

Epigenetic Regulation of Tamoxifen-Resistant Breast Cancer: An Update

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Abstract

Breast cancer is the most common cause of death in women around the world. Epigenetic changes modulate transcriptional activity in several diseases, including cancer. Cancer epigenetics explains gene expression changes without DNA mutations. Aberrant DNA methylation, histone modifications, and mRNA expression promote tumor growth and metastasis. In cancer cells, chemo-resistance occurs via Multidrug Resistance (MDR), apoptotic suppression, DNA damage response, epigenetic alterations, and competitive endogenous RNA. Owing to drug resistance, quiescence, and varied cancer cell production, Cancer Stem Cells (CSCs) are critical to tumor formation, metastasis, and recurrence after therapy. In addition, MDR promotes drug efflux, enhanced secretion of growth factors, and DNA modifications in cancer patients, thereby causing fatalities in cancer patients. Heterogeneity and epigenetic plasticity cause drug resistance due to various factors. However, the molecular mechanism of epigenetic drug resistance is still unravelled completely. Overexpressed c-MYC leads to cancer and tamoxifen resistance. Despite the molecular underpinning of cancer development, drug resistance is continued in a myriad number of cases. Epigenetic changes affect CSCs viability and tumor aggressiveness. These processes can be blocked by medicines. Tamoxifen is used widely for breast cancer treatment; however, latent treatments have emerged as a tamoxifen-resistant phenotype. Epigenetic modifications cause resistance by upregulating and altering the tumor microenvironment and deregulating the immune response. The knowledge of epigenetic pathways in clinical treatment resistance may enhance the outcome of cancer patients. Multifactorial heterogeneous resistance is common in many targeted therapies. Many resistance mechanisms to targeted therapy may converge, including route reactivation. This review summarizes the epigenetic alterations, MDR, and development of tamoxifen resistance in breast cancer.

Keywords: Breast Cancer, Cancer Epigenetics, Cancer Stem Cells, Multidrug Resistance, Tamoxifen Resistance

1. Introduction

Tamoxifen has been studied extensively and is used in the primary treatment of breast cancer. Although tamoxifen has been used to treat various ailments, it has garnered a preponderant preference for the treatment of early breast cancer since its inception. Tamoxifen, established in the early 1970s, is an anti-estrogen medication. Over the past four decades, tamoxifen has been used to treat breast cancer in women throughout the

globe, resulting in a 30% reduction in the death rate^{1,2}. Tamoxifen is the most successful therapy and care for breast cancer; however, the resistance of cancer cells is a key downside of its use³. Tamoxifen is an FDA-approved selective estrogen modulator hydrophobic drug that is administered primarily to patients with postmenopausal breast cancer^{4,5}. The success of tamoxifen in treating Ductal Carcinoma *In Situ* (DCIS) and the efficacy of new endocrine treatments as a tolerability profile with efficacy in both premenopausal and postmenopausal women have

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set a benchmark in cancer management. In addition, tamoxifen, administered orally, is a potential treatment for bipolar disorder^{6,7}. Tamoxifen has a mechanistic involvement in bipolar disorder due to its antagonistic effects on Estrogen Receptors (ERs) in breast tissue and the suppression of Protein Kinase C (PKC)^{8,9}. Tamoxifen has played a vital role in saving many lives in medical oncology and healthcare during the past few decades¹⁰⁻¹³.

Tamoxifen reduces ER function in hormone-sensitive breast cancers by competing with estrogen for the binding of ER, thereby limiting the estrogenic actions that cause cancer formation and proliferation^{14,15}. However, high-level resistance to hormone therapy has been discovered in most HIV-positive people, which results in *de novo* or acquired resistance. Therapies target estrogen synthesis or ER activation. Even though these treatments have helped a significant number of individuals, drug resistance continues to be a serious issue^{16,17}. The ability to develop resistance to established therapy characterizes the disease *viz.*, cancer, thereby resulting in higher cancer-related mortality rates¹⁸.

Resistance is a well-known phenomenon that arises when illnesses develop resistance to therapeutic interventions. This concept was first presented when bacteria developed resistance to specific antibiotics; however, comparable processes have been observed in other diseases, including cancer^{18,19}. Multidrug Resistance (MDR) is a recognized phenomenon clinically, where human tumours that develop resistance to one form of treatment develop resistance to numerous additional medications that are sometimes distinct in structure and mechanism of action²⁰. Efflux of the drug in microorganisms and drug-resistant cancers is disorder-specific¹⁸. The discovery of MDR pathways in human cancer led to the development of therapeutic medicines to combat MDR²⁰. Tamoxifen resistance is the cause of treatment failure in bone cancer cases²¹. Tamoxifen tolerance has unknown mechanisms²². Recently, various molecular pathways paved the way to develop tamoxifen resistance, such as receptor tyrosine kinase pathways and miRNAs²³. Although modern chemotherapy treatment has improved, it is still ineffective against disseminated tumours. Resistance to anticancer treatments is a complex process that begins with changes in drug targets, implying the need for more targeted therapies in the therapeutic arsenal²⁴.

