



# Metabolic Syndrome and Breast Cancer: Prevalence, Treatment Response, and Prognosis

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Metabolic syndrome is a type of multifactorial metabolic disease with the presence of at least three factors: obesity, diabetes mellitus, low high-density lipoprotein, hypertriglyceridemia, and hypertension. Recent studies have shown that metabolic syndrome and its related components exert a significant impact on the initiation, progression, treatment response, and prognosis of breast cancer. Metabolic abnormalities not only increase the disease risk and aggravate tumor progression but also lead to unfavorable treatment responses and more treatment side effects. Moreover, biochemical reactions caused by the imbalance of these metabolic components affect both the host general state and organ-specific tumor microenvironment, resulting in increased rates of recurrence and mortality. Therefore, this review discusses the recent advances in the association of metabolic syndrome and breast cancer, providing potential novel therapeutic targets and intervention strategies to improve breast cancer outcome.

**Keywords:** breast cancer, metabolic syndrome, obesity, incidence, treatment response, prognosis

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## INTRODUCTION

Breast cancer is a malignant tumor with the highest incidence in women of all ages in the world and is associated not only with hormones or factors related to reproduction but also with environmental factors in general (1). Epidemiological studies have shown that early menarche, postmenopausal weight gain, a high-fat diet, and long-term use of exogenous estrogen are associated with a high risk of breast cancer (2). Recent studies have also shown that a specific lifestyle characterized by reduced physical activity and fat-rich dietary habits, refined carbohydrates and animal protein, which consequently causes metabolic syndrome (MetS), plays a crucial role in breast cancer initiation (1, 3). Metabolic syndrome, also known as insulin resistance syndrome or syndrome X, is a type of multifactorial metabolic disease. The definition of MetS takes into account the presence of at least three factors, namely, abdominal obesity/high body mass index (BMI), insulin resistance, hypertension, hypertriglyceridemia, and low high-density lipoprotein (HDL) (3–5). The prevalence of obesity has increased rapidly in recent years with the number of overweight/obese people almost doubling since the 1980s, representing one-third of the world's population, and the proportion may reach 57.8 percent by 2030 (6, 7). In Western countries, the prevalence of MetS in the adult population is between 20% and 25% (8). Notably, the incidence rate increases significantly with age, resulting in the prevalence of people aged over 50 reaching 40–45% (8). Moreover, changes in the balance between insulinotropic and anti-inflammatory cytokines driven by abdominal obesity

may lead to insulin resistance, which is a core component of MetS. Asian women are particularly vulnerable to these diseases because they have greater abdominal and visceral fat than white women with similar BMI (9). Previous studies have confirmed that elderly and postmenopausal women are more susceptible to MetS (3). Several cohort studies and meta-analyses have also highlighted the link between MetS and its components and the prevalence, recurrence, and mortality of various cancers, including breast cancer (4, 10, 11).

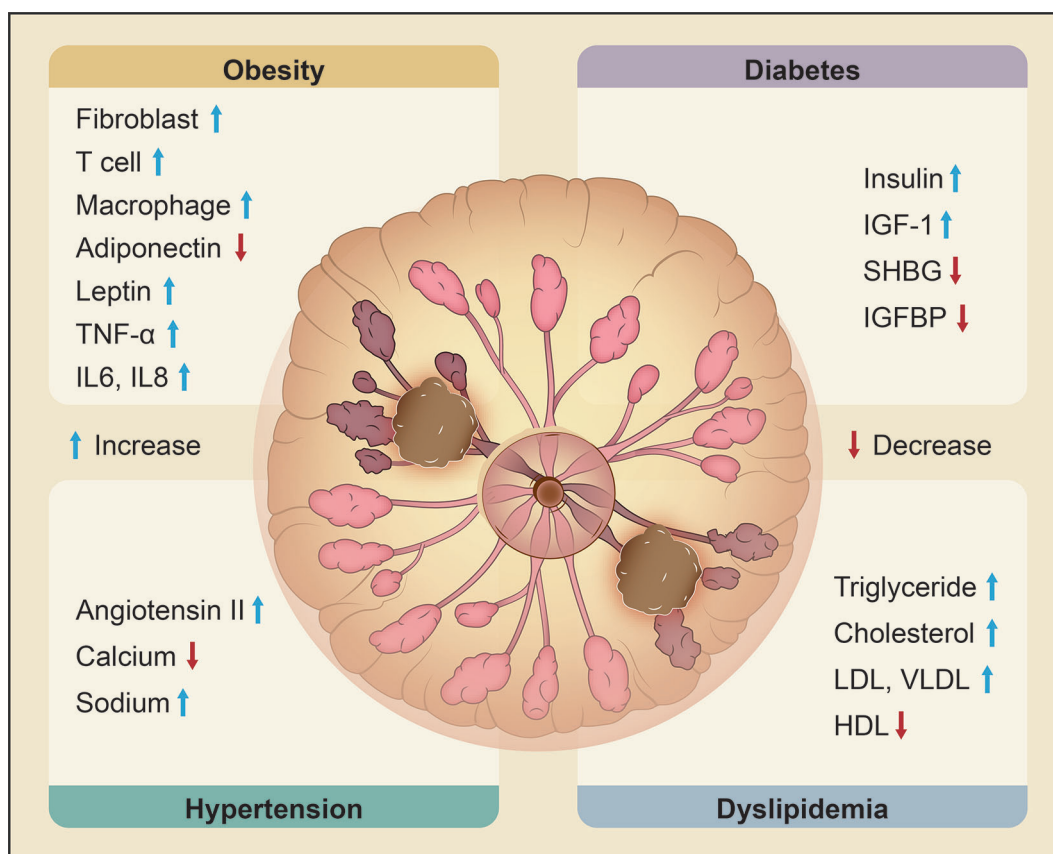
MetS has become a significant public health problem worldwide, and in-depth research on MetS and breast cancer is increasing. However, there is a paucity of systematic reviews focusing on breast cancer and metabolic syndrome and a comprehensive understanding of this topic. Thus, in this review, we will systematically discuss the latest research advances on MetS and their associations with the incidence, treatment response, prognosis and progression mechanism of breast cancer, which will help provide new therapeutic targets and strategies to improve the prognosis of breast cancer patients.

## METABOLIC SYNDROME AND BREAST CANCER RISK

A population-based study by Russo et al. (12) found that MetS and its components are associated with an increased risk of breast cancer. This connection has been proven in many other studies (13, 14), and it is more pronounced in postmenopausal women regardless of race (odds ratio [OR] = 1.75, 95% confidence interval [CI] 1.37-2.22) (15). With the increase in the number of MetS components, the risk of breast cancer increases for postmenopausal women ( $P = 0.01$ ) (16). Factors related to the risk of breast cancer are discussed in the following four sections (**Figure 1**). **Table 1** summarizes reports about the correlation between metabolic syndrome and its components and the risk of breast cancer with different subtypes.

### Obesity and Breast Cancer Risk

Obesity is associated with the risk of postmenopausal breast cancer (26, 27) as well as greater tumor burden and higher



**FIGURE 1** | Metabolic syndrome and its molecular changes related to the risk of breast cancer. Obesity, diabetes, hypertension and dyslipidemia are the main components of metabolic syndrome and are all significantly related to the risk of breast cancer. Obesity increases fibroblasts, T cells, macrophages, leptin, TNF- $\alpha$ , IL-6 and IL-8 and decreases adiponectin. Diabetes is characterized by upregulation of insulin and IGF-1 and downregulation of SHBG and IGFBP. Hypertension is associated with increased ANG II and sodium and decreased calcium. Dyslipidemia leads to high levels of TG, TC, LDL, and VLDL and low levels of HDL. Changes in the expression of these key molecules are correlated with an elevated risk of breast cancer. ('the blue arrow' means increase and 'the red arrow' means decrease).

