

Association of *Apolipoprotein E* Gene Polymorphism with Ischemic Stroke in Coronary Heart Disease Patients Treated with Medium-intensity Statins

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Objective: To study the association of *apolipoprotein E* (*APOE*) gene polymorphism with ischemic stroke (IS) in coronary heart disease (CHD) patients treated with medium-intensity statins.

Methods: The retrospective study was performed on 662 samples including 169 CHD subjects complicated with IS, 296 subjects with CHD, and 197 control subjects. The *APOE* gene was obtained from case files. Univariable and multivariable logistic regression analyses were utilized to recognize the possible risks of CHD and IS.

Results: The frequency of $\epsilon 3$ - $\epsilon 4$ genotype was increased in the CHD group ($p=0.013$) and CHD-IS group ($p=0.001$), the frequency of $\epsilon 4$ allele was also increased in the CHD group ($p=0.047$) and the CHD-IS group ($p=0.009$) compared with control group. $\epsilon 3$ - $\epsilon 4$ genotype was the independent risk for CHD and CHD-IS after adjusting for traditional risk factors with adjusted odds ratio (AOR) 2.210, 95%CI: 1.263–3.867, $p=0.005$ and (AOR 2.794, 95%CI: 1.539–5.072, $p=0.002$). The $\epsilon 4$ allele was also significantly associated with CHD (AOR 2.126, 95%CI: 1.265–3.575, $p=0.004$) and CHD-IS (AOR 2.740, 95%CI: 1.569–4.784, $p=0.001$).

Conclusion: These results demonstrated that $\epsilon 4$ allele influenced the development of CHD with or without IS, especially for the genotype of $\epsilon 3$ - $\epsilon 4$. CHD patients carrying the $\epsilon 3$ - $\epsilon 4$ genotype and the $\epsilon 4$ allele were significantly associated with the incidence of IS, even if medium-intensity statins had been used.

Keywords: *apolipoprotein E*, polymorphism, coronary heart disease, ischemic stroke, statins

Introduction

Coronary heart disease (CHD) and stroke cause immense health and economic burdens.¹ According to the research conducted by the China National Stroke Registry, ischemic stroke (IS) is a predominant stroke subtype, accounting for 66.7%.² CHD and IS are both based on atherosclerosis caused by lipid metabolism disorders. As one of the complex heterogeneous diseases of multiple etiologies and major clinical manifestations, IS is not only associated with various traditional risk factors, but also influenced by the gene polymorphism.

Apolipoprotein E (*APOE*) gene is a common candidate gene which exerts an important role in lipid metabolism. It includes three alleles ($\epsilon 2$, $\epsilon 3$, and $\epsilon 4$) which express three isoforms of proteins (E2, E3, and E4). These alleles of *APOE* gene

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constitute six genotypes ($\epsilon 2\text{-}\epsilon 2$, $\epsilon 2\text{-}\epsilon 3$, $\epsilon 2\text{-}\epsilon 4$, $\epsilon 3\text{-}\epsilon 3$, $\epsilon 3\text{-}\epsilon 4$, and $\epsilon 4\text{-}\epsilon 4$),³ among which the most common genotype is $\epsilon 3\text{-}\epsilon 3$, and $\epsilon 3$ is the most predominant allele in most races.⁴⁻⁶ Allele $\epsilon 2$ is associated with higher level triglyceride (TG), lower level low density lipoprotein cholesterol (LDL-C) in plasma and lower risk of coronary artery disease when compared with $\epsilon 3$ allele. Allele $\epsilon 4$ is associated with higher plasma levels of LDL-C, total cholesterol (TC), apolipoprotein B and increased risk of coronary artery disease compared with $\epsilon 3$ allele.⁷