Furthermore, there is a growing body of data showing that tamoxifen has immunomodulatory

effects, including findings from *in vitro* and *in vivo* trials, as well as findings from breast cancer patients who took the medication²⁵. These data suggest that tamoxifen may trigger a change in immunity from cellular (T-helper 1) to humoral (T-helper 2) levels²⁶. The recent crystallization of the estradiol and raloxifene ER complex provides intriguing insights with regard to anti-estrogen action and, for the first time, established a mechanism of tamoxifen drug resistance²⁷. Epigenetic processes influence cancer, chromosomal imprints, gene suppression, diversification, morphogenesis, and X chromosome inactivation²⁸. The resistance to cancer treatment may be influenced by epigenetic mechanisms, which are transmitted somatically during cell division. Owing to the rapid rate of epigenetic change in tumours, a diverse set of gene expression patterns may evolve as a result of medication selection throughout treatment, leading to acquired resistance²⁹. Simultaneous epigenetic control of numerous genes results in the development of drug resistance in the tumours, and this has crucial implications for biomarker investigations of clinical outcomes after chemotherapy and therapeutic treatments to avoid or regulate drug resistance. As a result, new knowledge about the molecular mechanisms of tamoxifen resistance, MDR, and epigenetic regulation will aid in the development of drug resistance-fighting regimens, the discovery of novel therapeutic agents with a lower risk of developing resistance, and the development of more effective treatment strategies³⁰.

2. Tamoxifen

2.1 History of Development

Tamoxifen is a synthetic non-steroidal anti-estrogen developed by a team of chemists, endocrinologists, and reproductive researchers at Imperial Chemical Industries Ltd., now known as AstraZeneca, in the late 1950s³¹. Beatson hypothesized, in the late 1800s, that oophorectomy may cure invasive breast cancer in premenopausal women. Tamoxifen was used originally to treat infertility in 1966³². Tamoxifen became the first targeted breast cancer agent in 1972³³⁻³⁷.

In 1978, tamoxifen was used mostly in postmenopausal individuals with metastatic breast cancer. Its remedial element now includes Stages I and II patients and early and late postmenopausal women³⁸. Tamoxifen was designed to treat contralateral breast cancer and is being at

present studied as a chemo-preventive therapy in healthy high-risk women²². Recently, several articles on putative tamoxifen-resistance pathways have been published. V.C. Jordan's technique of targeted ER-positive malignancies with long-term adjuvant tamoxifen saved a large number of breast cancer patients³⁹⁻⁴¹. Tamoxifen resistance is thought to be the result of several reasons. Knock-down of ER, upregulation of a particular growth factor receptor, stimulation of PI3K/AKT/mTOR pathway (in particular PTEN inactivation), and NF- κ B signaling play important roles in tamoxifen resistance^{4,42}.

2.2 Tamoxifen Resistance

In the 1950s, Ethamoxytriphetol (MER-25 or Merrell) and Clomiphene citrate (Clomid, Merrell) were brought up⁴³. Walpole and colleagues, in the 1960s, produced various triarylethylene derivatives, which include alkyl substitutes for chlorine in clomiphene for the treatment of hormone-dependent cancers. Animal research, at the time, had suggested that clomiphene and similar chemicals caused cataracts by accumulating desmosterol, a cholesterol precursor^{43,44}. Some breast cancer tumours may develop tamoxifen resistance despite its effectiveness. Clinical studies revealed that three years of intermittent treatment of tamoxifen leads to the development of tamoxifen resistance. Down-regulation, mutation, or deletion of ERs and de-escalated co-activator activation changed tamoxifen pharmacology^{45,46}. Tamoxifen-induced MCF-7 tumours in nude mice may be further accelerated by estrogen, which tamoxifen inhibits, thereby implying that tamoxifen has both aggressive and antagonistic qualities in these tumours⁴⁷.

2.3 Tamoxifen: Mechanism of Action

Tamoxifen has a dual mode of action: (i) it competes with 17 β -estradiol (E2) at the receptor site, reducing E2's breast cancer growth; and (ii) it binds DNA upon metabolic activation, beginning carcinogenesis. Tamoxifen competes with ER of breast cancer⁴⁸. Tamoxifen treatment reduces the breast tumour and is associated with decreasing the serum insulin-like growth factor (IGF-1)⁴⁹ while enhancing sex hormone binding globulin (SHBG)^{48,50,51}. An increase in SHBG inhibits free estradiol, which reduces tumour-promoting substances. Tamoxifen treatment kills the ER α -positive breast cancer cells⁵². Protein kinase C suppresses gene transcription, resulting in this condition⁵². Hypotheses explaining tamoxifen's

apoptotic activity include a threefold spike in cytoplasmic and mitochondrial calcium ions or TGF- β production^{52,53}.

