Abstract: Owing to improved early surveillance and advanced therapy strategies, the current death rate due to breast cancer has decreased; nevertheless, drug resistance and relapse remain obstacles on the path to successful systematic treatment. Multiple mechanisms responsible for drug resistance have been elucidated, and miRNAs seem to play a major part in almost every aspect of cancer progression, including tumorigenesis, metastasis, and drug resistance. In recent years, exosomes have emerged as novel modes of intercellular signaling vehicles, initiating cell–cell communication through their fusion with target cell membranes, delivering functional molecules including miRNAs and proteins. This review particularly focuses on enumerating functional miRNAs involved in breast cancer drug resistance as well as their targets and related mechanisms. Subsequently, we discuss the prospects and challenges of miRNA function in drug resistance and highlight valuable approaches for the investigation of the role of exosomal miRNAs in breast cancer progression and drug resistance.

Keywords: microRNA, exosome, breast cancer, drug resistance

Introduction

Breast cancer (BCa) is the most common invasive malignancy and the second leading cause of tumor-related death among women worldwide. Due to the improved early diagnosis and advanced therapeutic strategies, the mortality of BCa has been appreciably decreased. However, although the majority of patients were responsive to the initial treatment for certain time, the patients eventually developed more aggressive tumor forms that were generally resistant to the chemotherapy and radiotherapy. Thus, as a major challenge, metastasized recurrence leading to a poor prognosis remains to be conquered in BCa therapeutics. Current therapeutic strategies for treating BCa usually combine surgery with multiple adjuvant therapies, such as chemotherapy, radiotherapy, hormonal therapy, targeted therapy, or combination thereof. However, multiple drug resistance (MDR) is a major obstacle for successful systemic therapeutics. Multiple mechanisms involved in drug resistance have been elucidated in BCa treatment, including somatic mutations or epigenetic modifications within drug targets, deregulated apoptosis or survival, adapted signaling pathways or metabolic reprogramming, as well as drug reaching to the targets or drug transport modification within the tumor microenvironment. In this context, abnormal expression of miRNA leading to deregulation of gene expression has been implicated in the therapeutic resistance. miRNAs, a group of 20–25 nucleotides noncoding RNA that specifically bind to the 3′-untranslated region of mRNA of the target genes, negatively regulate gene expression at post-transcriptional level. It has been well recognized that miRNA-mediated downregulation of gene expression is involved in tumorigenesis, metastasis, and tumor response to treatment. Particularly, increasing evidences have demonstrated that the...
dysfunction of miRNA contributes to therapeutic resistance in BCa. As shown in Tables 1–4, a number of miRNAs are deregulated in BCa, which is associated with radioresistance and MDR including chemotherapy, endocrine therapy, and targeted therapy.\(^9\) However, individual miRNA can repress several hundred target genes, and single target gene is often regulated by multiple miRNAs. Importantly, exosomes function as vehicles delivering miRNAs from donating cells to accepting cells in the tumor microenvironment, which plays a pivotal role in the regulation of miRNA functions.\(^11\)

Exosomes, extracellular vesicles secreted from various types of cells, carrying small molecules including miRNAs and proteins.\(^12\) Initially, exosome was identified as cell debris containing metabolic products. However, the recent studies have verified that exosome is critical for cell–cell communication.\(^13\) As key components in the tumor microenvironment, exosomes shuttle bioactive molecules and lead to the exchange of genetic information and metabolic reprogramming of the recipient cells.\(^14\) Thus, exosome is thought to be critical for signal transduction between cells, and the function of exosome directly contributes to the promotion of proliferation, invasiveness, and metastasis of recipient cells, as well as transferring drug resistance, thereby enhancing tumor recurrence and metastasis.\(^15\) Our previous studies showed that tumor-derived exosomes transfer doxorubicin resistance to the susceptible BCa cells through exosomes shuttling miR-222.\(^16\)

In this review, we focus on specific miRNAs associated with BCa drug resistance. Since substantial evidences predict that exosome-mediated dynamic and reciprocal cross-talks are essential for sustaining specific tumor microenvironment, we particularly highlight the recent novel findings that revealed the aspects of exosomal miRNAs implicated in drug resistance, which may be beneficial for seeking

