Transition Patterns of Weight Status and Their Predictive Lipid Markers Among Chinese Adults: A Longitudinal Cohort Study Using the Multistate Markov Model

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Background: Obesity is well recognized as a risk factor for cardiometabolic diseases. The development of obesity is a dynamic process that can be described as a multistate process with an emphasis on transitions between weight states. However, it is still unclear what convenient biomarkers predict transitions between weight states. The aim of this study was to show the dynamic nature of weight status in adults stratified by age and sex and to explore blood markers of metabolic syndrome (MetS) that predict transitions between weight states.

Methods: This study involved 9795 individuals aged 18 to 56 at study entry who underwent at least two health check-ups in the eight-year period of study in the health check-up centre of our institution. Weight, height and biochemical indices were measured at each check-up. The participants were divided into four groups based on age and sex (young male, middle-aged male, young female and middle-aged female groups). A multistate Markov model containing 3 states (healthy weight, overweight and obesity) was adopted to study the longitudinal weight data.

Results: Young people were more likely to transit between weight states than middle-aged people, and middle-aged people were more resistant to recover from worse states. The mean sojourn time in obesity was greatest in the middle-aged male group (6.23 years), and the predicted rate of obesity beginning with healthy weight was greatest in the young male group (13.7%). In multivariate models, age group and triglyceride (TG) and high-density lipoprotein cholesterol (HDL) levels were significant for specific transitions in females, whereas age group and HDL levels were significant in males. In females, if HDL levels increased 1 mmol/L, the probability of progression from healthy weight to overweight decreased by 37.0% (HR= 0.63), and the probabilities of recovery (overweight to healthy weight and obesity to overweight) increased by 62.0% (HR= 1.62) and 1.23-fold (HR= 2.23), respectively. In males, if TG levels increased 1 mmol/L, the risk of progression from healthy weight to overweight increased by 24.0% (HR= 1.24). Each unit increase in HDL levels was associated with a 0.99-fold (HR= 1.99) increase in the chance of recovery from overweight to healthy weight and with a 0.37-fold (HR= 0.63) decrease in the risk of progression from healthy weight to overweight.

Conclusion: The weight status of young people was less stable than that of middle-aged people. Males were more likely to become overweight and more resistant to recover from worse states than females. Young males with healthy weight were more likely to develop obesity than other healthy weight groups. Blood lipid levels, especially HDL, were predictors of weight transitions in adults. Prevention and intervention measures should be applied early.

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Introduction

Overweight and obesity have been worldwide public health concerns. Obesity is responsible for many diseases, such as cardiovascular diseases (CVDs), stroke, and cancer. Obesity and obesity-related diseases have brought heavy social and economic burdens worldwide. With the rapid economic growth in China, Chinese people’s lifestyle has changed in many aspects, including diet, forms of transportation, physical activity and education, which have been accompanied by a rapid increase in the prevalence of overweight/obesity over the past two decades, especially in urban areas.

Many previous studies have focused on the prevalence of overweight/obesity at a certain time or described trends in prevalence over a period. However, weight change is a dynamic process in which individuals may experience successive weight states over a period, involving healthy weight, overweight and obesity. Therefore, the progression of obesity should be defined as a multistate process with a focus on transitions between weight states, which cannot be reflected by the prevalence of overweight/obesity alone. The flow rate of a population from one weight state to another and the mean sojourn time of individuals at each weight state are essential to understand the nature of the progression of overweight/obesity. Moreover, previous studies have shown that males and females have different metabolic characteristics, and the incidence rates of metabolic diseases associated with obesity differ across different age and sex groups. Therefore, it is important to explore the features of transitions between weight states by age and sex to develop targeted prevention and intervention strategies. To the best of our knowledge, there have not been many studies on dynamic weight changes in Chinese adults.

