

Meta Analysis

Journal of International Medical Research 49(11) I-II © The Author(s) 2021 Article reuse guidelines: sagepub.com/journals-permissions DOI: 10.1177/03000605211061033 journals.sagepub.com/home/imr



Association between serum lipids and breast cancer risk in premenopausal women: systematic review and meta-analysis

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Abstract

Objective: Associations between serum lipids and their individual components with premenopausal breast cancer risk are unclear. This meta-analysis summarized the literature on serum lipids and premenopausal breast cancer risk to elucidate their relationship.

Methods: Eligible studies were identified by searching the PubMed, Embase, China National Knowledge Infrastructure, and Wanfang databases until 31 December 2020. Standardized mean difference (SMD) scores with 95% confidence intervals (95%Cls) were used to assess the impact of serum lipids on premenopausal breast cancer risk. The I² statistic was calculated to measure the percentage of heterogeneity, and Egger's test was performed to measure publication bias.

Results: Thirteen studies were included. The SMD scores of triglycerides (TG) and low-density lipoprotein cholesterol (LDL-C) were 12.90 (95%CI: 7.19–18.61) and 31.43 (95%CI: 8.72–54.15), respectively. The SMD scores of total cholesterol (TC) and high-density lipoprotein cholesterol (HDL-C) were not significantly different between the groups. The included studies were highly heterogeneous. There were no publication biases found in TC, LDL-C, or HDL-C analyses, whereas publication bias was present in the TG analysis.

Conclusions: TG and LDL-C were higher in premenopausal breast cancer patients than in women without breast cancer. However, no significant differences were found in TC or HDL-C levels.

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Premenopausal breast cancer, serum lipid, meta-analysis, triglyceride, low-density lipoprotein cholesterol, tumor biomarker

Date received: 24 May 2021; accepted: 1 November 2021

Introduction

Carcinoma of the breast is the most common cancer in women worldwide.¹ Epidemiologic evidence has suggested that metabolic syndrome is associated with a higher risk of breast cancer.^{2–4} Metabolic syndrome represents a complex group of metabolic disorders including obesity, hyperglycemia, hypertension, and dyslipidemia.⁵ Although dyslipidemia is a component of metabolic syndrome, its relationship with breast cancer risk remains unclear.

Dyslipidemia is a cluster of lipoprotein abnormalities that is defined by the presence of one of the following: high levels of triglycerides (TG), total cholesterol (TC), low-density lipoprotein cholesterol (LDL-C), or low levels of high-density lipoprotein cholesterol (HDL-C). The primary metabolite of cholesterol. 27-Hydroxycholesterol (27-HC), is an endogenous selective estrogen receptor modulator (SERM) with estrogen-like effects; it can induce the proliferation of estrogen receptor-positive breast cancer cells and can facilitate metastasis.6,7 Laboratory studies have indicated that cholesterol is a component of lipid rafts, which act as platforms for cell signaling, invasion, and migration.^{8,9} LDL-C and HDL-C regulate cholesterol levels by mediating cholesterol transport between the liver and blood and the transport of 27-HC through esterification;¹⁰ thus, they may have an indirect effect on breast cancer progression. Meanwhile, LDL-C can trigger the proliferation and migration of cancer cells by activating the P38 signaling pathway, which is related to breast cancer development.¹¹

Furthermore, increased HDL-C content can lead to increased expression of antiinflammatory cytokines such as interleukin-10 and decreased expression of insulin-like growth factor-I; the former protects against breast cancer, whereas the latter promotes carcinogenesis.^{12–14} The diversity of lipid components and the complex mechanisms of each component has made elucidating the relationship between blood lipids and breast cancer difficult.

Menstrual status can also affect serum lipids. LDL-C and TG levels are lower in premenopausal women compared with in postmenopausal women, whereas HDL-C levels show the opposite trend.^{15,16} Therefore, the role of lipoproteins in breast cancer risk may vary according to menopausal status, and several studies support this conclusion.^{17,18} To our knowledge, this is the first meta-analysis to explore the relationship between serum lipids and breast cancer risk only in premenopausal women.

Materials and methods

This meta-analysis was registered with INPLASY in September 2021 (registration number ID: INPLASY202190011), and the DOI number is 10.37766/ inplasy2021.9.0011. This meta-analysis was performed in accordance with the PRISMA 2020 guidelines.¹⁹

Search strategy

We searched the PubMed, Embase, China National Knowledge Infrastructure, and Wanfang databases without restrictions for articles published until 31 December 2020. The following keywords were used for search: "breast cancer," "breast the "breast neoplasm," carcinoma," "serum lipid," "blood lipid," "dyslipidemia," "lipoprotein," "triglyceride," and "cholesterol." Furthermore, we also reviewed the reference lists of all retrieved articles to identify additional studies.