Statins are one of the most commonly prescribed medicines worldwide which are utilized to prevent and stabilize atherosclerotic plaque. Statin therapy could also reduce the risk of IS in CHD patients.⁸⁻¹⁰ In the current clinical practice in China, medium-intensity statins are widely used in the routine treatment of CHD. Whether the routine dose of statins could reduce IS risk in CHD patients was still poorly documented. There was no data demonstrating the relationship between *APOE* gene polymorphism and IS in CHD patients treated with medium-intensity statins. In this study, we aimed to investigate the association of *APOE* gene polymorphism with IS in CHD patients treated with medium-intensity statins, of whom the difference on plasma lipid levels between $\epsilon 4$ carriers and $\epsilon 3\text{-}\epsilon 3$ genotype.

Methods

Inclusion Criteria and Exclusion Criteria of the Study Population

The study complied with the Declaration of Helsinki for investigation in human beings and was approved by the Ethics Committee of The First Affiliated Hospital of Nanchang University. All patient data complied with relevant data protection and privacy regulations.

The single center retrospective study consisted of 662 subjects, all of whom were admitted to the First Affiliated Hospital of Nanchang University between September 2013 and August 2017. All subjects were divided into control group ($n=197$), CHD group ($n=296$) and CHD-IS group ($n=169$). The control group was collected from healthy individuals who visited the Department of Physical Examination Center. All control patients declared that they had no history of CHD and IS, and this was confirmed by electrocardiogram (ECG) and brain computerized tomography (CT). CHD group patients were checked by coronary angiography for at least 50% lumen obstruction in a major coronary artery or one of their branches.

CHD-IS group patients included CHD patients complicated with IS, which were diagnosed according to the criteria amended by the Chinese Medical Association in National Cerebrovascular Disease Conference in 2015, with confirmation by brain CT or magnetic resonance imaging (MRI). In the CHD-IS group, IS happened in CHD patients who insisted on taking medium-intensity statins since diagnosed as CHD. Telephone follow-up and checking out prescriptions for outpatients were used to determine if CHD patients were taking statins until the occurrence of IS, and CHD patients who took medium-intensity statins for less than one month when diagnosed as IS were excluded. Medium-intensity statins included atorvastatin (20 mg), rosuvastatin (10 mg), or pitavastatin (2 mg). Other CHD medicines had no special requirements. Exclusion criteria included heart valve disease, atrial fibrillation, hemorrhagic stroke or ischemic stroke before coronary heart disease, neurodegenerative disorders, hepatic and renal diseases, and any form of endocrine or metabolic diseases. Information were obtained from these case files which included age, gender, obesity, discharge diagnosis, history of smoking, dyslipidemia, diabetic mellitus, hypertension, homocysteine (Hcy), *APOE* genotypes, vital signs, laboratory examination, ECG, head imaging examination (CT or MRI), cardiac ultrasound, coronary arteriography, complications, and the dose of statins.

Definitions

Obesity was defined as body mass index (BMI) ≥ 25 kg/m². Cigarette smokers meant someone who had smoked continuously more than one year even if they were no longer a smoker. Hypertension was defined as blood pressure $\geq 140/90$ mmHg or the use of antihypertensive medications. Dyslipidemia were defined as when one had levels of TG ≥ 1.70 mmol/L, TC ≥ 5.72 mmol/L, LDL-C ≥ 3.12 mmol/L, high-density lipoprotein cholesterol (HDL-C) ≤ 1.0 mmol/L. Diabetic mellitus was defined as fasting plasma glucose ≥ 6.1 mmol/L and postprandial glucose ≥ 11.1 mmol/L, and/or diagnosed diabetes mellitus receiving treatment.