Lipid indices such as total cholesterol (TC), triglyceride (TG), high-density lipoprotein cholesterol (HDL) and low-density lipoprotein cholesterol (LDL) are biomarkers that can predict CVD and metabolic syndrome (MetS). In addition, the relationships between these biomarkers and weight have also been shown in many cross-sectional and cohort studies. However, most prospective studies have focused on the effects of weight gain on these biochemical indices, but whether changes in blood lipids predict weight gain remains unclear. In a limited number of studies, the role of blood lipids in predicting overweight/obesity has been inconsistent due to differences in subject characteristics and follow-up period. Lower HDL levels were found to be the only blood predictor of the development of metabolically healthy obesity in nonobese healthy adults in Iran after a mean follow-up period of 13 years. TC levels predicted a change in body mass index (BMI) over 12 months in Australian pre-pubertal children, while TG, HDL and insulin levels did not. It is well known that blood lipids are closely related to lifestyle, such as physical activities and dietary habits, so it is beneficial to seek convenient blood lipids to predict weight changes and, in turn, develop prevention lifestyle strategies. Therefore, it is urgent to explore easily measured blood lipids to predict overweight/obesity. Moreover, because weight change is a dynamic process that may include different weight states, it is necessary to identify convenient blood lipid biomarkers that affect transitions between weight states.

While an individual’s weight status changes continuously, only a specific state can be observed at a certain time. Therefore, the exact time of the change in weight state is often unclear. Furthermore, the time interval between two observations may vary by individual. To overcome such issues, the multistate Markov model has been adopted, which is a useful tool to describe a process in which an individual goes through a series of states. First, the transition intensity, which reflects the flow rate of transitions between any two states, can be obtained from the model. Second, classic logistic and Cox regression models only include a single outcome, but the multistate Markov model can analyse multiple outcomes in one system. Third, the multistate Markov model does not require equal lengths for the time intervals between two consecutive observations for different individuals, and therefore, this model makes full use of longitudinal data. Fourth, in the multistate Markov model, the covariate values at the preceding visit are used to predict the outcome state at the subsequent visit; thus, this model can powerfully determine predictors for transitions.

Previous studies have shown that the prevalence of obesity/overweight in China was higher in the North than in the South and in urban areas than in rural areas. Dalian is a modern and developed city in Northeast China. The aim of the current study was to build a multistate Markov model to describe the dynamic nature of weight status in adults by age and sex and to show how demographics and blood lipids affect transitions between weight states. We focused on young and middle-aged people for the early detection and prevention of obesity. To the best of our knowledge, this is the first longitudinal study to analyse weight change and their predictive lipid markers in Chinese adults by the multistate Markov model.
Materials and Methods

Patients and Data

This was a retrospective cohort study. The data were collected from health check-ups acquired from January 2010 to December 2017 at the Second Hospital Affiliated with Dalian Medical University. This study was approved by the ethics committee of Dalian Medical University (Ethics Approval No. 2020 006). The eligibility criteria for the study were age between 18 and 56 at the first check-up, two or more health check-ups in the eight-year study period, and no history of diabetes mellitus or CVD at the first check-up. The data for the first check-up were regarded as baseline data. Participants with a body mass index (BMI) <18.5 at baseline or whose measurement of any blood markers (Glu, TC, TG, HDL and LDL) were outside of the mean ± 3 standard deviations at baseline were excluded from the cohort. Finally, data from a total of 9795 participants were collected in this study. The participants were divided into four groups based on age and sex. The young group included individuals with baseline ages between 18 and 36, and the middle-aged group included individuals with baseline ages between 36 and 56. The maximum ages of the young and middle-aged groups were no more than 42 and 62 years, respectively, during the follow-up.

At each check-up, the subjects were asked for information on their personal medical history, especially a history of metabolic diseases such as diabetes, hyperlipidaemia, and hypertension, and their medication situation. Their weight with light clothing and barefoot height were measured. BMI was calculated as the body mass divided by the square of the body height, in units of kg/m². Venous blood was taken after 12 hours of overnight fasting. Biochemical measures included serum glucose (Glu), TG, TC, HDL and LDL, which were measured on a Siemens ADVIA 2400 automatic biochemical analyser. Glu levels were determined by the hexokinase method. TG levels were determined by the phosphoglyceride oxidase-peroxidase method. TC levels were determined by the visible absorption spectrophotometry method. HDL levels were measured by using the enzyme selective protection method. LDL levels were measured by the visible absorption spectrophotometry method.