**TABLE 1 |** Obesity and its measurement indexes with the risk of different breast cancer subtypes.

| No.                        | Author               | Patients(N) | Region    | Menopausal status  | Comparison                          | Molecular subtype          | Measurement     | 95%CI     | P         | Ref  |
|----------------------------|----------------------|-------------|-----------|--------------------|-------------------------------------|----------------------------|-----------------|-----------|-----------|------|
| 1                          | Agresti R. et al.    | 1779        | Italy     | Pre-menopausal     | BMI $\geq$ 25 vs. BMI<25            | LuminalB(HER2-)            | OR=1.30         | 0.79-2.18 | N/A       | (13) |
|                            |                      |             |           |                    |                                     | LuminalB(HER2+)            | OR=1.78         | 0.88-3.63 | N/A       |      |
|                            |                      |             |           |                    |                                     | HER2+(non-luminal)         | OR=1.89         | 0.78-4.60 | N/A       |      |
|                            |                      |             |           |                    |                                     | TNBC                       | OR=3.04         | 1.43-6.43 | N/A       |      |
|                            |                      |             |           |                    |                                     | WC $\geq$ 80cm vs. WC<80cm | LuminalB(HER2-) | OR=2.55   | 1.53-4.24 |      |
|                            |                      |             |           | LuminalB(HER2+)    | OR=2.11                             | 1.03-4.35                  | N/A             |           |           |      |
|                            |                      |             |           | HER2+(non-luminal) | OR=1.28                             | 0.50-3.27                  | N/A             |           |           |      |
|                            |                      |             |           | TNBC               | OR=1.03                             | 0.42-2.53                  | N/A             |           |           |      |
|                            |                      |             |           | Post-menopausal    | BMI $\geq$ 25 vs. BMI<25            | LuminalB(HER2-)            | OR=1.51         | 1.14-2.00 | N/A       |      |
|                            |                      |             |           |                    |                                     | LuminalB(HER2+)            | OR=1.20         | 0.76-1.92 | N/A       |      |
| HER2+(non-luminal)         | OR=1.43              | 0.79-2.57   | N/A       |                    |                                     |                            |                 |           |           |      |
| TNBC                       | OR=1.11              | 0.65-1.90   | N/A       |                    |                                     |                            |                 |           |           |      |
| WC $\geq$ 80cm vs. WC<80cm | LuminalB(HER2-)      | OR=1.17     | 0.75-1.81 |                    |                                     | N/A                        |                 |           |           |      |
| LuminalB(HER2+)            | OR=0.82              | 0.41-1.63   | N/A       |                    |                                     |                            |                 |           |           |      |
| HER2+(non-luminal)         | OR=1.36              | 0.50-3.69   | N/A       |                    |                                     |                            |                 |           |           |      |
| TNBC                       | OR=0.89              | 0.41-1.95   | N/A       |                    |                                     |                            |                 |           |           |      |
| 2                          | Chen H. et al.       | 4557        | USA       | Both               | Type 2 DM vs. non-DM                | ER+/HER2+                  | OR=0.77         | 0.40-1.48 | <0.05     | (17) |
|                            |                      |             |           |                    |                                     | TNBC                       | OR=1.38         | 1.01-1.89 | <0.05     |      |
| 3                          | Maskarinec G. et al. | 681         | USA       | Both               | Type 2 DM in subtypes               | H2E                        | OR=1.38         | 0.93-2.06 | <0.05     | (18) |
|                            |                      |             |           |                    |                                     | ER+/PR+                    | HR=1.17         | 1.05-1.29 | N/A       |      |
|                            |                      |             |           |                    |                                     | ER-/PR-                    | HR=1.03         | 0.83-1.26 | N/A       |      |
| 4                          | Michels K.B. et al.  | 116488      | USA       | Both               | Type 2 DM vs. non-DM                | ER+/PR- or ER-/PR+         | HR=1.01         | 0.81-1.28 | N/A       | (19) |
|                            |                      |             |           |                    |                                     | ER+                        | HR=1.22         | 1.01-1.47 | N/A       |      |
| 5                          | Millikan R.C. et al. | 3446        | USA       | Pre-menopausal     | WHR $\geq$ 0.84 vs. WHR<0.84        | ER-                        | HR=1.13         | 0.79-1.62 | N/A       | (20) |
|                            |                      |             |           |                    |                                     | LuminalB                   | OR=0.90         | 0.40-1.80 | N/A       |      |
| 6                          | Munsell M.F. et al.  | /           | /         | Pre-menopausal     | BMI $\geq$ 30 vs. BMI<25            | HER2+(non-luminal)         | OR=0.90         | 0.40-2.20 | N/A       | (21) |
|                            |                      |             |           |                    |                                     | TNBC                       | OR=1.90         | 1.00-3.60 | N/A       |      |
|                            |                      |             |           | Post-menopausal    | BMI $\geq$ 30 vs. BMI<25            | LuminalB                   | OR=0.50         | 0.20-0.90 | N/A       |      |
|                            |                      |             |           |                    |                                     | HER2+(non-luminal)         | OR=0.50         | 0.30-1.00 | N/A       |      |
|                            |                      |             |           |                    |                                     | TNBC                       | OR=1.40         | 0.70-2.70 | N/A       |      |
| 7                          | Palmer J.R. et al.   | 1851        | USA       | Both               | Type 2 DM vs. non-DM                | ER+/PR+                    | RR=0.78         | 0.67-0.92 | 0.67      | (22) |
|                            |                      |             |           |                    |                                     | ER-/PR-                    | RR=1.06         | 0.70-1.60 | 0.004     |      |
| 8                          | Pierobon M. et al.   | 3845        | USA       | Pre-menopausal     | BMI $\geq$ 30 vs. BMI<30            | ER+/PR+                    | RR=1.39         | 1.14-1.70 | 0.001     | (23) |
|                            |                      |             |           |                    |                                     | ER-/PR-                    | RR=0.98         | 0.78-1.22 | 0.02      |      |
| 9                          | Rosner B. et al.     | 77232       | USA       | Pre-menopausal     | Per 25 lbs weight gain              | ER+                        | HR=1.02         | 0.80-1.31 | N/A       | (24) |
|                            |                      |             |           |                    |                                     | ER-                        | HR=1.43         | 1.03-2.00 | N/A       |      |
| 10                         | Suzuki R. et al.     | 41594       | Japan     | Post-menopausal    | Per increment of 5kg/m <sup>2</sup> | TNBC                       | OR=1.43         | 1.23-1.65 | N/A       | (25) |
|                            |                      |             |           |                    |                                     | ER+/PR+                    | RR=1.13         | 0.89-1.43 | 0.32      |      |
|                            |                      |             |           |                    |                                     | ER+/PR-                    | RR=2.19         | 1.33-3.61 | 0.002     |      |
|                            |                      |             |           |                    |                                     | ER-/PR-                    | RR=1.61         | 1.09-2.38 | 0.016     |      |
|                            |                      |             |           |                    |                                     | ER+/PR+                    | RR=2.24         | 1.50-3.34 | N/A       |      |
| ER+/PR-                    | RR=0.63              | 0.31-1.27   | N/A       |                    |                                     |                            |                 |           |           |      |
| ER-/PR-                    | RR=0.67              | 0.38-1.17   | N/A       |                    |                                     |                            |                 |           |           |      |

BMI, Body mass index; ER, Estrogen receptor; HER2, Human epidermal growth factor 2; N/A, Not applicable; OR, Odds ratio; PR, Progesterone receptor; RR, Relative risk; TNBC, Triple negative breast cancer; WC, Waist circumference.

histopathological grade (28). BMI is routinely used to measure obesity but lacks information about actual body composition. Waist circumference (WC) and waist-to-hip ratio (WHR) are more commonly adopted for evaluating body fat distribution. Compared with patients with BMI < 25 kg/m<sup>2</sup>, patients with BMI  $\geq$  30 kg/m<sup>2</sup> have larger tumors, poorer differentiation, a higher frequency of lymph node invasion, and more advanced disease (29). According to The International Agency for Research on Cancer, ample evidence suggests that increasing weight gain is a risk factor for postmenopausal breast cancer (30). A small case-control study by Schapira et al. (31) found that breast cancer patients had 45% more visceral fat tissue than the control group. In addition, epidemiological studies have shown that WC/WHR increases are related to breast cancer (10, 32, 33).

However, some studies have also reported that obesity may have a protective effect on the risk of premenopausal breast cancer (8, 34), which requires further evidence.

Given that excessive estrogen stimulates breast tissue proliferation (35), the increase in estrogen levels in the body caused by obesity is considered to be one of the mechanisms associated with breast cancer because adipose tissue is the main source of estrogen in postmenopausal women (34). Chronic inflammation caused by the imbalance of fat homeostasis due to obesity also promotes the development of tumors. Aromatase is a kind of cytochrome P450 enzyme mainly located in the adipose tissue of breast, abdomen, thigh and buttocks, but it may also be present in tumor tissue. It can catalyze the formation of estrone and estradiol from androstenedione and testosterone (34).

Obese adipocytes upregulate the expression of pro-inflammatory factors such as TNF- $\alpha$  and IL-6 through the obesity-inflammation-aromatase axis (36), resulting in enhanced transcription of CYP19 gene encoding aromatase, thereby promoting the production of aromatase. Adipose tissue not only stores adipocytes, it is also an endocrine organ producing biologically active molecules called adipokines. These adipokines bind to specific receptors on the surface of target cells and affect the metabolism of tissues and organs (37). Among the adipokines, leptin increases the risk of disease (38), whereas adiponectin may have a protective effect. Some case-control studies have indicated that low adiponectin levels are associated with increased breast cancer risk and a more aggressive phenotype (39, 40).

For premenopausal women, obesity has a protective effect on hormone receptor-positive breast cancer but increases the risk of estrogen receptor (ER)+/progesterone receptor (PR)- and ER-/PR- breast cancer (24). A meta-analysis conducted by Munsell et al. (21) indicated a summary risk ratio (RR) of 0.78 (95% CI 0.67-0.92) for hormone receptor-positive breast cancer and 1.06 (95% CI 0.70-1.60) for hormone receptor-negative breast cancer in premenopausal women associated with obesity. A positive correlation is noted between obesity and triple-negative breast cancer (TNBC) (23). A multiple logistic regression analysis of 1,779 patients with primary invasive breast cancer in Italy showed that premenopausal women with WC  $\geq$  80 cm were prone to luminal B breast cancer, including HER2-negative (OR = 2.55, 95% CI 1.53-4.24) and HER2-positive women (OR = 2.11, 95% CI 1.03-4.35), whereas women with BMI  $\geq$  25 kg/m<sup>2</sup> (OR = 3.04, 95% CI 1.43-6.43) were significantly related to TNBC compared with other subtypes (13).

For postmenopausal women, obesity is strongly associated with the risk of ER+/PR+ breast cancer but has a weak association with PR- breast cancer (25). Munsell et al. (21) showed in their meta-analysis that obesity was significantly associated with hormone receptor-positive breast cancer in postmenopausal women (RR = 1.39, 95% CI 1.14-1.70), but the RR for hormone receptor-negative breast cancer in postmenopausal women was 0.98 (95% CI 0.78-1.22). A study on the association between BMI and breast cancer subtypes in postmenopausal women in the Mediterranean found that BMI > 25 kg/m<sup>2</sup> was positively correlated with the risk of luminal breast cancer but not TNBC (41). In a case-control study of a large population conducted by the Carolina Breast Cancer of the United States (20), women with a large WHR regardless of menopausal status (premenopausal OR = 1.9, 95% CI 1.0-3.6; postmenopausal OR = 1.4, 95% CI 0.7-2.7) were more prone to TNBC compared with other breast cancer subtypes.

Therefore, these results indicate that obesity plays a potential role in the prevalence of breast cancer. Obesity may be more likely to increase the risk of hormone receptor-negative breast cancer in premenopausal women and hormone receptor-positive breast cancer in postmenopausal women. In different phenotypes, the risk correlation also varies with measurements of obesity, among which WC and WHR were two important

values. Abdominal obesity is correlated with more aggressive molecular types regardless of menopausal status.

## Diabetes Mellitus and Breast Cancer Risk

Diabetes is one of the most frequent chronic diseases in the global population. Insulin resistance is the key factor in the pathogenesis of type 2 diabetes and the most typical and serious phenomenon (42). It is defined as decreased sensitivity to insulin-mediated glucose disposal and inhibition of hepatic glucose production (43) and presents as dysfunction of insulin transduction in glucose uptake and utilization in body skeletal muscles, adipocytes and hepatocytes (44), which leads to hyperglycemia, hyperinsulinemia and various disorders. Insulin resistance is the core part of MetS, and the corresponding increase in fasting blood glucose levels and the effects of hyperinsulinemia in breast cancer are also frequently studied. A meta-analysis (45) showed that compared with the nondiabetic group, the hazard ratio (HR) of the diabetes group was 1.23 (95% CI 1.12-1.34), which indicates that diabetes is a risk factor for breast cancer. In addition, hyperinsulinemia associated with insulin resistance is also correlated with a high risk of breast cancer. Zhu et al. (46) found that the OR for breast cancer associated with the highest quartile versus the lowest quartile of insulin was 1.45 (95% CI 1.20-1.75). Elevated insulin levels can cause high insulin growth factor-1 (IGF-1) bioavailability, leading to the occurrence and proliferation of breast cancer (10). The insulin receptor and IGF-1 receptor are widely expressed in breast cancer cells and promote cell proliferation mainly *via* the insulin receptor substrate (IRS)/phosphatidylinositol 3-kinase (PI3K) and Ras/mitogen-activated protein kinase (MAPK) pathways (8, 39).

