

Supplementary Table 1. Distribution of plasma sOB-R levels and other characteristics among the study population

Characteristics	N	Value ^a
Age (yr)	1,504	56.2 (6.9)
BMI (kg/m ²)	1,504	28.0 (5.6)
WHR	1,022	0.80 (0.07)
Physical activity (MET-hr/week)	1,487	15.8 (17.8)
Alcohol (g/day)	1,488	5.4 (8.7)
Smoking status (%)	1,504	
Never smoked		719 (47.8)
Past smoker		618 (41.9)
Current smoker		167 (11.1)
Menopausal status (%) ^b	1,504	
Pre-menopause		335 (22.3)
Post-menopause, never used hormone		467 (31.5)
Post-menopause, hormone past user		196 (13.0)
Post-menopause, hormone current user		506 (33.6)
Family history of diabetes (%)	1,504	477 (31.7)
Type 2 diabetes (%)	1,504	684 (45.5)
Fasting status (%)	1,504	968 (64.4)
Total energy (kcal/day)	1,488	1800 (518)
Glycemic load	1,488	96.1 (15.8)
Cereal fiber (g/day)	1,488	4.0 (1.7)
Polyunsaturated to saturated fat ratio	1,488	0.51 (0.13)
Trans fat (g/day)	1,488	1.9 (0.5)
sOB-R (ng/mL)	1,504	31.4 (10.5)
Leptin (ng/mL)	1,504	25.0 (14.3)

^aMean (SD) for continuous variables; n (%) for categorical variables. ^bHormone refers to hormone replacement therapy.

Supplementary Table 2. Top genotyped and imputed SNPs ($P < 5 \times 10^{-8}$) for plasma sOB-R^a

SNP	Position (bp) ^b	Gene	Gene region	Alleles (+/-)	MAF ^c	r^2	Beta ^d	s.e.	P^d
rs2767485	65751505	<i>LEPR</i>	Intron	T/C	0.17	0.9986	0.098	0.012	5.77E-15
rs12079231	65743743	<i>LEPR</i>	Intron	C/G	0.17	0.9999	0.097	0.012	6.88E-15
rs11208656	65743375	<i>LEPR</i>	Intron	A/G	0.17	1	0.097	0.012	7.11E-15
rs12062820	65743083	<i>LEPR</i>	Intron	T/C	0.17	1	0.097	0.012	7.11E-15
rs1782755	65759392	<i>LEPR</i>	Intron	C/A	0.17	1	0.097	0.012	8.66E-15
rs1627238	65758667	<i>LEPR</i>	Intron	G/A	0.17	0.9999	0.096	0.012	8.88E-15
rs1171275	65755221	<i>LEPR</i>	Intron	C/T	0.17	0.9995	0.096	0.012	1.13E-14
rs1180445	65755507	<i>LEPR</i>	Intron	T/C	0.17	0.9995	0.096	0.012	1.13E-14
rs1171278	65760734	<i>LEPR</i>	Intron	C/T	0.17	0.9786	0.096	0.012	2.42E-14
rs1171276	65760037	<i>LEPR</i>	Intron	A/G	0.19	0.9348	0.095	0.012	5.43E-14
