scientific reports

OPEN

Metabolic syndrome clustering patterns and the association with cardiovascular disease among post‑menopausal Korean women

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The metabolic syndrome (MetS) is a group of diseases conceptualized as a clustering of risk factors, with the risks of developing MetS in women varying signifcantly before and after menopause. This study investigated MetS clustering patterns and their association with cardiovascular disease (CVD) risk among post-menopausal women (n= 2479) using data from the Korean Genome Epidemiology Study. Using latent class analysis, three groups emerged: diabetic (5.6%), hypertensive (35.2%), and low-risk (59.2%). Relative to the low-risk group, the diabetic group demonstrated associations with older age, a family history of chronic disease, an increased Homeostatic Model Assessment for Insulin Resistance (HOMA-IR), an elevated white blood cell (WBC) count, experience with hysterectomy, being a non-drinker, more physical activity, and excessive sleep. The hypertensive group was associated with older age, lower monthly income, a family history of chronic disease, increased HOMA-IR, a higher WBC count, more physical activity, and excessive sleep. The diabetic and hypertensive groups had a signifcantly higher CVD risk than the low-risk group (diabetic: odds ratio [OR] = 2.41 [1.11, 5.27]; hypertensive: OR= 2.46 [1.33, 4.55]). This study identifed potential markers for MetS screening in post-menopausal women, highlighting the need for early intervention and personalized healthcare for middle-aged women to reduce CVD risk following menopause.

Keywords Metabolic syndrome, Latent class analysis, National Cholesterol Education Program Adult Treatment Panel III, Post-menopause, Cardiovascular disease

Abbreviations

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Metabolic syndrome (MetS) is diagnosed when an individual exhibits a cluster of conditions including hyperglycemia, abdominal obesity, dyslipidemia, and hypertension. Reaven frst coined the term 'Syndrome X' to describe this phenomenon, with a common cause among afected patients being a decline in beta cell function due to insulin resistance¹. The primary goal of preventing and treating MetS is to mitigate the morbidity and mortality associated with these conditions². The prevalence of MetS in the Republic of Korea has been rising steadily over time, affecting both men and women ^{[3](#page-9-2)}. Consequently, medical expenses and national economic losses due to MetS are expected to increase, highlighting the urgent need for its management 4 .

Although MetS is a predictor of various diseases, its pathophysiology remains elusive⁵. Several studies have utilized factor analysis to examine the structural connections between the components of MetS and identify common underlying causes^{6–15}. The literature review findings suggest that between two and four factors can explain the relationships between the components of MetS, indicating that there is no single pathophysiological pathway for its development. However, factor analysis has limitations such as the arbitrariness of factor extraction criteria and the requirement for continuous variables to follow a normal distribution^{[16](#page-10-1)}. Furthermore, it is essential to consider the normal range for each MetS component when applying these results to clinical practice. Latent class analysis (LCA) is a person-centered statistical method that examines non-continuous and categorical variables to classify individuals into distinct groups and create personalized interventions based on their response patterns^{[17](#page-10-2)}.

MetS is acknowledged as a lifestyle disease shaped by factors such as socioeconomic status and health behaviors, with its effects differing according to gender and menopausal status¹⁸. Middle-aged women, particularly those who are post-menopausal, face an increased risk of MetS owing to a decline in female sex hormones, lead-ing to heightened vulnerability throughout a considerable portion of their lives^{19,20}. Studies utilizing LCA^{[16](#page-10-1),[21](#page-10-6)-[24](#page-10-7)} have typically identifed between two and four groups of clustering patterns of MetS components. However, they did not account for the unique characteristics of post-menopausal women, who are particularly vulnerable to MetS owing to hormonal changes unlike women of other age groups. As the risk of developing MetS increases considerably after middle age, particularly in women owing to factors such as menopause^{19,25}, it is essential to closely monitor MetS occurrence and patterns in post-menopausal women.