Based on the criteria of the Working Group on Obesity in China, the subjects were classified into three weight status categories at each assessment: healthy weight, overweight and obesity. Healthy weight, overweight and obesity were defined as a BMI of 18.5–23.9 kg/m², a BMI of 24–27.9 kg/m² and a BMI ≥ 28.0 kg/m², respectively.

Markov Model

The Concept of Multistate Models

The development of a disease usually includes several stages. A subject may be followed up at an arbitrary time for different reasons. A multistate model has the advantage of investigating the evolution of a disease by considering continuous status changes.

\[ S(t) \] denotes an individual’s state at time \( t \). The transition intensity \( q_{rs} \) is the instantaneous risk of transitioning from state \( r \) to state \( s \) at time \( t \):

\[
q_{rs}(t) = \lim_{\delta t \to 0} \frac{P(S(t + \delta t) = s | S(t) = r)}{\delta t}
\]

All possible intensities form the transition matrix \( Q = (q_{rs}) \). The diagonal entries are defined by \( q_{rr} = - \sum_{r \neq s} q_{rs} \). All entries can be estimated by entering data into the Markov model. The covariate vector \( z(t) \) for transition intensity \( q_{rs} \) can be investigated by the following equation:

\[
q_{rs}(z(t)) = q_{rs}(0) \exp(\beta_{rs}^T z(t))
\]

where \( \beta_{rs}^T \) is the coefficient vector representing the effects of covariates on the transition from state \( r \) to state \( s \) and can be interpreted similarly to those in the ordinary Cox regression model in terms of hazard ratios.

\( q_{rs}(z(t)) \) is used to determine the transition probability matrix \( P(t-u) \), which includes all transition probabilities \( p_{rs}(t-u, z(u)) \), where \( u \) and \( t \) are the initial and ending points of the interval between two consecutive observations, respectively. Time-dependent covariates are assumed to be piecewise constant in the model, that is, the value of the covariate at initial point \( u \) is used to predict the state at the end point \( t \).

Three-State Markov Model

In this study, the Markov model included three states: healthy weight, overweight and obesity. These three states were mutually exclusive, and there was no absorbing state because the mortality rate among healthy people was very low. Note that the weight status of an individual changes continuously, so the individual may advance to or recover from the adjacent states in the continuous-time multistate model. Transitions were permitted from healthy weight to overweight, overweight to obesity, overweight to healthy weight and obesity to healthy weight.
weight and obesity to overweight. A model schematic is presented in Figure 1. Because of different characteristics in metabolism between males and females, Markov models were constructed by sex. Age group, Glu, TC, TG, HDL and LDL were exploratory variables to determine significant factors for weight transitions.

Statistical Analysis
Data were recorded and analysed with Excel 2016. Means and standard deviations were used to express continuous variables with a normal distribution, and medians and quartile were used to present variables with a non-normal distribution. Percentages were used to describe discrete variables. The MSM package R 3.2 was used to build the continuous multi-state Markov model. Transition intensities between states and mean sojourn time in each state were estimated by age and sex. The covariates in the Markov models included age group (middle-aged group vs young group) and Glu, TC, TG, HDL, and LDL levels. The effects of covariates on weight transitions were assessed by sex. First, univariate models were used to determine the effect of each factor. Second, significant factors in the univariate models were entered into the multivariate model. Excel 2016 was used to make a line graph with predicted rates of obesity in individuals beginning with healthy weight over 7.5 years.

Results
A total of 9795 individuals aged 20–56 years, 46.3% male and 53.7% female, were included in our study. A summary of the information of individuals at baseline is presented in Table 1. General information on weight states at baseline and for each follow-up year is presented in Table 2. The average length of the follow-up time was 3.83 years. The prevalence of overweight and obesity was much higher in males than in females. The prevalence of overweight and obesity increased with the length of follow-up in females, whereas it steadily fluctuated in males.