rs1751492	65765213	<i>LEPR</i>	Intron	T/C	0.30	1	-0.076	0.010	4.00E-13
rs1186403	65773121	<i>LEPR</i>	Intron	T/C	0.30	1	-0.076	0.010	4.40E-13
rs2154381	65768289	<i>LEPR</i>	Intron	A/G	0.30	1	-0.076	0.010	4.61E-13
rs1171266	65776412	<i>LEPR</i>	Intron	C/T	0.30	0.9925	-0.075	0.010	1.35E-12
rs1171274	65753426	<i>LEPR</i>	Intron	T/C	0.27	0.9984	-0.076	0.011	3.62E-12
rs7516840	65745183	<i>LEPR</i>	Intron	G/C	0.27	0.9992	-0.076	0.011	3.68E-12
rs10889556	65742140	<i>LEPR</i>	Intron	A/G	0.27	1	-0.076	0.011	3.74E-12
rs10889558	65749554	<i>LEPR</i>	Intron	G/A	0.27	0.9984	-0.076	0.011	3.79E-12
rs1327116	65748639	<i>LEPR</i>	Intron	C/A	0.27	0.9985	-0.076	0.011	3.80E-12
rs7533108	65750432	<i>LEPR</i>	Intron	G/C	0.27	0.9983	-0.076	0.011	3.83E-12
rs12068561	65746541	<i>LEPR</i>	Intron	C/T	0.27	0.9988	-0.076	0.011	3.84E-12
rs10889557	65743292	<i>LEPR</i>	Intron	G/A	0.28	0.9587	-0.076	0.011	3.94E-12
rs4655680	65814057	<i>LEPR</i>	Intron	G/T	0.29	0.9884	-0.074	0.011	4.03E-12
rs6697315	65766144	<i>LEPR</i>	Intron	T/C	0.36	1	-0.070	0.010	5.30E-12
rs3828039	65826144	<i>LEPR</i>	Intron	A/G	0.28	0.9764	-0.074	0.011	5.57E-12
rs11208674	65811905	<i>LEPR</i>	Intron	T/C	0.29	1	-0.073	0.010	5.66E-12
rs12033452	65811023	<i>LEPR</i>	Intron	C/A	0.29	1	-0.073	0.010	5.66E-12
rs4655518	65812446	<i>LEPR</i>	Intron	T/G	0.29	1	-0.073	0.010	5.66E-12
rs6657632	65811254	<i>LEPR</i>	Intron	A/G	0.29	1	-0.073	0.010	5.66E-12
rs2104564	65742018	<i>LEPR</i>	Intron	T/C	0.27	1	-0.076	0.011	5.87E-12
rs12042884	65825780	<i>LEPR</i>	Intron	C/G	0.28	0.9771	-0.074	0.011	5.89E-12
rs1782754	65765936	<i>LEPR</i>	Intron	A/G	0.27	0.9992	-0.075	0.011	6.71E-12
rs12042877	65825736	<i>LEPR</i>	Intron	C/T	0.28	0.9784	-0.074	0.011	6.76E-12
rs12028951	65825669	<i>LEPR</i>	Intron	G/A	0.28	0.9788	-0.073	0.011	7.08E-12
rs4370791	65824816	<i>LEPR</i>	Intron	A/G	0.28	0.9807	-0.073	0.011	7.29E-12
rs1475398	65755845	<i>LEPR</i>	Intron	G/C	0.27	1	-0.075	0.011	8.01E-12
rs11208657	65745941	<i>LEPR</i>	Intron	A/G	0.28	0.9610	-0.075	0.011	8.09E-12