MetS is associated with an increased risk of cardiovascular disease (CVD), particularly in women^{[26,](#page-10-9)27}. However, research on MetS has primarily focused on the presence or total number of MetS risk factors, which has not led to a profound understanding of the heterogeneous nature of the condition's development^{[28](#page-10-11)}. Consequently, comprehensive studies are necessary to understand the multifaceted nature of MetS and its intricate relationship with CVD. Given the elevated risk faced by post-menopausal women with MetS, this study aimed to (1) identify the clustering patterns of MetS components among women aged 40 to 69 who were post-menopausal at the time of the baseline survey conducted from 2001 to 2002; (2) pinpoint the characteristics of participants associated with MetS class membership; and (3) explore how the risk of CVD varies across MetS classes.

Methods

We performed a secondary analysis using the dataset from the Korean Genome Epidemiology Study (KoGES), an ongoing population-based cohort study of adults aged 40 to 69 living in Ansan and Anseong provinces in 2000.

Data source

We employed a secondary analysis using KoGES, which comprehensively represents the characteristics of the South Korean population. Initiated by the Korea Centers for Disease Control and Prevention Agency in 2001, KoGES aimed to create a framework and guidelines for monitoring and addressing risk factors for the development of non-communicable diseases in South Koreans. It commenced with a baseline survey from 2001 to 2002, which was subsequently administered every two years for eight waves. Focusing on non-communicable diseases, such as diabetes, hypertension, and CVD, KoGES encompasses a wide array of data, including physical examinations, laboratory analyses (e.g. blood and urine tests), and interviews detailing lifestyle habits and disease history. Given its comprehensive coverage of noncommunicable diseases that are closely linked to MetS and the depth of data pertaining to physiological measurements and lifestyle factors, KoGES is well-suited for fulflling the present study's objectives. The dataset is accessible to approved researchers, and detailed information on the cohort profile has been presented elsewhere^{[29](#page-10-12)}.

Participants and sample size

The initial sample consisted of 10,030 adults, of whom 5272 were women aged 40 to 69, living in both rural and urban areas in central South Korea. Written informed consent was obtained from all participants. All methods used in this study were performed by the guidelines and regulations of the Declaration of Helsinki. Tis study utilized a baseline survey conducted from 2001 to 2002, which had the least missing data and was currently

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available for research. Tis study was approved by the Institutional Review Board of Yonsei University Medical Center (IRB #4-2021-0373).

For this analysis, we included women aged 40 to 69 who were not menstruating at the time of the baseline survey. Among 5272 women aged 40 to 69, we classifed 3299 as menopausal as they answered 'No' to the question 'Are you currently menstruating?' Tus, we excluded participants who had been previously diagnosed with CVD at baseline $(n=63)$, reported current use of female sex hormones at baseline $(n=388)$, or were unresponsive to the questionnaire (n=369) from the analysis. Supplementary Table 1 presents the baseline characteristics of the excluded and included participants. Finally, we included 2479 samples in the analysis.

Measures

Measurement and defnition of MetS

MetS components were diagnosed using the defnition based on the criteria from the 2004 revised version of the National Cholesterol Education Program Adult Treatment Panel III, including the waist circumference (WC) cut-of point suggested by the Korean Society of Obesity to account for the typical features of South Korea's population. The cut-off points for MetS components were as follows: WC≥85 cm for women, diastolic/ systolic blood pressure (DBP/SBP)≥85/130 mmHg, high-density lipoprotein cholesterol (HDL-C)<50 mg/dL in women, triglycerides (TG)≥150 mg/dL, and fasting blood glucose (FBG)≥100 mg/dL. We considered the medication status for diabetes, hypertension, or hyperlipidemia to be a separate MetS component. Using these eight components, we statistically derived MetS clustering patterns using LCA.

Characteristics considered for LCA

We considered general, biological, fertility-related, and health-related behavioral characteristics to be correlates of MetS latent class membership.

General characteristics included age, marital status, family history, monthly income, profession, and education level. Marital status was a binary variable set based on whether the respondent was married or single. Family history was a binary variable based on whether the participant's parents or siblings had been diagnosed with hypertension, diabetes, or hyperlipidemia. We categorized monthly income as less or more than 2 million won. We classifed homemakers as unemployed, factory workers as manual laborers, farmers as agricultural workers, and people such as office workers, the self-employed, salespeople, and professionals as non-manual laborers. We classifed education level as their fnal level of schooling (high school and below or college and above).