The estimated transition intensities are given in Table 3. In both age groups, males were more likely to progress from healthy weight to overweight than females and were more resistant to recover from obesity. Young females were more likely to move to other states than

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**Table 1** Main Characteristics of Subjects at Baseline

| Variable                        | Males (n = 4537) | Females (n = 5258) |
|---------------------------------|------------------|--------------------|
| **Age group (baseline)**        |                  |                    |
| 18 ≤ age ≤ 36 years, n (%)     | 1840 (40.6)      | 2856 (54.3)        |
| 36 < age ≤ 56 years, n (%)     | 2697 (59.4)      | 2402 (45.7)        |
| **BMI, M (P_{25}, P_{75})**    | 25.13 (23.12,27.16) | 22.15 (20.55,24.12) |
| Healthy weight, n (%)          | 1591(35.1)       | 3874 (73.7)        |
| Overweight, n (%)              | 2088 (46.0)      | 1138 (21.6)        |
| Obesity, n (%)                 | 858 (18.9)       | 246 (4.7)          |
| Serum glucose, mmol/L, M (P_{25}, P_{75}) | 5.51 (5.23, 5.85) | 5.22 (4.96, 5.52) |
| TC, mmol/L, M (P_{25}, P_{75}) | 4.77 (4.25, 5.33) | 4.58 (4.09, 5.17)  |
| TG, mmol/L, M (P_{25}, P_{75}) | 1.31 (0.93, 1.87) | 0.83 (0.64, 1.13)  |
| HDL, mmol/L, M (P_{25}, P_{75})| 1.12 (0.98, 1.29) | 1.40 (1.22, 1.60)  |
| LDL, mmol/L, M (P_{25}, P_{75})| 2.85 (2.42, 3.33) | 2.56 (2.13, 3.05)  |

**Abbreviations:** Glu, glucose; TC, total cholesterol; TG, triglycerides; HDL, high-density lipoprotein cholesterol; LDL, low density lipoprotein cholesterol; HW, healthy weight; M, median; OW, overweight; OB, obesity; P, percentile.
middle-aged females (transition intensity from healthy weight to overweight: 0.11 vs 0.09; overweight to healthy weight: 0.37 vs 0.18; overweight to obesity: 0.12 vs 0.06; obesity to overweight: 0.42 vs 0.27). Young males and middle-aged males had similar intensities of transitions between states, while only the probability of progressing from overweight to obesity seemed higher in young males than in middle-aged males (0.10 vs 0.06). Table 4 shows the mean sojourn time in each state specified by sex and age. In both age groups, the mean sojourn times in overweight and obesity were greater in males than in females. In both males and females, middle-aged individuals had greater sojourn times in the same state than young individuals. The greatest sojourn time in overweight and obesity appeared in the middle-aged male group (5.90 and 5.38 years, respectively).

The results regarding the progression from healthy weight to obesity in each group are presented in Figure 2. The rate of obesity in individuals beginning with healthy weight increased with time. Young groups were more likely to develop obesity than middle-aged groups, and males were more likely to develop obesity than females. The predicted rates of obesity varied greatly between groups. After 7.5 years, obesity developed in an estimated 13.7% of young males and 4.5% of middle-aged females.

Table 5 displays the effects of covariates on each transition in females. The 95% confidence interval (CI) indicated that age group and Glu, TC, TG, HDL and LDL levels were significant factors for some particular transitions in the univariate models. Compared with young females, middle-aged females tended to maintain their preceding weight state. Increases in TC, TG and LDL levels predicted the transition from healthy weight to overweight, and increases in Glu and LDL levels also made females less likely to recover from obesity. Females with higher HDL levels were more likely to recover from overweight or obesity and less likely to progress from healthy weight to overweight. However, only age group and HDL levels were significant for some particular transitions in the multivariate model. If HDL levels increased 1 mmol/L, the probability of progression from healthy weight to overweight decreased by 37.0% (HR = 0.63), and the probabilities of recovery (overweight to healthy weight and obesity to overweight) increased by 62.0% (HR = 1.62) and 1.23-fold (HR = 2.23), respectively.