Statistical Analysis

All analyses were conducted using the statistical software SPSS statistics 23 (IBM Corporation, Armonk, NY, USA), and followed the Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) statement. Continuous variables were expressed as mean \pm SD

or median (with IQR), and categorical variables as numbers and percentages. One-way ANOVA followed by post-hoc Bonferroni multiple comparison test was used to evaluate the mean difference of the data among the three groups (control, CHD, and CHD-IS). The chi-squared tests and Fisher-F were used for categorical variables. The difference of allele and genotype among the three groups and deviations from Hardy–Weinberg equilibrium were tested by chi-squared tests or Fisher-F. The univariable logistic regression analysis with unadjusted OR and 95% CI were used to explore the association between *APOE* polymorphism and diseases. Multivariable binary logistic regression analysis with adjusted OR and 95%CI, which followed adjusting for age, sex, smoking history, SBP and Hcy, were used to determine the independent risk factor for development of IS after CHD. A value of $p < 0.05$ was considered statistically significant.

Results

The Baseline Characteristics of the Population in This Study

The demographic and clinical data were shown in Table 1. There was a significant increase in age of the CHD-IS and CHD groups compared with the control group ($p = 0.009$). The proportion of males in the CHD-IS and CHD groups was increased when compared with the control group ($p < 0.001$).

Other clinical parameters such as systolic blood pressure (SBP) and Hcy were significantly higher in both CHD-IS and CHD groups when compared with the control group ($p < 0.001$ and $p = 0.020$, respectively). CHD-IS group and CHD group had significantly higher frequencies of smoking and dyslipidemia compared with control group ($p = 0.003$ and $p < 0.001$, respectively). Despite of taking medium-intensity statins, the levels of TC and LDL-C in the CHD-IS and CHD groups still showed significant increase when compared with the control group ($p = 0.015$ and $p < 0.001$, respectively). The lower levels of TC and LDL-C were observed in the CHD-IS group compared with CHD group, but no significant difference was shown ($p = 0.241$ and $p = 0.139$, respectively). The levels of TG and HDL-C showed no significant difference among three groups ($p = 0.211$ and $p = 0.339$, respectively).

The Frequencies of *APOE* Genotype and $\epsilon 3$, $\epsilon 4$, $\epsilon 2$ Alleles

The genotype distribution of control, CHD, and CHD-IS groups were in Hardy–Weinberg equilibrium (Table 2). The frequency of $\epsilon 3$ - $\epsilon 3$ genotype was significantly decreased in the CHD-IS and CHD groups ($p < 0.001$ and $p = 0.029$, respectively) compared with the control group, while $\epsilon 3$ allele was only observed to have a lower prevalence in the CHD-IS group ($p = 0.002$). The frequency of $\epsilon 3$ - $\epsilon 4$ genotype was significantly increased in the CHD-IS

Table 1 Demographic and Clinical Data of Different Groups

Variables	Controls (n=197)	CHD-IS (n=169)	CHD (n=296)
Age (years)	61.23±9.97	64.92±12.22**	63.42±12.27*
Sex (male)	107	119***	225***
HR (bpm)	80.21±17.71	81.21±19.25	80.43±20.04
SBP (mmHg)	124.09±20.38	130.22±22.61**	128.23±22.13*
DBP (mmHg)	81.19±16.56	77.25±14.51	77.97±14.51
Smoking (%)	37.60	51.50***	53.00***
Hypertension (%)	32.60	56.20***	55.00***
Dyslipidemia (%)	2.50	9.50**	3.40*
Diabetes mellitus (%)	12.70	20.70	17.20
Hcy (μmol/L)	16.97±7.13	19.84±10.07****	20.51±10.66***
Glucose (mmol/L)	5.78±1.40	6.07±2.04	5.95±1.68
TC (mmol/L)	3.73±0.88	4.29±0.98***	4.31±1.21***
TG (mmol/L)	1.42±0.86	1.54±0.98	1.39±0.99
HDL-C (mmol/L)	1.17±0.81	1.15±0.38	1.23±0.63
LDL-C (mmol/L)	2.68±0.86	2.87±1.11**	3.03±1.39*
Statin (–%)	–	95.40	18.30

Notes: Data are presented as mean ±SD, or numbers (n) and percentage of subjects. * p -value < 0.05 in comparison between control group and CHD group, ** p -value < 0.05 in comparison between control group and CHD-IS group, *** p -value < 0.001 in comparison between control group and CHD group, **** p -value < 0.001 in comparison between control group and CHD-IS group.

