

Association between rs155971 in the *PCSK1* gene and the lipid profile of obese Thai children: a family-based study

S. Kulanuwat¹, J. Santiprabhob², B. Phonrat³, C. Limwongse⁴, A. Tungtrongchitr⁵, N. Chongviriyaphan⁶ and R. Tungtrongchitr¹

¹Department of Tropical Nutrition & Food Science, Faculty of Tropical Medicine, Mahidol University, Bangkok, Thailand

²Division of Endocrinology & Metabolism, Department of Pediatrics, Faculty of Medicine Siriraj Hospital, Mahidol University, Bangkok, Thailand ³Department of Clinical Tropical Medicine, Faculty of Tropical Medicine, Mahidol University, Bangkok, Thailand

⁴Division of Molecular Genetics, Department of Research & Development, Faculty of Medicine Siriraj Hospital, Mahidol University, Bangkok, Thailand ⁵Department of Parasitology, Faculty of Medicine Siriraj Hospital, Mahidol University, Bangkok, Thailand

⁶Department of Pediatrics, Faculty of Medicine Ramathibodi Hospital, Mahidol University, Bangkok, Thailand

Corresponding author: R. Tungtrongchitr E-mail: rungsunn.tun@mahidol.ac.th

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ABSTRACT. Genetic variants of the *POMC* and *PCSK1* genes cause severe obesity among patients in the early stages of childhood. This family-based study analyzed the links between single nucleotide polymorphisms (SNPs) in either the *POMC* or *PCSK1* genes and obesity, as well as obesity-related traits among obese Thai children and their families. The variants rs1042571 and rs6713532 in the *POMC* gene in a sample of 83 obese children and their family members were investigated using polymerase chain reaction (PCR)-restriction

fragment length polymorphism. In addition, the SNPs rs6232, rs155971, rs3762986, rs3811942, and rs371897784 of PCSK1 were analyzed in all samples using PCR and gene sequencing methods. Participants with the homozygous variant genotype in rs155971 had significantly elevated cholesterol and low-density lipoprotein cholesterol (LDL-C) levels (P = 0.011, OR = 1.025, 95%CI = 1.006-1.045; and P = 0.006, OR = 1.030, 95%CI = 1.009-1.053, respectively) after adjustment for age, gender, and body mass index (BMI). In addition, patients with the heterozygous variant genotype in rs371897784 of PCSK1 had a 1.249fold higher risk (95%CI = 1.081-1.444, P = 0.027) of increased waist circumference than patients with the normal genotype, after adjustment for age, gender, and BMI. However, this analysis did not find any correlation between obesity and SNPs in PCSK1 and POMC. Therefore, these common variants in PCSK1 and POMC were not the major cause of obesity in the Thai subjects sampled. However, variants in PCSK1 did affect cholesterol level, LDL-C level, and waist circumference.

Key words: *POMC* gene; *PCSK1* gene; Genetic variant; Obesity; Body mass index; Single nucleotide polymorphism