rs1171270	65770429	<i>LEPR</i>	Intron	A/G	0.27	0.9995	-0.075	0.011	8.12E-12
rs1171269	65769390	<i>LEPR</i>	Intron	C/T	0.27	0.9996	-0.075	0.011	8.18E-12
rs1171271	65771378	<i>LEPR</i>	Intron	T/C	0.27	0.9995	-0.075	0.011	8.19E-12
rs1177681	65771707	<i>LEPR</i>	Intron	A/G	0.27	0.9996	-0.075	0.011	8.19E-12
rs1022981	65772622	<i>LEPR</i>	Intron	A/G	0.27	0.9997	-0.075	0.011	8.20E-12
rs1171272	65772399	<i>LEPR</i>	Intron	T/G	0.27	0.9996	-0.075	0.011	8.20E-12
rs6676495	65824705	<i>LEPR</i>	Intron	G/A	0.28	0.9845	-0.073	0.011	8.88E-12
rs1171262	65774334	<i>LEPR</i>	Intron	T/A	0.27	0.9984	-0.075	0.011	9.86E-12
rs1171263	65774521	<i>LEPR</i>	Intron	C/T	0.27	0.9968	-0.075	0.011	1.09E-11
rs1171264	65775602	<i>LEPR</i>	Intron	G/C	0.27	0.9945	-0.074	0.011	1.48E-11
rs1171267	65776442	<i>LEPR</i>	Intron	G/T	0.36	0.9857	-0.069	0.010	1.59E-11
rs12038998	65821716	<i>LEPR</i>	Intron	C/A	0.28	0.9946	-0.072	0.011	1.59E-11
rs4655528	65822077	<i>LEPR</i>	Intron	A/G	0.28	0.9946	-0.072	0.011	1.62E-11
rs11208676	65820711	<i>LEPR</i>	Intron	A/G	0.28	0.9945	-0.072	0.011	1.69E-11
rs11208677	65821005	<i>LEPR</i>	Intron	A/G	0.28	0.9944	-0.072	0.011	1.72E-11
rs10493379	65818515	<i>LEPR</i>	Intron	G/A	0.28	1	-0.072	0.011	1.83E-11
rs11208679	65822326	<i>LEPR</i>	Intron	G/A	0.28	0.9998	-0.072	0.011	1.83E-11
rs6691346	65818985	<i>LEPR</i>	Intron	G/A	0.28	0.9999	-0.072	0.011	1.83E-11
rs11208675	65818012	<i>LEPR</i>	Intron	T/G	0.28	0.9999	-0.072	0.011	1.83E-11
rs1343981	65817916	<i>LEPR</i>	Intron	A/G	0.28	0.9999	-0.072	0.011	1.83E-11
rs1343982	65817839	<i>LEPR</i>	Intron	C/T	0.28	0.9999	-0.072	0.011	1.83E-11
rs3790424	65816601	<i>LEPR</i>	Intron	A/G	0.28	0.9999	-0.072	0.011	1.83E-11
rs1938487	65819855	<i>LEPR</i>	Intron	T/C	0.28	0.9998	-0.072	0.011	1.84E-11
rs3790425	65815700	<i>LEPR</i>	Intron	A/G	0.28	1	-0.071	0.011	1.92E-11
rs1171261	65773990	<i>LEPR</i>	Intron	C/T	0.27	1	-0.074	0.011	2.08E-11
rs6696954	65827583	<i>LEPR</i>	Intron	G/T	0.37	0.8718	0.072	0.011	2.30E-11
rs1409802	65793939	<i>LEPR</i>	Intron	G/A	0.27	0.9861	-0.074	0.011	3.01E-11
rs1171265	65775840	<i>LEPR</i>	Intron	G/A	0.37	0.8990	-0.070	0.010	3.70E-11
rs6676419	65824797	<i>LEPR</i>	Intron	C/T	0.50	0.9662	-0.063	0.010	4.48E-11
rs6673324	65803651	<i>LEPR</i>	Intron	A/G	0.50	0.9932	0.063	0.009	4.55E-11
rs7519977	65806084	<i>LEPR</i>	Intron	G/A	0.27	0.9975	-0.072	0.011	6.97E-11
rs4655517	65812013	<i>LEPR</i>	Intron	T/C	0.50	0.9987	0.062	0.009	7.13E-11
rs10158279	65806284	<i>LEPR</i>	Intron	G/T	0.50	1	0.062	0.009	7.46E-11
rs10789184	65809550	<i>LEPR</i>	Intron	G/A	0.27	1	-0.072	0.011	9.24E-11
rs1137100	65809029	<i>LEPR</i>	Missense (Lys109Arg)	A/G	0.27	1	-0.072	0.011	9.24E-11
rs3790427	65815338	<i>LEPR</i>		T/C	0.27	0.9982	-0.071	0.011	1.02E-10
rs10789186	65823065	<i>LEPR</i>	Intron	G/A	0.50	0.9779	-0.062	0.009	1.06E-10
rs10889563	65821755	<i>LEPR</i>	Intron	A/G	0.49	0.9795	-0.061	0.009	1.11E-10