The effects of covariates on transitions in males are shown in Table 6. Age group and TG and HDL levels were
### Table 3: Transition Intensities Estimated by the Multistate Model \( q_{rs}(95\% CI) \)

| Original States | Follow-Up States (Young Group) | Follow-Up States (Middle-Aged Group) |
|-----------------|---------------------------------|-------------------------------------|
|                 | Healthy Weight                  | Overweight                          | Obesity | Healthy Weight | Overweight | Obesity |
| **Female**      |                                 |                                    |         |               |            |        |
| Healthy weight  | \(-0.11 (-0.12, -0.10)\)        | \(0.11 (0.10, 0.12)\)              | 0       | \(-0.09 (-0.10, -0.08)\) | \(0.18 (0.16, 0.20)\) | 0       |
| Overweight      | \(0.37 (0.34, 0.41)\)           | \(-0.50 (-0.54, -0.45)\)          | \(0.12 (0.10, 0.15)\) | \(-0.24 (-0.26, -0.20)\) | \(0.27 (0.22, 0.33)\) | \(0.06 (0.05, 0.08)\) |
| Obesity         | 0                               | \(-0.42 (-0.51, -0.34)\)          | 0       | 0              | 0.09 (0.08, 0.10) | \(-0.27 (-0.33, -0.22)\) |
| **Male**        |                                 |                                    |         |               |            |        |
| Healthy weight  | \(-0.17 (-0.19, -0.15)\)        | \(0.17 (0.15, 0.20)\)              | 0       | \(-0.16 (-0.18, -0.15)\) | \(0.09 (0.08, 0.10)\) | 0       |
| Overweight      | \(0.11 (0.10, 0.13)\)           | \(-0.21 (-0.23, -0.19)\)          | \(0.10 (0.08, 0.11)\) | \(0.16 (-0.17, -0.14)\) | \(0.16 (0.14, 0.19)\) | \(0.06 (0.05, 0.07)\) |
| Obesity         | 0                               | \(-0.17 (-0.20, -0.15)\)          | 0       | 0              | 0.09 (0.08, 0.10) | \(-0.16 (-0.19, -0.14)\) |

**Note:** \( q_{rs} \) is the instantaneous risk of transitioning from state \( r \) to state \( s \).

### Table 4: Estimated Mean Sojourn Time in Each State

| States            | Young Group | Middle-Aged Group |
|-------------------|-------------|-------------------|
|                   | Estimate (Years) | Standard Errors | 95% CI | Estimate (Years) | Standard Errors | 95% CI |
| **Female**        |              |                  |        |                |                |        |
| Healthy weight    | 9.04        | 0.41             | 8.28, 9.87 | 10.67 | 0.54 | 9.67, 11.77 |
| Overweight        | 2.02        | 0.09             | 1.85, 2.20 | 4.15 | 0.20 | 3.78, 4.55 |
| Obesity           | 2.41        | 0.24             | 1.98, 2.94 | 3.68 | 0.36 | 3.05, 4.45 |
| **Male**          |              |                  |        |                |                |        |
| Healthy weight    | 5.75        | 0.34             | 5.13, 6.46 | 6.15 | 0.33 | 5.53, 6.84 |
| Overweight        | 4.81        | 0.25             | 4.34, 5.33 | 6.44 | 0.29 | 5.90, 7.03 |
| Obesity           | 5.79        | 0.46             | 4.95, 6.77 | 6.23 | 0.46 | 5.38, 7.21 |
significant factors for some transitions in the univariate and multivariate models. Middle-aged males were less likely to change their weight state than young males. Males with higher TG or lower HDL levels had a greater risk of progressing from healthy weight to overweight, while higher HDL levels predicted recovery from overweight. If TG levels increased 1 mmol/L, the risk of progression from healthy weight to overweight increased by 24.0% (HR= 1.24). Each unit increase in HDL levels was associated with a 0.99-fold (HR= 1.99) increase in the chance of recovery from overweight to healthy weight and with a 0.37-fold (HR= 0.63) decrease in the risk of progression from healthy weight to overweight.