Tamoxifen functions as an estrogen agonist and antagonist in different parts of the body. It binds to ERs preferentially, producing both estrogenic and anti-estrogenic effects; as a specific ER modulator, it is patient-specific due to its binary conditioning⁵⁴. It competes with estrogen in breast tissue, causing anti-estrogenic and anticancer effects. Intracellular processes impede cell cycle progression. It boosts rather than inhibits ER in bone, which may avoid fractures in postmenopausal women⁵⁵. It also works as an estrogen agonist in premenopausal women's hypothalamus, thereby increasing gonadotropin levels and causing ovulation. Tamoxifen is metabolized in the liver by a role to CYP450 enzymes such as 2B6, 2C9, 2C19, 2D6, and 3A4. The half-life of tamoxifen is 5–7 days and N-desmethyl tamoxifen is 14 days^{56,57}.

The estrogen-ER complex homodimerizes and attaches to estrogen response elements in estrogen-sensitive genes. AF1 and AF2 (transcriptional co-activators) bind with other molecules to boost RNA Pol II activity and control gene activity⁵⁸. Tamoxifen competes with estrogen for ER binding⁵⁹. Tamoxifen-ER complexes bind to estrogen-sensitive genes and form homodimers⁵⁸. However, only if AF1 is operational while on tamoxifen-ER binding⁶⁰. When AF2 is inactive, estrogen-responsive gene transcription and co-activator binding are diminished. As a consequence, tamoxifen slows cell growth by blocking the G1 phase of the cell cycle. Tamoxifen may have several modes of action, tamoxifen may affect breast cancer epithelial cells indirectly by altering cytokine levels both locally and systemically⁶¹. It increases the generation of TGF- β , which inhibits breast cancer cells^{61,62}. Tamoxifen suppresses IGF-1, an anti-angiogenic breast cancer mitogen^{63,64}. Tamoxifen binds to non-specific sites as well as particular binding sites⁶⁵.

3. Mechanisms of Resistance: Endocrine Therapy

Significant advancements in the treatment of early breast cancer have been observed during the past 25 years. One of the most significant achievements in respect of hormone receptor-positive breast cancers is the widespread use of Endocrine Therapy (ET) similar to tamoxifen, a specific selective ER modulator, or Aromatase Inhibitors (AIs),

both nonsteroidal analogous (such as letrozole) and anastrozole or steroidal comparable (such as exemestane), which block estrogen production⁶⁶. The resistance that develops during therapy might be inherent, which means that it exists before the commencement of the treatment or is acquired during the treatment^{46,67}.

4. Estrogen Receptor

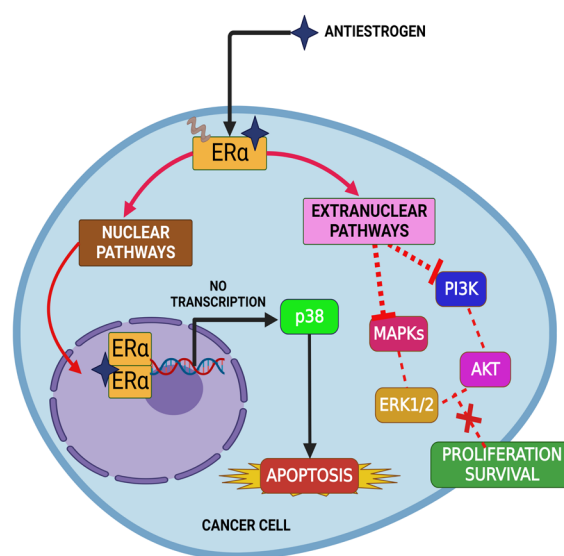
Estradiol (E2) binds to ER, a ligand-activated key transcription that interacts with estrogen response elements (EREs) in nonsupervisory regions of its target genes^{68,69}. ER may homo- (ER α -ER α) or heterodimerize (ER α -ER β) to interfere with transcriptional activity, which is encoded by genes on chromosomes 6 and 14. Each ER α /ER β subtype regulates genes differently, expresses differently in different cells and tissues, and influences various downstream signalling cascades⁷⁰. ER activation enhances tumorigenesis by proliferating and invading breast cancer cells. ER reduces cell proliferation, decreases epithelial-to-mesenchymal transition, and boosts tamoxifen sensitivity^{71,72}. High ER levels in individuals without ER are related to higher survival and tamoxifen response, whereas low ER levels may lead to endocrine treatment resistance^{73,74}. ER amplification in tamoxifen-resistant women's pre-invasive metastatic breast cancer seems to inhibit ER-stimulated transcription.

