Author’s response to reviews

Title: Association of rs662799 in APOA5 with CAD in Chinese Han population

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Dear Editor,

We have studied the valuable comments from you carefully, and revised the manuscript. The point to point responds to your comments are listed as following:

(i) Page 5, lines 1-15 (Sequencing Data Analysis and SNP calling).

All of the text in this section has been taken directly from the following paper: A novel three-round multiplex PCR for SNP genotyping with next generation sequencing. This section needs to be re-written and attribution to this paper referenced accordingly.

The sentences have been revised in Methods page 5(Sequencing Data Analysis).

(ii) Page 6, lines 37-43 (Despite the merits of our analysis, there are limitations that must be considered........diagnosis may vary due to differences in the inclusion of CHD cases and non-CHD controls). These lines have been taken Positive Association between APOA5 rs662799 Polymorphism and Coronary Heart Disease: A Case-Control Study and Meta-Analysis. Please re-write this portion. In particular, some of the text in that section is not
correct with respect to your manuscript, as publication and language bias is only an issue with meta analyses (which the above paper is). Your paper is not a meta analysis.

We have re-written this portion in Discussion page 6 (There are limitations to the study. Our analysis only includes the study of Chinese Han population, and the sample size is small. So, it might not apply to other populations.)