Research data have yielded inconsistent reports on the relationship between diabetes mellitus (DM) and the risk of different types of breast cancer. Michels et al. (19) and Maskarinec et al. (18) found that type 2 DM was strongly associated with ER-positive breast cancer (HR for Michels et al. = 1.22, 95% CI 1.01-1.47; HR for Maskarinec et al. = 1.17, 95% CI 1.05-1.29). The main mechanism may be due to crosstalk between estrogen and the insulin/IGF-1 signaling pathway. The activation of ER $\alpha$  is influenced by the insulin/IGF-1 pathway, and estrogen may increase the expression and activity of certain proteins in the pathways that promote signal transduction (47). Palmer et al. (22) observed opposite results. Specifically, a medical history of diabetes was positively correlated with the risk of ER- breast cancer (HR = 1.43, 95% CI 1.03-2.00) rather than ER+ breast cancer. Clinical data indicate that the diabetic state may promote a more aggressive cancer subtype. A retrospective study (17), including 4,557 cases, showed that women diagnosed with type 2 DM have a higher risk of TNBC (OR = 1.38, 95% CI 1.01-1.89) or human epidermal growth factor receptor 2 (HER2)-overexpressing breast cancer (OR = 1.38, 95% CI 0.93-2.06) than patients without a history of diabetes. Several studies also supported that type 2 DM is associated with the risk of TNBC. Regarding antidiabetes treatment, metformin has been indicated to block breast cancer cell cycle progression and selectively induce apoptosis. Previous

studies showed that patients treated with metformin had a better breast cancer prognosis (48, 49). Liu et al. (49) reported that metformin has a tumor suppressive effect on breast cancer through a variety of molecular effects, especially on TNBC. They observed that compared with the controls, tumor growth ( $P = 0.0066$ ) and cell proliferation ( $P = 0.0021$ ) in tumor-bearing nude mice treated with metformin were significantly inhibited. Metformin-induced apoptosis, proteolysis of polyadenosine diphosphate-ribose polymerase (PARP) and reduction of epidermal growth factor receptor (EGFR) (a key receptor in TNBC cells) were not observed in phenotypes other than TNBC. Chen et al. (17) found that compared with nondiabetic people, the use of metformin was associated with an increased risk of TNBC (OR = 1.54, 95% CI 1.07-2.22). This may be due to the relatively small sample of women with DM enrolled in the study, and the limited collection of data on history of DM and drug use was only two years before the diagnosis of breast cancer, which limits the evaluation of the impact of long-term use of metformin on the risk of TNBC.

DM and its related hyperinsulinemia and insulin resistance are risk factors for breast cancer. However, the correlation between DM and breast cancer subtypes has inconsistent conclusions among studies and requires further exploration and research. The use of metformin may improve the prognosis of breast cancer patients with diabetes or a prediabetic state, but solid clinical evidence is needed.

## Dyslipidemia and Breast Cancer Risk

Dyslipidemia, including high total triglyceride (TG) levels, high total cholesterol (TC) levels and low serum HDL levels, is also considered to be associated with the occurrence of breast cancer, but the results are inconsistent. A prospective study launched by His et al. (50) showed that TC (HR 1 mmol/L increment = 0.83, 95% CI 0.69-0.99,  $P = 0.04$ ) and HDL (HR 1 mmol/L increment = 0.48, 95% CI 0.28-0.83,  $P = 0.009$ ) were inversely associated with the risk of breast cancer. Low serum HDL was independently correlated with an increased risk of breast cancer, especially after menopause (51, 52). However, Kitahara et al. (53) found that TC was positively correlated with breast cancer risk (HR = 1.17, 95% CI 1.03-1.33) in the Korean population. Katzek et al. (54) found that TGs (HR for highest vs. lowest quartile = 0.65, 95% CI 0.46-0.92) were negatively associated with breast cancer risk, but HDL (HR for highest vs. lowest quartile = 1.39, 95% CI 1.01-1.93) was positively associated with breast cancer risk, which differed from previous results. A significant increase in TGs and a decrease in HDL have been observed in TNBC patients (4). Therefore, more clinical data and reliable meta-analyses are needed to confirm the relationship between lipid metabolism and breast cancer risk. A case-control study on Chinese women (33) showed that among all MetS components, the hypertriglyceridemia waist circumference phenotype (HTWC), namely, elevated waist circumference and triglycerides, significantly increased the risk of breast cancer (OR = 1.56, 95% CI 1.02-2.39) regardless of menopausal status. Pelton et al. (55) found that mice fed a high-fat/high-cholesterol diet had

significantly higher percentages of tumor cell proliferation and higher microvessel density in preclinical models. Furberg et al. (56) suggested that low serum HDL in overweight and obese women was associated with higher levels of breast mitogens and estrogens; thus, HDL might represent a biological marker of breast cancer risk. In particular, Boyd et al. (57) observed a positive relationship between low HDL levels and atypical hyperplasia of the mammogram.

These results suggest that lipid disorders are a risk factor for breast cancer and may promote tumor cell proliferation and blood vessel formation. However, the relationship between different lipids and the risk of different subtypes of breast cancer still needs to be confirmed through more clinical evidence, so these lipids may be used as biologically reasonable markers to identify and intervene in high-risk individuals. Regarding the possible mechanism by which triglycerides, cholesterol and lipoprotein affect the occurrence and progression of breast cancer, further exploration is needed. Then, new lipid-related strategies can be launched for the prevention and treatment of breast cancer based on the findings.

## Hypertension and Breast Cancer Risk

Regarding hypertension and the risk of breast cancer, the results of studies are not consistent. A cohort study in Finland (58) found no correlation between hypertension and breast cancer (standardized incidence ratio = 0.94, 95% CI 0.84-1.04). Some studies also showed that hypertension was not associated with TNBC (4). A case-control study by Pereira et al. (59) found that women with hypertension (blood pressure  $\geq 140/90$  mmHg) had a higher risk of breast cancer (OR = 4.18; 95% CI 1.81-9.64), and a significant association was observed in postmenopausal women (OR = 2.84, 95% CI 1.09-7.39). A meta-analysis (60) also showed a significant association between hypertension and increased breast cancer risk (risk ratio [RR] = 1.15, 95% CI 1.08-1.22), especially for postmenopausal women. In contrast, there was no significant correlation with premenopausal women (RR = 0.97, 95% CI 0.84-1.12) or the Asian population (RR = 1.07, 95% CI 0.94-1.22).

Furthermore, Largent et al. (61) additionally studied the impact of the use of antihypertensive drugs on the risk of breast cancer among people with hypertension. Compared to people not receiving antihypertensive treatment, those who used antihypertensive drugs for more than 5 years had a significantly increased risk of invasive breast cancer (RR = 1.18, 95% CI 1.02-1.36), especially ER+ breast cancer (RR = 1.21, 95% CI 1.03-1.42). Among the drugs, people who used diuretics for more than 10 years showed a significant association with the occurrence of breast cancer (RR = 1.16, 95% CI 1.01-1.33), especially ER+ breast cancer (RR = 1.21, 95% CI 1.03-1.42). The possible mechanism is that breast cancer and hypertension have a common pathophysiological pathway mediated by adipose tissue causing chronic inflammation to form metabolic syndrome (62), and hypertension may increase the risk of disease by blocking and changing apoptosis (63). More research is needed to clarify the relationship between hypertension and breast cancer.

## METABOLIC SYNDROME AND BREAST CANCER PROGRESSION

### Molecular Changes in Patients With Metabolic Syndrome

As the body's metabolic state changes, the corresponding molecular level also changes, leading to proliferation, invasion and metastasis of breast cancer *via* various signaling pathways and target genes (Figure 2). Under an obese state, the tumor microenvironment changes and produces more fibroblasts and immune cells, such as T cells, macrophages and endothelial cells (64). In the mammary gland, the interaction between obese adipocytes and breast cancer cells leads to the transformation of mammary adipocytes into cancer-associated adipocytes (CAAs) (3), which secrete more leptin and reduce the production of adiponectin. Obese adipose tissue is associated with chronic inflammation, which promotes the production of proinflammatory factors, such as TNF- $\alpha$ , IL-6, and IL-8, and inhibits the secretion of anti-inflammatory factors, such as adiponectin (3, 8). In addition, obese adipose tissue is also associated with increased aromatase activity, which promotes the conversion of androgens to estrogen (65).

Diabetes, especially type 2 diabetes, is often accompanied by insulin resistance, leading to hyperinsulinemia. Diabetes also promotes the production of IGF-1 in the liver (3) and inhibits the secretion of sex hormone-binding globulin (SHBG) (51) and IGF-binding protein (IGFBP) (66).

The renin-angiotensin system (RAS) is the main pathway for regulating blood pressure, and an increase in ANG II (67) leads to hypertension. Serum hypocalcemia, one of the characteristics of hypertension, is also related to the occurrence of tumors.

Dyslipidemia is mainly characterized by high TG and high TC. Elevated cholesterol and increased very-low-density lipoprotein (VLDL) and LDL promote tumor proliferation through different pathways (68).

We reviewed the main findings recent years on the principal molecular actors that are involved in the interactions between MetS and breast cancer biology, including leptin, adiponectin, insulin and IGF-1, Angiotensin II and Calcium and, cholesterol and lipoprotein.

### Leptin

Leptin (Figure 2A) is a hormone secreted by adipose cells that acts on the hypothalamus to suppress appetite, increase basal metabolism and inhibit the synthesis of adipocytes by binding to leptin receptors. Leptin is also considered an important biomarker of metabolic syndrome. Elevated levels of leptin are related to obesity, and obese people may exhibit leptin resistance (35, 69). Leptin levels are also proportional to the degree of insulin resistance.