### Table 1 miRNAs involved in the regulation of chemotherapeutic resistance in breast cancer

<table>
<thead>
<tr>
<th>miRNA</th>
<th>Drug</th>
<th>Target(s)</th>
<th>Expression level*</th>
<th>Mechanism</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>miR-505</td>
<td>Doxorubicin</td>
<td>Akt3 (indirect)</td>
<td>↓</td>
<td>Not investigated</td>
<td>19</td>
</tr>
<tr>
<td>miR-181a</td>
<td>Doxorubicin</td>
<td>Bcl-2</td>
<td>↓</td>
<td>Increase apoptosis</td>
<td>24</td>
</tr>
<tr>
<td>miR-663</td>
<td>Doxorubicin</td>
<td>HSPG2</td>
<td>↑</td>
<td>Inhibit apoptosis</td>
<td>7</td>
</tr>
<tr>
<td>miR-25</td>
<td>Doxorubicin</td>
<td>ULK1</td>
<td>↑</td>
<td>Inhibit autophagy</td>
<td>6</td>
</tr>
<tr>
<td>miR-145</td>
<td>Doxorubicin</td>
<td>MRPI</td>
<td>↓</td>
<td>Induce intracellular doxorubicin accumulation</td>
<td>8</td>
</tr>
<tr>
<td>miR-644a</td>
<td>Doxorubicin</td>
<td>CTBP1</td>
<td>↓</td>
<td>Inhibit EMT</td>
<td>95</td>
</tr>
<tr>
<td>miR-128</td>
<td>Doxorubicin</td>
<td>Bmi-1, ABCC5</td>
<td>↓</td>
<td>Increase apoptosis</td>
<td>23</td>
</tr>
<tr>
<td>miR-30c</td>
<td>Doxorubicin</td>
<td>TWF1 (PTK9)</td>
<td>↓</td>
<td>Reverse EMT</td>
<td>32</td>
</tr>
<tr>
<td></td>
<td>Paclitaxel</td>
<td>VIM, IL-11</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>miR-326</td>
<td>Doxorubicin</td>
<td>MDR-1</td>
<td>↓</td>
<td>Downregulate MRP-1</td>
<td>36</td>
</tr>
<tr>
<td>miR-181a</td>
<td>Doxorubicin</td>
<td>Bax</td>
<td>↑</td>
<td>Inhibit apoptosis</td>
<td>86</td>
</tr>
<tr>
<td>miR-106b–25 cluster</td>
<td>Doxorubicin</td>
<td>EP300</td>
<td>↑</td>
<td>Activate EMT</td>
<td>25</td>
</tr>
<tr>
<td>miR-34a</td>
<td>Doxorubicin</td>
<td>HDAC1, HDAC7</td>
<td>↑</td>
<td>Inhibit autophagic cell death</td>
<td>30</td>
</tr>
<tr>
<td>Cisplatin</td>
<td>Cisplatin</td>
<td>HDAC7</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>miR-345</td>
<td>Cisplatin</td>
<td>MRPI</td>
<td>↓</td>
<td>Uncharacterized</td>
<td>27</td>
</tr>
<tr>
<td>miR-7</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>miR-302b</td>
<td>Cisplatin</td>
<td>E2F1 (direct)</td>
<td>↓</td>
<td>Inhibit cell cycle progression</td>
<td>28</td>
</tr>
<tr>
<td></td>
<td></td>
<td>ATM (indirect)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>miR-24</td>
<td>Cisplatin</td>
<td>BimL, F1H1</td>
<td>↑</td>
<td>Promote EMT and CSCs</td>
<td>79</td>
</tr>
<tr>
<td>miR-129-3p</td>
<td>Docetaxel</td>
<td>CP100</td>
<td>↑</td>
<td>Reduce cell cycle arrest and apoptosis</td>
<td>33</td>
</tr>
<tr>
<td>miR-34a</td>
<td>Docetaxel</td>
<td>BCL-2, CCND1</td>
<td>↑</td>
<td>Inhibit apoptosis</td>
<td>87</td>
</tr>
<tr>
<td>miR-125b</td>
<td>Paclitaxel</td>
<td>Sema4C</td>
<td>↓</td>
<td>Reverse EMT</td>
<td>35</td>
</tr>
<tr>
<td>miR-100</td>
<td>Paclitaxel</td>
<td>mTOR</td>
<td>↓</td>
<td>Enhance cell cycle arrest and apoptosis</td>
<td>31</td>
</tr>
<tr>
<td>miR-125b</td>
<td>Taxol</td>
<td>Bak1</td>
<td>↑</td>
<td>Inhibit apoptosis</td>
<td>34</td>
</tr>
<tr>
<td>miR-484</td>
<td>Gemcitabine</td>
<td>CDA</td>
<td>↓</td>
<td>Promote proliferation and cell-cycle redistribution</td>
<td>37</td>
</tr>
<tr>
<td>miR-200</td>
<td>Carboplatin</td>
<td>Zeb</td>
<td>↓</td>
<td>Reverse EMT</td>
<td>99</td>
</tr>
<tr>
<td>miR-218</td>
<td>MDR</td>
<td>Survivin</td>
<td>↓</td>
<td>Enhance apoptosis</td>
<td>38</td>
</tr>
<tr>
<td>miR-205</td>
<td>Doc, Dox</td>
<td>VEGFA, FGF2</td>
<td>↓</td>
<td>Inhibit PI3K/mTOR pathway and CSCs</td>
<td>37</td>
</tr>
<tr>
<td></td>
<td>plus CTX</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

Note: *Upregulation (↑) or downregulation (↓) of miRNAs in drug-resistant cell lines/tissues.