rs6673591	65820977	<i>LEPR</i>	Intron	A/G	0.49	0.9827	-0.061	0.009	1.18E-10
rs10789185	65818452	<i>LEPR</i>	Intron	G/C	0.49	0.9907	-0.061	0.009	1.49E-10
rs10889568	65832033	<i>LEPR</i>	Intron	T/C	0.26	0.9970	-0.070	0.011	1.56E-10
rs12405556	65835705	<i>LEPR</i>	Intron	G/T	0.26	1	-0.070	0.011	1.58E-10
rs4655539	65832251	<i>LEPR</i>	Intron	A/T	0.44	0.9692	-0.061	0.010	1.77E-10
rs10732836	65832048	<i>LEPR</i>	Intron	C/T	0.44	0.9705	-0.061	0.010	1.78E-10
rs6669117	65833660	<i>LEPR</i>	Intron	T/C	0.44	0.9684	-0.061	0.010	1.78E-10
rs4655537	65831389	<i>LEPR</i>	Intron	G/A	0.39	0.9086	0.066	0.010	1.85E-10
rs1137101	65831101	<i>LEPR</i>	Missense (Gln223Arg)	A/G	0.44	0.9752	-0.061	0.010	1.91E-10
rs2154380	65830029	<i>LEPR</i>	Intron	T/C	0.44	0.9777	-0.061	0.010	2.04E-10
rs11208682	65829184	<i>LEPR</i>	Intron	G/A	0.44	0.9766	-0.061	0.010	2.06E-10
rs10736402	65830588	<i>LEPR</i>	Intron	C/T	0.44	0.9759	-0.061	0.010	2.06E-10
rs10449758	65829705	<i>LEPR</i>	Intron	A/G	0.44	0.9786	-0.061	0.010	2.08E-10
rs12564626	65829130	<i>LEPR</i>	Intron	G/A	0.44	0.9740	-0.061	0.010	2.13E-10
rs10749754	65827228	<i>LEPR</i>	Intron	G/A	0.44	0.9660	-0.061	0.010	2.34E-10
rs1475397	65755746	<i>LEPR</i>	Intron	C/T	0.26	1	0.069	0.011	2.92E-10
rs10889567	65829638	<i>LEPR</i>	Intron	T/C	0.45	1	-0.059	0.009	3.10E-10
rs10157275	65739091	<i>LEPR</i>	Intron	C/T	0.13	0.9917	0.087	0.014	5.09E-10
rs1782763	65780488	<i>LEPR</i>	Intron	T/C	0.33	0.9858	-0.065	0.010	5.77E-10
rs10158579	65722644	<i>LEPR</i>	Intron	T/C	0.13	0.9996	0.086	0.014	5.95E-10
rs11808888	65725369	<i>LEPR</i>	Intron	G/A	0.13	0.9988	0.086	0.014	6.17E-10
rs9436302	65669149	<i>LEPR</i>	Intron	G/A	0.20	0.8588	0.080	0.013	6.75E-10
rs17127673	65728313	<i>LEPR</i>	Intron	A/G	0.14	1	0.085	0.014	6.80E-10
rs10128072	65729684	<i>LEPR</i>	Intron	A/C	0.13	1	0.086	0.014	7.64E-10
rs7602	65670539	<i>LEPR</i>	Intron	G/A	0.20	0.9984	0.072	0.012	8.57E-10
rs17127677	65730862	<i>LEPR</i>	Intron	G/T	0.13	0.9989	0.085	0.014	8.59E-10
rs11800275	65736292	<i>LEPR</i>	Intron	T/A	0.13	0.9979	0.085	0.014	9.49E-10
rs6694528	65735604	<i>LEPR</i>	Intron	C/T	0.13	0.9979	0.085	0.014	9.49E-10
rs17127690	65735537	<i>LEPR</i>	Intron	A/G	0.13	0.9980	0.085	0.014	9.52E-10
rs9436303	65669262	<i>LEPR</i>	Intron	A/G	0.24	0.8981	0.070	0.012	2.53E-09
rs1171279	65761081	<i>LEPR</i>	Intron	C/T	0.26	0.9859	0.065	0.011	2.96E-09
rs6690625	65850178	<i>LEPR</i>	Intron	T/G	0.19	0.9591	-0.073	0.013	5.98E-09
rs12040007	65852748	<i>LEPR</i>	Intron	G/A	0.19	0.9589	-0.073	0.013	6.78E-09
rs4655556	65853151	<i>LEPR</i>	Intron	G/A	0.19	0.9591	-0.073	0.013	8.0E-09
rs1938484	65853870	<i>LEPR</i>	Intron	C/A	0.19	0.9590	-0.073	0.013	8.0E-09
rs9436300	65667852	<i>LEPR</i>	Intron	G/A	0.33	0.7046	-0.069	0.012	1.5E-08
rs7534511	65667718	<i>LEPR</i>	Intron	G/A	0.34	0.6901	-0.070	0.012	1.6E-08
rs12567606	65914472	NA	NA	A/G	0.22	0.9645	-0.067	0.012	1.7E-08