Abbreviations: Hcy, homocysteinemia; TC, total cholesterol; HDL, high-density lipoprotein cholesterol; TG, triglyceride; LDL, low-density lipoprotein cholesterol; SBP, systolic blood pressure; DBP, diastolic blood pressure.

Table 2 Genotype and Allele Frequencies Distribution of APOE Gene Polymorphism in the CHD and CHD-IS Groups Compared with the Control Group

Genotype	Groups		
	Control (n=197)	CHD (n=296)	CHD-IS (n=169)
ε2-ε2	4 (2.00%)	2 (0.70%)	2 (1.20%)
ε2-ε3	9 (4.60%)	16 (5.40%)	13 (7.70%)
ε2-ε4	0 (0%)	2 (0.70%)	2 (1.20%)
ε3-ε3	157 (79.70%)	210 (70.90%)	106 (62.60%)
ε3-ε4	22 (11.20%)	58 (19.6%)	42 (24.90%)
ε4-ε4	5 (2.50%)	8 (2.70%)	4 (2.40%)
p-value	0.130	0.729	0.983
Allele ε2 (95%CI)	0.043 (0.029–0.047)	0.037 (0.029–0.041)	0.056 (0.042–0.069)
Allele ε3 (95%CI)	0.876 (0.841–0.892)	0.834 (0.803–0.891)	0.790 (0.767–0.796)
Allele ε4 (95%CI)	0.081 (0.068–0.087)	0.129 (0.114–0.132)	0.154 (0.141–0.167)

and CHD groups ($p=0.001$ and $p=0.013$, respectively), and the allele of ε4 had a higher prevalence in the CHD-IS and CHD groups ($p=0.002$ and $p=0.014$, respectively). In addition, the frequencies of ε2-ε3 genotype and ε2 allele only had an increased tendency in the CHD-IS group compared with the control group, but no significant difference was observed ($p=0.210$ and $p=0.415$, respectively). The frequencies of ε2-ε2, ε4-ε4, and ε2-ε4 genotypes had no statistically significant differences observed among three groups (Table 3).

The Association Between APOE Gene Polymorphism and the Development of CHD and IS

The ε4 allele was associated with low lipid clearance property. Therefore, the ε3-ε4 genotype could be the risk factor for the development of CHD and IS. The ε4 allele

increased the risk of both CHD-IS and CHD groups with unadjusted OR 1.964 (95%CI: 1.228–3.142, $p=0.005$) and 1.666 (95%CI: 1.079–2.572, $p=0.021$). The ε3-ε4 genotype increased the risk of CHD with unadjusted OR 1.939 (95%CI: 1.143–3.287, $p=0.014$) and showed the same tendency about CHD-IS with unadjusted OR 2.631 (95%CI: 1.496–4.626, $p=0.001$) (Table 4). The multivariate binary logistic regression analysis showed that ε3-ε4 genotype was the independent risk factor for development of CHD-IS and CHD after being adjusted for age, sex, smoking history, SBP and Hcy ($p=0.002$, OR 2.794, 95%CI: 1.539–5.072 and $p=0.005$, OR 2.210, 95%CI: 1.263–3.867, respectively). Furthermore, the ε4 allele was found to be independent risk factor for development of CHD-IS and CHD ($p=0.001$, OR 2.740, 95%CI: 1.569–4.784 and $p=0.004$, OR 2.126, 95%CI: 1.265–3.575, respectively). In the present study, age, male, SBP, and Hcy were also independent risks for CHD-IS and CHD, but smoking

Table 3 Genotype and Allele Frequencies Distribution of APOE Gene Polymorphism in the CHD Group and CHD-IS Group Compared with the Control Group