Abbreviations: EMT, epithelial–mesenchymal transition; CSC, cancer stem cell.
Table 2 miRNAs involved in the regulation of endocrine resistance in breast cancer

<table>
<thead>
<tr>
<th>miRNA</th>
<th>Drug</th>
<th>Target(s)</th>
<th>Expression level</th>
<th>Mechanism</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>miR-451</td>
<td>Tamoxifen</td>
<td>14-3-3ζ, ERα</td>
<td>↓</td>
<td>Inhibit PI3K/Akt/mTOR pathway and enhance apoptosis</td>
<td>77</td>
</tr>
<tr>
<td>miR-451a</td>
<td>Tamoxifen</td>
<td>14-3-3ζ, ERα</td>
<td>↓</td>
<td>Inhibit autophagy</td>
<td>47</td>
</tr>
<tr>
<td>miR-320a</td>
<td>Tamoxifen</td>
<td>ARPP-19, ERα, c-Myc, Cyclin</td>
<td>↓</td>
<td>Activation of the c-Myc and Cyclin D1 pathways</td>
<td>46</td>
</tr>
<tr>
<td>miR-15a</td>
<td>Tamoxifen</td>
<td>Cyclin E1, Bcl-2</td>
<td>↓</td>
<td>Promote cell growth arrest and apoptosis</td>
<td>44, 91</td>
</tr>
<tr>
<td>miR-320a</td>
<td>Tamoxifen</td>
<td>ARPP-19, eRRc, c-Myc, Cyclin</td>
<td>↓</td>
<td>Activation of the c-Myc and Cyclin D1 pathways</td>
<td>46</td>
</tr>
<tr>
<td>miR-15a</td>
<td>Tamoxifen</td>
<td>Cyclin E1, Bcl-2</td>
<td>↓</td>
<td>Promote cell growth arrest and apoptosis</td>
<td>44, 91</td>
</tr>
<tr>
<td>miR-378a-3p</td>
<td>Tamoxifen</td>
<td>GOLT1A</td>
<td>↓</td>
<td>Uncharacterized</td>
<td>51</td>
</tr>
<tr>
<td>miR-342</td>
<td>Tamoxifen</td>
<td>Uncharacterized</td>
<td>↓</td>
<td>Promote apoptosis and cell cycle progression</td>
<td>49</td>
</tr>
<tr>
<td>miR-574-3p</td>
<td>Tamoxifen</td>
<td>CLTC</td>
<td>↓</td>
<td>Not investigated</td>
<td>52</td>
</tr>
<tr>
<td>miR-873</td>
<td>Tamoxifen</td>
<td>CDK3</td>
<td>↓</td>
<td>Inhibit ERα activity; restore anti-proliferative effect</td>
<td>43</td>
</tr>
<tr>
<td>miR-375</td>
<td>Tamoxifen</td>
<td>MTDH</td>
<td>↓</td>
<td>Reverse EMT</td>
<td>50</td>
</tr>
<tr>
<td>miR-214</td>
<td>Tamoxifen</td>
<td>UCP2</td>
<td>↓</td>
<td>Inhibit inhibiting PI3K/Akt/mTOR pathway and enhance apoptosis</td>
<td>48</td>
</tr>
<tr>
<td>miR-519a</td>
<td>Tamoxifen</td>
<td>PTEN, RB1, CDKN1A</td>
<td>↑</td>
<td>Promote proliferation, inhibit apoptosis</td>
<td>17</td>
</tr>
<tr>
<td>miR-101</td>
<td>Tamoxifen</td>
<td>MAGI-2</td>
<td>↑</td>
<td>Promote PI3K pathway</td>
<td>53</td>
</tr>
<tr>
<td>miR-301</td>
<td>Tamoxifen</td>
<td>FOXF3, BBC3, PTEN, COL2A</td>
<td>↑</td>
<td>Uncharacterized</td>
<td>56</td>
</tr>
<tr>
<td>miR-221/222</td>
<td>Fulvestrant</td>
<td>β-Catenin, TGF-β</td>
<td>↑</td>
<td>Deregulate multiple signaling pathways including β-catenin and TGF-β</td>
<td>78</td>
</tr>
<tr>
<td>miR-221/222</td>
<td>4-hydroxytamoxifen</td>
<td>P27kip1</td>
<td>↑</td>
<td>Promote cell cycle arrest</td>
<td>42</td>
</tr>
</tbody>
</table>

*Note: *↑upregulation (↑) or downregulation (↓) of miRNAs in drug-resistant cell lines/tissues.