Inclusion and exclusion criteria

Relevant studies were selected if they fulfilled the following criteria: 1) all subjects in the case group were histopathologically diagnosed as having primary breast cancer before menopause and had not started any antitumor treatments; and 2) the risk factors explored included at least one of the selected lipid components (TG, TC, LDL-C, and HDL-C). The study was excluded if included patients had primary hyperlipidemia, metabolic syndrome, or were taking drugs or hormones that affect lipid metabolism.

Data extraction and quality assessment

Two authors (JW and WL) independently assessed study eligibility and extracted data, any disagreements were resolved by consensus. If the study met the inclusion criteria, the following data were extracted: author, year of publication, country, study design, sample size, and mean \pm standard deviation (SD) of serum lipids in the case and control groups.

The nine-star Newcastle–Ottawa Scale (NOS)²⁰ for case–control studies was used to assess methodological quality. NOS consists of three major aspects: selection, comparability, and exposure/outcome. High-quality studies were defined as those with seven or more stars.

Statistical analysis

We used STATA version 15.1 (STATA Corp., College Station, TX, USA) for all

analyses. As the explored exposure factors (serum lipids) were measurement data, standardized mean difference (SMD) scores with 95% confidence intervals (CIs) were calculated. The I² statistic was calculated to measure the percentage of heterogeneity, with I^2 values of 25%, 50%, and 75% corresponding to low, medium, and high data heterogeneity, respectively. When substantial heterogeneity was detected, random effect models were used. Otherwise, fixed effect models were used to assess the effect of the size of the included studies in the meta-analysis. Egger's tests were performed to measure publication bias. A p-value <0.05 was considered statistically significant.

Ethical approval

This study was a meta-analysis, and all data were extracted from previously published articles. Thus, obtaining informed consent was not required.

Results

Literature search results

The process for the literature search and study selection is displayed in a flow diagram in Figure 1. A total of 2027 articles were identified through the search strategy, among which 1842 were excluded because their contents did not meet the purpose of our study. After screening the titles and abstracts of 185 potentially relevant articles, 24 full texts were reviewed for eligibility. Ultimately, after excluding articles whose data did not contain the mean \pm standard deviation or did not report the blood lipid data of premenopausal and postmenopausal women separately, 13 studies were included in this systematic review.

Study characteristics

Characteristics of the 13 included articles²¹⁻³³ are listed in Table 1. NOS

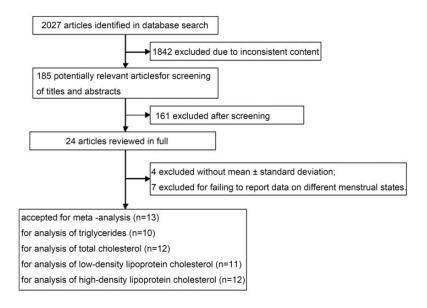


Figure 1. Flow diagram depicting the literature search and study selection.

scores of all included articles were seven stars: thus, they were all considered to be of high quality. Four articles^{21-23,25} were published in Chinese, and the remaining seven were published in English. The 10 articles that analyzed TG included 915 premenopausal women with breast cancer as the case group and 1779 healthy premenopausal women as the control group. The 12 articles that analyze TC included 1202 women in the case group and 1943 women in the control group. The 11 studies that analyzed LDL-C included 1142 and 1913 women in the case and control groups, respectively. For HDL-C analysis, there were 1279 cases enrolled in 12 studies with 2051 control subjects.

Relationship between serum lipids and premenopausal breast cancer risk

Figure 2 shows the relationships between the four serum lipid components and the risk of premenopausal breast cancer from our meta-analyses. The SMD of TG was 12.90 (95% CI: 7.19–18.61) (Figure 2a), indicating that TG levels of premenopausal breast cancer patients were significantly higher than those of healthy premenopausal women. Similarly, a significant increase in LDL-C (SMD 31.43, 95% CI: 8.72-54.15) was detected in the case group compared with the control group (Figure 2b). Although premenopausal women with breast cancer tended to have lower TC and higher HDL-C than those without breast cancer, no significant differences were found in TC (SMD 31.10, 95% CI: -9.98-70.18) or HDL-C (SMD -1.39, 95% CI: -2.90-0.12) between the two groups (Figure 2c and 2d).

