

Review Article

Endocrine-Resistant Breast Cancer: Underlying Mechanisms and Strategies for Overcoming Resistance

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Estrogen plays important roles in the development and progression of breast cancer. However, one-third of breast cancers fail to respond to endocrine therapy and most endocrine-responsive breast cancers subsequently become resistant to endocrine therapy. A tremendous effort has been made to elucidate the mechanisms responsible for the development of endocrine-resistance in breast cancer. Since the main target molecule of estrogen in breast cancer is estrogen receptor (ER)- α , most studies have focused on investigating quantitative and qualitative changes in ER- α in endocrine-resistant breast cancer. Breast cancers expressing no ER- α fail to respond to endocrine therapy. Some breast cancers expressing ER- α also fail to respond to endocrine therapy and most breast cancers with acquired endocrine resistance retain ER- α expression, which suggests that the disappearance of ER- α in breast cancer cells is not a common cause of resistance to endocrine therapy. Recent molecular biological studies have shown evidence that qualitative and functional changes, such as gene mutations and phosphorylation of ER- α , cause endocrine resistance in breast cancer. In addition, it has been suggested that endocrine resistance could be induced by epigenetic changes, such as hypoxia, in breast cancer tissues. Understanding the precise mechanisms that underlie endocrine resistance may enable clinicians to develop new strategies for retarding or overcoming endocrine resistance in breast cancer.

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Key words: Endocrine resistance, Breast cancer, Growth factor, Signaling pathway, Hypoxia

Accumulated knowledge provided from the results of clinical studies has suggested that endocrine therapy is useful for chemoprevention and treatment for patients with clinical breast cancer in either adjuvant or metastatic settings. However, the clinical benefit provided by endocrine therapy is not available to patients with estrogen receptor (ER)-negative and progesterone receptor (PgR)-negative breast cancer. In addition, one-third of ER-positive recurrent breast cancer cases fail to respond to endocrine therapy. Furthermore, even if recurrent breast tumors respond to first- and/or second-line endocrine therapies, they fre-

quently develop resistance to these therapies after varying durations of response. Most endocrine-responsive breast tumors ultimately become endocrine-resistant.

Breast cancer with acquired resistance to first-line endocrine therapies, such as the antiestrogen tamoxifen frequently responds to second-line endocrine therapies, such as aromatase inhibitors or progestins. In addition, third-line endocrine therapy sometimes provides clinical benefit to patients that suffer from endocrine-refractory breast cancer. These findings support the clinical strategy of treating ER- and/or PgR-positive breast cancer with as many endocrine therapies as possible, and subsequently applying cytotoxic chemotherapy. These findings also suggest that there are several steps or complicated mechanisms causing endocrine resistance.

It should be noted that endocrine resistance is a term sometimes used to designate resistance to a certain endocrine therapeutic agent, such as tamoxifen. Tamoxifen resistance is a common problem in clinics and the underlying mecha-

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Abbreviations:

ER, Estrogen receptor; PgR, Progesterone receptor; LH-RH, Luteinizing hormone-releasing hormone; AP-1, Activating protein-1; EGF-R, Epidermal growth factor receptor; PI3K, Phosphatidylinositol-3-OH-kinase; HAT, Histone acetyltransferase; ERBF1, Estrogen receptor promoter B-associated factor 1

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nisms have been extensively investigated for the last two decades. Endocrine resistance in breast cancer was divided into four categories by possible mechanisms in the present review as follows: 1) specific resistance to antiestrogens, 2) ligand-independent ER activation, 3) modified ER-interacting proteins, and 4) loss of ER expression. In addition, possible clinical approaches to retard or overcome endocrine resistance are discussed.