Table 3 miRNAs involved in regulation of target therapy resistance in breast cancer

<table>
<thead>
<tr>
<th>miRNA</th>
<th>Drug</th>
<th>Target(s)</th>
<th>Expression level</th>
<th>Mechanism</th>
<th>Reference</th>
</tr>
</thead>
<tbody>
<tr>
<td>miR-21</td>
<td>Trastuzumab</td>
<td>PTEN, PDCD4</td>
<td>↑</td>
<td>Promote PI3K/Akt pathway</td>
<td>71–73</td>
</tr>
<tr>
<td>miR-211</td>
<td>Trastuzumab</td>
<td>PTEN</td>
<td>↑</td>
<td>Promote PI3K/Akt pathway</td>
<td>75</td>
</tr>
<tr>
<td>miR-210</td>
<td>Trastuzumab</td>
<td>Uncharacterized</td>
<td>↑</td>
<td>Uncharacterized</td>
<td>76</td>
</tr>
<tr>
<td>miR-205-5p</td>
<td>Trastuzumab</td>
<td>P63</td>
<td>↑</td>
<td>Promote EGFR pathway</td>
<td>78</td>
</tr>
<tr>
<td>miR-200c</td>
<td>Trastuzumab</td>
<td>ZNF17 and ZEB1</td>
<td>↓</td>
<td>Inhibit TGF-β signaling</td>
<td>79</td>
</tr>
<tr>
<td>miR-542-3p</td>
<td>Trastuzumab</td>
<td>AKT</td>
<td>↓</td>
<td>Inhibit PI3K pathway</td>
<td>80</td>
</tr>
<tr>
<td>miR-375</td>
<td>Trastuzumab</td>
<td>IGF1R, AKT</td>
<td>↓</td>
<td>Inhibit PI3K/Akt pathway</td>
<td>81</td>
</tr>
<tr>
<td>miR-630</td>
<td>Lapatinib</td>
<td>IGF1R</td>
<td>↓</td>
<td>Uncharacterized</td>
<td>82</td>
</tr>
<tr>
<td>miR-16</td>
<td>Trastuzumab</td>
<td>Cyclin and FUBPI</td>
<td>↓</td>
<td>Enhance chromatin accessibility</td>
<td>83</td>
</tr>
<tr>
<td>miR-7</td>
<td>Trastuzumab</td>
<td>EGFR</td>
<td>↓</td>
<td>Suppress EGFR expression</td>
<td>84</td>
</tr>
<tr>
<td>miR-515</td>
<td>Trastuzumab</td>
<td>CD44</td>
<td>↓</td>
<td>Inhibit CSC marker</td>
<td>103</td>
</tr>
</tbody>
</table>

*Note: *↑upregulation (↑) or downregulation (↓) of miRNAs in drug-resistant cell lines/tissues.

Abbreviations: EMT, epithelial-mesenchymal transition; CSC, cancer stem cell; TGF-β, transforming growth factor-β.

more effective therapeutic targets for the improvement of BCa treatment.

miRNAs involved in BCa drug resistance

The function of miRNA influences a variety of physiological and pathological processes by regulation of the target genes expression, which is generally implicated in tumor progression and resistance to treatment. A number of miRNAs were identified to be involved in BCa drug resistance by screening the established resistant cell or tissue models using either high-throughput sequencing or miRNA microarray. Subsequently, functional validation was conducted to assess the role of miRNA in drug resistance by ascertaining the expression of target genes and the alteration of related pathways. Finally, the selected candidate miRNAs were evaluated by using animal experimental models and/or clinical samples, which provided promising approaches to discover the
potential drug targets for improving BCa therapeutics and prognosis. Clinically used drugs for treating BCa mainly involve in chemotherapy, hormone therapy, and targeted therapy. In this study, we summarize drug resistance and relevant miRNA-mediated mechanisms in BCa.

Chemotherapy resistance

BCa patients with high risk of recurrence are generally recommended for adjuvant chemotherapy after definitive surgery. Doxorubicin (adriamycin) was the first-line anticancer drug to treat BCa which blocks DNA replication. Correspondingly, miR-505, miR-128, and miR-145 function as tumor suppressors, which are downregulated in doxorubicin-resistant BCa, and upregulation of these miRNAs led to reversal of doxorubicin resistance. In contrast, miR-663, miR-181a, and miR-106b–25 clusters serve as oncogenic miRNAs; downregulation of them resulted in enhancement of the sensitivity of doxorubicin-resistant cells.

Similar to doxorubicin, cisplatin is used to treat multiple types of cancer by inhibiting DNA replication. A microarray screening for miRNA expression signatures in cisplatin-resistant BCa cells versus sensitive BCa cells has shown upregulation of 46 miRNAs but downregulation of 57 miRNAs. In particular, miR-146a, miR-10a, miR-221/222, miR-345, miR-200b, and miR-200c were significantly deregulated in the resistant cells. Functional validation experiments indicated that miR-345 and miR-7 endow the cisplatin-resistant phenotype, while miR-302b can sensitize BCa cells to cisplatin.