Genotype	Groups		
	Control (n=197)	CHD (n=296)	CHD-IS (n=169)
ε2-ε2	0.020	0.007	0.012
ε2-ε3	0.046	0.054	0.077
ε2-ε4	0	0.007	0.012
ε3-ε3	0.797	0.709*	0.626**
ε3-ε4	0.112	0.196*	0.249**
ε4-ε4	0.025	0.027	0.024
Allele ε2	0.043	0.037	0.056
Allele ε3	0.876	0.834	0.790**
Allele ε4	0.081	0.129*	0.154**

Notes: *p-value <0.05 compared between control group and CHD group, **p-value <0.05 compared between control group and CHD-IS group.

Table 4 Associations of *APOE* Gene Polymorphism with the Risk of the CHD Group and CHD-IS Group Compared with the Control Group Represented as Unadjusted OR

Genotype	Controls (n=197)	CHD (n=296)	Unadjusted OR (95%CI)	p-value	CHD-IS (n=169)	Unadjusted OR (95%CI)	p-value
ε2-ε2	4	2	0.328 (0.060–1.809)	0.201	2	0.578 (0.105–3.195)	0.530
ε2-ε3	9	16	1.194 (0.517–2.757)	0.679	13	1.741 (0.725–4.180)	0.215
ε2-ε4	0	2			2		
ε3-ε3	157	210	0.622 (0.405–0.955)	0.030	106	0.422 (0.265–0.672)	0.001
ε3-ε4	22	58	1.939 (1.143–3.287)	0.014	42	2.631 (1.496–4.626)	0.001
ε4-ε4	5	8	1.067 (0.344–3.309)	0.911	4	0.931 (1.097–3.194)	0.916
Allele ε2	17	22	0.735 (0.377–1.433)	0.366	19	1.321 (0.675–2.584)	0.416
Allele ε3	345	494	0.716 (0.495–0.1036)	0.076	267	0.527 (0.354–0.783)	0.002
Allele ε4	32	78	1.666 (1.079–2.572)	0.021	52	1.964 (1.228–3.142)	0.005

was only found to be the independent risk for CHD (Table 5).

The Correlation of ε4 Allele Carriers with Lipid Profiles

The significant differences of *APOE* distribution among these groups mainly resulted from different frequencies of the ε4 allele. The distributions of ε4 allele frequencies could cause the differences of plasma lipid levels. Even though CHD-IS group patients continuously took medium-intensity statins, the differences of plasma lipid levels were still observed between ε4 carriers and ε3-ε3 genotype in all three groups (Table 6). In the control group and CHD group, the values of TC and LDL-C showed significant elevation in ε4 carriers compared with the ε3-ε3 genotype. However, the values of TG and HDL-C level were observed with no significant difference in ε4 carriers when compared with the ε3-ε3 genotype in the control and CHD groups. In the CHD-IS group, there was significant elevation in TC level but not in LDL-C level in ε4 carriers compared with the ε3-ε3 genotype.

Discussion

Atherosclerosis is a systemic disease caused by lipid metabolism disorders in which coronary arteriosclerosis and cerebral arteriosclerosis can occur simultaneously. Clinical trials had confirmed the beneficial effects of statin treatment in primary and secondary prevention of cardiovascular disease and stroke.¹¹ The statins affected the development of arteriosclerosis by decreasing cholesterol concentration and other mechanisms, which could further prevent the occurrence of IS in CHD patients. Statins could reduce the risk of IS in CHD patients.^{9–11} IS is the result of complex interactions involving different