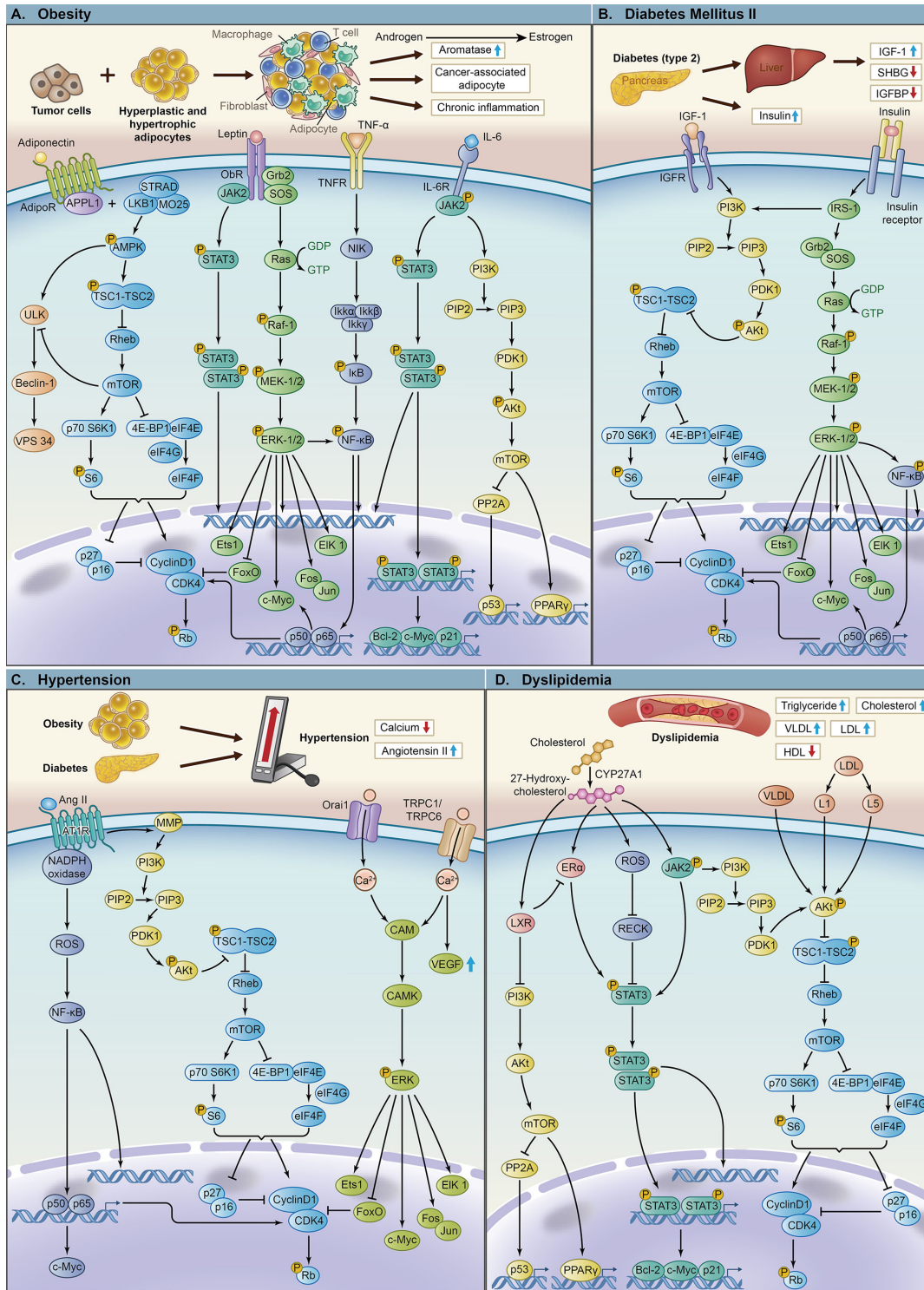
Sieminska et al. (70) found that the leptin level in postmenopausal women was positively correlated with the number of metabolic syndrome components. Leptin is expressed in normal breast tissue, breast cancer cells and solid tumors. Recent research shows that leptin can upregulate aromatase expression by regulating the p53-hypoxia inducible

factor 1 $\alpha$ /pyruvate kinase M2 axis in breast adipose stromal cells, increasing the risk of breast cancer (71) and promoting the growth of breast tumors. Pan et al. (72) demonstrated that serum leptin levels were related to overall breast cancer risk (standardized mean difference = 0.46, 95% CI 0.31-0.60), especially in Chinese women (standardized mean difference = 0.61, 95% CI 0.44-0.79). Some *in vitro* experiments have shown that leptin can increase the expression of cox-2 and members of the PI3K/Akt pathway in cocultures of leptin and breast cancer cells, such as MCF-7, MDA-MB-231 and BT474, to promote tumor growth (73–75). Leptin has also been found to promote breast cancer invasion and metastasis through the upregulation of procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2 (PLOD2) and IL-18 (76, 77). In *in vivo* experiments, tumor-bearing mouse models showed that leptin increased the volume of tumors and promoted lung metastasis (76). A case-control study by Maccio et al. (78) showed that leptin levels are associated with higher TNM staging and an increased risk of distant metastasis ( $P < 0.05$ ) in postmenopausal breast cancer patients. Regarding potential mechanisms (79), leptin can activate the JAK2/STAT3, PI3K/Akt/mTOR, and extracellular regulated protein kinase (ERK) pathways by binding to its own receptors expressed in tumor cells and stromal components, including immune cells, endothelial cells and tumor-associated fibroblasts. Moreover, it may stimulate the epithelial mesenchymal transition (EMT) (80), which has been confirmed to activate the EMT of breast cancer cells by upregulating pyruvate kinase M2 to activate the PI3K/Akt pathway (81), increasing the activity of matrix metalloproteinases (MMPs), promoting the formation and maintenance of breast cancer stem cell (BCSC) survival, inducing the activation and proliferation of endothelial cells and modulating tumor-immune cell cross talk. Sabol et al. (82) found through molecular experiments that adipose stem cells modified by obesity promoted the secretion of leptin and upregulated and activated the IL-6 and Notch signaling pathways in ER+ breast cancer cells, resulting in ER+ breast cancer radiation resistance. Leptin signaling affects the progression of ER- breast cancer (83). Recent *in vitro* experiments have also found that the expression of leptin receptor and leptin-targeting genes is associated with reduced survival rate of ER- breast cancer and chemotherapy resistance, indicating that the coexpression of leptin receptor and leptin targeting genes can be used as a prognostic indicator of ER-breast cancer patients (84).

Numerous *in vivo* and *in vitro* studies confirmed the role of leptin in recurrence and metastasis. However, relevant clinical evidence is relatively lacking, so more research is needed to investigate leptin as a predictive biomarker and a novel therapeutic target in the clinic.

### Adiponectin

Adiponectin (Figure 2A) is a protein synthesized and secreted by white adipocytes. Obese patients have lower levels of adiponectin compared with normal patients. In contrast to leptin, the concentration of adiponectin decreases as the number of metabolic syndrome components increases (85–88). Adiponectin activates various signaling pathways (AMPK,



**FIGURE 2** | The key pathways and molecular mechanisms of components of metabolic syndrome leading to tumor proliferation, invasion and metastasis. **(A)** The mechanism of obesity and its related key molecules leading to breast cancer with important signaling pathways, including LKB1/AMPK/mTOR, AMPK/ULK, JAK2/STAT3, Ras/Raf/MEK/ERK, NF-κB and PI3K/Akt/mTOR. **(B)** The mechanism of diabetes mellitus type 2 and its related key molecules leading to breast cancer with important signaling pathways, including PI3K/Akt/mTOR, Ras/Raf/MEK/ERK and NF-κB. **(C)** The mechanism of hypertension and its related key molecules leading to breast cancer with important signaling pathways, including NF-κB, PI3K/Akt/mTOR and CAM/CAMK/ERK. **(D)** The mechanism of dyslipidemia and its related key molecules leading to breast cancer with important signaling pathways, including PI3K/Akt/mTOR, JAK2/STAT3 and ERα/STAT3.

PI3K/Akt, MAPK, PPAR- $\alpha$ , STAT3 and NF- $\kappa$ B) (39) and exerts various cellular functions by binding to two receptors, adiponectin receptor 1 (AdipoR1) and 2 (AdipoR2). Adiponectin improves insulin sensitivity *in vivo* by activating PPAR- $\alpha$  (89). Based on the role of adiponectin in the abovementioned signaling pathways, the short peptide ADP355 based on adiponectin can exert antiproliferative effects in breast cancer cells through the STAT3 and ERK1/2 signaling pathways (39, 90). Adiponectin also increases insulin sensitivity in muscles and liver through the AMPK pathway and improve insulin resistance (91) as an endogenous insulin sensitizer. Adiponectin deficiency causes downregulation of the activity of PTEN and activation of the PI3K/Akt signaling pathway, thereby promoting breast tumors (92). In addition, adiponectin controls inflammation, inhibits angiogenesis (93), reduces proliferation, and promotes apoptosis. It inhibits the expression of adhesion molecules in endothelial cells, suppresses the growth of macrophage precursors and downregulates TNF- $\alpha$  and IL-6/8 to control inflammation (94). Adiponectin potentially inhibits angiogenesis through the AMPK pathway to enhance nitric oxide production, activate endothelial nitric oxide synthase (95), and suppress the expression of pro-angiogenesis factors, such as vascular endothelial growth factor (VEGF) and IL-8. *In vitro* experiments further demonstrated that in MCF-7 and MDA-MB-231 cell lines, the proliferation of endothelial cells expressing adiponectin receptors is inhibited by adiponectin, thereby reducing VEGF expression and inhibiting metastasis, invasion, and angiogenesis (96). Moreover, adiponectin may inhibit TNF- $\alpha$ -induced activation of the NF- $\kappa$ B pathway to promote apoptosis.

Adiponectin is negatively related to the risk of breast cancer. Many studies have shown that low adiponectin levels are associated with the risk of breast cancer and the progression of more aggressive subtypes (40), which is more common in ER-/PR- breast cancer. *In vivo* experiments showed that in animals receiving ER- MDA-MB-231 cells, the tumor volume was significantly reduced after adiponectin pretreatment (97). Moreover, Oh et al. (98) found that adiponectin may protect against recurrence in ER-/PR- breast cancer patients. Although adiponectin exhibits antiproliferative effects on cell growth in both ER+ and ER- cell lines, the main mechanisms may be different. For ER+ breast cancer, a low adiponectin concentration can amplify ER signaling to increase the proliferation of ER+ breast cancer cells, leading to the progression of breast tumors (40, 99, 100). Low adiponectin levels may mediate the proliferation of ER- breast cancer cells by regulating genes involved in the cell cycle (such as p53, Bax, Bcl-2, c-myc and Cyclin D1) (101), which indicates that adiponectin can be used as adjuvant therapy for ER- breast cancer, and its local and systemic application can reduce tumor size and inhibit tumor metastasis.

The possible mechanism by which adiponectin exerts antitumor growth and proautophagy effects is the LKB1-AMPK-mTOR/ULK pathway (102). *In vitro* experiments have shown that adiponectin increases the expression of the tumor suppressor gene LKB1 in breast cancer cells and subsequently reduces tumor adhesion, migration and invasion through the

AMPK-p70S6 kinase (S6K) axis (103). Adiponectin binds to the C-terminus of AdipoR, and its N-terminus binds to the PTB domain of signal adaptor protein (APPL1), affecting LKB1 expression. In obese patients, adiponectin production is reduced such that binding to AdipoR is inhibited. The binding of the domain of APPL1 to LKB1 is suppressed, so AMPK and ULK expression is consequently inhibited (104). Some studies have summarized that the *in vitro* data of adiponectin's antitumor proliferation effect in breast cancer are mainly limited to full-length adiponectin (fAd) (105, 106), one of the forms of adiponectin. The other form of adiponectin, globular adiponectin (gAd), may have the opposite effect. It can activate AMPK through its high-affinity receptor AdipoR1 acting on aggressive tumor cells, and AMPK upregulates autophagy by activating ULK1 to enhance tumor metastasis (106). Mauro et al. further found that gAd inhibits the growth of ER+ breast cancer but promotes ER+ breast cancer proliferation (107).

