

**Supplementary Table 1.** Surrogates used in this study (derived from the CEU HapMap)

CHR	SNP on our Illumina Infinium™ II HumanHap550 BeadChip	SNP used in other system	r <sup>2</sup>	Genes Nearby
1	rs2793831	rs10923931	1	NOTCH2
3	rs4411878	rs4607103	1	ADAMTS9
6	rs4712523	rs10946398	1	CDKAL2
7	rs1635852	rs864745	0.965	JAZF1
9	rs2383208	rs10811661	1	CDKN2A/B
10	rs11257622	rs12779790	0.833	CDC123-CAMK1D
11	rs729287	rs3740878	1	EXT2
11	rs163171	rs151290	0.951	KCNQ1
12	rs1353362	rs7961581	0.973	TSPAN8-LGR5

**Supplementary Table 2.** Effect of the G allele rs7923837 at the *HHEX-IDE* locus on pediatric BMI by gender.

N: number of individuals tested; Effect Size: regression coefficient for the test SNP; SE: standard error of the regression coefficient; Test statistic: additive model; P: unadjusted two-sided trend test P-value.

	N	Effect Size	SE	Test Statistic	P
<b>Discovery Cohort</b>					
<b>MALE</b>	1857	0.04092	0.03993	1.025	0.306
<b>FEMALE</b>	1735	0.1472	0.04051	3.633	0.000288
<b>Replication Cohort</b>					
<b>MALE</b>	1946	0.08304	0.03964	2.095	0.0363
<b>FEMALE</b>	1646	0.0436	0.04138	1.054	0.292
<b>Combined Cohort</b>					
<b>MALE</b>	3803	0.06225	0.02813	2.213	0.0270
<b>FEMALE</b>	3381	0.09741	0.02895	3.365	0.000773

**Supplementary Table 3.** Effect of the G allele rs7923837 at the *HHEX-IDE* locus on pediatric BMI by age bin in the combined cohort.

N: number of individuals tested; Effect Size: regression coefficient for the test SNP; SE: standard error of the regression coefficient; Test statistic: additive model; P: unadjusted two-sided trend test P-value.

	N	Effect Size	SE	Test Statistic	P
<b>Age Bin (yrs old)</b>					
<b>2 to 6</b>	2244	0.1205	0.03616	3.332	0.000877
<b>7 to 11</b>	1847	0.07482	0.04192	1.785	0.07446
<b>12 to 18</b>	3093	0.04473	0.02896	1.545	0.1225

**Supplementary Table 4.** Effect of the G allele rs7923837 at the *HHEX-IDE* locus on pediatric BMI by individual ages in the combined cohort.

N: number of individuals tested; Effect Size: regression coefficient for the test SNP; SE: standard error of the regression coefficient; Test statistic: additive model; P: unadjusted two-sided trend test *P*-value.

<b>AGE</b>	<b>N</b>	<b>Effect Size</b>	<b>SE</b>	<b>Test Statistic</b>	<b>P</b>
<b>2</b>	555	0.04397	0.0743	0.5917	0.554
<b>3</b>	540	0.1594	0.06834	2.332	0.0201
<b>4</b>	419	0.1585	0.08553	1.853	0.0646
<b>5</b>	351	0.1216	0.09438	1.289	0.198
<b>6</b>	379	0.1269	0.08714	1.456	0.146
<b>7</b>	374	0.2323	0.09286	2.502	0.0128
<b>8</b>	367	0.1742	0.09794	1.778	0.0762
<b>9</b>	340	-0.09863	0.09608	-1.027	0.305
<b>10</b>	369	-0.01515	0.09316	-0.1626	0.871
<b>11</b>	397	0.08216	0.08903	0.9228	0.357
<b>12</b>	420	0.00608	0.08068	0.07536	0.94
<b>13</b>	480	-0.08932	0.0769	-1.162	0.246
<b>14</b>	503	0.1643	0.06957	2.362	0.0186
<b>15</b>	547	0.02947	0.06593	0.447	0.655
<b>16</b>	508	0.1319	0.06776	1.947	0.0521
<b>17</b>	386	0.06315	0.08567	0.7371	0.462
<b>18</b>	249	-0.07207	0.1037	-0.695	0.488