According to demographic characteristics described in the 13 included articles, four studies included an African population,^{28,29,31,32} eight included an Asian population,^{21–27,30} and one had a European population.³³ Subgroup analyses were performed based on the African and Asian populations (Table 2). In the African population, only TG levels of premenopausal

Author Number of suby Number of cases controls Exposure TG TC IDL-C HDL-C Yang LN (2020) china case-control 9 9 TG, TC, LUC 12050±5050 190.21±33.25 102.45±41.37 56.44±12.76 Jing GM (2019) China case-control 38 37 TG, TC, LUC 183.17 ±46.07 247.04 ±23.58 111.34 ±11.21 62.63 ±10.05 Hang Z (2019) China case-control 39 76, TC, LDLC NR 1736.61 ±36.34 100.90 ±17.78 39.05 ±11.21 Adalanka HMK Sri case-control 39 76, TC, DDLC NR 1736.61 ±36.34 100.90 ±17.78 39.05 ±11.21 Adalanka HMK Sri case-control 39 76, TC, DDLC NR 1736.61 ±36.34 103.90 ±17.78 39.05 ±11.21 Adalanka HMK Sri case-control 50 76, TC, DDLC NR 1736.61 ±36.34 103.90 ±17.78 39.05 ±11.21 Adalanka HMK Sri case-control 50 76, TC, DDLC NR 20.56 ±40.48	Case group mean \pm SD (mg/dL)		Control group mean \pm SD (mg/dL)	ean \pm SD (mg/dL			
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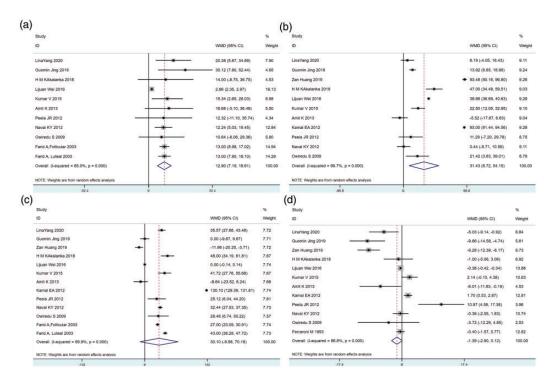


Figure 2. Forest plot depicting the standardized mean difference (SMD) scores and 95% confidence intervals for studies comparing the serum lipid levels of premenopausal breast cancer patients with those of healthy women: (a) triglycerides (TG), (b) low-density lipoprotein cholesterol (LDL-C), (c) total cholesterol (TC), and (d) high-density lipoprotein cholesterol (HDL-C).

Subgroup	Number of studies	Number of patients			Heterogeneity	
		Case	Control	SMD (95% CI)	l ² (%)	Р
TG						
African	3	128	91	12.92 (9.84–16.01)	0.00	0.996
Asian	7	787	1683	13.54 (5.57–21.51)	76.0	<0.001
LDL-C						
African	3	165	126	42.40 (-18.89-103.69)	98.5	<0.001
Asian	8	977	1787	27.26 (2.49–52.03)	99.4	<0.001
тс						
African	4	225	156	50.98 (-7.08-109.04)	99.9	<0.001
Asian	8	977	1787	16.94 (2.00–31.87)	97.9	<0.001
HDL-C				× ,		
African	3	165	126	3.22 (-3.47-9.91)	78.8	0.009
Asian	8	977	1787	-3.18 (-5.640.72)	88	<0.001

Table 2. Subgroup analysis based on different ethnic groups.

TG, triglycerides; TC, total cholesterol; LDL-C, low-density lipoprotein cholesterol; HDL-C, high-density lipoprotein cholesterol; SMD, standardized mean difference; CI, confidence interval.

breast cancer patients were significantly higher than in the healthy premenopausal population. However, in the Asian population, TG, TC and LDL-C levels were all significantly higher in the case group than in the control group, while HDL-C levels showed the opposite trend.

Percentage of heterogeneity and publication bias

The studies selected in this analysis were performed in different countries and with different populations. The I^2 values of all four blood lipid components were >75% (TG: 85.0%, LDL-C: 99.9%, TC: 99.9%, and HDL-C: 86.8%) (Figure 2). Thus, the studies included in this meta-analysis were highly heterogeneous, and random effect models were used to estimate summary hazard ratios (HRs) and 95% CIs. Considering the small study effect, publication bias was tested using Egger's test. No evidence of publication bias was found in the TC (P = 0.189), LDL-C (p = 0.107), or HDL-C (p = 0.575) analyses, whereas evidence of publication bias was found in the TG (p = 0.001) analysis.

Discussion

We performed this meta-analysis to determine whether serum lipids are a risk factor for premenopausal breast cancer. The results support the hypothesis that a significant positive association exists between dyslipidemia (high levels of TG and LDL-C) and premenopausal breast cancer risk. In Asian populations, all four indicators of dyslipidemia (high levels of TG, LDL-C, TC, and low levels of HDL-C) were considered risk factors for premenopausal breast cancer. Most of the studies included in this meta-analysis included Asian or African populations, probably because the peak age of breast cancer varies in different regions. Epidemiological evidence has shown that the peak age of breast cancer in Asia and Africa is earlier than in western countries.^{34,35} For example, in Asian countries, the peak age of breast cancer ranges from 40 to 50 years old, whereas in western countries, the peak age ranges from 60 to 70 years old.³⁶

Compared with the other lipid components, relatively few studies have been conducted on the relationship between breast cancer risk and TG. Some studies have indirectly reflected the relationship between TG and breast cancer risk by confirming that a high-fat diet can increase breast cancer risk.^{37,38} As early as the 1990s, Goodwin et al.³⁹ found that elevated triglyceride levels were an independent risk factor for premenopausal breast cancer. Han et al.⁴⁰ also reported that elevated serum TG levels might be correlated with an increased risk of breast cancer. Four studies included in our meta-analysis reached the same conclusion.^{22,26,30,31} A statistically significant difference was found in the TG concentrations between premenopausal breast cancer patients and healthy women, with patients having higher TG levels; this was consistent in African and Asian populations.