Possible Mechanisms of Action Underlying Endocrine Resistance (Table 1)

1) Specific Resistance to Antiestrogens

Tamoxifen has been used as a chemoprevention agent (unfortunately, not in Japan) as well as a therapeutic agent for patients with breast cancer in the preoperative, postoperative and metastatic settings. Other antiestrogens, such as toremifene, have also been used as alternatives to tamoxifen. Therefore, antiestrogen resistance is very common in clinics. In addition, tamoxifen-withdrawal response and tamoxifen-stimulated tumor growth have been observed in clinics and experimental studies¹⁻³. Tamoxifen might be deleterious in some antiestrogen-resistant tumors. Several hypothetical mechanisms responsible for antiestrogen resistance have been advocated.

1) Modified tamoxifen metabolism is suspected to cause tamoxifen resistance. Approximately one decade ago, tamoxifen metabolites, a less potent antiestrogenic cis-isomer of 4-hydroxytamoxifen and an estrogenic metabolite, metabolite E, were detected in both animal models and tamoxifen-resistant breast tumors⁴. However, the causal relation between the increased levels of these metabolites and tamoxifen resistance remains to be clarified.

2) Tamoxifen has been known to stimulate ovarian estrogen production and an elevated level of serum estrogen may interfere with the antiestrogenic activity of tamoxifen⁵. Actually, combined tamoxifen and luteinizing hormone-releasing hormone (LH-RH) agonists has been suggested to be superior to either agent alone in clinics⁶.

3) Enhancement of estrogenic activity of antiestrogens by a certain mutational change in the ER- α gene or the prominent activating protein (AP)-1-mediated transactivation might induce antiestrogen resistance as well as antiestrogen-stimulated growth^{7,8}. However, no common mutation in the ER- α gene has been detected in antiestrogen-resistant breast tumors. A series of exper-

Table 1. Action Mechanisms Underlying Endocrine-Resistance in Breast Cancer

Specific Resistance to Antiestrogens
Modified metabolism of antiestrogens
Stimulated estrogen synthesis in ovaries
Enhanced agonistic activity by ER mutations or AP-1 activation
Ligand-independent ER Activation
Phosphorylation of ER by growth factor signaling pathway
ER mutations or ER mRNA variants
Modified ER-interacting Proteins
Quantitative or qualitative changes in cofactors
Interaction with ER- β and its isoforms
Loss of ER Expression
Hypermethylation in the ER promoter
Loss of trans factors, such as ERBF-1
Epigenetic changes, such as hypoxia

imental and clinical results suggests a positive correlation between up-regulation of AP-1 activity and acquired tamoxifen resistance^{9,10}. It has been suggested that cellular oxidative stress induced by tamoxifen might activate signaling pathways leading to the activation of AP-1⁸.

2) Ligand-Independent ER Activation

It has been classically presumed that transactivation of target genes by ER requires a ligand, such as estrogen, binding to ER. Recently, experimental studies have revealed that several growth factors and their intracellular signaling pathways can stimulate ER activity in the absence of a ligand. In addition, it has been suggested that this ligand-independent ER activation is caused by the phosphorylation of different sites in the ER protein. These mechanisms may explain the development of endocrine-resistant breast cancer.

1) Overexpression of HER1 (also known as epidermal growth factor receptor, EGF-R) and/or HER2 (also known as ErbB-2) has been detected in 20-30% of primary breast cancers. Experimental and clinical studies have suggested that overexpression of HER1 and/or HER2 causes tamoxifen-resistance^{11,12}. Recent fundamental studies also revealed that the intracellular signaling pathway activated by HER1 or HER2, RAS/RAF/MEK/ERK1/2, phosphorylates the serine 118 residue located in ER and results in ligand-independent transactivation of target genes of ER¹³⁻¹⁵. It was also reported that increased ERK1/2 activity correlates with endocrine resistance and shorter survival in patients with breast cancer¹⁶. Moreover, experimental studies have suggested that the

blockade of these signaling pathways by EGF-R tyrosine kinase inhibitors, MEK inhibitors and the anti-HER2 monoclonal antibody trastuzumab may restore the anti-tumor effect of antiestrogens¹⁷⁻¹⁹.

