

Apolipoprotein E: A Genetic Bridge Between Premature Atherosclerosis and Alzheimer Disease

Apolipoprotein E (APOE) represents one of the most important common genetic determinants linking coronary artery disease (CAD) and neurodegeneration. ApoE links lipid metabolism, vascular health, and brain function. The three common APOE alleles, $\epsilon 2$, $\epsilon 3$, and $\epsilon 4$, encode functionally different isoforms with important cardiovascular and neurological consequences.^{1,2} Table 1 depicts the ApoE alleles and related clinical consequences.

Peripheral and brain ApoE biology should be considered separately. In the circulation, ApoE is present mainly on triglyceride-rich lipoproteins and their remnants and on a subset of high-density lipoprotein (HDL) particles. It facilitates receptor-mediated lipoprotein clearance and influences plasma lipid concentrations. Large-scale human data show a graded relationship between APOE genotype, low-density lipoprotein (LDL) cholesterol, and coronary risk; $\epsilon 2$ is generally associated with lower LDL-cholesterol and lower coronary risk, whereas $\epsilon 4$ is associated with higher LDL-cholesterol and a modest increase in coronary risk across the genotype spectrum.^{1,2}

In this issue, Dogsom et al³ report an exploratory hospital-based case-control study of 63 Mongolian patients with angiographically confirmed CAD and 61 controls. The study provides data from a population poorly represented in cardiovascular genetic research. The strongest finding was lower odds of CAD among individuals with the $\epsilon 2/\epsilon 3$ genotype (OR 0.29, 95% CI 0.12-0.71). Although $\epsilon 4/\epsilon 4$ was more frequent among cases, its association with CAD was not statistically significant (OR 2.68, 95% CI 0.68-10.55). Thus, the study does not establish $\epsilon 4$ as a CAD risk factor specifically in Mongolians, but provides hypothesis-generating data requiring confirmation.³ The small genotype groups also limit multivariable precision and preclude important sex-specific analyses. In addition, non-HDL-cholesterol or ApoB would also have provided a more complete assessment of atherogenic lipoprotein burden.

Population-specific evaluation is particularly important for APOE. Allele frequencies and effect sizes vary across ancestries. In Alzheimer disease (AD), the effect associated with $\epsilon 4$ differs substantially across ancestry groups and is influenced by global and local genetic ancestry and gene-environment interactions.⁴ This supports generating APOE data from Mongolian and other understudied populations.

The study also illustrates why lipid values obtained after CAD has developed require cautious interpretation. Total cholesterol and triglycerides were lower among cases, while LDL-cholesterol was higher.³ Detailed lipid-lowering treatment data were unavailable, so these differences should not simply be attributed to therapy. More importantly, a single lipid value does not represent lifetime exposure. This is relevant in premature atherosclerosis, where cumulative atherogenic exposure and inherited susceptibility are central considerations.⁵

EDITORIAL COMMENT

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Table 1. Cardiovascular and Neurological Effects of Common APOE Alleles

Feature	APOE ϵ 2	APOE ϵ 3	APOE ϵ 4
Peripheral lipid phenotype	Generally lower LDL-C; altered remnant lipoprotein metabolism	Reference phenotype	Generally higher LDL-C and total cholesterol
Coronary atherosclerosis	Generally lower coronary risk than ϵ 3; ϵ 2/ ϵ 2 may predispose to dysbetalipoproteinemia in susceptible individuals	Reference phenotype	Modestly higher coronary risk in large population studies
Premature CAD	Possible protective association, but evidence is limited	Reference phenotype	ϵ 3/ ϵ 4 associated with increased premature CAD risk in a recent observational case-control study
Brain ApoE and lipid transport	ApoE-containing HDL-like particles participate in CNS cholesterol transport; isoform-specific effects differ	Reference ApoE function	Altered ApoE lipidation and cellular cholesterol handling
Alzheimer disease	Associated with lower AD risk than ϵ 3	Reference phenotype	Strongest common genetic risk allele for late-onset AD; clear gene-dose effect
Selected brain effects	Generally associated with lower AD risk; mechanistic effects remain context dependent	Reference phenotype	Amyloid- β pathology, altered astrocyte lipid handling, microglial lipid accumulation, neuroinflammation, BBB dysfunction, and tau-related pathways
Effect of ancestry	Allele frequency and effect size vary among populations	Allele frequency varies among populations	Allele frequency varies across populations; APOE-related AD risk is strongly modified by global and local genetic ancestry, while cardiovascular associations also show population heterogeneity
Clinical interpretation	Context dependent; generally associated with lower LDL-C and AD risk, but ϵ 2/ ϵ 2 may predispose to dysbetalipoproteinemia	Most common reference isoform	Risk allele, not a deterministic diagnosis; effect depends on ancestry and other risk factors