Biological characteristics included an increased Homeostatic Model Assessment for Insulin Resistance (HOMA-IR), white blood cell (WBC) count, and C-reactive protein (CRP). HOMA-IR is a reliable marker for MetS as it can be calculated from the FBG and insulin blood concentrations, as proposed by Matthews et al.³⁰. We assessed CRP levels through a blood test and categorized them as normal if they were below 1 mg/dL and moderately increased if they were 1 mg/dL or above following the standard proposed by Nehring et al[.31.](#page-10-14) We measured WBC count as a continuous variable, determined using a blood test.

Fertility-related characteristics included age at menarche, gravidity, age at frst delivery, and history of hysterectomy. In the Republic of Korea, with an average age at menarche of 12.4 years^{[32](#page-10-15),[33](#page-10-16)}, it was categorized as <13 years old or 13 years and older. Previous studies^{[34](#page-10-17)–39} that demonstrated a significant relationship between gravidity and MetS used diferent methods to process the gravidity variable. To better refect the distribution of gravidity in our study population (median 5, mean 5.39), we categorized them into three groups: no births, one to fve births, and six or more births. A birth was defned as either a live birth or a stillbirth, excluding miscarriages. The age at first delivery, sourced from open-ended baseline survey responses, denotes the age at which a participant frst gave birth; we coded it as 0 if they had never been pregnant. We assessed experience with hysterectomy using a 'yes' or 'no' answer to the question, 'Have you ever had surgery to remove your uterus (i.e. hysterectomy)?.

Health behavioral characteristics included smoking, alcohol consumption, physical activity, unrecommended dietary habits, stress, and sleep duration. We categorized smoking status as a non-smoker ('Never smoked'), former smoker ('Used to smoke but do not currently smoke'), and current smoker ('Occasionally smoke' or 'Habitually smoke'). Based on a meta-analysis study 40 , which found that the prevalence of the metabolic syndrome is lowest in women consuming less than 20 g of alcohol per day, we classifed individuals who consume less than 20 g of alcohol per day as 'moderate drinkers' and those who consume 20 g or more per day as 'heavy drinkers'. We quantifed physical activity using the following:

the metabolic equivalent of task value of activity \times the number of physical activities per week

 \times the minutes of exercise per day \times 3.5 \times body weight (in kg)/200.

We assessed unrecommended dietary habits based on three criteria: (1) consuming more than 130 g of carbohydrates, (2) consuming more than 2000 mg of sodium, and (3) consuming less than 20 g of fber per day, each earning one point⁴¹. We measured stress using a 30-question self-assessment tool developed by the Ministry of Health and Welfare, with scores ranging from 0 to 30 and higher scores signifying greater stress 42 . We assessed sleep duration based on self-reported average daily sleep over the past year, including naps, and categorized it as adequate (6–8 h), insufficient (<6 h), or excessive (>8 h) based on National Sleep Foundation's guidelines^{[43](#page-10-22)} and research findings⁴³⁻⁴⁷.

Risk of CVD

For this study, we defned CVD as including myocardial infarction, congestive heart failure, coronary artery disease, peripheral vascular disease, and cerebrovascular disease (stroke, hemorrhage, etc.); CVD occurs in the heart and cerebral blood vessels. We considered CVD to have occurred if a participant responded 'No' to the question 'Have you ever been diagnosed with CVD by a doctor in the past?' in the 2001–2002 baseline survey and also responded 'Yes' at least once during the eight follow-ups to the question 'Have you been diagnosed with CVD since the last survey [in the past 2 years]?'.