2) Another intracellular signaling pathway, which is also stimulated by receptor tyrosine kinases, such as HER1 and HER2, is phosphatidylinositol-3-OH-kinase (PI3K)/AKT (also known protein kinase B) pathway. It was reported that phosphorylation of the serine 167 residue located in ER by AKT results in ligand-independent activation of ER^{20, 21}. In addition, a negative regulator of AKT, PTEN, which acts as a tumor suppressor, is frequently inactivated in breast cancer²².

3) Other intracellular signaling pathways, such as protein kinase A induced by G-protein-coupled receptors and the 90-kDa ribosomal S6 kinase RSK, are also known to be involved in ER phosphorylation^{23, 24}. However, the roles of these pathways in the development of endocrine resistance remain to be elucidated.

There are other mechanisms that result in ligand-independent ER activation. Certain mutations or mRNA variants of the ER gene may produce a constitutively active form of ER protein.

1) A missense mutation of the tyrosine 537 residue in the ligand binding region of ER was detected in metastatic breast cancer tumors. This mutant ER was reported to be equally active in the absence of ligand or in the presence of estradiol or tamoxifen. Substitution of this residue by different amino acids has been shown to result in the ligand-independent activation of ER together with co-activator recruitment^{25, 26}. However, other studies suggest that only 1% of primary breast cancers carry gene mutations of ER in the coding region, most of which did not result in alteration at the protein level²⁷. Therefore, the clinical significance of the mutation in endocrine resistance is questionable.

2) Alternatively spliced mRNA lacking specific exons, which encode truncated and potentially constitutively active forms of ER, has been discovered²⁸. However, analysis of breast tumors failed to confirm a significant role of the variants in the development of endocrine-resistance. The pathological significance of the ER mRNA variants remains to be determined²⁹⁻³¹.

3) Modified ER-Interacting Proteins

A series of recent studies have revealed that ER-interacting proteins, called cofactors, which include co-activators and co-repressors, have

important roles in mediating transcriptional activation of target genes by the ER. These findings suggest that quantitative or qualitative changes of the ER-related cofactors could contribute to the development of endocrine-resistant breast cancer. However, no significant relation has been confirmed between alterations of co-activators or co-repressors and endocrine resistance in breast tumors³². However, some experimental findings support these attractive mechanisms.

1) Overexpression of nuclear receptor co-activator protein (NCOA)-1 was reported to increase the agonist activity of tamoxifen in experimental cell systems³³. This suggests that co-activator overexpression could contribute to endocrine-resistance. In addition, it might be possible that increased histone acetyltransferase (HAT) activity caused by the co-activator up-regulation is also involved in the development of endocrine resistance. Therefore, newly developed HAT inhibitors might be useful in the treatment of endocrine-resistant breast tumors³⁴.

2) In a mouse model system, it was reported that expression of one of the co-repressors, nuclear co-repressor protein (NCOR)-1, was decreased in tamoxifen-resistant tumors³⁵. However, no other studies support this interesting finding in human endocrine-resistant tumors.

3) A novel co-activator, L7/SPA, was discovered and the complex of this co-activator and tamoxifen was reported to lead to increase transcription activation in target genes³⁶. Overexpression of this kind of co-activator may contribute to endocrine resistance. However, no study has so far clarified this hypothetical mechanism.

4) A recently discovered ER subtype, ER- β , may play a role in the development of endocrine resistance. A series of studies suggested that the expression of wild-type ER- β in breast cancer cells predicts a favorable prognosis, but an isoform of ER- β gives a poorer prognosis³⁷⁻³⁹. Actually, experimental studies suggested that this isoform reduces ER- α -mediated transactivation in a dominant negative fashion⁴⁰. Further studies are needed to clarify the role of ER- β in the development of endocrine resistance.

