

Association of Ghrelin Gene Polymorphisms with Obesity in Japanese Children

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Background

Recently, ghrelin (GHRL) has attracted attention as a hormone connected with energy metabolism and appetite. The relationship between ghrelin gene polymorphisms and obesity has been analyzed in adults, but the influence of these SNPs on childhood obesity is uncertain.

Objective

We performed SNPs analysis of the GHRL gene and examined its relationship with childhood obesity.

			Re	esul	ts						
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		Mean ± SD	Range								
	n (M / F)	44 (31 / 13)			A-604G		TT, 35			CT, S	9
	Age (years)	11.4 ± 2.9	(5.9-17.8)								
	Total cholesterol (mg/dL)	180.8 ± 33.4	(123-266)								
	LDL-cholesterol (mg/dL)	109.6 ±31.2	(55.8-191.6)								
	HDL-cholesterol (mg/dL)	52.7 ±13.4	(24-91.7)								
	Triglycerides (mg/dL)	129.6 ±76.4	(29-346)		C-501A		TT, 37			GT,	7
	obesity index (%)	49.1 ±13.4	(21.9-79.3)								
	BMI (kg/m²)	27.3 ± 4.3	(19.4-37.2)								
	BMI Z- score (SD)	2.2 ± 0.4	(1.5-3.0)							TT.	2
	Fasting glucose (mg/dL)	95.9 ±12.4	(82-133)						ОТ	10	
	Fasting insulin (µU/mL)	32.9 ±45.5	(2.2-244)		C247A		GG, 26		GT,	16	
	HOMA-IR	8.8 ±14.5	(0.5-80.1)								
	HbA1c (NGSP) (%)	5.6 ±0.6	(4.1-8.4)		0%	10% 20%	30% 40% 50%	60%	70% 80)% 90%	100%

Population and Methods

We analyzed 44 patients (31 boys, 13 girls) treated in our clinic from 2010 to 2014. The average age at the first visit was 11.4 years old (range, 5.9–17.8 years) old) (Table 1).

We selected three SNPs of the ghrelin gene: g.A-604G (rs27647), g.C-501A (rs26802), and g.C247A (rs696217, Leu72Met) (Fig.1). The SNPs were genotyped using the ABI 7500 Fast Real-Time PCR System and Taqman[®] SNP Genotyping Assays.

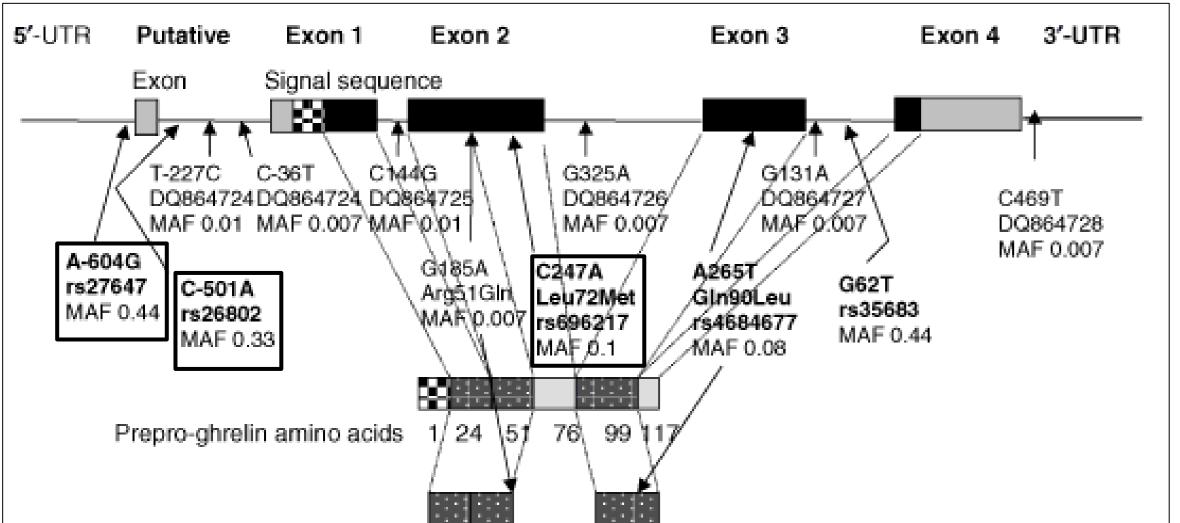


Table 1. Clinical characteristics of obese children in Japan (Kyoto).

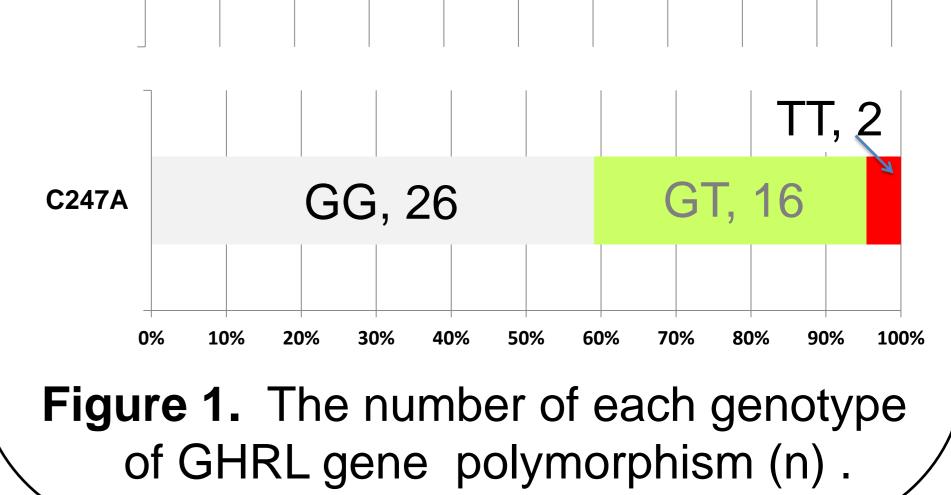


Table 2. Association between GHRL gene polymorphisms and clinical parameters on obese children (n=44) in Japan (P-value).