Both ER α and ER β subtypes may vary in the regulation of gene expression, cellular location, and cancer pathogenicity⁷⁵. When compared with the full-length ER protein, at least two docked ER isoforms (ER36 and ER46), which result from alternative splicing, exon deletion, and promoter activity, act in a dominant negative manner^{76,77}. In endometrial cells, ER36 was limited to the mitochondrial matrix, had no genetic activities, and was associated with tamoxifen resistance; however, ER46 increased tamoxifen perceptivity in tamoxifen-resistant MCF-7 cells. On the basis of the removal of exon 8, ER comprises many isoforms similar to ER1 (full-length), known as ER2 to ER5. Nuclear ER2 and ER5 have been shown to decrease ER function while promoting ER1 transactivation. As they are co-expressed consistently with ER⁷⁸, their relevance to endocrine resistance is being investigated at present. As ER subtypes push and pull, their percentage is being examined as a diagnostic tool for endocrine drug response⁷⁹. Owing to a paucity of adequate antibodies, subtype data are questionable⁸⁰. Large quantities of tyrosine kinase receptors, signalling

proteins, and ER are predicted in caveolae and lipid rafts^{81,82}. In tamoxifen-resistant cells, ER relocation increases epidermal growth factor receptor (EGFR) binding and downstream signalling. Another endocrine resistance media is characterized most probably in this manner⁸³ (Figure 1).

4.1 Estrogen Receptor Alteration

A possible tamoxifen resistance mechanism for ER protein structural alterations results in the altered affinity of the receptor. Defects in the ER gene encoding may cause aberrant receptors physically. These mutations may inactivate the ER, giving the tumour an ER-negative look. If mutations affect critical receptor amino acids, functionally active ER species with distinct estrogen and tamoxifen specificities may arise²². There are innumerable studies reporting that mutations and mRNA splice variants in ER α result in tamoxifen resistance⁸⁴⁻⁸⁶.



ANTIESTROGEN MEDIATED ER α PATHWAY INHIBITION

Figure 1. Anti-estrogens, which are estrogen blockers, function by inhibiting the Estrogen Receptor (ER) either by inhibiting its nuclear pathways or by inhibiting the extranuclear pathways. In the nuclear pathway, anti-estrogen binds to the ER α dimer preventing it from transcribing, thereby leading to cell apoptosis. In the extranuclear signaling pathways, the cell survival of the cancer cells inhibited by blocking via MAPKs, PI3K, ERK1/2, and AKT signaling molecules.

4.2 Estrogen Receptor Cofactors

ER is a modulator protein containing an N-terminal transactivation domain (AF1), and a C-terminal Ligand Binding Domain (LBD). Co-activators/co-repressors, and transcriptional and histone modifiers comprise the bound ER⁸⁷. When ligand is bound, helix 5 and 12 create a hydrophobic pocket and protein surface for interacting with co-activator LXXLL motifs⁸⁷. Tamoxifen binds to the same regulatory sites as estrogen-bound ER, but shifts helix 12 away from the ligand binding pocket, which leads tamoxifen-bound ER to employ a co-repressor group rather than a co-activator group to counterbalance hormone response^{88,89}.

High levels of ER co-activators, such as amplification in ER co-activator (AIB1), have been established to augment tamoxifen's agonist effect and contribute to tamoxifen resistance^{87,90}. Tamoxifen's agonistic actions on the ER were exacerbated by a prolonged decrease in nuclear receptor co-repressor 1, and co-repressor activity during tamoxifen therapy, which increases tamoxifen resistance⁹¹. ER's non-classical genomic effects include indirect DNA binding. Membrane-bound non-classical signalling activates MAPK and G-protein-coupled receptor activation leading to the induction of mitogenesis⁹². Tamoxifen-ER antagonist activity is transformed into an aggressive action as a result of these complexes, which influences breast cancer cell proliferation⁹³. An increase in the expression of these factors has also been linked to endocrine treatment resistance⁹⁴. SRY-box 9 (SOX9) is a stem cell factor that promotes mitosis, migratory phenotype, and endocrine treatment resistance⁹⁵. FOXA1 co-expresses with ER α throughout mammary gland development and early breast cancer formation^{69,96}. Over-expressed FOXA1 stimulates tumorigenesis and endocrine resilience proteins, including IL-8⁹⁷.

Endocrine resistance is also fueled by non-ER-dependent reprogramming chromatin landscapes. EZH2, a chromatin-altering enzyme, confers endocrine treatment resistance by reducing GREB1, an ER cofactor. EZH2 is high in tamoxifen-resistant samples, and low GREB1 changes the ER transcriptional machinery and transcript, which leads to resistive phenotypes in hormone-positive cells⁹⁵. In another research, drug-sensitive MCF-7 cell lines have different open chromatin landscapes⁹⁸. Traditional ER signalling had no discernible benefit in these cell lines. Nonetheless, chromatin remodelling enriches and over-expresses the NOTCH network and some other molecular targets in resistant cells, with NOTCH3 responsible for

resistance. NOTCH3-activated PBX1 induces endocrine resistance genes⁹⁸. The crosstalk of PKC- α and Notch-4 signalling induces tamoxifen resistance in breast cancer⁹⁹. Recently, the over-expression of Notch-1 has been associated with the development of tamoxifen resistance and MDR in breast cancer¹⁰⁰.