Statistical analysis

LCA is a statistical method used to estimate unobserved heterogeneity within a population; it classifes participants with similar response patterns into groups and identifies their characteristics and differences⁴⁸. In this study, we employed the bias-adjusted 3-step approach proposed by Asparouhov and Muthén⁴⁹. Previously, researchers included covariates directly in the same model to identify the class solution, a method known as the one-step approach, but this often led to flawed models^{[50,](#page-11-0)51}. The one-step approach has been replaced with newer methodologies, with the three-step approach now widely recommended^{49,52}. This approach involves first identifying the measurement model, adding covariates separately, and fxing measurement parameters to those from the initial model to achieve more unbiased results^{51,53}. This model consists of the following sequential steps.

Step 1: We determined the optimal number of latent classes by considering various indices, including information criteria, quality of classifcation, model ft comparisons, latent class proportions, and their theoretical/ clinical meaningfulness. We explored the number of latent classes, ranging from two to fve; covariates were not included in this step. We used the Akaike information criterion (AIC) , the Bayesian information criterion (BIC), and the sample size-adjusted Bayesian information criterion (SABIC) to assess model ft. We evaluated classifcation quality using entropy, which ranged from 0 to 1, with higher values denoting lower classifcation errors[54.](#page-11-4) Additionally, we employed the Lo-Mendell-Rubin-adjusted likelihood ratio test (LMR LRT) and the bootstrap-adjusted likelihood ratio test (BLRT) to determine whether a model with *k* latent classes performed better than a model with *k* − 1 latent classe[s55.](#page-11-5) Best practices advise careful consideration of each of these indices when selecting the best-ft model.

Step 2: Given the latent class distribution identifed in Step 1 and the response patterns of each individual, we estimated the posterior probabilities of membership in specific latent classes using Bayes' rule⁴⁹. This approach calibrates the classifcation uncertainty rate, which is essential when examining covariates to predict latent class membership in Step 3. We assigned individuals to the class classifed in Step 1 to which they had the highest estimated probability of belonging.

Step 3: We performed a multinomial logistic regression analysis by incorporating the covariates and correcting the uncertainty classifcation errors that occurred in Step 2. We performed this step to identify the characteristics that determine class membership. Multinomial regression involves confguring binomial logistic regression models, which predict the probability of an individual belonging to each category relative to the reference group[56](#page-11-6).

Step 4: We conducted a discrete-time survival analysis to explore the diferences in the hazard probability of a CVD event among classes, which is the conditional probability of an event occurring at a particular time for a specific individual⁵⁷.

We placed more than half of the participants (59.2%) in the low-risk group; they exhibited the lowest risk of MetS. Thus, we set the low-risk group as the reference group. The missing data rate in this study was 1.1%, with the FBG levels and HOMA-IR having the most missing data (5.1%). We used full-information maximum likelihood estimation to minimize the bias caused by missing values⁵⁸. We used R 4.2.0 and Mplus 7.3 to carry out the analyses mentioned above.

Results

Number of latent clusters

Table [1](#page-3-0) lists the statistical criteria used to evaluate model ft, which yielded conficting outcomes. Comparing the criteria as the number of latent classes increased from 1 to 5, the AIC was lowest when there were 5 latent

Table 1. Comparison of diferent latent classes based on model selection statistics. *AIC* Akaike information criterion, *BIC* Bayesian information criterion, *SABIC* sample-adjusted Bayesian information criterion, *BLRT* bootstrap adjusted likelihood ratio test, *LMR-LRT* Lo-Mendell-Rubin adjusted likelihood ratio test. **p*<0.05, ***p*<0.01, ****p*<0.001.

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Fig. 1. Information criterion based on the number of latent classes. *AIC* Akaike information criterion, *BIC* Bayesian information criterion, *SABIC* sample size-adjusted BIC.

classes, while the BIC and SABIC were lowest when there were 3 latent classes (Fig. [1](#page-4-0)). Furthermore, the frst insignifcant LMR LRT and BLRT appears for 4 and 5 class respectively, indicating that no more than 4 latent classes are required for the entire sample. The 4-class model also had the highest entropy value, which is ideal when close to 1. Researchers are encouraged to examine and report entropy, but not to solely rely on the value for determining the final class solution^{[52](#page-11-2)}. While there is no definitive consensus on the best criteria for determining the optimal number of latent clusters, it is accepted that: (a) the BIC is one of the most reliable measures and should be included^{50,52}, and (b) the final model should be selected based on theoretical interpretability. We used substantive reasons to guide our decision. First, the BIC was the lowest for this model, indicating a better ft and parsimony^{49,[50,](#page-11-0)52}. Second, in similar prior studies^{16[,22](#page-10-26)}, the configuration of MetS components by classes was most similar when using three classes. Terefore, the three-class model was deemed the best ft for this analysis.