SNP	Age	T cho	LDL- cho	HDL- cho	TG	obesity index	BMI	BMI z	BS	insulin	HbA1c	HOMA-IR
A-604G,												
rs27647 ^a	0.1	0.3	0.3	0.4	0.8	0.4	0.7	0.3	0.8	0.4	0.4	0.4
C-501A,												
rs26802 ^a	0.2	0.4	0.8	0.1	0.7	0.2	0.6	0.1	0.2	0.4	0.7	0.4
C247A,												
rs696217 ^b	0.7	0.002	0.03	0.2	0.8	0.6	0.7	0.9	0.6	0.5	0.4	0.5



Obestatin peptide

Ghrelin gene (GHRL, chromosome 3p25.3) (#1)

Discussion

- Our study suggested that g.C247A (rs696217, Leu72Met) of GHRL gene polymorphism was related to hypercholesterolemia in obese Japanese children (Table 2, Fig. 2).
- The relationship between the SNPs of GHRL gene and HOMA-IR index was not seen in Japanese childhood obesity, although g.A-604G and g.C247A (Leu72Met) of GHRL polymorphism were related to insulin resistance in Caucasian adults (#2,#3,#4, Fig.3). The probability that childhood obesity could translate to adulthood obesity is known to be 40-85%. The patient having the specified SNPs of GHRL gene may show insulin resistance when they grow up. The ratio of these SNPs almost accorded with the ratio of East Asian people (#5, Fig1). But because there is a race difference, we cannot discuss these

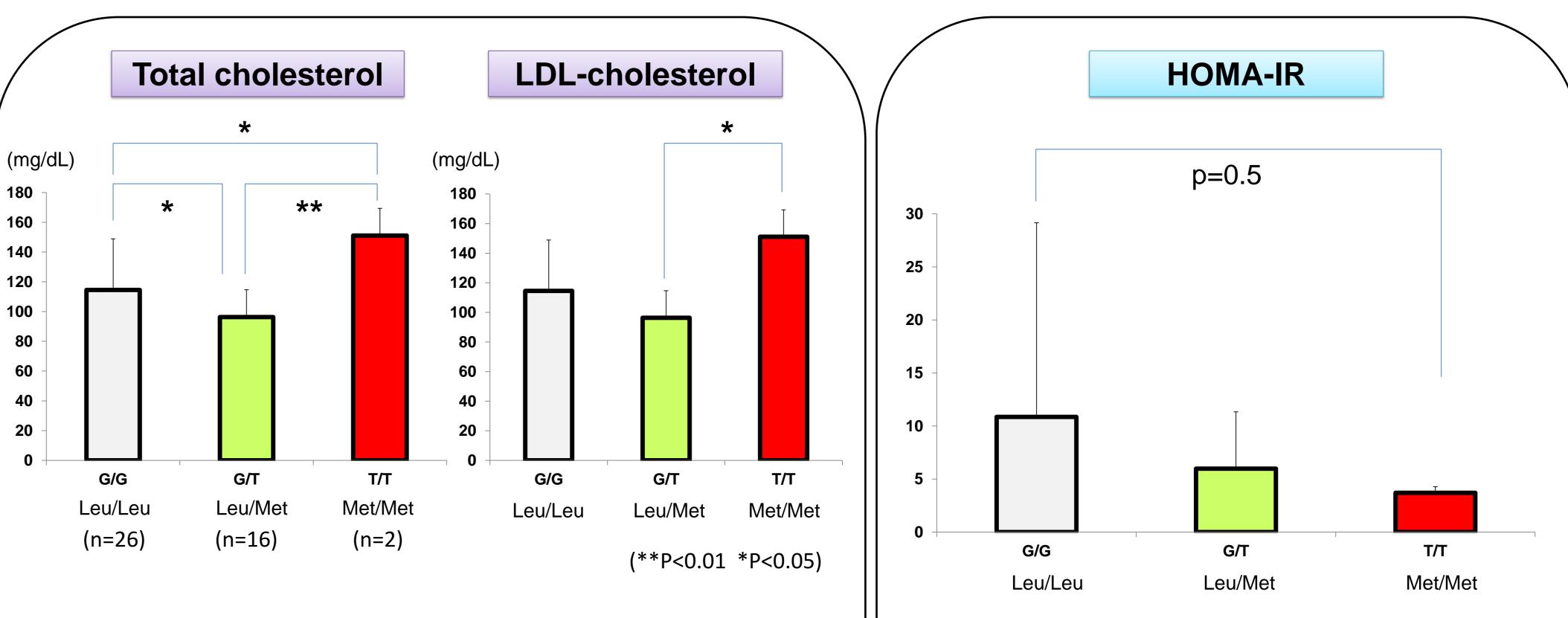


Figure 2. Total cholesterol and LDL- cholesterol according to g.C247A (Leu72Met) genotypes.

Figure 3. HOMA-IR according to g.C247A (Leu72Met) genotypes.

SNPs among all races.

Conclusions

Our study suggests that Met/Met of g.C247A (rs696217, Leu72Met) could be a risk factor for hypercholesterolemia in childhood obesity in Japan. • We could not find the relationship between BMI, BMI-Z score and GHRL gene polymorphisms. In addition, the relationship between insulin resistance and GHRL gene polymorphisms was not observed. • We are planning to examine the relationship between diet and GHRL gene polymorphisms in childhood obesity on a larger sample size for definitive conclusion.

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References

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