4.3 Estrogen Receptor and its Signaling Pathways

Post-translational modifications to the wild-type ER protein result in ligand-independent ER activation consistently. Growth factor signalling protein over-expression alters the ER, making the cell tamoxifen resistant. The phosphorylation of the ER α is due to the over-expression of receptor tyrosine kinases corresponding to HER2, EGFR, and IGF1R¹⁰¹⁻¹⁰³. Phosphorylation activates the ER in the absence of a ligand, rendering cells tamoxifen resistant. Endocrine resistance has also been connected to MAPK or the PI3K/AKT signalling pathway activation¹⁰⁵. Furthermore, HER2 transfection in hormone receptor-positive cells down-regulates ER α , thereby providing resistance to anti-hormone treatment^{105,106}. c-Jun/AP1 exerted over-expression and reduced the sensitivity of tamoxifen treatment in ER α -positive breast cancers¹⁰⁷. Interleukin-6 (IL-6) potentiates the ER α signalling and activates cell proliferation in MCF-7 breast cancer cells¹⁰⁸.

5. Acquired Tamoxifen Resistance: Potential Mechanisms

Several pathways have been connected to the development of acquired resistance to tamoxifen. Many of the proposed processes, on the other hand, have no conclusive evidence to back them up. Tamoxifen has been found in *in vitro* and *in vivo* studies to promote cell proliferation after prolonged exposure^{65,109}. The ability of tamoxifen to select a sub-clone of tamoxifen-stimulated cells changes tamoxifen in such a way that the cells release a stimulatory signal or generate a genetic mutation, which increases unknown drug sensitivity. Tamoxifen has been discovered to form DNA adducts in the liver of rats, implying that it may have genotoxic qualities that cause mutations¹¹⁰.

Even though the mechanisms which cause tamoxifen resistance are unknown, recent research suggests that many pathways are implicated. Changes in ER levels, lower ER affinity, enhanced cellular mechanisms for avoiding

Tamoxifen cytotoxicity, reduced cellular tamoxifen level, enhanced levels of antagonizing metabolites, and other routes have helped to understand tamoxifen resistance²².

6. Mechanisms: Anti-Estrogen Resistance

Several processes have been linked to the development of tamoxifen resistance. However, many of the postulated processes have little solid evidence to back them up. Drugs were once thought to be responsible for causing metabolic changes in cells that resulted in drug resistance.

Tamoxifen is thought to limit tumour cell proliferation primarily by interacting with ER²². Recent data from breast cancer biopsies link tamoxifen resistance to altered growth factor receptor expression and downstream signalling cascades¹⁷. Kinases downstream of these receptors, such as ERK1/2, p38, AKT, and p21-activated kinase-1, are active consistently in nonresponsive cancers with *de novo* or acquired resistance^{111–115}. In tamoxifen-resistant tumors, BCAR1, AIB1, and ER α expression are all elevated significantly. Tamoxifen reduces estrogen-dependent cancer cell growth by preventing G0/G1 phase¹¹⁶. Interaction tamoxifen with the receptor is thought to result in the formation of a complex that, when linked with estrogen-responsive regions, stops target genes from being transcribed. The ensuing blockage is thought to be cytostatic mostly, and it can be reversed by adding estradiol. It is still unclear whether tamoxifen causes apoptosis or cell death.

Tamoxifen's anti-estrogenic actions differ with species and target tissues. Rats and humans have tamoxifen sensitivity. Tamoxifen has estrogen-agonist effects in uterine tissues; however, it is an estrogen inhibitor in most instances¹¹⁷. Tamoxifen affects postmenopausal gonadotropin levels, plasma proteins, and vaginal epithelium^{116–118}. It is not known if the variation in the anti-estrogenic effect is due to species or tissue-specific tamoxifen metabolism or different transcription factors that modify signal interpretation by the cell after the anti-estrogen interacts with the ER. Numerous other factors would still influence the cellular response of tamoxifen.

Tamoxifen was also found to bind to non-ER locations¹¹⁹. The digestive system, uterus, ovaries, brain,

and kidneys have the most estrogen-binding sites and Anti-Estrogen Binding Sites (AEBS)¹²⁰. The AEBS are different from the ER and emerge only after estradiol therapy¹¹⁹. Anti-estrogen affinity for AEBS does not appear to be related to the biological efficacy of anti-estrogen, which implies that AEBS does not mediate anti-estrogen activity directly^{121,122}. Many investigations have linked AEBS binding to anti-estrogen-related physiological activities, such as protein kinase C inhibition¹²³, calmodulin inhibition¹²⁴, and interactions with histamine¹²⁵, dopamine¹²⁶, and muscarinic receptors¹²⁷. AEBS have generated interest consistently, although their role in the antitumor efficacy of tamoxifen is uncertain.