AD, Alzheimer disease; ApoE, apolipoprotein E protein; APOE, apolipoprotein E gene; BBB, blood-brain barrier; CAD, coronary artery disease; CNS, central nervous system; HDL, high-density lipoprotein; LDL-C, low-density lipoprotein cholesterol.

The relationship between APOE ϵ 4 and premature CAD also deserves attention. Li et al⁶ reported that ϵ 3/ ϵ 4 was independently associated with premature CAD in a case-control study. This is consistent with broader evidence linking ϵ 4 to higher LDL-cholesterol and coronary risk.^{1,2} APOE genotyping is not part of routine premature CAD assessment, but unexpectedly early or severe atherosclerosis should prompt evaluation for inherited lipid disorders and other genetic contributors.⁵

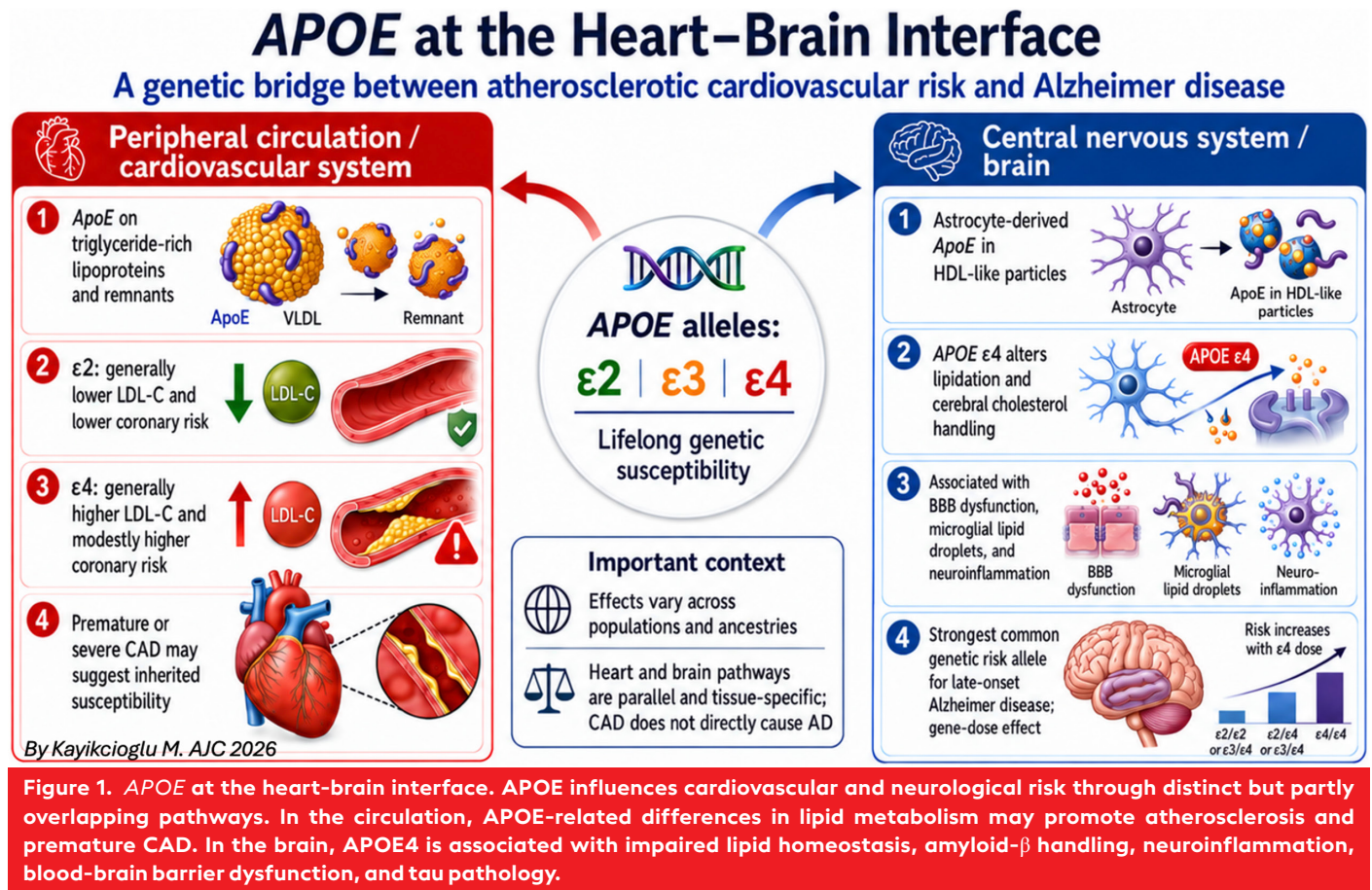
Brain cholesterol metabolism is largely compartmentalized from plasma lipoprotein metabolism. Under physiological conditions, ApoE is produced predominantly by astrocytes and secreted in HDL-like lipoprotein particles that redistribute cholesterol and phospholipids among brain cells.^{1,7}

