. Surprisingly, p62 responsible for the sequestration of the misfolded and non-functional proteins for degradation, along with sequestered proteins by the autolysosomes, was upregulated upon AOEE treatment. Of note, p62 accumulation indicates the induction of abortive autophagy. The use of the autophagy inhibitor 3- MA caused partial reversal of the viability of AOEE treated MDA-MB-231 cells,

which suggests that autophagy induced by AOEE is not protective mechanism and contributes to the growth inhibitory effect induced by AOEE. Ginkgolide B, a natural derived compound extracted from *Ginkgo biloba* leaves with anti-cancer potential, was reported to inhibit the proliferation of A549 and H1975 lung cancer cell lines by inducing Beclin-1 dependent autophagy accomplished by the upregulation of p62 protein (Wang et al., 2020).

In addition to the role of autophagy in the growth inhibitory effect of AOEE in MDA-MB-231 cells, the results of the present study demonstrated that AOEE induces cellular senescence in concentration- and time- dependent manner. Cellular senescence has emerged as a potent target for inhibiting the proliferation of cancer cells with low doses of radiotherapy or chemotherapy to minimize the side effects of both treatment choices, by inducing irreversible, or at least prolonged, cell cycle arrest (Zeng et al., 2018). Interestingly, AOEE has significantly increased the expression of SA-β-gal; a biomarker for the induction of senescence, in MDA-MB-231 cells. The results also demonstrated upregulation of p16 protein, which is highly expressed in senescent cells and known to inhibit the Cyclin D-CDK4/6 complex required for the cell cycle progression from G1 to S phase. G1/S cell cycle arrest and alteration in the cell cycle regulatory proteins were detected by AOEE. While the cell cycle regulatory protein, Cyclin D1, which plays a role in cell cycle progression from G1 to S phase, was downregulated in the treated cells, the CDKs inhibitors  $p21^{WAF1}$  and  $p27^{Kip1}$  were upregulated upon treating MDA-MB-231 cells with AOEE for 48 hours. Even though p53, which regulates the expression of  $p21^{WAF1}$  and  $p27^{Kip1}$  (Philipp-Staheli et al., 2004), is mutated in MDA-MB-231 cells, the upregulation of both proteins in the treated cells suggests p53-independent pathway for the induction of senescence and cell cycle arrest. PCNA, a marker for cell proliferation, plays major roles in the DNA replication and replication-associated processes such as mismatch repair and chromatin assembly in the S phase of cell cycle (Boehm et al., 2016). Interestingly,  $p21^{WAF1}$  found to regulate the function of PCNA, by inhibiting its binging with DNA polymerase δ (Wang et al., 2021). A clear reduction in the PCNA levels was detected at AOEE concentration of 200 μg/mL, which further confirm the antiproliferative effect of AOEE on MDA-MB-231 cells. Rb is an important downstream target for Cyclin D1 and Cyclin E1 that regulates the G1/S transition in the cell cycle. Phosphorylation of Rb; initiated by the effect of Cyclin D/CDKs complexes, followed by the hyperphosphorylation effect mediated by Cyclin E/CDK2 results in the release of E2F transcription factors required for the G1/S transition (Beasley et al., 2003). The results of this study demonstrated hypophosphorylation of Rb upon AOEE treatment, which indicates that the p16/Cyclin D1/Rb pathway plays a major role in the G1/S cell cycle arrest. The anti-cancer potential of a novel indenone derivative was also reported to induce senescence and G1 cell cycle arrest in MDA-MB-231 cells (Priyanga et al., 2020).

Double-strand breaks in DNA can result mainly from the exposure to exogenous agents such as radiation and certain chemicals or during DNA replication and repair (Cannan & Pederson, 2016). DNA double-strand breaks can highly affect the cell cycle progression, where a single double-strand break reported to be sufficient to induce cell cycle arrest in G1 or G2 (van den Berg et al., 2018). Additionally, DNA damage was reported to be linked to the induction of senescence via the activation of ATM/ATR/p53/p16 pathway (Von Zglinicki et al., 2005). The levels of the phosphorylated form of Hisotone-H2AX, which is used as biological marker for double-strand breaks in the cells, were increased starting at 200 μg/mL of AOEE. Thus, the results obtained from the present study suggest that DNA damage is the initiator for the cell cycle arrest starting at 200 μg/mL. On the other hand, the absence of significant upregulation of p16 at 200 μg/mL of AOEE can be explained by the minimal concurrent DNA double-strand breaks at the same concentration.