Several studies have shown that ER β is lost during carcinogenesis, which indicates that it may play a tumour-suppressing role^{128,129}. When both receptors are expressed, ER β has been found to reduce ER α 's function as a transcriptional activator while promoting anti-proliferative and pro-apoptotic activities^{128,130}. Numerous genes controlled by ER β , but not ER α , have been found; and researchers are determining whether these transcripts have anti-proliferative or pro-apoptotic features that might explain ER β 's tumour suppressor function^{128,131,132}.

7. Multidrug Resistance

Drug inactivation and inhibition are caused by various indigenous and foreign factors, including cellular reprogramming, neoplastic stimulation, drug efflux due to the over-expression of MDR genes, and metabolic changes. Changes in indigenous and external stimuli result in the DNA damage repair mechanism, evasion of programmed cell death, and epithelial–mesenchymal translocation^{133–135}. The unpredictability of the acquired cell death evasion strategies of the cancer cells may also compromise immune surveillance¹³⁶. Furthermore, the link between cancer and oxidative stress has been investigated thoroughly, which indicates that ROS plays a crucial role in cancer progression¹³⁷. An imbalance in redox homeostasis is a key element in the development of cancer treatment resistance. As oxidative stress is important for cell survival, it may also contribute to cancer medication resistance¹³⁸.

Single-Agent Resistance (SAR) or MDR might be confined to the medicine that is used to treat a patient.

Owing to its ability to disrupt the expected medication response in cancer patients, resistance to several treatments during cancer therapy has been a “clinician’s nightmare.” As a result, dealing with drug resistance in a cancer treatment program is difficult^{100,139}.

8. Mechanisms of Drug Resistance

An increase in the expression of ATP-Binding Cassette (ABC) carrier proteins, which export substrates across the cell membrane through ATP hydrolysis, drives cancer therapy resistance¹⁴⁰. These drug efflux transporters decrease the cellular concentration of the drug and compromise treatment responsiveness¹⁴¹. Research links 48 ABC transporters to humans¹⁴². Many of them protect the kidney, pancreas, liver, GI tract, testes, and brain endothelial arteries¹⁴³. There are 13 ABC transporters are linked to cancer therapy resistance. ABCB1 (permeability glycoprotein/MDR1)¹⁴⁴, ABCC1 (MDR related protein-1, MRP1), and ABCG2/BCRP (breast cancer resistance protein) have been studied carefully to better understand multidrug resistance. ABC transporters are involved in the removal of xenobiotics and harmful endogenous chemicals from cells and organs in order to maintain interstitial equilibrium. These membrane-bound transporters are used by cancer cells to develop drug resistance¹⁴⁵. Domains were constructed by utilizing Illustrater of Biological Sequences, and these sequences were acquired from Uniprot Database^{146,147}.

In the processes of these ABC transporters, ATP plays a significant role. ATP levels in drug-resistant cancer cells are higher than in parental cancer cells¹⁴⁸. When ATP levels are low, cancer cells become much more susceptible to treatment. Drug-sensitive cells become drug-resistant as their intracellular ATP concentrations increase¹⁴⁹. Extracellular ATP also promotes the expression of ABC transporters, resulting in increased drug efflux¹⁵⁰. Extracellular ATP concentrations are known to increase the tumour microenvironment¹³³. With the help of a mechanism known as “macropinocytosis,” cancer cells consume this extracellular ATP. Numerous chemotherapeutic drugs become resistant as a result of the considerable increase in intracellular ATP concentration¹⁵¹. Accumulating evidence has shown the combination of therapies in tamoxifen-resistant cancer, a higher opportunity to develop MDR, and stem-like phenotypes in breast cancer^{100,139,152}.

9. Drug Resistance and Epigenetics

Epigenetics determines cell destiny and pathogenic provenience. Nongenetic heterogeneity causes tumour-initiating cells and/or therapy resistance. Epigenetic changes produce inadequate gene expression, which lasts across cell cycles and contributes to nongenetic variation and resistance to treatment¹⁵³.

Precision oncology benefits from epigenetic-based diagnostic and prognostic techniques substantially. Numerous genetic diagnostic tests are in clinical studies or usage¹⁵⁴. Epidrugs, or drugs that target epigenetic modulators, were developed as a result of precision oncology efforts to address dysregulated epigenetic pathways¹⁵⁵.

Epigenetic modifications govern gene transcription via altering chromatin packaging and regulate DNA accessibility to sequence-specific transcription factors. Genetic variation, chromatin rearrangement, histone modification, and non-coding RNA alterations are linked to cancer chemoresistance^{155,156} (Figure 2).

