Association of the Common Genetic Variant Upstream of *INSIG2* Gene with Obesity Related Phenotypes in Chinese Children and Adolescents¹

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Objective To study the association between the rs7566605 variant of *INSIG2* and obesity-related phenotypes in Chinese children and adolescents. **Methods** The study sample consisted of two independent cohorts of Chinese children and adolescents. Anthropometric indices, lipids, blood pressure, fasting glucose, insulin and percentage of fat mass were determined. PCR with restriction fragment length polymorphism analysis was performed for genotyping the rs7566605 variant. **Results** In each of the two independent cohorts, no significant association was observed between rs7566605 and obesity under additive, dominant or recessive model. We also did not detect any difference in the genotype frequency between all the obese children and controls. Furthermore, we did not find evidence of an association between body composition indices and metabolic phenotypes in all children. However, the triglyceride level of CC homozygotes was significant trend of severe obesity in a *post-hoc* test. **Conclusion** *INSIG2* rs7566605 variant is not associated Chinese children.

Key words: INSIG2 gene; Obesity-related phenotypes; Children and adolescents

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