The ERK1/2 and p38 pathways are members of the MAPK signaling pathway responsible for several cellular processes, including, cell proliferation, differentiation, migration, senescence and apoptosis, by delivering extracellular signals to the nucleus (Sun et al., 2015). The phosphorylation of ERK1/2 at threonine and tyrosine is required to acquire its biological functions. Importantly, ERK1/2 pathway was reported to have dual effect on cellular senescence; where it can promote or inhibit the induction of senescence via various mechanisms and under different dose- and durationcircumstances (Zou et al., 2019). Furthermore, ERK1/2 was previously shown to have a role in the induction of autophagy (Lee et al., 2013; Yuan et al., 2018). Similar to ERK1/2 pathway, p38 pathway was also reported to be involved in the regulation of the cell cycle and induction of cell cycle arrest in several phases. Surprisingly, p38 activation was reported to be reduced in many types of tumors, and the reduction of p38 activation is linked to proliferation in those tumors. Therefore, the activation of p38 pathway can activate senescence (Zarubin & Jiahuai, 2005). The complex relationship between p38 and autophagy is not clear yet and needs further investigation, but several reports found dual regulatory effect of p38 activation on autophagy, and autophagy induction effect on p38 activation (Sui et al., 2014; Webber, 2010). The upregulation in the phosphorylated p38 and ERK1/2, in AOEE treated MDA-MB-231, demonstrated in the present work suggests that both pathways might be related to the induction of senescence and autophagy.

In summary, AOEE inhibited the viability of three human breast cancer cell lines (MDA-MB-231, Hs578t and MCF-7) and the effect seems to be independent of the ER expression status and p53 status. Different cellular mechanisms and pathways were associated with the growth inhibitory effect of AOEE in the high invasive triplenegative breast cancer cell line MDA-MB-231. The antiproliferative activity and cell cycle arrest induced by AOEE in MDA-MB-231 cells were accomplished by the downregulation of cyclin D1, PCNA, p-Rb and the upregulation of  $p21^{WAF1}$  and p27<sup>Kip1</sup>. Additionally, AOEE induced prolonged autophagy accomplished by the upregulation of LC3-II, Beclin-1 and p62. Moreover, AOEE induced DNA double stranded breaks and cellular senescence characterized by the upregulation of p-H-H2AX, SA-β-gal and p16 respectively in MDA-MB-231 cells. Finally, AOEE induced activation of ERK and p38 pathways, which might be related to the induction of autophagy and senescence. Further investigation on the exact role of the ERK and p38 pathways in the autophagy and senescence induced by AOEE is warranted. Figure 26 represents a hypothetical model summarizes the molecular events induced by AOEE in MDA-MB-231 cells.



Figure 26: Proposed model demonstrating the underlying mechanism of action in AOEE-induced anti-breast cancer activity in MDA-MB-231 cells

## **Chapter 5: Conclusion**

In conclusion, the present study reveals a negative action of AOEE on the viability of breast cancer cells *in vitro*. The study showed, for the first time, that AOEE inhibits MDA-MB-231, human breast cancer, cells proliferation and growth. The antiproliferation effect of AOEE on MDA-MB-231 cells was associated with the cell cycle arrest, through modulation of cell cycle regulatory proteins, autophagy and cellular senescence induction, and DNA double stranded breaks. MAPK signaling pathway was also involved in the anti-breast cancer activity of AOEE. Therefore, AOEE could be a rich source for active compounds. Further identification and analysis of the active compounds responsible for the observed anti-cancer activity of AOEE is warranted. *In vivo* studies are also required to validate the *in vitro* AOEE anti-breast cancer effect.

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