rs1046011	65671584	<i>LEPR</i>	Intron	C/T	0.30	0.9014	-0.062	0.011	2.0E-08
rs4655555	65852857	<i>LEPR</i>	Intron	T/A	0.22	0.8930	-0.069	0.012	2.3E-08
rs6667615	65935218	NA	NA	G/A	0.21	0.8965	-0.070	0.013	2.6E-08
rs6588156	65933021	NA	NA	G/A	0.21	0.9100	-0.069	0.012	2.6E-08
rs6667616	65935219	NA	NA	C/A	0.21	0.8877	-0.070	0.013	2.7E-08
rs4655583	65927995	NA	NA	C/A	0.22	0.9463	-0.067	0.012	2.7E-08
rs6664534	65932663	NA	NA	A/T	0.21	0.9192	-0.069	0.012	3.0E-08
rs9436301	65668515	<i>LEPR</i>	Intron	T/C	0.24	0.8046	0.069	0.012	3.0E-08
rs4655795	65931956	NA	NA	G/T	0.21	0.9222	-0.069	0.012	3.2E-08
rs4394621	65938424	NA	NA	A/G	0.21	0.8792	-0.070	0.013	3.4E-08
rs4655792	65931852	NA	NA	T/G	0.20	0.9456	-0.068	0.012	3.6E-08
rs12130476	65897438	NA	NA	G/C	0.20	0.9761	-0.068	0.012	3.7E-08
rs2186245	65890652	NA	NA	C/T	0.19	0.9787	-0.068	0.012	3.8E-08
rs4655793	65931866	NA	NA	C/A	0.21	0.9378	-0.068	0.012	3.9E-08
rs7518632	65875938	NA	NA	C/A	0.19	0.9894	-0.068	0.012	4.1E-08
rs3762274	65836701	<i>LEPR</i>	Intron	T/C	0.37	0.9232	-0.055	0.010	4.4E-08
rs1892535	65869769	<i>LEPR</i>	Intron	G/A	0.19	0.9936	-0.068	0.012	4.5E-08
rs12145237	65897470	NA	NA	C/T	0.20	0.9762	-0.067	0.012	4.6E-08
rs12043772	65907622	NA	NA	T/A	0.20	0.9860	-0.066	0.012	4.8E-08