environmental and genetic factors.³ *APOE* gene is a common candidate gene which plays an important role in lipid metabolism. The relationship of *APOE* polymorphism and coronary artery disease had been explored in several ethnic groups, including Caucasian in the USA,¹² Italian,¹³ Turkish,¹⁴ Chinese¹⁵ and Indian.¹⁶ The association between *APOE* polymorphism and the risk of different subtypes of stroke had been evaluated, the ε4 allele might predict an increased risk for different subtypes of stroke, including IS and hemorrhagic stroke.¹⁷ The *APOE* ε4+ genotype was related to more severe intracranial atherosclerosis score.¹⁸ The ε4 allele was associated with higher plasma levels of LDL-C, TC, apolipoprotein B and increased the risk of coronary artery disease when compared with ε3 allele.¹⁹ ε4 allele was an independent risk factor for myocardial infarction.²⁰ At present clinical practice, medium-intensity statins are widely used in China. However, there is no research demonstrating the relationship between *APOE* gene polymorphism and IS in CHD patients treated with medium-intensity statins, so it is of great interest for us to study the relationship.

In present study, higher prevalence of elderly, male, higher level of SBP, TC, and LDL-C were observed in CHD group and CHD-IS group, all of which were the established risk factors for CHD and IS. Higher level of SBP damaged the intima of the artery, which promoted the deposition of cholesterol and lipids, and aggravated the formation of atherosclerotic plaque. In addition, our results were consistent with several epidemiological studies in which Hcy was an independent risk factors for stroke.^{21–23}

In the Chinese population, the predominant genotype is ε3-ε3, whereas the most common allele is ε3.^{5,24} In this study, the ε3-ε3 genotype accounted for 79.7% in the control group, 70.9% in the CHD group and 62.6% in

Table 5 Association of *APOE* Gene Polymorphism with the Risk of the CHD Group and CHD-IS Group Compared with the Control Group Represented as Adjusted OR

	CHD		CHD-IS	
	Adjusted OR (95%CI)	p-value	Adjusted OR (95%CI)	p-value
Age	1.028 (1.010–1.047)	0.002	1.036 (1.015–1.057)	0.001
Sex	0.400 (0.261–0.615)	0.001	0.542 (0.333–0.883)	0.014
Smoking	1.628 (1.083–2.446)	0.019	1.458 (0.911–2.334)	0.116
SBP (mmHg)	1.011 (1.001–1.020)	0.026	1.013 (1.002–1.024)	0.016
Hcy (μmol/L)	1.056 (1.026–1.090)	0.001	1.048 (1.018–1.080)	0.002
ε3-ε4 genotype	2.210 (1.263–3.867)	0.005	2.794 (1.539–5.072)	0.002
ε4 allele	2.126 (1.265–3.575)	0.004	2.740 (1.569–4.784)	0.001

Table 6 The Comparison of Fasting Lipid Concentration Between *APOE* ε4 Carriers and ε3/ε3 Genotypes

Group	Control		CHD		CHD-IS	
	ε3-ε3 (n=157)	ε4 carriers (n=27)	ε3-ε3 (n=210)	ε4 carriers (n=68)	ε3-ε3 (n=106)	ε4 carriers (n=48)
TC (mmol/L)	3.807 (0.765)	4.407 (0.916)*	4.023 (0.945)	5.377 (1.808)**	4.195 (0.886)	4.700 (1.024)***
TG (mmol/L)	1.395 (0.838)	1.615 (0.602)	1.311 (0.834)	1.390 (0.919)	1.467 (0.779)	1.599 (0.810)
HDL (mmol/L)	1.224 (0.794)	1.161 (0.354)	1.345 (0.863)	1.201 (0.347)	1.310 (0.885)	1.252 (0.386)
LDL (mmol/L)	2.722 (0.877)	3.564 (0.810)*	2.563 (0.796)	3.625 (1.297)**	2.513 (1.299)	2.903 (0.790)