4) Loss of ER Expression

Endocrine-resistant breast tumors do not always lose ER- α expression. Therefore, loss of ER- α expression is only one of the possible mechanisms of endocrine-resistance in breast cancer. However, this mechanism of action has long been suggested

Table 2. Clinical Strategies for Overcoming Endocrine-Resistance in Breast Cancer

Inhibition of Growth Factor Signaling Pathway
Purpose: Blocking ligand-independent ER activation
Candidates: trastuzumab, ZD1839 and other tyrosine kinase inhibitors
Retardation of the Development of Resistance
Intermittent endocrine therapy
Alternative endocrine therapy
Combined endocrine therapy
Hypoxic cytotoxins

in Japan as follows: 1) ER-positive breast cancer could consist of so-called mosaic cell mixtures with ER-positive and negative tumor cells. 2) When endocrine therapy effectively kills ER-positive tumor cells, ER-negative tumor cells could selectively grow. 3) Finally, this breast cancer loses ER expression and acquires total endocrine resistance⁴¹.

Another possible mechanism leading to loss of ER expression is that some genetic or epigenetic changes in breast tumors gradually cause a reduction in ER expression. For example, hypermethylation of the promoter region of the ER gene causes transcriptional down-regulation and decreases the expression of ER protein⁴². Another cause of transcriptional down-regulation is decreased expression of transcription-promoting trans elements for ER, such as estrogen receptor promoter B-associated factor (ERBF) 1⁴³. However, it is not yet known whether these changes directly cause loss of ER or a loss of ER caused by other mechanisms subsequently induces these changes.

Recently, the inverse relation between the expression of ER and hypoxia-inducible factors such as hypoxia-inducible factor-1 α and carbonic anhydrase IX, was reported in breast cancer tissues^{44, 45}. In addition, our previous experimental study suggested that hypoxic conditions lead to the post-transcriptional down-regulation of ER- α and its function⁴⁴. These findings suggest that hypoxic microenvironments, which are frequently observed in breast tumors, cause a decrease in ER expression. It has been suggested that hypoxia drives the malignant progression of many types of human cancers⁴⁶. Therefore, it might be possible that hypoxic microenvironments caused by an increase in tumor size or induced by hormonal agents provide a selective pressure for the outgrowth of ER-poor and more aggressive clones and finally cause a loss of ER expression in breast

tumors.

New Strategies against Endocrine-Resistant Breast Cancer (Table 2)

As described above, there are many hypothetical action mechanisms underlying endocrine resistance in breast cancer. Some appear to be irreversible and impossible to overcome. However, others appear to be reversible and might be overcome by new strategies. In the last part of this review, new clinically testable strategies are discussed.

1) Inhibition of Growth Factor Signaling Pathway

It was suggested that the up-regulation of growth factor-mediated signaling pathways causes ER phosphorylation, induces ligand-independent ER activation and leads to endocrine-independent and -resistant growth of breast cancer^{11-15, 20, 21}. Some recent studies also provided clinical evidence suggesting that breast cancer overexpressing HER1 and/or HER2 together with ER expression is likely to be resistant to endocrine therapy^{47, 48}. These findings suggest the following: 1) Inhibitors of the growth factor signaling pathway, such as tyrosine kinase inhibitors, may decrease ER phosphorylation and ligand-independent ER activation. 2) This may induce the estrogen-dependent growth pathway. 3) Then, endocrine therapeutic agents effectively inhibit the growth of breast cancer.