In addition, docetaxel and paclitaxel are often used to treat cancers by interfering the function of microtubules during cell division. It has been reported that miR-34a, miR-100, and miR-30c were downregulated in paclitaxel-resistant BCa cells. By contrast, miR-129-3p was upregulated in docetaxel-resistant BCa cells and contributed to docetaxel resistance. However, two studies present controversial results regarding the role of miR-125b in the regulation of paclitaxel resistance in BCa. Zhou et al primarily demonstrated that miR-125b was increased in paclitaxel-resistant BCa cells as an oncogenic miRNA. While Yang et al reported that miR-125b presumably serves as a tumor suppressor miRNA (Table 1).

Moreover, it is recognized that miR-326 and miR-484 can reverse MDR phenotype; nevertheless, miR-218 and miR-205 contribute to the acquisition of MDR in BCa cells.

Hormone therapy resistance

Since ~70% of BCa patients exhibit high levels of estrogen receptor-α (ER-α), targeting ER-α seems to be the most valuable endocrine-based therapy. Several modulators of ER were selected to treat ER-positive BCa patients. In particular, tamoxifen, a well-documented hormone therapeutic drug, is able to bind to ER and block the interaction between estrogen and ER. Unfortunately, resistance and disease relapse have limited its long-term therapeutic efficacy. Manavalan et al initially screened miRNA expression profiles and identified multiple miRNAs deregulated in tamoxifen-resistant (TamR) cells versus the sensitive control cells.
The downregulation of miRNAs in TamR BCa cells was further confirmed to consist of miR-15a, miR-16, miR-320, miR-451, miR-214, miR-342, miR-873, miRNA-375, miR-378a-3p, and miR-574-3p. Subsequently, mechanistic studies have ascertained that these miRNAs play the roles of tumor suppressors due to the inhibition of multiple cell survival signaling pathways.43,45–52 Conversely, miR-101, miR-221/222, miR-301, and miRNAs-C19MC clusters were found to highly express in TamR BCa cells.17,53–56 (Table 2).

**Targeted therapy resistance**

Overexpression of human epidermal growth factor receptor 2 (HER2) is highly correlated to BCa metastasis and poor prognosis.57 HER2 targeted therapy has successfully achieved excellent results for treating HER2-positive BCa.58 Trastuzumab, a humanized monoclonal antibody targeting HER2, lapatinib, a small-molecule tyrosine kinase inhibitor dually targeting HER2 and epithelial growth factor receptor (EGFR), have largely improved therapy outcome and prognosis.59–65 However, despite their initial efficacy achieved for the early patient response, most of the patients eventually developed resistance after 1 year.66–70 Several mechanistic studies have revealed that deregulation of miRNA is involved in BCa trastuzumab resistance. Correlation of upregulation of miR-21 with trastuzumab resistance was validated in vitro and in vivo.71–73 However, one study asserted that the results from clinical research did not display the association between miR-21 expression and recurrent disease after adjuvant trastuzumab treatment.74

In addition to miR-21, miR-221 has been reported to promote trastuzumab resistance and metastasis of HER2-positive BCa.75 Also, circulating miR-210 levels in plasma from HER2-positive and trastuzumab-resistant BCa patients were determined to be correlated to BCa growth and metastases, suggesting that plasma miR-210 may serve as a predictive biomarker in surveillance of the therapeutic responsiveness.76 Furthermore, miR-205 was identified as a tumor suppressor, whose expression is downregulated in BCa cells, but is upregulated in patient-derived cancer stem cells (CSCs).77,78 miR-200c seemed to restore trastuzumab sensitivity and suppress invasion of BCa cells.79 Downregulation of miR-542-3p and miR-375 was also reported to contribute to trastuzumab resistance.80,81 In addition, miR-630 was found to be downregulated in lapatinib-resistant cell lines,82 while miR-16 expression was upregulated by trastuzumab and lapatinib.83 In the case of HER2Δ16, an oncogenic isoform of HER2 has been reported to prompt metastasis and MDR including trastuzumab and tamoxifen.84 The results from miRNA array have indicated that 16 miRNAs were abnormally expressed in HER2Δ16-expressed cells, particularly miR-7 appeared to have a 4.8-fold reduction.84

**Mechanisms involved in miRNA-mediated therapeutic resistance of BCa**

The function of miRNA in drug resistance has been ascribed to the alteration of drug transporters leading to efflux of anticancer agents,113,114,115 modification of autophagy/apoptosis to enhance survival,24,28,47,48,55,86 promotion of growth factors to disturb associated signal pathways57,71,77,87–88 and activation of epithelial–mesenchymal transition (EMT) process and augment of CSC populations to promote metastasis25,32,50,79,80 (Figure 1).