Distribution of MetS components among the participants

The participants exhibited a mean WC of 84.41 ± 9.39 cm and a mean FBG level of 85.70 ± 18.08 mg/dL. SBP averaged 126.61 ± 19.84 mmHg, while DBP averaged 81.66 ± 11.59 mmHg. The HDL-C levels had a mean of 45.04 ± 10.07 mg/dL, and the TG levels had a mean of 162.00 ± 95.72 mg/dL. Regarding medication use, 5.8%

Table 2. Distribution of metabolic syndrome components among the participants (N=2479). *DBP* diastolic blood pressure, *FBG* fasting blood glucose, *HDL-C* high-density lipoprotein cholesterol, *M* mean, *SBP* systolic blood pressure, *SD* standard deviation, *TG* triglycerides, *WC* waist circumference.

Table 3. Response probability of metabolic syndrome components for each latent class. *BP* blood pressure, *FBG* fasting blood glucose, *HDL-C* high-density lipoprotein cholesterol, *TG* triglycerides, *WC* waist circumference. Values in bold indicate response probabilities higher than 0.5.

Fig. 2. Response probability of metabolic syndrome components in each latent class.

(143) of participants were taking anti-diabetic medications, 19.2% (476) were using anti-hypertensive medications, and 0.5% (12) were taking lipid-lowering medications (Table [2](#page-4-1)).

Probability of MetS components in each latent class

Table [3](#page-5-0) and Fig. [2](#page-5-1) present the distribution and characteristics of each class based on the conditional probabilities for each component. The first class (diabetic) comprised 5.6% of participants and was the most likely to take anti-diabetic medication (100.0%), followed by a decreasing likelihood of low HDL-C levels (84.9%), high FBG (84.6%), high TG (77.6%), high WC (61.6%), and high BP (54.5%). The second class (hypertension) comprised 35.2% of participants. Individuals in this class were most likely to have high BP (83%), followed by a decreased likelihood of having low HDL-C (80.7%), high TG (71%), and high WC (62.8%). The largest class (low-risk)

Table 4. Characteristics associated with metabolic syndrome class membership (ref. low-risk group). *CRP* C-reactive protein, *HOMA-IR* homeostatic model assessment for insulin resistance, *WBC* white blood cell, *RRR* relative risk ratio.

Table 5. Comparing the risk of cardiovascular disease by metabolic syndrome latent classes. *CI* confdence interval, *OR* odds ratio. Class 1=diabetic group, Class 2=hypertensive group, Class 3=low-risk group. *Reference group.

Fig. 3. Odds ratio for cardiovascular disease by metabolic syndrome latent class.

included 59.2% of participants who were most likely to have low HDL-C levels (65.9%), followed by a declining likelihood of having high TG levels (61.3%).

Characteristics associated with MetS class membership

Table [4](#page-6-0) shows the results of the conditional model, incorporating covariates and employing a 3-step bias correction method. Setting the low-risk group as the reference, older age (Relative risk ratio [RRR]=1.12, *p*<0.001), a family history of chronic disease (RRR=4.01, *p*<0.001), a higher HOMA-IR (RRR=4.76, *p*<0.001), a higher WBC count (B=1.19, $p=0.007$), experience with hysterectomy (RRR = 2.59, $p=0.003$), being a non-drinker (moderate: RRR = 0.48, $p=0.041$; heavy: RRR=0.06, *p*<0.001), increased physical activity (RRR=1.01, *p*=0.006), and excessive sleep (RRR=1.89, *p*=0.009) led to greater odds of being in the diabetic group than the low-risk group.

