Letter to the Editor

The Association between Apolipoprotein E Polymorphism and Diabetic Nephropathy in Iranian Patients

To the Editor,

We read with great interest the recently published article by Karimoei et al evaluating the association between apolipoprotein E (ApoE) polymorphism and diabetic nephropathy (DN) in a selected population of Iranian patients with diabetes mellitus (DM).¹ DN is the main cause of end-stage renal disease which affects approximately 35% of patients with DM.² While several predisposing risk factors including hyperglycemia, duration of DM, smoking, hypertension, age, 3,4 and a positive family history have been identified,⁵ much of the mechanism by which DN develops in patients with DM remains unknown.⁶ As research in this area continues, much focus has been devoted to the genetic basis in recent vears.⁷ ApoE is a plasma protein which contributes in lipid metabolism. ApoE gene is located on chromosome 19q and has 3 major alleles (2, 3, and 4) which are located on exon 4 of this gene.^{7,8} Several studies have investigated the association between ApoE polymorphism and development of DN; however, their findings were conflicting.⁷ There have been multiple studies showing association between 2 allele and development of DN;⁸⁻¹⁵ whereas, other studies have shown no correlation.¹⁶⁻¹⁹ Even more controversy surrounded 4 allele, as some studies have indicated 4 allele as a protective factor;^{8,14,20,21} while other studies have suggested this allele to be a risk factor of DN.^{17,19,22} Karimoei et al

in their study have shown protective effects of 4 allele on DN and found no association between 2 allele and DN in Iranian patients.¹ However, we believe that before making such a conclusion an important methodological issue should be considered.

In the index study, the age and duration of DM were significantly higher in patients with DN than patients without DN (61.5 ± 8.5 vs. 57.3 ± 8.1 , P < 0.05 and 12.01 \pm 7.6 vs. 10.2 \pm 6.9, P < 0.05, respectively).¹ It is well-established that both age and duration of DM are two risk factors of DN.^{3,4} In fact when two study groups were not matched for these two factors the final conclusion may be misleading. Because by passing the time, probably, a number of patients with DM may eventually develop DN and this issue can make bias on allele frequency between the two groups. In other words, it is possible that some patients initially categorized as DM group without DN will eventually transfer to DN group over the ensuing years; hence, both age and DM duration adjustment are required to minimize the "bias of group transformation."

In other similar studies in the literature, this issue has been appreciated and the two study groups are consistently matched for age and duration of DM.^{8,11,16,18,20,22} Given these concerns, the results reported by Karimoei et al will be more convincing after they recalculate the association between ApoE polymorphism and DN, after age and duration of DM are adjusted.

Conflict of interest: None declared.

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