**Autophagy/apoptosis**

Autophagy is a lysosome-mediated intracellular self-catabolic degradation process for maintaining cellular homeostasis.91 Recent studies suggest that miRNA affects drug resistance by the regulation of autophagy that is considered to be a surviving mechanism.92,93 miR-451a is involved in BCa resistance to tamoxifen through significantly reduced autophagosomes.87 A novel compound isoliquiritigenin can reverse chemoresistance of BCa cells by negatively regulating miR-25, and overexpression of miR-25 was validated to stop isoliquiritigenin-induced autophagy and trigger chemo-sensitization by targetingULK1.6

Nevertheless, as the procedure of programmed cell death, apoptosis is considered to be involved in sensitization of cancers to treatment, which is regulated by pro-apoptotic regulators like caspases and/or antiapoptotic regulators including Bcl-2 family.94,95 As a tumor suppressor, miR-505 induces cell apoptosis via targeting Akt3 that participates in metastasis and drug resistance.19–22 miR-100 sensitizes BCa cells to paclitaxel by inducing cell apoptosis mainly by targeting mTOR.35 Bak1 was validated to be a direct target of miR-125b, which enhanced docetaxel-induced apoptosis.34 In addition, NF-κB-dependent STAT3 activation contributes to the upregulation of miR-181a to suppress BAX, resulting in decreased apoptosis and increased invasion of triple-negative breast cancer (TNBC).95 Inhibition of miR-181a significantly enhanced the sensitivity of TNBC to doxorubicin.96 Uregulation of miR-34a converted docetaxel resistance of BCa cells by downregulation of Bcl-2.97 In addition to miR-34a, miR-15a, miR-16, and miR-214 were also found to target Bcl-2 family in TamR cells.45,48,49 Moreover, miR-218 was found to contribute to the development of MDR in BCa cells by targeting survivin.38
Cell cycle arrest

It has been shown that there is a balance between cell cycle arrest and cell death in cancer therapy. CDKs act as dominators in mammalian for the regulation of cell cycle through phosphorylation of their selected cyclins by maintaining exit or entry to the different phases of the cell cycle. Several miRNAs participate in cell cycle regulation by targeting CDKs, cyclins, and E2F. Overexpression of miR-302b can sensitize BCa cells to cisplatin by targeting E2F1, a major regulator of the G1/S transition. In TamR BCa cells, miR-221/222 has been shown to target cell cycle inhibitor p27Kip1 that contributes to antagonizing cell death and promoting hormone-independent cell growth. Restoration of miR-320a expression is able to sensitize TamR cells to tamoxifen through the downregulation of c-Myc and Cyclin D1. In addition, the forced expression of miR-16 reduced BCa resistance to trastuzumab and lapatinib, and the underlying mechanism involves inhibition of c-Myc-enhanced chromatin assessibility.

CSCs and EMT

The intractable tumor relapses suggest that not all cells have been targeted, for example, CSCs may be a typical “escaping cells” that are capable of regenerating the tumor and do not respond to therapy drugs. CSCs are thought to be major contributors for drug resistance by activating HIF1α under hypoxic condition. It has been reported that miR-24 increases the expression of CSC markers, conferring cisplatin...
resistance. Although miR-205 was identified as a tumor suppressor, it has appeared to be upregulated in patient-derived CSCs. Similarly, miR-205 suppresses ErbB3 expression in BCa cells, whereas it inhibits EFGFR by targeting p63 in BCa CSCs. In addition, CD44 is a putative marker of CSCs, highly expressed in trastuzumab-resistant HER2-positive BCa. Knockdown of CD44 can efficiently restore the trastuzumab sensitivity of the resistant cells. Several miRNAs have been shown to regulate CD44 expression in CSCs, particularly miR-515 is essential for the regulation of CD44 in response to trastuzumab resistance.

Increasing evidences suggest that the function of miRNA is particularly involved in the EMT process that plays a crucial role in MDR by the promotion of tumor metastasis. E-cadherin transcriptional activator EP300 is suppressed by miR-106b–25 cluster, leading to acquired doxorubicin resistance by activating EMT. In addition, miR-24 confers cisplatin resistance by enhancing EMT process by targeting BimL and F1H1. In contrast, miR-644a suppresses a transcription repressor CTBP1, resulting in enhanced p53-mediated inhibition of EMT. Furthermore, miR-30c increases chemoresistance by targeting EMT-related cytokines in BCa, such as vimentin, stress fiber F-actin, twinfilin 1, Interleukin-6, and interleukin-1. Loss of miRNA-375 in TamR cells resulted in the activation of EMT, and ectopic expression of miRNA-375 in TamR cells was able to restore tamoxifen sensitivity by reversing EMT. Intriguingly, miR-125b serves as a tumor suppressor miRNA through the inhibition of EMT by targeting Sema4C, regardless of its effect on apoptosis.

It is worth noting that cells undergoing EMT often obtain properties associated with CSCs. Thus, EMT can cause drug resistance by inducing CSCs. TNBC accounts for ~15%–20% of all BCa cases, which is often documented as clinical chemoresistance due to the lack of molecular targets, particularly claudin-low tumors with extremely rich CSCs and EMT markers. As a central regulator of EMT, miR-200 has been shown to inhibit self-renewal capabilities of CSCs and keep cells in an epithelial state by suppressing the function of Zeb transcription factors. Induction of miR-200 in claudin-low tumors resulted in the suppression of tumor growth by inhibiting the functions of CSCs and reversal of EMT.