^an = 1,504; sOB-R levels were transformed on a natural logarithm scale. ^bPosition based on NCBI build 36.3. ^cMinor allele frequency among study participants. ^dRegression coefficients and P values for every one copy of minor allele were estimated from linear regression models adjusted for age at blood draw, diabetes case-control status, fasting status, body mass index, menopausal status, and postmenopausal hormone use..

Supplementary Table 3. Top genotyped and imputed SNPs ($P < 1 \times 10^{-6}$) for plasma sOB-R^a among controls

SNP	Position (bp) ^b	Gene	Gene region	Alleles (+/-)	MAF ^c	Beta ^d	s.e.	P^d
rs2767485	65751505	<i>LEPR</i>	Intron	T/C	0.17	0.104	0.017	5.97E-10
rs12079231	65743743	<i>LEPR</i>	Intron	C/G	0.17	0.104	0.017	6.13E-10
rs11208656	65743375	<i>LEPR</i>	Intron	A/G	0.17	0.104	0.017	6.17E-10
rs12062820	65743083	<i>LEPR</i>	Intron	T/C	0.17	0.104	0.017	6.17E-10
rs1171275	65755221	<i>LEPR</i>	Intron	C/T	0.17	0.101	0.017	1.20E-09
rs1180445	65755507	<i>LEPR</i>	Intron	T/C	0.17	0.101	0.017	1.20E-09
rs1627238	65758667	<i>LEPR</i>	Intron	G/A	0.17	0.101	0.016	1.20E-09
rs1782755	65759392	<i>LEPR</i>	Intron	C/A	0.17	0.101	0.016	1.20E-09
rs1171278	65760734	<i>LEPR</i>	Intron	C/T	0.17	0.100	0.017	2.45E-09
rs1171276	65760037	<i>LEPR</i>	Intron	A/G	0.19	0.100	0.017	2.75E-09
rs2104564	65742018	<i>LEPR</i>	Intron	T/C	0.27	-0.091	0.016	8.69E-09
rs7516840	65745183	<i>LEPR</i>	Intron	G/C	0.27	-0.090	0.016	1.11E-08
rs10889556	65742140	<i>LEPR</i>	Intron	A/G	0.27	-0.090	0.016	1.13E-08
rs1171274	65753426	<i>LEPR</i>	Intron	T/C	0.27	-0.090	0.016	1.13E-08
rs10889558	65749554	<i>LEPR</i>	Intron	G/A	0.27	-0.090	0.016	1.14E-08
rs7533108	65750432	<i>LEPR</i>	Intron	G/C	0.27	-0.090	0.016	1.14E-08
rs1327116	65748639	<i>LEPR</i>	Intron	C/A	0.27	-0.090	0.016	1.14E-08
rs12068561	65746541	<i>LEPR</i>	Intron	C/T	0.27	-0.090	0.016	1.14E-08
rs1475398	65755845	<i>LEPR</i>	Intron	G/C	0.27	-0.089	0.016	1.80E-08
rs6697315	65766144	<i>LEPR</i>	Intron	T/C	0.36	-0.081	0.014	1.81E-08
rs1751492	65765213	<i>LEPR</i>	Intron	T/C	0.30	-0.084	0.015	1.85E-08
rs11208657	65745941	<i>LEPR</i>	Intron	A/G	0.28	-0.088	0.016	2.07E-08
rs10889557	65743292	<i>LEPR</i>	Intron	G/A	0.28	-0.088	0.016	2.52E-08
rs1186403	65773121	<i>LEPR</i>	Intron	T/C	0.30	-0.083	0.015	2.80E-08
rs2154381	65768289	<i>LEPR</i>	Intron	A/G	0.30	-0.083	0.015	2.80E-08
rs1171265	65775840	<i>LEPR</i>	Intron	G/A	0.37	-0.082	0.015	3.89E-08
rs1171267	65776442	<i>LEPR</i>	Intron	G/T	0.36	-0.079	0.014	4.92E-08
rs1171266	65776412	<i>LEPR</i>	Intron	C/T	0.30	-0.082	0.015	5.36E-08
rs1782754	65765936	<i>LEPR</i>	Intron	A/G	0.27	-0.086	0.016	5.76E-08
rs1171270	65770429	<i>LEPR</i>	Intron	A/G	0.27	-0.085	0.016	7.82E-08
rs1171269	65769390	<i>LEPR</i>	Intron	C/T	0.27	-0.085	0.016	7.84E-08
rs1171271	65771378	<i>LEPR</i>	Intron	T/C	0.27	-0.085	0.016	7.88E-08
rs1177681	65771707	<i>LEPR</i>	Intron	A/G	0.27	-0.085	0.016	7.88E-08
rs1022981	65772622	<i>LEPR</i>	Intron	A/G	0.27	-0.085	0.016	7.89E-08
rs1171272	65772399	<i>LEPR</i>	Intron	T/G	0.27	-0.085	0.016	7.89E-08
rs1171262	65774334	<i>LEPR</i>	Intron	T/A	0.27	-0.085	0.016	8.21E-08
rs1171263	65774521	<i>LEPR</i>	Intron	C/T	0.27	-0.085	0.016	8.59E-08