The humanized anti-HER2 monoclonal antibody, trastuzumab, is the only agent that inhibits the growth factor-mediated signaling pathway and has been introduced into clinics⁴⁹. Our previous experimental study suggested that combined treatment with antiestrogen and this antibody effectively inhibits the growth of breast cancer cells expressing both ER and HER2¹⁹. Several clinical studies launched worldwide tested the clinical usefulness of this combination therapy⁵⁰. Howev-

er, a recent clinical study, in which the anti-tumor effects of the antiestrogen tamoxifen and the aromatase inhibitor letrozole were compared in the neoadjuvant setting, suggested that tamoxifen was significantly less effective than letrozole in ER-positive breast cancer overexpressing HER1 and/or HER2⁵¹. It was reported that tamoxifen might be deleterious or ineffective in the adjuvant setting for patients with breast cancer overexpressing HER2⁵². These findings suggest that combined treatment with an aromatase inhibitor and this antibody is more suitable for clinical studies recruiting patients with breast cancer overexpressing HER2.

Very recently, an EGF-R-specific tyrosine kinase inhibitor, ZD1839, was introduced into clinics for the treatment of non-small cell lung cancer⁵³. This agent effectively inhibits the signaling pathway mediated by EGF-R⁵⁴. It was reported that approximately 20% of breast cancers overexpress EGF-R, and breast cancer overexpressing EGF-R is more aggressive and resistant to endocrine therapy⁵⁵. Our preliminary study also suggested that ER-positive breast cancer overexpressing EGF-R more rapidly acquires endocrine-resistance (unpublished data). Moreover, experimental studies have suggested that ZD1839 is not only effective in breast cancer cells overexpressing EGF-R but is also effective in breast cancer overexpressing HER2^{54, 56}. These findings suggest that ZD1839 is also a promising agent for combined use with endocrine therapeutic agents.

2) Retardation of the Development of Endocrine-Resistance

Even if it is difficult to overcome endocrine resistance, it may be possible to prolong the anti-tumor effect of endocrine therapy or retard the occurrence of endocrine resistance. Two hypothetical strategies are introduced and discussed below.

1) Acquired endocrine-resistance has been recognized as an adaptive response of breast cancer cells to endocrine therapy. If so, it can be suggested that intermittent treatment with a single endocrine therapeutic agent, combined treatment with two different agents or alternating use of two different agents may interfere with the adaptation process and retard the occurrence of endocrine-resistance. Prostate-specific antigen is a very sensitive and specific tumor marker, and a new strategy, in which intermittent treatment with an LH-RH agonist is given to patients with advanced prostate

cancer, has been tested in clinical trials⁵⁷. In addition, a recent clinical study suggests that alternating treatment with tamoxifen and the aromatase inhibitor aminoglutethimide was more effective than tamoxifen alone in the adjuvant setting for patients with breast cancer⁵⁸. These findings prompted us to start large-scaled prospective clinical trials to test these new treatment modalities.

2) Our recent experimental study suggested that an epigenetic change caused by a hypoxic microenvironment may play an important role in the development of endocrine resistance in breast cancer⁴⁴. If so, hypoxic cytotoxins, such as tirapazamine⁵⁹, which specifically kill cancer cells under hypoxic conditions, may be useful for retarding the occurrence of endocrine resistance. Our preliminary findings suggest that the administration of hypoxic toxins restores ER expression in human breast cancer xenografts in athymic nude mice (unpublished data). Further experimental studies are needed to clarify this novel treatment strategy.

Conclusions

Although there are many attractive mechanisms of action possibly responsible for the development of endocrine resistance, a common cascade has not yet been discovered. In other words, many mechanisms may be involved. From a clinical point of view, endocrine therapy appears to be less toxic and provides better quality of life for patients with breast cancer compared with cytotoxic chemotherapy. Therefore, clinicians and researchers have to make a serious effort to develop new strategies for enhancing or prolonging the efficacy of endocrine therapy. With regard to clinical implications, strategies for retarding endocrine resistance, such as the intermittent, alternating or combined endocrine therapy, are testable in clinical trials. In addition, the administration of inhibitors for the growth factor signaling pathway may not only prolong the efficacy of endocrine therapy but also overcome a certain type of endocrine resistance in the near future. Well-designed clinical trials supported by scientific evidence are essential to clarify new promising strategies.

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