The ϵ 4 allele alters several aspects of central nervous system lipid homeostasis. Human induced pluripotent stem cell-derived astrocytes carrying ϵ 4/ ϵ 4 show altered ApoE lipidation and reduced neurotrophic support.⁸ APOE ϵ 4 has also been associated with blood-brain barrier dysfunction and pericyte injury.⁹ Human brain and cellular studies have identified an APOE4-associated microglial phenotype characterized by triglyceride-rich lipid-droplet accumulation and enhanced inflammatory and neurotoxic responses.¹⁰ These findings connect disturbed lipid handling, inflammation, and neurodegeneration.

Clinically, APOE ϵ 4 is suggested as the strongest common genetic risk allele for late-onset AD, with a clear gene-dose effect.⁴ Fortea et al¹¹ showed that ϵ 4 homozygotes have a highly predictable age-related pattern of AD biomarker abnormalities and proposed that ϵ 4 homozygosity may represent a distinct genetic form of AD. This does not mean that every ϵ 4 homozygote will inevitably develop clinical dementia.

Thus, APOE links cardiovascular and neurological disease through parallel, tissue-specific pathways rather than through a simple sequence in which atherosclerosis causes AD (Figure 1). For cardiologists, the practical message remains prevention. Genetic susceptibility conferred by APOE is lifelong, while many cardiovascular risk factors are modifiable. Early and sustained control of LDL cholesterol and other atherogenic lipoproteins, blood pressure, and diabetes, together with smoking avoidance, weight management, and regular physical activity, remains fundamental to the prevention of premature atherosclerosis. Whether these interventions specifically attenuate APOE4-related AD risk remains uncertain. Importantly, higher physical activity has been associated with lower AD risk across APOE ϵ 4 carrier status and polygenic risk levels, suggesting that lifestyle remains relevant even with increased genetic susceptibility.¹²

The study by Dogsom et al³ opens a broader discussion than its modest size might suggest. APOE is not simply another



lipid-related gene; it is a biological link between lifelong lipid metabolism, premature atherosclerotic susceptibility, brain lipid homeostasis, and neurodegeneration.

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