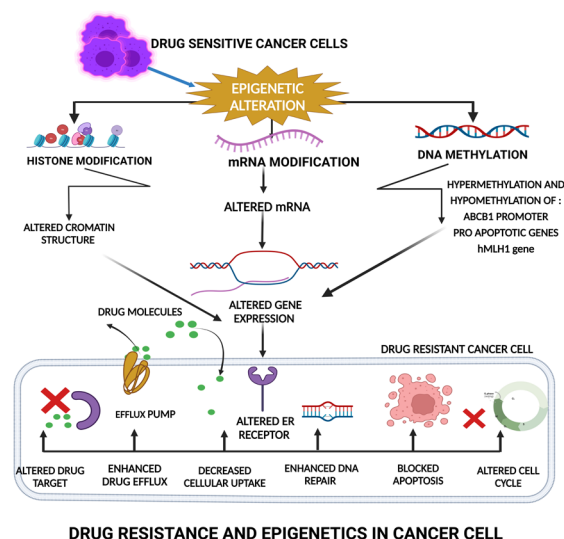


Figure 2. Drug-sensitive cancer cells upon epigenetic alterations lead to histone modification, methylation, and mRNA modification, which lead to altered gene expression. These alterations in gene expression lead to drug target site alteration, cell cycle and ER receptor alteration, decreased cellular uptake, enhanced drug efflux, blocked apoptosis, and enhanced cellular repair; thereby, making the drug-sensitive cancer cell resistant to the drug.

10. Molecular Mechanism

Aberrant modification of CpG islands around gene promoter regions induces gene silencing during tumour development, as governed by molecular mechanisms¹⁵⁷. DNA methylation is associated with Histone Deacetylases (HDACs), chromatin condensation, and gene silencing¹⁵⁸. The frequency of epimutations is far larger than that of genetic mutations. According to the study, 61 unusual mutations were detected during metastasis, 15 of which were driver genes and the rest were altered passenger genes. As a result, these modifications have a smaller impact on sub-population selection related to tumour formation and medication resistance. According to the accumulated evidence, demethylation of the MDR1 protagonist appears to be associated with (MDR) in a range of tumour types^{159,160}. Epigenetic alterations improve DNA damage repair in cancer cells by reviving the DNA repair molecule MGMT, which enhances the survival of malignant cells. Cancer cells affect the sensitivity to modification and epigenetic suppression of proapoptotic genes, such as *APAF1* and *hMLH1*, and oncogenes and tumour suppressor genes, such as *BRCA1* and *E-cadherin*. Exosomes are also thought to have a role in epigenetic alterations, according to recent research¹⁶¹. According to another recent study, exosomes are considered to play a role in tumour growth, cell proliferation, and metastasis. Extracellular vesicles carry proteins and nucleic acids to target cells, affecting chromatin alteration, chromatin structure, and RNA post-transcriptional control¹⁶². Tumour microenvironment fibroblasts and immunocytes release exosomes where they transport a range of loadings and microRNAs (miRNAs). Exosomal miRNAs intervene in drug resistance mechanisms, such as medicine efflux, medication metabolism changes, drug target mutations, DNA repair, metabolic alterations, cancer stem cells, and epigenetic modifications¹⁶³. Exosomal miRNAs promote resistance to medication.

11. ER α Signaling and Epigenetics

Estradiol activates ER target sites, which recruit hundreds of ER co-regulators to chromatin to ensure optimum transcriptional and repressive activity. Even though each ER molecule resides on chromatin for only seconds, it cycles on and off for minutes and hours after estradiol stimulation. ER receptor coactivators such as P300/CBP, SWI/SNF and PRMTs, are activates the epigenetic

changes associated with them. SRC-1, SRC-2, and SRC-3 attach to ER physically and serve as a platform for ER to recruit additional stimulating kinases and chromatin remodelling complexes¹⁶⁴. P300, a histone acetyltransferase (HAT), communicates with ER-bound enhancers through SRC-3 to acetylate histone H3 lysine 27 (H3K27ac), thereby activating enhancers^{165,166}. Estradiol elevates H3K27ac concentrations at loci in which the ER has a regulatory effect¹⁶⁷⁻¹⁶⁹. An increase in the H3K27ac at ER-bound enhancers corresponds with BRG1, the SWI/SNF catalytic element, which shows that ER promotes the SWI/SNF complexes to redesign and activate enhancers¹⁷⁰. NCoR1, NCoR2, and LCoR engage with an epigenetic suppressor to down-regulate estradiol-repressed genes¹⁷¹. The ER corepressor BRCA1 is well-known. After binding to the AF2 area, BRCA1 monoubiquitinates the ER, thereby decreasing its transcriptional activity¹⁶⁸. The transcriptional and oncogenic effects of these coregulators have been well-studied¹⁷²⁻¹⁷⁴. Estradiol-dependent ER recruitment requires colonial transcription factors. Pioneer transcription factors in ER breast cancer have been studied¹⁶⁹. GRHL2 shows numerous functional similarities to FOXA1 and GATA3, including estradiol-independent chromatin assembly and ER target gene regulation^{170,171,175}. PRC1 and PRC2 increase estradiol-induced ER corresponding transcriptional activation in breast cancer cells by recruiting chromatin^{168,176-178}. Polycomb complexes have also been discovered to have both repressive and activating effects in stem cells, embryonic development, and cancer cells, as well as the activities that take place inside these cells¹⁷⁶. The roles of GRHL2 and polycomb-group proteins as ER signalling regulators in ER+ breast cancers require further investigation (Figure 3).