Based on adiponectin-mediated cytotoxic autophagy, we can reasonably hypothesize that adiponectin usage represents a new adjuvant therapy strategy for obese breast cancer patients and that combining adiponectin with chemotherapeutic drugs may therefore reduce the dose of chemotherapeutic drugs (108). Studies have concluded (109) that leptin and adiponectin receptors seem to be the most promising molecular targets for the treatment of metabolic syndrome-related cancers. In the future adjuvant therapy of breast cancer, for patients with metabolic syndrome, it may be possible to improve the efficacy by adjusting the circulating levels of the above molecules to obtain a better prognosis.

## Insulin and IGF-1

Insulin (**Figure 2B**) is an anabolic hormone. It has mitogenic, antiapoptotic and angiogenic effects, which are partially related to cancer progression and mortality. Hyperinsulinemia can also affect the prognosis of breast cancer mainly through the following mechanisms. Insulin itself promotes the synthesis of DNA, RNA, and ATP, induces mitosis and angiogenesis, and inhibits apoptosis. High insulin levels induce aromatase activity (110), increase the chance of conversion of androgen to estrogen, and promote mitosis. Moreover, high insulin levels also reduce the production of SHBG synthesized by the liver (111) and increase the proportion of circulating estradiol, leading to the growth of breast cancer. *In vitro* experiments proved that insulin induces EMT, invasion and migration of tumor cells (112). In a chronic hyperinsulinemic state, the ER is activated to promote tumor growth by regulating the cell cycle, apoptosis factors, and nutrient metabolism (113). This mechanism provides a basis for using metformin to treat ER+ breast cancer patients with diabetes. In addition, hyperinsulinemia mediates the production of IGF-1 (**Figure 2B**), inhibits the production of IGFBP in the liver, increases the bioavailability of IGF-1 and stimulates further tumor growth through excessive activation of the PI3K-Akt-mTOR pathway and Ras-MAPK pathway (114).

MetS patients with insulin resistance have high circulating insulin levels. Insulin binds to the insulin receptor on the cell membrane and activates IRS1 and subsequently activates the Ras/Raf/MEK/ERK pathway to regulate cell proliferation and



differentiation. Insulin also stimulates the production of IGF-1 in the liver and downregulates the secretion of IGFBP1 and 2 in the liver, leading to increased bioavailability of circulating IGF-1 (66, 115). IGF-1 receptor (IGF-1R) is a transmembrane receptor tyrosine kinase that is similar in structure to the insulin receptor. It is often overexpressed in breast cancer cells and regulates cell proliferation, survival, differentiation, and transformation (116–119). IGF-1 combines with IGFR, which activates PI3K and thereby activates Akt. Activated Akt further induces the mTOR pathway, leading to cell proliferation and protein synthesis in tumor cells.

Therefore, improving insulin resistance through the above pathways can inhibit tumor growth and proliferation directly or indirectly, and it may also relieve other metabolic abnormalities that can improve the curative effect of adjuvant therapy, leading to better disease prognosis.

## Angiotensin II and Calcium

RAS and calcium (Figure 2C) are hypertension-related molecules. Studies have reported the relationship and possible mechanism between these molecules and breast cancer. RAS is an important physiological mechanism of hypertension, and studies have shown that it also plays an important role in the progression of breast cancer. It has been observed in some *in vitro* and *in vivo* experiments that all components of the RAS are overexpressed in breast cancer cells (67). Among them, angiotensin II (Ang II) (Figure 2C) is a well-known hypertension hormone that binds to angiotensin II type 1 receptor (AT1R) and angiotensin II type 2 receptor (AT2R). Ang II-AT1R is related to breast cancer proliferation, and its mechanism has been extensively studied. After Ang II binds to AT1R, AT1R interacts with nicotinamide adenine dinucleotide phosphate (NADPH) oxidase to activate AT1R and promotes the infiltration of macrophages and the production of VEGF and reactive oxygen species (ROS). ROS further promote members of the downstream NF- $\kappa$ B pathway to bind to target genes and mediate cell transcription. AT1R also promotes the secretion of matrix metalloproteinases and then activates the PI3K/Akt/mTOR pathway to promote tumor growth and inhibit cell apoptosis.

Calcium ions are also an important part of the pathogenesis of hypertension. Studies have found that calcium ion channels are associated with tumor proliferation, angiogenesis, apoptosis and metastasis. In hypertensive patients, calcium ions flow through calcium channels, resulting in intracellular high calcium and extracellular low calcium. A meta-analysis by Wulaningsih et al. (120) found that serum calcium had a protective effect on breast cancer (RR 0.80, 95% CI 0.66–0.97). Deliot et al. (121) showed that calcium ions flood into cells through the Orai1 channel (Figure 2C), phosphorylating ERK *via* the calcium/calcium-dependent calmodulin (CAM)/calmodulin kinase II (CAMK) pathway, which regulates cell proliferation. In addition, Orai1 levels in TNBC were increased compared with that in non-TNBC (122). Orai1 is a 33kDa protein with four transmembrane domains and its functional channel is a hexamer composed of 6 Orai1 subunits (123). Orai1 forms a highly selective calcium ion channel in the plasma membrane to mediate the transmission of calcium ions (124). The antiproliferative and antimigratory effects of Orai1 silencing were observed in the MDA-MB-231 basal breast

cancer cell line. This finding reveals that the Orai1 channel may represent a therapeutic target for TNBC. Calcium can also stimulate breast cancer cell proliferation and activate VEGF through the (transient receptor potential-canonical) TRPC6/TRPC1 channel (Figure 2C). The use of calcium channel blockers may inhibit the growth of breast cancer tumors. The use of calcium-channel blockers for 10 or more years was associated with a greater than twofold increase in the risk of ductal breast cancer (OR 2.4, 95% CI 1.2–4.9) and lobular breast cancer (OR 2.6, 95% CI 1.3–5.3) according to Li et al. (125).

## Cholesterol and Lipoprotein

High cholesterol (Figure 2D) is an important feature of dyslipidemia in obese patients and one of the side effects of adjuvant therapy in patients. It has a certain impact on the pathophysiology and progression of breast cancer. Nelson et al. (126) found that 27-hydroxycholesterol (27HC), a primary metabolite of cholesterol and an ER and liver X receptor (LXR) ligand, might play an important role in the progression of breast cancer. Cholesterol is converted into 27HC by cytochrome P450 oxidase. Then, 27HC activates ER and LXR to promote breast cancer cell growth and metastasis. Nelson et al. further confirmed that in mouse models of breast cancer, 27HC significantly reduced tumor latency ( $P < 0.05$ ) and accelerated tumor growth ( $P < 0.05$ ). Increases in markers, such as cell proliferation, angiogenesis, and macrophage infiltration, were also observed in mice treated with 27HC. In higher grade tumors, increased expression of cytochrome P450 oxidase was observed (OR 6.7, 95% CI 1.7–27,  $P = 0.0007$ ), indicating that cancer cells could increase 27HC levels through autocrine signaling. In addition, 27HC activates the STAT3 pathway by reactive oxygen species (ROS) methylation and activation of JAK and extracellular ER $\alpha$ . LXR suppresses the PI3K/Akt pathway to inhibit cell proliferation and downregulate ER expression (127), and ER enhances STAT3 activation (128).

Epidemiologically, abnormal lipoprotein levels (Figure 2D) have also been shown to be significantly associated with breast cancer. Data from Lu et al. (68) provide a new mechanism by which lipoproteinemia promotes tumor development. They found that the VLDL, L1 and L5 subfractions of LDL (Figure 2D) activated Akt to promote cell migration by Ser473 phosphorylation. VLDL, L1, and L5 also increase mesenchymal markers, such as vimentin and  $\beta$ -catenin, to induce EMT and enhance angiogenic factors in breast cancer to promote angiogenesis. Therefore, reducing circulating cholesterol or inhibiting its conversion to 27HC and abnormal lipoprotein levels may represent new strategies for the prevention and/or treatment of breast cancer. However, the effect of lipid composition on the growth of breast cancer cells and its mechanism still require more research.

## METABOLIC SYNDROME AND TREATMENT EFFECT ON BREAST CANCER

### Metabolic Syndrome and Adjuvant Therapy Efficacy

Healy et al. (129) reported that MetS is associated with more aggressive tumor biology, such as later tumor stage ( $P = 0.022$ )

and lymph node invasion ( $P = 0.028$ ). Compared with breast cancer patients who do not have metabolic abnormalities, those with MetS and its components show worse treatment responses to various therapies.

Regarding chemotherapy, a study by Stebbing et al. (130) on the relationship between MetS and response to chemotherapy for breast cancer found that among patients with metabolic syndrome, the proportion of progression, stability and response to treatment after chemotherapy ranged from 61.1% to 42.9% to 33.3% ( $P$  for trend = 0.03). This finding indicated that the MetS-mediated risk of nonresponse to chemotherapy was higher and that treatment efficacy was worse. Litton et al. (131) found that overweight (BMI = 25–29 kg/m<sup>2</sup>) and obese (BMI  $\geq$  30 kg/m<sup>2</sup>) groups exhibited more difficulties than normal or underweight (BMI < 25) groups in achieving pathologic complete response (pCR) (OR = 0.67, 95% CI 0.45–0.99) with neoadjuvant chemotherapy. This unsatisfactory treatment response may be due to underdose in obese patients (132). In the guidelines issued by the American Society of Clinical Oncology, full-dose chemotherapy based on body weight was recommended to obese patients because their worsening survival may be related to the insufficient dose of cytotoxic therapy (41, 133). However, considering safety and therapeutic benefits, overweight and obese patients worldwide are still receiving reduced doses of chemotherapy (23, 134).

DM can also affect the efficacy of chemotherapy. The elevated concentration of IGF-1 under hyperglycemia can specifically inhibit cell death induced by anticancer drugs in MCF-7 cells, which suggests its involvement in the mechanism of drug resistance. Zeng et al. (135) showed that hyperglycemia-induced resistance to chemotherapy was only observed in ER+ breast cancer cells, indicating that antiestrogen may promote the effectiveness of chemotherapy in such patients.