Those who were older (RRR=1.14, *p* < 0.001) had an average monthly income of less than 2 million won (RRR=0.46, *p*=0.017), a family history of chronic disease (RRR=3.49, *p*<0.001), a higher HOMA-IR (RRR=4.37, *p*<0.001), a higher WBC count (RRR=1.22, *p*=0.003), increased physical activity (RRR=1.01, *p*=0.006), and excessive sleep (RRR=1.59, $p=0.043$) were more likely to be in the hypertensive group than the low-risk group.

Risk of CVD by MetS class

Of the 2479 participants,121 participants (4.9%) developed CVD. We calculated the probability of developing CVD based on the MetS classes while accounting for classifcation errors. Table [5](#page-6-1) displays the risk of CVD according to the MetS class. CVD was signifcantly more likely to occur in the diabetic group (odds ratio [OR]=2.46, $p = 0.027$) and hypertensive group (OR=2.41, $p = 0.004$) than in the low-risk group. However, there was no significant difference in the risk of CVD between the diabetic and hypertensive groups (OR = 0.98, *p* = 0.959) (Fig. [3](#page-7-0)).

Discussion

We identifed the clustering patterns of MetS components in post-menopausal women in South Korea and explored the characteristics associated with MetS class membership. Moreover, we examined the hazard probability of CVD according to three MetS classes. Our fndings highlight the need for intervention programs targeting MetS in post-menopausal women and underscore the importance of CVD prevention initiatives.

We identified three MetS classes: diabetes (5.6%), hypertension (35.2%), and low-risk (59.2%). Studies^{[16](#page-10-1),[18](#page-10-3)-[21](#page-10-6)} that identifed similar MetS clustering patterns reported varying numbers of classes, ranging from two to four. These discrepancies likely stem from differences in the study populations, MetS definitions, and the factors considered. Although our study provides valuable insights into efective strategies for preventing MetS in postmenopausal women, further research is required to validate whether the characteristics of the derived classes hold in diverse populations.

Unlike previous studies, we did not identify a latent class of healthy participants who did not meet the MetS criteria. Among the 1524 participants in the low-risk group, only 100 had no components that ft the MetS diagnostic criteria. Te diferentiation observed may be linked to menopause-related hormonal shifs, such as reduced estrogen and heightened progesterone, which lead to changes in fat distribution, muscle loss, and a rise in metabolic risk factors⁵⁹

The diabetic group, comprising only 5.6% of participants, is of particular note because every individual in this group was actively prescribed anti-diabetic medications. Ahanchi, et al.²² also recognized a congruent clustering pattern based on the use of anti-diabetic medications. The diabetic group showed clear metabolic changes associated with insulin resistance, which leads to conditions such as hyperglycemia and diabetes, concurrently resulting in suboptimal absorption of free fatty acids and consequently fostering increased TG production within hepatocytes^{[5,](#page-9-4)60}. Furthermore, the accumulation of surplus TG impairs lipoprotein lipase activity, hindering the hydrolysis of TG and impeding the production of active $HDL-C⁵$. The emergence of hyperinsulinemia, an adaptive response to insulin resistance, contributes to the development of hypertension by intensifying salt retention and elevating cytosolic Ca²⁺ levels in the kidneys^{[9](#page-9-6)}. Consequently, the diabetic group served as a clear representation of the metabolic changes associated with insulin resistance.

The hypertensive group (comprising 35.2% of participants) exhibited a higher prevalence of hypertension, which previous studies identified as an independent factor for MetS²⁰. Menopause—marked by decreased estrogen levels and increased arterial stifness, sodium sensitivity, and fat accumulation—leads to higher blood pressure in middle-aged women 61 . Women should be made aware of the risk of developing post-menopausal hyperglycemia or hypertension and encouraged to make lifestyle changes well before menopause.