**Discussion**

The current study describes the dynamic evolution of obesity by sex and age and explores the blood lipids that can predict weight transitions based on a large longitudinal dataset.

Our results indicated that males were more likely to transit from healthy weight to overweight and more resistant to recover from worse states than females. Moreover, males had a higher prevalence and more sojourn time in overweight and obesity than females. This might be due to differences in lifestyles, genetics or behaviour between males and females. The wheat staple pattern and the rice staple pattern are two traditional dietary patterns in China. A recent study in Shanghai showed that males with these two dietary patterns had a greater risk of obesity than females. Moreover, traditional Chinese cultures favour females with slender body shapes, so females may be more sensitive to their weight and more active in weight control than males. In the current study, the predicted rates of obesity with time were greater in males than in females beginning at healthy weight. In a study based on a large population in Northeast China, males had a higher prevalence of overweight or obesity than females. Some studies in other countries also obtained similar results, with a higher prevalence of obesity in males. These results on differences in prevalence between males and females were partly similar to those we found. However, the prevalence of obesity was greater in females than in males in some populations. These opposite results may be due to differences in the geographical and cultural background of the population.

Many studies have reported that the prevalence of obesity increases with age until 60 years in both males and females. In our study, age was also responsible for the weight transitions. Young people were more likely to transit from one weight state to another, regardless of progression or recovery. Middle-aged people with overweight or obesity tended to maintain their preceding weight state. These results were partly consistent with one of our previous studies, in which the numbers of transitions from obesity to healthy weight were greater in young people than in middle-aged people. One reason may be a decline in metabolic capability in middle-aged people, which makes it difficult for those who have been obese to lose weight. There is evidence that weight gain is associated with metabolic disorders in menopausal females. Middle-aged people with healthy weight might be likely to have had a healthy lifestyle from a young age,
### Table 5 Effects of Different Covariates on Weight Transitions in Females [Hazard Ratios (95% CIs)]

| Variable | Univariate Model | Multivariate Model |
|----------|-----------------|--------------------|
|          | HW → OW | OW → HW | OW → OB | OB → OW | HW → OW | OW → HW | OW → OB | OB → OW |
| Age group" | 0.85 (0.74, 0.97)* | 0.48 (0.41, 0.55)* | 0.52 (0.40, 0.67)* | 0.65 (0.50, 0.86)* | 0.77 (0.67, 0.88)* | 0.45 (0.39, 0.52)* | 0.52 (0.40, 0.68)* | 0.73 (0.55, 0.97)* |
| Glu      | 1.08 (0.96, 1.22) | 0.92 (0.82, 1.03) | 0.91 (0.75, 1.10) | 0.81 (0.68, 0.98)* | 1.03 (0.90, 1.17) | 1.01 (0.91, 1.12) | 0.99 (0.83, 1.19) | 0.91 (0.76, 1.10) |
| TC       | 1.09 (1.02, 1.18)* | 1.06 (0.98, 1.15) | 0.92 (0.80, 1.06) | 0.90 (0.77, 1.04) | 1.02 (0.79, 1.32) | 1.15 (0.89, 1.48) | 1.13 (0.72, 1.77) | 1.09 (0.69, 1.70) |
| TG       | 1.19 (1.04, 1.36)* | 1.00 (0.84, 1.19) | 0.99 (0.83, 1.21) | 0.92 (0.76, 1.11) | 1.12 (0.95, 1.31) | 1.08 (0.91, 1.27) | 1.05 (0.81, 1.37) | 1.03 (0.79, 1.35) |
| HDL      | 0.59 (0.47, 0.75)* | 1.70 (1.30, 2.14)* | 1.21 (0.76, 1.91) | 2.47 (1.44, 4.23)* | 0.63 (0.43, 0.93)* | 1.62 (1.10, 2.37)* | 1.17 (0.58, 2.36) | 2.23 (1.07, 4.62)* |
| LDL      | 1.19 (1.09, 1.30)* | 1.00 (0.91, 1.11) | 0.86 (0.72, 1.02) | 0.76 (0.63, 0.92)* | 1.18 (0.90, 1.55) | 0.93 (0.71, 1.23) | 0.81 (0.50, 1.31) | 0.73 (0.45, 1.19) |

**Notes:** "Age group was middle-aged group vs young group; *P<0.05.

**Abbreviations:** Glu, glucose; TC, total cholesterol; TG, triglycerides; HDL, high-density lipoprotein cholesterol; LDL, low density lipoprotein cholesterol; HW, healthy weight; OW, overweight; OB, obesity.