For endocrine therapy, Ewertz et al. (29) followed up on 18,967 Danish women with BMI information who received early breast cancer treatment found that after one decade, in contrast to the patients with BMI < 25 kg/m<sup>2</sup>, endocrine therapy in patients with BMI  $\geq$  30 kg/m<sup>2</sup> had a worse response. Compared to normal weight women, the disease-free survival (DFS) rate (HR = 1.78, 95% CI 1.12–2.83) and overall survival (OS) rate (HR = 2.28, 95% CI 1.16–4.51) are worse in obese women receiving tamoxifen (TAM) combined with aromatase inhibitors (AIs) (136). In this regard, they further studied the role of fulvestrant and TAM in chemotherapy resistance. Fulvestrant is a selective estrogen receptor degrader that can not only competitively bind to ER, but also induce ER degradation and down-regulate ER levels. Fulvestrant blocked hyperglycemia-mediated chemotherapeutic resistance, but TAM did not (135). Diabetes is another major factor affecting endocrine therapy, which may lead to a worse treatment response (8, 137). For DM patients, IGF-1 receptor or IRS-1 overexpression increases the resistance of breast cancer cells to antiestrogen therapy (138).

Trastuzumab is a widely used targeted medicine. It is a monoclonal antibody against HER2, affecting the transmission of growth signals through specific binding to the HER2 receptor, and at the same time kill tumor cells through antibody-dependent cell-mediated cytotoxicity pathway. For targeted

therapy, obese adipocytes secrete more IGF-1, thereby increasing drug resistance, such as trastuzumab resistance (139), but the underlying mechanism remains unclear. Lee et al. (140) demonstrated in their research that in HER2+ breast cancer patients receiving trastuzumab treatment, DM was a significant unfavorable prognostic factor for DFS ( $P = 0.006$ ) and OS ( $P = 0.017$ ), which was consistent with previous reports (141).

For radiation therapy, in the study of Sabol et al. (82), when breast cancer cells were cocultured with obesity-altered adipose stem cells, breast cancer cell apoptosis decreased and survival increased after radiotherapy. Research has suggested that obesity-altered adipose stem cells upregulate IL-6 and activate the Notch signaling pathway to induce radiation tolerance. Fang et al. (142) found that higher BMI was associated with worse quality of life for breast cancer patients before, during, and after radiotherapy even after adjusting for other factors. DM patients also showed worse response and radiation tolerance through elevation of IGF-1R and IRS-1 (47).

## Adjuvant Therapy and Metabolic Changes

A published observational study (143) showed that after patients completed corresponding chemotherapy, metabolic syndrome and its components were significant risk factors ( $P < 0.01$ ). Among them, the fasting blood glucose level changed by 20.3% and the triglyceride level changed by 18.4%, the deterioration of which were more significant. Several studies have demonstrated that breast cancer patients who have received chemotherapy are more likely to gain weight than those who have not (144–146). Fredslund et al. (147) also observed consistent results. Changes were evident in body fat ( $P = 0.01$ ), triglycerides ( $P = 0.03$ ), WC ( $P = 0.008$ ), glucose ( $P = 0.02$ ) and diastolic blood pressure ( $P = 0.04$ ) of premenopausal women, whereas changes in WC ( $P = 0.03$ ), HDL ( $P = 0.05$ ) and glucose ( $P = 0.02$ ) were observed in postmenopausal women. These findings suggested that chemotherapy may have greater adverse effects in premenopausal breast cancer patients because this group may be induced to undergo either transient or permanent early menopause that promotes the rapid decline of estradiol levels in the body, leading to weight gain and a rapid and continuous increase in fat mainly in the abdomen, which causes a series of changes in metabolism-related components. Such weight gain may also increase the risk of chemotherapy-related diabetes. Glucocorticoids used in combination with chemotherapeutic drugs or chemotherapeutic drugs themselves, such as platinum and cyclophosphamide, can cause abnormal glucose metabolism through direct or indirect mechanisms and worsen the pre-existing DM state in susceptible individuals (148). Among the three breast cancer treatments, including radiotherapy, chemotherapy, and endocrine therapy, evaluated in the analysis conducted by Bordelea et al. (149), only chemotherapy ( $P = 0.03$ ) was related to new-onset diabetes.

Studies have shown that endocrine therapy, such as estrogen inhibitors, may also lead to worse metabolic status. TAM is a commonly used antiestrogen drug that can cause dyslipidemia, a well-known side effect with a worsening HDL level and circulating TG level, in women with breast cancer (150).

However, TAM was reported to lower circulating TC and (low-density lipoprotein) LDL (151). Fatty liver ( $P = 0.000$ ) and visceral adipose tissue ( $P = 0.000$ ) are also significant side effects of receiving TAM (152).

Furthermore, the use of TAM can lead to increased fasting blood glucose levels and insulin resistance even when administered in low doses. A population-based cohort study launched by Sun et al. (153) demonstrated that TAM users showed a significantly increased risk of DM compared with non-TAM users among breast cancer patients (adjusted HR = 1.31, 95% CI 1.19-1.45). TAM also leads to a decrease in insulin sensitivity. Johansson et al. (150) observed a 7-fold decrease in insulin sensitivity among breast cancer patients using TAM (OR = 0.15, 95% CI 0.03-0.88) compared with nonusers. The underlying mechanism of the link between TAM and diabetes is still unclear. A possible explanation is that estrogen maintains steady blood glucose, and TAM may affect the interaction between estrogen and insulin by inhibiting estrogen. Additionally, hypertriglyceridemia and fatty liver caused by TAM are features of insulin resistance and glucose intolerance (153).

AIs, including letrozole, anastrozole and exemestane, are another important hormone therapy. Similar to TAM, abnormal lipid metabolism is one of the most significant side effects of AI, but the specific impact is different. Compared with TAM, AI may not change TG levels but increase TC and LDL-C levels. Among AIs, anastrozole and letrozole increase TC and LDL-C levels, whereas exemestane decreases TC, LDL-C, TG and HDL-C levels (151, 154). An early *in vitro* study showed that 17-hydroxy exemestane, a metabolite of exemestane, may elicit androgenic effects by binding to the androgen receptor. Bell et al. (154) hypothesized that this may be a potential mechanism by which exemestane could reduce HDL cholesterol. However, less is known about nonsteroidal aromatase inhibitors, such as anastrozole and letrozole. Hong et al. (155) showed in their cohort study that the incidence rate of fatty liver was increased in the TAM group than in the AI group ( $P = 0.021$ ). The reason may be that TAMs increase circulating TG levels and promote insulin resistance, thereby promoting susceptibility to fatty liver. These findings also suggested that TAMs have direct liver toxicity, but more evidence is needed to confirm this hypothesis (155).

In addition, hypertension is a characteristic adverse event of bevacizumab therapy. The incidence was 17.9% in a clinical study (156). The possible mechanism is that bevacizumab inhibits vascular endothelial growth factor (VEGF), which enhances endothelial nitric oxide synthase activity. These effects lead to decreased production of vasodilator nitric oxide, which causes hypertension (157).

Thus, an interaction exists between adjuvant therapy and human body metabolic status. Patients with unstable metabolic status will receive unfavorable results from adjuvant therapy. Conversely, adjuvant therapy will also cause deterioration of the human body's metabolic status.

## Obesity and Cardiotoxicity of Trastuzumab and Anthracyclines

Compared with chemotherapy alone, the combination of targeted medicine and chemotherapy has clinical advantages in

improving the DFS and OS rates of breast cancer patients (158–160). Anthracyclines and trastuzumab are widely used in the treatment of breast cancer patients, and both can induce acute or chronic dose-dependent cardiotoxicity, especially acute cardiac insufficiency, which is the most serious side effect. This condition is characterized by reduced left ventricular ejection fraction (LVEF), which is typically asymptomatic or associated with heart failure (161). A meta-analysis of 8,754 breast cancer patients treated with anthracyclines, sequential anthracyclines (anthracyclines followed by trastuzumab), and trastuzumab launched by Guenancia et al. (162) showed that in an unadjusted analysis, overweight plus obesity was obviously related to the risk of cardiotoxicity of the above therapies. The OR was 1.38 (95% CI 1.06-1.80) for overweight plus obesity, 1.47 (95% CI 0.95-2.28) for obese individuals and 1.15 (95% CI 0.83-1.58) for overweight individuals. Subgroup analysis showed that the risk of cardiotoxicity gradually increased with increasing BMI ( $P = 0.05$ ). For different administrations, the rate of cardiotoxicity of obese patients treated with anthracyclines only was 20% (95% CI 5%-43%), and obese patients treated with trastuzumab with or without anthracyclines had a 16% (95% CI 10%-24%) rate of cardiotoxicity.

Gunaldi et al. (163) found an association between postmenopausal women ( $P = 0.01$ ) and cardiotoxicity caused by trastuzumab. Obesity ( $P = 0.0001$ ) and hypertension ( $P = 0.002$ ) were related to lower LVEF in patients, whereas diabetes ( $P = 0.766$ ) was not statistically significant. However, a retrospective study (164) observed a correlation between a history of diabetes and trastuzumab-related cardiotoxicity ( $P = 0.01$ ). Therefore, evaluation of heart function or appropriate dose reduction is needed for obese breast cancer patients before treatment with anthracycline or trastuzumab. During and after treatment, LVEF should also be evaluated regularly to prevent heart insufficiency and heart failure. The relationship between people with DM and the incidence of cardiotoxicity caused by trastuzumab requires further exploration.

Obese patients or mice fed a high-fat diet were more sensitive to the cardiotoxicity caused by anthracyclines (165, 166). The mechanism may be due to downregulation of cardiac peroxisome proliferator-activated receptor- $\alpha$ , decreased mitochondrial adenosine monophosphate (AMP)- $\alpha 2$  protein kinase and a reduction in cardiac adenosine triphosphate (ATP) levels after doxorubicin administration according to Mitra et al. (166). Therefore, high-fat diet-induced obese rats are highly sensitized to anthracycline-induced cardiotoxicity by downregulating cardiac mitochondrial ATP generation, increasing oxidative stress and downregulating the Janus kinase (JAK)/signal transducers and activators of transcription 3 (STAT3) pathway. Additionally, Maruyama et al. (167) found that adiponectin-knockout mice showed aggravated left ventricular systolic dysfunction after injection of doxorubicin, whereas exogenous adiponectin improved this condition in wild-type and adiponectin-knockout mice. These results indicate that downregulation of adiponectin levels partly affects adverse cardiac reactions, suggesting that adiponectin may be used as a therapy to prevent cardiotoxicity caused by anthracyclines in

obese breast cancer patients. More evidence is needed to confirm the underlying mechanism of trastuzumab-induced cardiotoxicity in obese patients.