The low-risk group, representing 59.2% of participants, exhibited notably elevated occurrences of low HDL-C $(65.9%)$ and high TG $(61.3%)$. These observations are associated with decreased estrogen levels during menopause, leading to unfavourable changes in HDL-C subtypes, specifcally a decrease in HDL-2 and an increase in HDL-3, along with elevated TG levels. Furthermore, a substantial percentage of individuals with diabetes and hypertension displayed coexisting dyslipidemia, which is consistent with the high prevalence of low HDL-C and high TG levels across all groups. Post-menopausal women are especially susceptible to dyslipidemia due to hormonal changes. However, awareness and treatment rates for dyslipidemia lag behind those for other chronic illnesses. It is imperative to emphasize early dyslipidemia prevention in middle-aged women through communitybased health services and regular screenings to prevent CVD. Implementing a mandatory lipid screening system or providing accessible lipid testing devices for middle-aged women could improve the awareness, treatment, and control of dyslipidemia.

Among general characteristics, age, monthly income, and family history distinguished the low-risk group from the diabetic and hypertensive groups. Biological aging lowers glucose tolerance and insulin secretion, which can lead to atherosclerosis; however, this can be mitigated by addressing obesity and promoting a healthy lifestyle⁶². It is also essential to proactively identify patients with a family history of chronic disease and consider those with such a history as potential targets for MetS management interventions—even if they are currently healthy. Monthly income distinguished low-risk individuals from the hypertensive group, which is consistent with studies that demonstrated an inverse linear relationship between household income and the prevalence of hypertension^{63,64}. These findings imply that there may be socioeconomic disparities in access to primary prevention interventions among middle-aged women.

Among the biological characteristics, HOMA-IR and WBC counts were correlated with MetS classes, indicating a link between insulin resistance, inflammation, and MetS. Oda and Kawai⁶⁵ suggested that cytokine secretion from mast cells mediates systemic infammatory responses and contributes to MetS development by stimulating the synthesis of infammatory markers. Given that HOMA-IR and WBC count are fairly easy to measure, these markers could serve as cost-efective indicators for early MetS diagnosis.

Sleep duration, a health-related behavioral characteristic, distinguished the low-risk group from the other two groups. Studies have shown varying efects of sleep duration on MetS, particularly in women, although the fndings regarding the relationship between sleep duration and MetS have been inconsistent^{45,[46](#page-10-28),66}. Prolonged sleep can have adverse health outcomes including sleep fragmentation, increased fatigue due to poor sleep quality, reduced exposure to sunlight, and physical inactivity. Women are more susceptible to sleep disruption than men owing to fuctuations in estrogen and estradiol levels during menopause, which are ofen accompanied by hypersomnia. Incorporating sleep duration assessments into MetS management could help identify individuals at high risk of developing MetS and determine the need for sleep-related interventions.

Among fertility-related characteristics, experience with hysterectomy diferentiated the low-risk and diabetic groups. A study⁶⁷ showed that women with a history of hysterectomy had higher abdominal fat tissue volumes and weights than those without such a history. Including fertility-related questions in women's health questionnaires could help identify individuals at risk of diabetes and guide preventive interventions. Compared to men, women experience more severe outcomes of MetS³⁹, and post-menopausal women show an increase in prevalence owing to hormonal changes and pregnancy experiences⁶¹. Specifically, fertility-related characteristics such as age of menarche, gravidity, age of first delivery, and hysterectomy status have been reported as risk factors for MetS in previous studies [34–](#page-10-17)[38](#page-10-29)[,67,](#page-11-17)[68](#page-11-18). In this study, only hysterectomy status infuenced MetS patterns, aligning with fndings that other fertility-related characteristics showed no significant correlation with MetS after adjusting for confounders like health behaviors^{[39](#page-10-18)}.

Alcohol consumption was used to distinguish between the low-risk and diabetic groups, with heavy drinkers being more likely to belong to the low-risk group. While a meta-analysis⁴⁰ indicated a lower prevalence of MetS among drinkers, especially those with moderate alcohol intake, we found that heavy drinkers had the lowest likelihood of being in the diabetic group.

Physical activity also served as a distinguishing factor between the low-risk and other groups; those who were physically active were more likely to be in the diabetic or hypertensive groups. Tis could be owing to reverse causation, as those at a higher risk of these conditions might engage more ofen in health management behaviors, leading to increased physical activity, which is commonly the initial recommended lifestyle change for managing diabetes and hypertension^{[68](#page-11-18)}.

