Additional file 2. Association within or nearby linkage region[a] (P<10^{-6} & LOD≤1.9)

	Ethnic Group	EA: European Americans		AA: African Americans	MA: Mexican Americans		JA: Japanese Americans
PUBLISHED LINKAGE RESULTS	1-LOD chromosomal region (Trait & LOD of linkage analysis)	2q12.1-2q13 (Weight, LOD=2.91)		16p13.2-12.1 (BPfactor, LOD=2.25)	3p26 (Insulin-Glucose, Blood Pressure factors, LOD=2.20)(1) (Serum uric acid, LOD=4.9)(2)		5q33.1-5q34 (log-Waist, LOD=1.98)
ASSOCIATION RESULTS	trait	Weight	Weight	log(TG)	log(HDL)	log(HDL)	Fasting Glucose
	Chromosomal Region	2q14.2		16p13.13	3p26.3-p26.2		5q23.3-5q31.1[b]
	Chr	2	2	16	3	3	5
	Bp	121836875	121843978	11562798	2004251	2001175	130581195
	rsid	rs139940998[c]	rs144756634[c]	rs551107164	rs17005939	rs12631510	rs6875363
	Gene	GLI2-TFCP2L1		AK126539	CNTN6-CNTN4		LYRM7-CDC42SE2
	Function	intergenic	intergenic	ncRNA_intronic	intergenic	intergenic	intergenic
	A1	A	A	G	C	C	T
	A2	G	T	A	T	T	C
	Freq[§]	0.0087	0.0089	0.011	0.4186	0.4375	0.6857
	Freq GENNID[‖]	0.01	0.01	0.01	0.35	0.35	0.4421
	β	1.84	1.84	1.41	-0.08	-0.08	-0.54263
	SE	0.32	0.32	0.26	0.02	0.02	0.109372
	P	7.34E-09	7.34E-09	7.61E-08	8.02E-08	9.13E-08	1.38E-07
	I^2 (% heterogeneity)	-	-	-	80.7%	79.2%	0%[h]

[a]Candidate linkage regions for EA, AA and JA were from publication analyses (3-7)
[b]Region with evidence of association is very close to previously nominated 1-LOD interval but not within it
[c]Bolded variants were significantly associated with weight (P<$5x10^{-8}$).
[f]Frequency of A1 allele in gnomAD (8) : NFE (Non-Finnish Europeans) for EA, EAS (East Asians) for JA, AMR (Latinos) for MA, and AFR (African/African Americans) for AA
[g] Frequency of A1 allele in GENNID as estimated by GCTA (9)
[h] I^2 = -51.6% and was to set 0% based on Higgins et al **(**10**)**

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