### Table 6 Effects of Different Covariates on Weight Transitions in Males [Hazard Ratios (95% CIs)]

| Variable | Univariate Model | Multivariate Model |
|----------|-----------------|--------------------|
|          | HW → OW | OW → HW | OW → OB | OB → OW | HW → OW | OW → HW | OW → OB | OB → OW |
| Age group" | 0.94 (0.80, 1.10) | 0.83 (0.69, 0.99)* | 0.65 (0.53, 0.79)* | 0.93 (0.75, 1.15) | 0.91 (0.78, 1.07) | 0.82 (0.68, 0.99)* | 0.64 (0.52, 0.79)* | 0.92 (0.74, 1.13) |
| Glu      | 1.02 (0.91, 1.14) | 1.05 (0.94, 1.18) | 0.92 (0.79, 1.07) | 1.03 (0.93, 1.14) | – | – | – | – |
| TC       | 1.06 (0.97, 1.17) | 1.04 (0.93, 1.16) | 0.91 (0.81, 1.03) | 0.94 (0.83, 1.07) | – | – | – | – |
| TG       | 1.30 (1.19, 1.41)* | 0.93 (0.83, 1.04) | 0.93 (0.83, 1.05) | 0.94 (0.84, 1.06) | 1.24 (1.13, 1.37)* | 1.01 (0.89, 1.13) | 0.91 (0.80, 1.03) | 0.95 (0.84, 1.08) |
| HDL      | 0.49 (0.36, 0.69)* | 1.97 (1.37, 2.82)* | 0.76 (0.49, 1.19) | 1.32 (0.78, 2.23) | 0.63 (0.45, 0.88)* | 1.99 (1.35, 2.94)* | 0.67 (0.41, 1.10) | 1.24 (0.70, 2.17) |
| LDL      | 1.07 (0.96, 1.20) | 0.99 (0.88, 1.13) | 0.99 (0.86, 1.15) | 0.95 (0.81, 1.10) | – | – | – | – |

**Notes:** "Age group was middle-aged group vs young group; *P<0.05.

**Abbreviations:** Glu, glucose; TC, total cholesterol; TG, triglycerides; HDL, high-density lipoprotein cholesterol; LDL, low density lipoprotein cholesterol; HW, healthy weight; OW, overweight; OB, obesity.
which helped to maintain their healthy weight. Our results suggested that prevention and intervention should be implemented early for young people due to their weight instability and easier recovery.

In the current study, TC, TG and LDL levels were found to positively predict the onset of overweight in males and females, whereas they were not significant predictors for transitioning from overweight to obesity. Moreover, HDL levels were a sensitive predictor of weight transitions in both males and females. Few cohort studies have investigated the probability of lipid indices predicting weight gain. A cohort study in Sydney showed that TC levels at baseline had a positive association with changes in BMI over 12 months in pre-pubertal children,\(^1\) which demonstrated that TC levels were a predictor of weight gain. The Tehran Lipid and Glucose Study identified lower HDL levels as a predictor of the development of metabolically healthy obesity in nonobese healthy adults.\(^2\) The findings in these studies were partly consistent with ours. However, a cohort study in Finnish adults showed that higher TC, higher Glu and lower HDL levels at baseline predicted weight loss over 6.5 years.\(^3\) This result was contrary to previous results and our results. In that study, the time between the baseline and the end point was relatively long, and fluctuations in the indicator before the end point could not be measured, which may have made the association between baseline biomarkers and outcome events slightly weak. Our study was based on longitudinal data with multiple repeated measurements, and the interval between any two consecutive follow-up years was approximately one year.