## METABOLIC SYNDROME AND PROGNOSIS OF BREAST CANCER

### Recurrence

Current evidence suggests that metabolic syndrome and its components are associated with an increased risk of breast cancer recurrence (3, 168). A retrospective study conducted by Ewertz et al. (29) assessed the events of locoregional recurrence and distant metastasis among 53,816 women with early-stage breast cancer up to 10 years after diagnosis and showed that obesity may have no effect on the risk of local recurrence ( $P > 0.05$ ) but had a significant association with distant metastasis 5 to 10 years after diagnosis. The HR of distant metastasis in patients with a BMI of 25 to 29 kg/m<sup>2</sup> was 1.42 (95% CI 1.17-1.73,  $P < 0.001$ ). The HR in patients with a BMI of 30 kg/m<sup>2</sup> or more was 1.46 (95% CI 1.11-1.92,  $P = 0.007$ ). In particular, adiponectin is an important adipocytokine, and its expression is reduced in obese people. Oh et al. (98) found that compared to the highest quartile of serum adiponectin levels, the lowest quartile showed a 2.82-fold increased risk.

Goodwin et al. (169) demonstrated that, regardless of menopausal status, fasting plasma insulin levels are associated with high tumor grade, axillary lymph node involvement, and risk of recurrence. Patients in the highest quartile of insulin level had an increased risk of distant recurrence (HR 2.0, 95% CI 1.2-3.3,  $P = 0.007$ ). However, in different molecular types of breast cancer, the correlation between insulin resistance and recurrence risk shows different results (98). In contrast with ER-/PR- patients, a negative association is noted between insulin resistance and tumor recurrence in ER+/PR+ people, which is consistent with serum insulin levels. Moreover, ER+/PR+ patients with hyperglycemia exhibited a lower risk of recurrence (HR = 0.48, 95% CI 0.26-0.89,  $P = 0.020$ ) (Table 2).

The impact of dyslipidemia on the prognosis of breast cancer has demonstrated conflicting results. In general, breast cancer patients with high levels of TC, low density lipoprotein-cholesterol (LDL-C) and TG and low levels of HDL-cholesterol (HDL-C) show poor prognosis (170, 172). However, a retrospective study by Jung et al. (173) observed opposite results. In the study, compared to high levels of LDL-C (quartile IV) and TGs (quartile IV), lower levels of LDL-C (quartile III) (HR 1.88, 95% CI 1.09-3.27,  $P = 0.02$ ) and abnormally low levels of TGs (quartile I) (HR 1.88, 95% CI 1.07-3.29,  $P = 0.03$ ) showed obviously higher risks of recurrence. Ozdemir et al. (174) also found that the risk of recurrence was increased in patients with normocholesterolemia compared with patients with hypercholesterolemia ( $P = 0.001$ ). However, Bahl et al. (172) found no statistically significant relationship between lipids and breast cancer outcome other than a trend toward a risk of recurrence with higher TC (HR 1.62 for the fourth to first quartile, 97.5% CI 0.98-2.69,  $P = 0.03$ ) in multivariate analysis.

Therefore, the potential role of lipids in the recurrence of breast cancer should be further studied.

Regarding hypertension, Braithwaite et al. (175) showed that hypertension was another independent predictor of prognosis in breast cancer. Interestingly, hypertension was associated with recurrence among African Americans (HR 1.60, 95% CI 1.07-2.40) but not their white counterparts. However, the reason was not clear, and more studies are needed to explain the distinction. The association among other ethnic groups requires more solid results.

The above findings indicate that MetS and its composition can be regarded as predictors of recurrence. Among them, evaluating adiponectin and insulin concentrations can help determine the prognosis in ER-/PR- patients, and corresponding interventions can be implemented to improve the prognosis and reduce the risk of recurrence. In addition, more research is needed to prove the impact of serum lipid levels and hypertension on the recurrence of breast cancer and the relationship between MetS and its components and breast cancer subtypes. Therefore, it could provide new and specific prognostic detection methods to monitor and prevent recurrence in breast cancer patients.

### Mortality

For patients with early-stage breast cancer, abnormal metabolic status will bring unsatisfactory outcomes. MetS is widely thought to be associated with a risk of mortality for breast cancer patients, and the risk sharply increases as the number of its components increases. Compared with individuals without any metabolic syndrome-related components, patients with 1-2 components have a 5-fold higher risk of death (HR = 4.90, 95% CI 1.47-16.35,  $P = 0.01$ ) and a 6-fold higher risk of breast cancer-specific death (HR = 6.07, 95% CI 1.41-26.21,  $P = 0.02$ ) (Table 2); patients with 3-5 components have a 12-fold higher risk of death (HR = 12.2, 95% CI 3.49-43.01,  $P < 0.0001$ ). In addition, the risk of breast cancer-specific death is 16-fold higher (HR = 15.97, 95% CI 3.49-73.16,  $P < 0.0001$ ) (3).

Epidemiologists show that the mortality rate of breast cancer among Asians is the lowest in the world, but it has rapidly increased in recent years due to the sharp increase in obesity and consequent metabolic disorders (176, 177). It was proven that obese women show a worse survival rate regardless of menopausal status. A study from the United States in 2009 found that after diagnosis with breast cancer, every gained 5 kg of body weight could increase breast cancer-specific mortality by 13% (66). Cho et al. (146) found that in all patients, BMI  $\geq 25$  kg/m<sup>2</sup> was an unfavorable factor for OS ( $P = 0.030$ ). In a Danish retrospective study of 18,967 early breast cancer patients (29), patients with BMI  $\geq 30$  kg/m<sup>2</sup> at the time of diagnosis had a 38% increase in breast cancer mortality compared with patients with a BMI  $< 25$  kg/m<sup>2</sup> (HR = 1.38, 95% CI 1.11-1.71). In addition, studies have also found that obesity was significantly associated with adverse outcomes in women with ER+ tumors, and obesity had an impact on ER- or HER2+ tumors (40, 178). The possible mechanisms may be due to obesity-related molecules, such as leptin, adiponectin, TNF- $\alpha$ , and IL-6, and their subsequent signal pathways. These findings also suggested that obese patients often have tumors detected later along with more aggressive tumor biological characteristics; therefore, fewer

**TABLE 2 |** The effect of metabolic syndrome and its components on recurrence and survival of breast cancer.

| No. | Author              | Pts (N) | Region  | Study design                    | Molecular subtype | Comparison  | Outcome                                      | HR    | 95%CI       | P         | Ref   |
|-----|---------------------|---------|---------|---------------------------------|-------------------|---|--|-------|-------------|-----------|-------|
| 1   | Buono G. et al.     | 717     | Italy   | prospective observational study | All               | 1-2MetS components vs. 0MetS component                  | OS   | 4.90  | 1.47-16.35  | 0.01      | (3)   |
|     |                     |         |         |                                 |                   | 3-5MetS components vs. 0MetS component                  | BCSS   | 6.07  | 1.41-26.21  | 0.02      |       |
|     |                     |         |         |                                 |                   |   | OS   | 12.20 | 3.49-43.01  | <0.0001   |       |
| 2   | Cho W. K. et al.    | 5668    | Korea   | retrospective cohort study      | All               | BMI $\geq$ 25 vs. BMI<25                                | BCSS   | 15.97 | 3.49-73.16  | <0.0001   | (146) |
|     |                     |         |         |                                 |                   | Non-hyperlipidemia vs. Hyperlipidemia                   | OS   | 1.356 | 1.038-1.773 | 0.03      |       |
|     |                     |         |         |                                 |                   |   | DFS  | 1.248 | 1.038-1.502 | 0.076     |       |
| 3   | Emaus A. et al.     | 1364    | Norway  | retrospective cohort study      | All               | BMI $\geq$ 30 vs. BMI=18.5-25                           | OS   | 1.47  | 1.08-1.99   | N/A       | (170) |
|     |                     |         |         |                                 |                   | highest tertile of cholesterol vs. lowest               | OS   | 1.29  | 1.01-1.64   | N/A       |       |
|     |                     |         |         |                                 |                   |   | highest tertile of blood pressure vs. lowest | OS    | 1.41        | 1.09-1.83 |       |
| 4   | Ewertz M. et al.    | 18967   | Denmark | retrospective cohort study      | All               | BMI $\geq$ 30 vs. BMI<25                                | BCSS   | 1.38  | 1.11-1.71   | 0.003     | (29)  |
| 5   | Minicozzi P. et al. | 1607    | Italy   | retrospective cohort study      | ER/PR+            | High glucose(>94.0mg/dl) vs. reference (84.1-94.0mg/dl) | BCSS   | 5.49  | 1.56-19.31  | N/A       | (171) |
| 6   | Oh S. W. et al.     | 747     | Korea   | retrospective cohort study      | ER-/PR-           | Hyperglycemia vs. non-hyperglycemia                     | Recurrence                                   | 0.77  | 0.15-4.17   | N/A       |       |
|     |                     |         |         |                                 | ER-/PR-           | Serum adiponectin                                       | Recurrence                                   | N/A   | N/A         | 0.009     |       |

BCSS, Breast cancer specific survival; BMI, Body mass index; DFS, Disease free survival; ER, Estrogen receptor; HR, Hazard ratio; MetS, Metabolism Syndrome; N/A, Not applicable; OS, Overall Survival; PR, Progesterone receptor; Pts, patients.

treatment opportunities with poorer therapeutic effects are available, which leads to an increased risk of death.

Diabetes is also associated with high breast cancer mortality, and a meta-analysis showed a significantly higher all-cause mortality risk (HR 1.49, 95% CI 1.35-1.65) of patients with breast cancer and diabetes (179). Another meta-analysis also indicated that HR was 1.51 (95% CI 1.34-1.70) for OS and 1.28 (95% CI 1.09-1.50) for DFS in breast cancer patients with diabetes compared to those without (180). However, its effects differ among subtypes. ER+/PR+ women with hyperglycemia are more likely to die of breast cancer (HR = 5.49, 95% CI 1.56-19.31), whereas ER-/PR- patients show no significant association (171). The underlying reasons may be that patients with diabetes receive less aggressive treatment because they are vulnerable to related comorbidities and may have a greater risk of chemotherapy-related toxicity. Additionally, diabetes can directly affect breast cancer by altering related molecules, such as insulin, IGF-1 and inflammatory markers. Bozcuk et al. (181) and Pasanisi et al. (182) found that fasting serum insulin levels are an independent predictor of OS in breast cancer patients, which may be related to the high expression of insulin receptors in breast cancer tissues.