**Growth factor-related pathways**

Numerous studies have revealed multiple signaling pathways involved in therapeutic resistance of BCa and suggested many important molecular targets for the development of new drugs to treat aggressive BCa that are resistant to transitional chemotherapy and radiotherapy. In particular, growth factors, cytokines including chemokines, serve as major contributors for promoting BCa metastasis and leading to therapeutic resistance.

**PI3K/Akt/mTOR**

PI3K/Akt/mTOR-mediated amplification of growth factors is well recognized to play a pivotal role in BCa metastasis and drug resistance. Multiple miRNAs were identified to participate in BCa MDR through the regulation of PI3K/Akt/mTOR signaling pathway. For example, miR-205 and miR-214 enhance chemoresistance and TamR by inhibiting PI3K/Akt/mTOR pathway. However, phosphorylation of Akt and mTOR are reduced by miR-451, resulting in restored sensitivity to tamoxifen. Downregulation of miR-542-3p contributes to trastuzumab resistance in BCa partially through inhibiting PI3K. In addition, PTEN, an inhibitor of PI3K, is a key regulator for PI3K/Akt/mTOR pathway. Reduction or loss of PTEN has been involved in BCa progression and drug resistance. PTEN was verified to be a target gene of miR-21 and miR-221. In addition, miR-101 and miRNA-519a confer TamR and chemoresistance by targeting PTEN. Reducing miR-301 expression was validated to increase sensitivity to tamoxifen.

**NF-κB**

NF-κB, stress responsible transcription factor, plays a crucial role in cancer progression and therapeutic resistance. In IKK family identified as standard upstream signaling to activate NF-κB, PI3K and STAT3 can also trigger NF-κB activation. Through NF-κB transcriptional activation, miR-21 is uniquely highly expressed in HER2-positive BCa and is responsible for the resistance to trastuzumab. In addition to PTEN, PDCD4 was found as another direct target of miR-21. In contrast, miR-31 represses NF-κB activation and increases apoptosis in TNBC cells by targeting protein kinase C epsilon.

**EGFR**

Overexpression of EGFR family containing ErbB1, HER1, HER2, HER3, and HER4 imperatively increases drug resistance of BCa. Notably, the activation of EGFR finally triggers PI3K/Akt/mTOR and STAT3 signaling pathways, resulting in increased cell survival by activating antiapoptotic proteins. miR-451 was reported to specifically target 14-3-3ζ, leading to enhanced apoptosis and sensitized BCa cells to endocrine drugs via inactivation of EGFR. miR-7 was identified to suppress EGFR expression and sensitize...
HER2Δ16 to the targeted therapy. In addition, miR-205 was also found as a tumor suppressor, sensitizing targeted therapy by suppressing HER3 and Zeb. However, it is upregulated in BCa CSCs and increases the therapeutic resistance through the regulation of P63-mediated EGFR activation.78,102

Cytokines
Cytokines including chemokines are important inflammatory factors and highly related to BCa progression and drug resistance. Several miRNAs have been involved in cytokine-mediated drug resistance of BCa due to the activation of EMT. Overexpression of miR-221/222 contributes to drug resistance through deregulation of multiple signaling pathways including β-catenin and transforming growth factor-β (TGF-β). In addition, miR-200c suppresses TGF-β signaling and counteracts trastuzumab resistance of BCa by targeting ZNF217 and ZEB1. Both miR-630 and miR-375 have been found to enhance patient responses to HER-targeting drugs by targeting IGF1R. Furthermore, miR-30c inhibits chemoresistance of BCa by targeting TWF1, an activator of EMT, leading to suppression of IL-6 and IL-11 production.32

ATP binding cassette (ABC) transporter
Overexpression of ABC transporters in cancer cells is considered to be a common mechanism of MDR due to efflux of drug substrates to maintain their concentrations below effective levels. Human ABC transporters are a large group of membrane protein complexes, including MDR1 (P-gp) and MRP1 (ABCC1). ABCC5, another member in ABC family, was targeted by miR-128, leading to sensitization of BCa to doxorubicin. In addition, miR-145 was able to inverse doxorubicin resistance in BCa by suppressing MRP1 and triggering drug efflux transporter. Targeting MRP1 by miR-326 was also validated to reverse MDR phenotype.36

Commentary
However, it is intriguing that there are some controversial results reported in response to the regulation of different sets of target genes. For example, downregulation of miR-181a was previously reported to inhibit BCa cell apoptosis induced by doxorubicin through the upregulation of Bcl-2, while miR-181a was upregulated in TNBC cells in response to chemotherapy, which was activated by STAT3 to target BAX. Similarly, miR-125b was upregulated in paclitaxel-resistant BCa cells and target Bak1. Nevertheless, miR-125b was also shown to be downregulated in paclitaxel-resistant MCF-7 cells, leading to reverse EMT by targeting Sema 4C. Despite one miRNA targets different genes which may exert the same or opposite effects, the roles of miRNAs in drug resistance of BCa finally depend on their targets for achieved dominant effects in particular cellular and tissue types corresponding to the particular treatment. In addition, it also needs to be noted that multiple miRNAs may interact with one another or may interplay on gene expression through either cooperative effects or antagonist competition, which forms a complicated network to regulate all the target genes in cancers. However, to date, most of studies cited in this review described only the role of miRNA in drug resistance dependent upon the effect of single miRNA, but the concert action of multiple miRNAs contributes to the resistance remains obscure.