Cholesterol plays an important role in the etiology of cardiovascular disease. High levels of TC and LDL and low levels of HDL are adverse factors that lead to cardiovascular disease.41,42 Changes in these three lipids are also risk factors for breast cancer. However, a review of related studies still cannot provide a clear picture because the results are inconsistent. A large prospective study showed a correlation between serum TC and breast cancer. Women with TC levels > 240 mg/dL reportedly had a higher risk of developing breast cancer than those with TC levels <160 mg/dL.⁴³ In contrast, another large prospective study that included 288,057 female participants suggested that low serum TC was a favorable factor for the prevention of breast cancer.44 A study of 38,823 Norwegian women found a 25% increased risk of breast cancer in the low HDL-C group compared with in the high HDL-C group.⁴⁵ A meta-analysis conducted by Melvin et al.⁴⁶ showed that increasing TC and decreasing HDL were associated with an increased cancer risk (including breast cancer) of 18% and 15%, respectively.

In 2015, two meta-analyses arrived at different conclusions regarding the relationship between these three lipids and breast cancer risk. One observed a modest inverse association between TC and HDL-C and the risk of breast cancer after considering studies that eliminated preclinical bias, and no association was found between LDL-C and breast cancer risk.47 The other meta-analysis suggested that serum HDL-C may protect against breast carcinogenesis among postmenopausal women but not premenopausal women, whereas serum TC and LDL-C levels may be not associated with breast cancer risk.⁴⁸ Unfortunately, most of these articles did not separately discuss the premenopausal and postmenopausal populations. In this meta-analysis, elevated LDL-C was found to be an adverse factor for premenopausal women, but higher TC and lower HDL levels were not related to the risk of premenopausal breast cancer. Notably, our subgroup analyses confirmed that TC and LDL-C were modest but statistically significant unfavorable factors, whereas HDL-C was a favorable factor for premenopausal breast cancer patients in the Asian population. No significant relationship was found between TC, LDL-C, or HDL-C and premenopausal breast cancer risk in the African population.

Our meta-analysis had several limitations. First, all studies included were of high quality according to the NOS assessment; however, the outcomes in the four components of serum lipids were moderate either because of high heterogeneity or the limited number of case-control trials

available for analysis. Additionally, the populations studied were mostly from Asia, followed by Africa. Data from European and American countries were lacking. Second, because studies with positive results tended to be published, especially case-control studies, publication bias might be a concern. In particular, when the number of articles included in the meta-analysis was small, tests for publication bias tended to lack statistical power. Finally, some relevant articles might not have been included in the analysis due to the different ways data are evaluated in various articles. For example, studies were excluded if the data did not report the mean \pm SD or failed to separately list the premenopausal and postmenopausal women.

The etiology of breast cancer is multifactorial. Thus, an increase or decrease in isolated cholesterol may not be the decisive factor for carcinogenesis. The results of our meta-analysis should be interpreted with caution, because the number of studies was extremely limited. Many studies have focused on postmenopausal women with breast cancer. We hope to find more large-scale prospective clinical studies that are focused on breast cancer in premenopausal women in the future.

Conclusion

To our knowledge, no previous metaanalysis has specifically evaluated the role of dyslipidemia in assessing breast cancer risk in premenopausal women. TG and LDL-C levels were higher in premenopausal breast cancer patients than in healthy premenopausal women. High serum TG and LDL-C levels may be associated with increased premenopausal breast cancer risk. Among Asian women, serum HDL-C may play a protective role, whereas TG, TC, and LDL-C may promote carcinogenesis in premenopausal breast.

Declaration of conflicting interest

The authors declare that there is no conflict of interest.

Funding

The authors disclosed receipt of the following financial support for the research, authorship, and publication of this article: This work was financially supported by the Chongqing Municipal Education Commission Science and Technology Research Project (KJQN201 800109), Chongqing Science and Health Joint Medical Research Project (2021MSXM222), National Cancer Center Climbing Fund (NCC201922B01), Central University's basic scientific research business fee medical integration project (2019CDYGYB004), and Chongqing Science and Technology Bureau Project (cstc2018jxjl130044).

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