Notes: *APOE* ε4 carriers = ε3-ε4, ε4-ε4, ε2-ε4. Independent sample t-test test was applied to compare between ε3/ε3 and *APOE* ε4 carriers. Data were expressed as mean ±SD. *p-value <0.05 in comparison between ε3/ε3 and ε4 carriers in the control group. **p-value <0.05 in comparison between ε3/ε3 and ε4 carriers in the CHD group. ***p-value <0.05 in comparison between ε3/ε3 and ε4 carriers in the CHD-IS group.

the CHD-IS group respectively. Our results were consistent with previous studies. The ε3-ε4 genotype had an increasing trend, accounting for 11.2% in the control group, 19.6% in the CHD group and 24.9% in the CHD-IS group, respectively. While the frequency of ε2-ε2 genotype and the ε2 allele was lower in the three groups. The ε3-ε4 genotype and ε4 allele had significantly higher prevalence in the CHD group compared with control group ($p=0.006$ and $p=0.007$, respectively). Compared with the control group, the ε3-ε4 genotype and ε4 allele in the CHD-IS group had significantly higher prevalence ($p=0.001$ and $p=0.005$, respectively). After being adjusted for age, sex, smoking history, SBP and Hcy, the multi-variable binary logistic regression analysis showed that ε3-ε4 genotype was 2.21-fold higher risk ($p=0.005$), while ε4 allele caused 2.126-fold higher risk ($p=0.004$) for the development of CHD. ε3-ε4 genotype demonstrated 2.794-fold higher risk ($p=0.002$) while ε4 allele caused 2.74-fold higher risk ($p=0.001$) for the development of IS in CHD patients treated with medium-intensity statins. These data suggested that ε3-ε4 genotype and ε4 allele were the independent risk factors for the development of CHD and CHD-IS. Medium-intensity statins therapy did

not completely prevent IS in CHD patients with the ε3-ε4 genotype or the ε4 allele.

The TC and LDL-C levels were significantly elevated in the CHD and CHD-IS groups compared with the control group ($p<0.001$ and $p=0.015$, respectively) (Table 1). The levels of TC and LDL-C were lower in the CHD-IS group than that in the CHD group, which could have resulted from more use of medium-intensity statins in the CHD-IS group, but there was no significant difference ($p=0.241$ and $p=0.139$, respectively). *APOE* gene polymorphism could play important role in lipid metabolism.²⁵ It was mainly the different frequencies of the ε4 allele that led to significant differences in *APOE* distribution among these groups. These distributions of the ε4 allele frequencies could cause the differences in plasma lipid levels. Although patients of the CHD-IS group continuously took medium-intensity statins when they were diagnosed with IS, the differences of plasma lipid levels still existed because of ε4 carriers and ε3-ε3 genotype among the three groups (Table 6). In CHD group, the levels of TC and LDL-C of ε4 carriers were significantly higher than those of ε3-ε3 genotype. In the CHD-IS group, there was no significant difference in LDL-C level between ε4 carriers and ε3-ε3 genotype, but TC level of ε4 carriers

increased significantly. In this study, the patients of CHD-IS group were limited to taking medium-intensity statins, and $\epsilon 4$ allele was still associated with higher plasma TC level when compared with $\epsilon 3$ - $\epsilon 3$ genotype.

There were also some limitations in this study. Firstly, the sample size in this study was limited. Secondly, in order to more truly study the effect of CHD treatment on the incidence rate of IS, we chose the most routine dose of statin in practical application. While other studies suggested that intensive-dose statin treatment could be more favorable for reducing the incidences of all strokes.²⁶

Our study supported that the $\epsilon 4$ allele was an independent risk factor for the development of CHD by affecting the plasma lipid levels, and the main genotype was $\epsilon 3$ - $\epsilon 4$. The $\epsilon 4$ allele and $\epsilon 3$ - $\epsilon 4$ genotype were firstly found to be the major predictors for development of IS in CHD patients treated with medium-intensity statins. Therefore, high-intensity statins would be needed for CHD patients who carried $\epsilon 4$ allele to prevent the development of IS more effectively.

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