rs4655680	65814057	<i>LEPR</i>	Intron	G/T	0.29	-0.081	0.015	1.04E-07
rs1171264	65775602	<i>LEPR</i>	Intron	G/C	0.27	-0.084	0.016	1.05E-07
rs11208674	65811905	<i>LEPR</i>	Intron	T/C	0.29	-0.080	0.015	1.07E-07
rs12033452	65811023	<i>LEPR</i>	Intron	C/A	0.29	-0.080	0.015	1.07E-07
rs4655518	65812446	<i>LEPR</i>	Intron	T/G	0.29	-0.080	0.015	1.07E-07
rs6657632	65811254	<i>LEPR</i>	Intron	A/G	0.29	-0.080	0.015	1.07E-07
rs1171261	65773990	<i>LEPR</i>	Intron	C/T	0.27	-0.084	0.016	1.27E-07
rs3828039	65826144	<i>LEPR</i>	Intron	A/G	0.28	-0.080	0.015	2.00E-07
rs12042884	65825780	<i>LEPR</i>	Intron	C/G	0.28	-0.080	0.015	2.10E-07
rs1782763	65780488	<i>LEPR</i>	Intron	T/C	0.33	-0.078	0.015	2.27E-07
rs1409802	65793939	<i>LEPR</i>	Intron	G/A	0.27	-0.082	0.016	2.38E-07
rs12042877	65825736	<i>LEPR</i>	Intron	C/T	0.28	-0.079	0.015	2.39E-07
rs12028951	65825669	<i>LEPR</i>	Intron	G/A	0.28	-0.079	0.015	2.44E-07
rs4370791	65824816	<i>LEPR</i>	Intron	A/G	0.28	-0.079	0.015	2.46E-07
rs7519977	65806084	<i>LEPR</i>	Intron	G/A	0.27	-0.081	0.016	2.75E-07
rs6676495	65824705	<i>LEPR</i>	Intron	G/A	0.28	-0.079	0.015	2.78E-07
rs12405556	65835705	<i>LEPR</i>	Intron	G/T	0.26	-0.081	0.016	2.81E-07
rs10789184	65809550	<i>LEPR</i>	Intron	G/A	0.27	-0.081	0.016	2.98E-07
rs1137100	65809029	<i>LEPR</i>	Missense (Lys109Arg)	A/G	0.27	-0.081	0.016	2.98E-07
rs3790427	65815338	<i>LEPR</i>	Intron	T/C	0.27	-0.081	0.016	3.23E-07
rs10889568	65832033	<i>LEPR</i>	Intron	T/C	0.26	-0.080	0.016	3.25E-07
rs10493379	65818515	<i>LEPR</i>	Intron	G/A	0.28	-0.076	0.015	5.12E-07
rs11208675	65818012	<i>LEPR</i>	Intron	T/G	0.28	-0.076	0.015	5.12E-07
rs11208679	65822326	<i>LEPR</i>	Intron	G/A	0.28	-0.076	0.015	5.12E-07
rs1343981	65817916	<i>LEPR</i>	Intron	A/G	0.28	-0.076	0.015	5.12E-07
rs1343982	65817839	<i>LEPR</i>	Intron	C/T	0.28	-0.076	0.015	5.12E-07
rs1938487	65819855	<i>LEPR</i>	Intron	T/C	0.28	-0.076	0.015	5.12E-07
rs3790424	65816601	<i>LEPR</i>	Intron	A/G	0.28	-0.076	0.015	5.12E-07
rs3790425	65815700	<i>LEPR</i>	Intron	A/G	0.28	-0.076	0.015	5.12E-07
rs6691346	65818985	<i>LEPR</i>	Intron	G/A	0.28	-0.076	0.015	5.12E-07
rs12038998	65821716	<i>LEPR</i>	Intron	C/A	0.28	-0.077	0.015	5.28E-07
rs4655528	65822077	<i>LEPR</i>	Intron	A/G	0.28	-0.077	0.015	5.28E-07
rs11208676	65820711	<i>LEPR</i>	Intron	A/G	0.28	-0.076	0.015	5.39E-07
rs11208677	65821005	<i>LEPR</i>	Intron	A/G	0.28	-0.076	0.015	5.44E-07
rs17127673	65728313	<i>LEPR</i>	Intron	A/G	0.14	0.091	0.018	7.94E-07
rs7602	65670539	<i>LEPR</i>	Intron	G/A	0.20	0.079	0.016	8.80E-07
rs10157275	65739091	<i>LEPR</i>	Intron	C/T	0.13	0.091	0.018	9.50E-07

^asOB-R levels were log-transformed. n = 820. ^bPosition based on NCBI build 36.3. ^cMinor allele frequency among study participants. ^dRegression coefficients and P values for every one copy of minor allele were estimated from linear regression models adjusted for age at blood draw, fasting status, body mass index, menopausal status, and postmenopausal hormone use.

Supplementary Table 4. Top genotyped and imputed SNPs (combined $P < 5 \times 10^{-8}$) for plasma sOB-R^a in a meta-analysis pooling results of cases and controls.