12. Future Research Directions of Tamoxifen in Cancer Treatment

Tamoxifen is a potent anticancer drug that plays a vital role in ER-positive breast cancer patients. The main drawback of this drug is its drug resistance, which most probably limits its efficacy, thereby emphasizing the importance of overcoming tamoxifen resistance in breast cancer. The methods to overcome breast cancer resistance are based on the major mechanism of drug resistance. Mechanistic pathways are inhibited, such

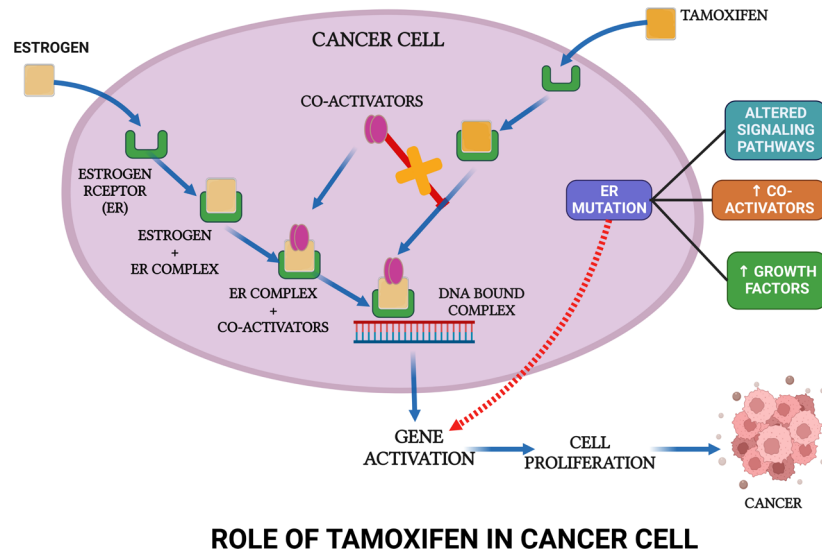


Figure 3. Tamoxifen is an estrogen competitor that competes for the active site on the Estrogen Receptor (ER) and blocks estrogen from binding to its receptor. Tamoxifen functions as an anti-estrogen by inhibiting the co-activators that bind with the ER complex. Upon tamoxifen binding, transcription of the ER complex is inhibited, as a result of which cancer cell growth and proliferation cease. However, an increase in the number of growth factors, co-activators, and altered pathways lead to alterations in ER and, ultimately, led to cancer cell formation.

as the RTK pathway, blocking protective autophagy, upregulation of ER α 36, cell cycle regulators, and EMT-like specific phenomenon. Some novel methods have broadened our vision to overcome the drug resistance of tamoxifen totally, by using combinational drug therapy with other drugs that can inhibit the development of drug resistance. In recent years, most of the research on the relationship among the major areas of autophagy, cell cycle regulators, and tamoxifen resistance has made great progress. Hence, these methods to overcome drug resistance by inducing autophagy mechanisms are limited most probably in the current research due to the inhibition of autophagy by its inhibitors.

The continued search for better autophagy inhibitors to overcome the resistance has thrown light to hypothesize that tamoxifen combined with other drugs may protect the mitochondrial function which in turn enhanced autophagy and overcome the drug resistance of tamoxifen. This view is a new arena to improve the drug resistance of tamoxifen, and more research studies are needed to explore a clear view in this field. The main target to overcome tamoxifen resistance is to target LEM4, which is a feasible direction for research in the future. It is already evident that the high expression of LEM4 in

drug-resistant cells is an important biological mechanism involved in the attenuation of the inhibitory effect of tamoxifen on the G1-S transition. Hence, targeting LEM4 will play a potent role in overcoming tamoxifen resistance in breast cancer and will be an important tool in the future.

13. Conclusion

Tamoxifen treatment epigenetically altered gene expression associated with changes in coregulators, epigenetics and post-translational changes, and genetic variations in the ER pathway may cause endocrine resistance. “Epigenetic treatment,” which appears to be promising, could be a solution to the problem of medication resistance in the cancer field. HDAC inhibitors, for example, have shown the potential to lower antiestrogen-resistant cell proliferation.

Finally, new information has emerged on the possible routes that contribute to tamoxifen resistance. Tamoxifen-resistant animals or patients show ER loss or alterations, decreased intracellular tamoxifen concentrations, and altered metabolite profiles, according to various scientific investigations. According to recent studies, breast cancer

cells become MDR and stem-like due to the treatment with tamoxifen. Research into epigenetic regulation in MDR and tamoxifen resistance can potentially lead to new cancer treatment regimens, which will improve patient outcomes.

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15. References

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