We compared the rates at which individuals in the three MetS classes experienced cardiovascular events. The diabetic and hypertensive groups were considerably more likely to experience a cardiovascular event than the low-risk group; however, there was no statistically signifcant diference between the diabetic and hypertensive groups.

Riahi et al.[16](#page-10-1) identifed three distinct groups for both genders and found considerable diferences in CVD risk between the healthy and MetS groups but not between the healthy and low-risk groups. Tis comparison suggests that MetS risk factor combinations vary based on the characteristics of the population, implying that a MetS management manual should consider both the specifc traits of individuals and combinations of their risk factors.

We found that dyslipidemia, marked by low HDL-C and high TG levels, was common across all classes, denoting an increased CVD risk when coupled with other factors such as diabetes and hypertension. However, neither low HDL-C nor high TG levels alone increased this risk considerably, underlining the role of early dyslipidemia in reducing CVD risk²⁴. Additionally, as the prevalence of dyslipidemia is notably higher in post-menopausal women, who face a greater CVD risk from dyslipidemia than men²⁰, lifestyle interventions and frequent screenings for the prevention of concurrent manifestation of dyslipidemia, diabetes, and hypertension afer menopause could help lower the incidence of CVD.

It is important to recognize this study's strengths and limitations. Utilizing the most comprehensive cohort dataset in the Republic of Korea enhanced the representativeness of our results and allowed for the identifcation of MetS clustering patterns and related factors in an understudied population. However, the limitations of LCA should be acknowledged, including its challenge in detecting rare latent classes in small sample sizes and its data-driven approach. Moreover, relying on baseline survey data for MetS onset identifcation and related factors, while minimizing missing data, hinders the examination of pre- and post-menopausal transitions and changes in MetS classes over time. Caution is warranted when estimating temporal relationships and when carefully considering potential reverse causality. Additionally, there exist a potential recall or social desirability bias from using self-reported questionnaires. Future research should supplement these with objective measures like electronic medical records. Moreover as this study utilized secondary data, variables such as abortion and breastfeeding that were not included in the dataset could not be analyzed, making it difcult to control for residual confounding. It is crucial to consider these limitations when interpreting the results. Future research should address these constraints to deepen our understanding of the complex relationships between MetS, its components, and related health risks.

Conclusion

We utilized LCA to delineate MetS clustering patterns in post-menopausal women, categorizing them into diabetic, hypertensive, and low-risk groups based on a range of characteristics, including general, biological, fertilityrelated, and health-related behaviors. We employed a bias-adjusted 3-step approach to ascertain features linked to class membership. Tis study highlights the need for specialized MetS management programs in healthcare that cater to post-menopausal women. By comparing CVD risk across MetS classes and identifying high-risk groups, this study provides a framework for targeted screenings. These findings suggest that programs designed to reflect the characteristics and CVD risk factors in post-menopausal women can play a vital role in helping middleaged women recognize health issues and establish actionable goals to enhance their self-management abilities.

Data availability

The data supporting the findings of this study are available from the Korean Genome Epidemiology Study (KoGES). Restrictions apply to the availability of the data, which were used under a licence for this study. Any additional information required to reanalyze the data reported in this paper is available from KoGES epidemiological data online sharing system [\(https://nih.go.kr/ko/main/contents.do?menuNo=300566\)](https://nih.go.kr/ko/main/contents.do?menuNo=300566) with permission.

Received: 22 May 2024; Accepted: 6 September 2024 Published online: 30 September 2024

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Author contributions

GSK supervised the research process and developed the concept and design of the study. YJC acquired the participants and/or data, analysed the data, and prepared the original draft. GSK, YJC, SHC, KHL, CGP, and MSS reviewed and edited the manuscript. All authors read and approved the fnal version of the manuscript.

Competing interests

The authors declare no competing interests.

Additional information

Supplementary Information The online version contains supplementary material available at [https://doi.org/](https://doi.org/10.1038/s41598-024-72403-1) [10.1038/s41598-024-72403-1](https://doi.org/10.1038/s41598-024-72403-1).

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