Function of exosome in BCa chemoresistance
Exosome is widely recognized to be an important transfer vehicle for cell–cell communication within the tumor microenvironment. Emerging evidences have demonstrated that exosome-mediated miRNA transfer between host–tumor cells plays a pivotal role in tumor progression and resistance to treatment. To confirm the effect of exosome-derived miRNA transferring on drug resistance transmitting, we further enlarged the significance of the miRNA delivered by exosomes. The results suggested that the miRNA regulated drug resistance due to exosomes involving more infused microenvironment within transmitting resistant capacity through miRNA-mediated alteration of gene expression. Nevertheless, in addition to doxorubicin, Wei et al have demonstrated distinct differences in the exosome concentration and size distribution between TamR cells and the sensitive cells. TamR cells seem to have an increased amount of exosomal RNA and proteins. Exosome secreted from TamR cells was able to enter into the sensitive cells and release miR-221/222, leading to TamR by targeting P27 and Era.6 Nevertheless,
TNBC cells secrete miRNA-134-enriched exosomes and contribute to increased sensitivity to anti-Hsp90 drugs through the reduction of STAT5B and Hsp90, suggesting that TNBC-derived exosomes may have therapeutic potential as miRNAs delivery vesicles.\textsuperscript{137}

In addition to BCa, exosomal miRNAs involving drug resistance have been studied in other types of cancer. Exosomal miR-155 contributes to chemoresistance of neuroblastoma by targeting TERF1.\textsuperscript{138} Increase of exosomal miR-155 was also found to enhance gemcitabine resistance in pancreatic ductal adenocarcinoma cells through the enhancement of anti-apoptotic activity.\textsuperscript{139} Exosomal miRNAs from lung cancer A549 cells were increased after exposure to cisplatin, which can endow the resistant capacity of the untreated cells.\textsuperscript{140} Exosomal miR-222-3p has been shown to enhance gemcitabine resistance by targeting SOCS3 in A549 cells.\textsuperscript{141} While exosomal miR-100-5p was decreased in cisplatin-resistant A549, leading to increase in cisplatin resistance through reverse regulation of mTOR.\textsuperscript{142} Exosomal miRNAs including miR-598, miR-34a, miR-146a, and miR-148a were significantly increased in docetaxel-resistant prostate cancer cells.\textsuperscript{143} Exosomal transfer of stromal-derived miR-21 can transmit paclitaxel resistance into ovarian cancer cells by targeting APAF1.\textsuperscript{13} In addition, exosomes derived from tumor-associated macrophages can mediate cisplatin resistance by directly transferring miR-21 to target PTEN in gastric cancer cells.\textsuperscript{144} Interestingly, Lou et al successfully sensitized hepatocellular carcinoma to targeting TERF1.

Exosome in the tumor microenvironment
Tumor microenvironment is often likened to the soil where tumor cells grow, which refers to the variety of normal cells, signaling molecules, blood vessels, and extracellular matrix surrounding the tumor cells.\textsuperscript{151} As a key component in the tumor microenvironment, exosomes shuttle bioactive molecules and mediate cellular communication in the tumor microenvironment, facilitating a more malignant and refractory tumor environment. The major function of exosome in the tumor microenvironment in response to drug resistance is considered to be a molecular shuttle. As described above, exosomes can enrich and transfer a variety of small molecules including miRNAs between donor cells and recipient cells. Recent studies suggest two main categories, namely from resistant cells to sensitive cells for transferring the resistant capacity as well as from stromal cells to their surrounded cancer cells for survival and resistance.\textsuperscript{13,132,136,140} The relevant stromal cells have been defined as cancer-associated fibroblasts, mesenchymal stem/stroma, and adipocytes.

Beyond its encapsulated abundant functional molecules, exosomes also participate in the regulation of drug resistance by acting as an efflux pump. The drugs and their metabolites can be encapsulated and exported by exosomes.\textsuperscript{152-154} Therefore, it is very promising to apply exosomes for targeted therapeutics for their unique advantages, because they have a good tolerance for their wide distribution in various human body fluids. Overall, exosomes can serve as an optimal carrier of drugs and functional molecules including diagnostic biomarkers and therapeutic molecules, which may provide a promising approach for the improvement of anticancer therapy.

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References


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