In the current study, HDL level was the strongest biomarker in both males and females. HDL is well recognized as a potent atheroprotective biomarker due to the role of HDL in the reverse cholesterol transport pathway. Cholesterol can be combined with HDL and transported from peripheral tissues to the liver and then excreted in the faeces.\(^4\) This process might be helpful to control weight by maintaining the balance of cholesterol in the body.

From the perspective of prevention, our findings showed that increases in HDL levels benefit transitions from worse weight states to better states or staying at better weights, which provided the direction for prevention and control of obesity. Several studies have reported that physical activities and dietary habits can significantly influence cholesterol. A study based on a large sample of Taiwanese adults showed that aerobic exercise and regular badminton were associated with higher levels of HDL.\(^5\) Another large-sample study in Taiwan found that even after adjusting for smoking and BMI, weekly exercise durations of < 2.5 and ≥ 2.5 hours both significantly increased HDL levels compared to exercise durations of 0 hours.\(^6\) Similar results were obtained in other studies.\(^7-10\) On the other hand, dietary habits also have a strong influence on HDL levels. A Mediterranean diet, which is characterized by high consumption of vegetables, fruits, legumes, wheat-based cereals, nuts, fish, and especially olive oil and low consumption of red and processed meat, is recognized as a dietary pattern that helps to increase HDL levels.\(^11\) However, a low-fat, plant-based eating pattern was linked to a decrease in HDL levels.\(^12\) Although there was a lack of individual information on physical activities and dietary habits in the current study, the association of HDL levels with lifestyle may partly explain our results. The principle of the Markov model in our study indicates that the transitions between weight states were influenced by HDL levels, which is changeable with lifestyle. Lifestyle factors such as dietary habits and physical activities may affect changes in blood lipids, leading to changes in body weight. In addition, finding convenient blood lipids that can predict weight changes benefits the development of corresponding lifestyle prevention strategies. For example, certain types of physical activity can be taken to increase or decrease the HDL concentration. The right food choices can lower HDL levels.\(^13,11,12\)

This study has some limitations. First, behavioural factors, such as dietary habits, physical activity and sleep time, were not included in the analysis. However, lifestyle may be an underlying factor affecting blood indicators. After identifying blood biomarkers that can predict weight state transitions, a corresponding lifestyle change would also be advocated to reduce the risk of obesity and cardiovascular disease. Future research will aim to investigate more variables to explore the important factors for the progression and regression of weight states. Second, the “natural” transition probability might not be completely natural because some individuals might have received interventions through health services.

**Conclusion**

The weight status of young people was less stable than that of middle-aged people. Males were more likely to progress to overweight from healthy weight and more resistant to recover from worse states than females. Young males with healthy weight were more likely to develop obesity than other healthy weight groups. Blood lipid levels, especially HDL, were the...
strongest predictors of weight transitions. Prevention and intervention measures should be adopted early due to the instability of weight states for the group of young people.

**Abbreviations**
Glu, serum glucose; TC, total cholesterol; TG, triglyceride; HDL, high-density lipoprotein cholesterol; LDL, low-density lipoprotein cholesterol; MetS, metabolic syndrome; BMI, body mass index.

**Data Sharing Statement**
The datasets used and analysed in the present study are available from the corresponding author upon reasonable request.

**Ethics Approval and Consent to Participate**
All participants involved in the study were informed about the purpose of the study, and all research procedures were conducted in accordance with the 1964 Declaration of Helsinki. This study was approved by the ethics committee of Dalian Medical University (Ethics Approval No. 2020 006). Since we excluded all identifiable personal information, the ethics committee agreed to waive the requirement for informed consent of the study.

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**Author Contributions**
XT is the first author. GS is the corresponding author. XT and GS designed the study. HZ, YZ, FL, QL, DH and GL screened and extracted the data. All authors contributed to data analysis, drafting or revising the article, have agreed on the journal to which the article will be submitted, gave final approval of the version to be published, and agree to be accountable for all aspects of the work.

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**Disclosure**
The authors declare that they have no competing interests.

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