Notably, in the study of Cho et al. (146), the absence of hyperlipidemia is an unfavorable prognostic factor for DFS (HR 1.447, 95% CI 1.080-1.937,  $P = 0.013$ ) and OS (HR 3.085, 95% CI 1.836-5.183,  $P < 0.001$ ) in breast cancer patients, which is perhaps related to the use of statins in patients with hyperlipidemia. A Norwegian research team found (170) that breast cancer patients with high total cholesterol levels (HR for OS is 1.29, 95% CI 1.01-1.64,  $P = 0.03$ ) had a high risk of overall mortality. This finding may

be attributed to the fact that cholesterol contributes to the progression and metastasis of tumors *via* several pathways, such as Akt and EGFR. Fan et al. (14) found that in the TNBC group, patients with low HDL showed worse relapse-free survival (RFS) (HR 3.266, 95% CI 2.087-5.112,  $P < 0.0001$ ) and OS (HR 3.071, 95% CI 1.732-5.445,  $P < 0.0001$ ). The possible mechanism is that HDL negatively correlates with angiotensin (ANG) II, which is positively associated with VEGF pathways in TNBC cells.

Therefore, MetS and its components play a profound role in the survival of breast cancer patients. In individuals, positive associations are observed between obesity as well as diabetes and mortality. Hyperlipidemia shows a protective effect on the mortality of breast cancer. The exact association between MetS and its related components and different phenotypes and underlying mechanisms remain unclear and require further study.

## Anti-MetS and Prognosis

Healy et al. (129) reported that MetS is associated with more aggressive tumor biology, such as later tumor stage ( $P = 0.022$ ) and lymph node invasion ( $P = 0.028$ ). Interventions, including medications, such as metformin and statins, low-calorie and low-fat diets and appropriate exercise to improve metabolic disorders, are needed to reduce the risk of comorbidities in breast cancer patients. The measures to treat MetS may consequently have a positive impact on the prognosis of patients to a certain extent.

Metformin is widely used in the treatment of hyperglycemia. Jiralersong et al. (183) showed that during neoadjuvant chemotherapy for breast cancer, patients with diabetes who

received metformin had a higher pathologic complete response (pCR) rate (24%, 95% CI 13%-34%) compared with those who did not (8%, 95% CI 2.3%-14%). These researchers additionally indicated that metformin use during neoadjuvant chemotherapy was an independent predictor of pCR (OR 2.95, 95% CI 1.07-8.17,  $P = 0.04$ ). Given the differences in insulin usage among the enrolled population, they further analyzed the effect of insulin usage on pCR. In the metformin group, the rate of pCR was not different for insulin use vs. no insulin use (27% vs. 23%,  $P = 0.75$ ). Metformin plays a role in inhibiting the proliferation, invasion and angiogenesis of tumor cells by reversing hyperinsulinemia (183) and improving insulin resistance (34). Some *in vitro* studies have shown that the antitumor effect of metformin *via* epidermal growth factor receptor (EGFR)-mediated pathways (49) is most prominent in TNBC cell lines. In addition, metformin may target the immune microenvironment of tumors to inhibit tumor proliferation (184).

It was reported that breast cancer patients without hyperlipidemia had worse DFS (HR 1.447, 95% CI 1.080-1.937,  $P = 0.013$ ) and OS (HR 3.085, 95% CI 1.836-5.183,  $P < 0.001$ ) (146), which may be due to the use of statins. The retrospective analysis of Li et al. (185) found that long-term use of statins ( $> 5$  years) was associated with improvements in OS (HR = 0.38, 95% CI 0.17-0.85,  $P < 0.018$ ) and DFS (HR = 0.15, 95% CI 0.05-0.48,  $P < 0.001$ ), even after adjusting for metabolic comorbidities. There was no significant difference in OS between patients taking statins for less than 5 years and those who did not take statins. A randomized phase III trial (186) conducted by the International Breast Organization on 8,010 patients with early postmenopausal hormone receptor-positive invasive cancer indicated that compared with patients who did not use cholesterol-lowering drugs, patients who received cholesterol-lowering drugs before endocrine therapy had better DFS (HR = 0.82, 95% CI 0.68-0.99). *In vitro*, statins inhibit the proliferation of the breast cancer cell line MCF-7, which may be due to the blockade of hydroxy methyl glutaryl coenzyme A (HMG-CoA) reductase (187). Ghosh-Choudhury et al. (188) found that simvastatin significantly inhibits the phosphorylation of Akt kinase in MDA-MB-231 breast cancer cells and further inhibits the mammalian target of rapamycin (mTOR) pathway. Statins also induce apoptosis in a variety of cancer cell lines, including colon, prostate, and breast cancer cells (185). Additionally, statins are well tolerated, and their drug interactions are limited. Among different statin types, lipophilic drugs show direct inhibition of breast cancer cell growth *in vitro* and *in vivo*, whereas, hydrophilic drugs such as pravastatin, have no effect (187, 189, 190).

For a long time, aerobic exercise has been widely believed to be effective in improving the abnormal metabolic state of the body, such as reducing fasting blood sugar, HDL, TGs, and WC. Resistance exercise can induce changes in insulin sensitivity by maintaining and/or increasing lean body mass, increasing glucose storage, reducing circulating glucose levels, and promoting a decrease in the amount of insulin required by obese people. A prospective study of 1,490 women with breast cancer conducted by Women's Healthy Eating and Living reported that the equivalent of walking 30 minutes a day for 6 days a week plus eating at least 5 servings of fruit and vegetables a week significantly benefited survival (HR = 0.56, 95% CI 0.31-0.98). However, in the analysis

of subtypes, this lifestyle intervention can only reduce the mortality of ER+ tumors ( $P < 0.05$ ).

Adhering to the Mediterranean diet is another option to reduce the occurrence of metabolic syndrome, preventing the prevalence of breast cancer and improving its prognosis. Some cohort studies have observed a corresponding risk reduction (191–193). The main characteristics of the Mediterranean diet are extensive consumption of fruits, vegetables, unrefined grains, legumes, fish, cereals, nuts, olive oil, and moderate drinking of wine during the main meal (194). Shifting to a Mediterranean diet can improve the imbalance of body metabolism (195) and prevent intractable diseases related to insulin resistance, such as obesity and breast cancer (196). A systemic review published in 2008 confirmed that the Mediterranean diet has a significant negative correlation with the risk of postmenopausal ER- breast cancer (197). Reducing inflammation may be a possible mechanism for the anticancer effect of the Mediterranean diet (1).

Therefore, medications for metabolic disorders and specific lifestyles may be effective strategies to improve the outcome of breast cancer patients, especially those with MetS, after diagnosis to obtain a better prognosis. These research conclusions may provide us with innovative treatments.

## CONCLUSION

Metabolic syndrome and its components have been widely considered to be correlated with the initiation and progression of breast cancer, which is due to obesity and its related adipokines, insulin and IGFs, abnormal serum lipids and lipoproteins and the molecules leading to hypertension. These molecular changes partly exert a profound influence on the tumor and its microenvironment. Metabolic syndrome is significantly associated with an increased risk, worse treatment response, invasive progression and poor prognosis of breast cancer. Notably, we systematically reviewed the mechanisms and pathways of the highlighted molecules affecting disease progression and summarized several potentially novel treatment targets. In the future, new treatment strategies can be prospectively performed based on the above findings to improve prognosis and improve quality of life for breast cancer patients.

## AUTHOR CONTRIBUTIONS

SD, ZW, and XC designed this study. SD performed the search and analysis. SD and ZW wrote the manuscript. XC and KS helped to revise the manuscript. All authors contributed to the article and approved the submitted version.

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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## GLOSSARY

|         |   |
|---------|---|
| 27HC    | 27-hydroxycholesterol   |
| AdipoR  | adiponectin receptor  |
| AI      | aromatase inhibitor   |
| AMP     | adenosine monophosphate   |
| AMPK    | adenosine monophosphate kinase  |
| ANG     | angiotensin   |
| APPL1   | adaptor protein containing the pleckstrin homology domain, phosphotyrosine-binding domain, and leucine zipper motif 1 |
| AT1R    | angiotensin II type 1 receptor  |
| AT2R    | angiotensin II type 2 receptor  |
| ATP     | adenosine triphosphate  |
| BMI     | body mass index   |
| BCSC    | breast cancer stem cell   |
| CAA     | cancer-associated adipocytes  |
| CAM     | calmodulin  |
| CAMK    | calmodulin kinase   |
| CI      | confidence interval   |
| DFS     | disease-free survival   |
| DM      | diabetes mellitus   |
| DNA     | deoxyribonucleic acid   |
| EGFR    | epidermal growth factor receptor  |
| EMT     | epithelial mesenchymal transition   |
| ER      | estrogen receptor   |
| ERK     | extracellular regulated protein kinases   |
| fAd     | full-length adiponectin   |
| gAd     | globular adiponectin  |
| HDL     | high-density lipoprotein  |
| HER2    | human epidermal growth factor receptor 2  |
| HMG-CoA | hydroxy methyl glutaryl coenzyme A  |
| HTWC    | hypertriglyceridemia waist circumference  |
| IGF-1   | insulin growth factor-1   |
| IGFBP   | insulin growth factor binding protein   |
| IL      | interleukin   |
| IRS     | insulin receptor substrate  |
| JAK     | Janus kinase  |

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|----------------|--|
| LDL            | low density lipoprotein                              |
| LKB1           | liver kinase B1                                      |
| LVEF           | left ventricular ejection fraction                   |
| LXR            | liver x receptor                                     |
| MAPK           | mitogen-activated protein kinase                     |
| MEK            | mitogen-activated protein kinase                     |
| MetS           | metabolic syndrome                                   |
| MMP            | matrix metalloproteinases                            |
| mTOR           | mammalian target of rapamycin                        |
| NADPH          | nicotinamide adenine dinucleotide phosphate          |
| NF- $\kappa$ B | nuclear factor kappa-B                               |
| OR             | odds ratio   |
| OS             | overall survival                                     |
| PARP           | poly adenosine diphosphate-ribose polymerase         |
| pCR            | pathologic complete response                         |
| PI3K           | phosphatidylinositol 3-kinase                        |
| PLOD2          | procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2   |
| PPAR           | peroxisome proliferators-activated receptors         |
| PR             | progesterone receptor                                |
| PTB            | phosphotyrosine binding                              |
| RAS            | renin-angiotensin system                             |
| RFS            | relapse-free survival                                |
| RNA            | ribonucleic acid                                     |
| ROS            | reactive oxygen species                              |
| RR             | risk ratio   |
| S6K            | p70S6 kinase   |
| Ser            | serine   |
| SHBG           | sex hormone binding globulin                         |
| STAT3          | signal transducers and activators of transcription 3 |
| TAM            | tamoxifen  |
| TC             | total cholesterol                                    |
| TG             | total triglyceride                                   |
| TNBC           | triple negative breast cancer                        |
| TNF            | tumor necrosis factor                                |
| TRPC           | transient receptor potential-canonical               |
| ULK            | unc-51-likekinase                                    |
| VEGF           | vascular endothelial growth factor                   |
| VLDL           | very-low-density-lipoprotein                         |
| WC             | waist circumference                                  |
| WHR            | waist-to-hip ratio                                   |