SNP	Gene	Gene region	Left Gene	Right Gene	Alleles (+/-)	z score ^b	P ^c	Direction ^d
rs2767485	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	7.883	3.19E-15	++
rs12079231	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/G	7.861	3.83E-15	++
rs12062820	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	7.856	3.98E-15	++
rs11208656	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	7.856	3.98E-15	++
rs1782755	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/A	7.838	4.56E-15	++
rs1627238	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	7.835	4.68E-15	++
rs1171275	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	7.800	6.20E-15	++
rs1180445	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	7.800	6.20E-15	++
rs1171278	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	7.708	1.27E-14	++
rs1171276	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	7.574	3.61E-14	++
rs1751492	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.524	5.30E-14	--
rs2154381	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.483	7.27E-14	--
rs1186403	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.480	7.45E-14	--
rs1171266	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-7.323	2.42E-13	--
rs6697315	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.274	3.50E-13	--
rs7516840	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/C	-7.202	5.93E-13	--
rs10889556	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.200	6.02E-13	--
rs1171274	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.199	6.07E-13	--
rs12068561	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-7.196	6.21E-13	--
rs10889558	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-7.194	6.29E-13	--
rs1327116	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/A	-7.194	6.30E-13	--
rs7533108	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/C	-7.192	6.38E-13	--
rs10889557	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-7.167	7.67E-13	--
rs4655680	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/T	-7.165	7.76E-13	--
rs1782754	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.135	9.66E-13	--
rs2104564	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.132	9.87E-13	--
rs12033452	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/A	-7.110	1.16E-12	--
rs4655518	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/G	-7.110	1.16E-12	--
rs6657632	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.110	1.16E-12	--
rs11208674	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.110	1.16E-12	--
rs1171269	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.097	1.28E-12	--
rs1171270	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-7.096	1.28E-12	--
rs1022981	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-7.096	1.29E-12	--
rs1171271	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.096	1.29E-12	--
rs1171272	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-7.095	1.29E-12	--
rs1177681	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/G	-7.095	1.29E-12	--

rs3828039	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-7.094	1.31E-12	--	
rs12042884	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/G	-7.089	1.35E-12	--	
rs11208657	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-7.09	1.35E-12	--	
rs1475398	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/C	-7.083	1.41E-12	--	
rs1171267	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/T	-7.082	1.43E-12	--	
rs1171262	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/A	-7.068	1.57E-12	--	
rs12042877	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/T	-7.067	1.58E-12	--	
rs12028951	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-7.062	1.65E-12	--	
rs4370791	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-7.059	1.68E-12	--	
rs1171263	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-7.053	1.75E-12	--	
rs6676495	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-7.035	1.99E-12	--	
rs1171264	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/C	-7.008	2.41E-12	--	
rs6696954	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/T	-6.987	2.81E-12	--	
rs12038998	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/A	-6.965	3.27E-12	--	
rs4655528	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.964	3.31E-12	--	
rs11208676	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.957	3.48E-12	--	
rs11208677	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.955	3.52E-12	--	
rs1171265	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-6.954	3.54E-12	--	
rs1171261	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-6.945	3.79E-12	--	
rs10493379	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.936	4.02E-12	--	
rs11208679	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.936	4.02E-12	--	
rs6691346	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.936	4.02E-12	--	
rs11208675	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/G	-6.936	4.03E-12	--	
rs1343981	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-6.936	4.03E-12	--	
rs1343982	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.936	4.03E-12	--	
rs1938487	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>PDE4B</i>	C/T	-6.936	4.03E-12	--	
rs3790424	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.936	4.03E-12	--	
rs3790425	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>PDE4B</i>	A/G	-6.927	4.30E-12	--	
rs1409802	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-6.898	5.29E-12	--	
rs6673324	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	6.861	6.85E-12	++	
rs6676419	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/T	-6.819	9.15E-12	--	
rs4655517	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	6.787	1.14E-11	++	
rs10158279	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/T	6.783	1.18E-11	++	
rs7519977	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-6.760	1.38E-11	--	
rs10789184	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	-6.716	1.87E-11	--	
rs1137100	<i>LEPR</i>	Missense	(Lys109Arg)	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	-6.716	1.87E-11	--
rs3790427	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>PDE4B</i>	T/C	-6.713	1.91E-11	--	
rs10889563	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.711	1.93E-11	--	

rs10789186	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.701	2.07E-11	--
rs6673591	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.699	2.10E-11	--
rs10789185	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/C	-6.653	2.87E-11	--
rs12405556	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/T	-6.648	2.96E-11	--
rs10889568	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-6.648	2.98E-11	--
rs4655537	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.645	3.04E-11	++
rs4655539	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/T	-6.596	4.23E-11	--
rs6669117	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-6.595	4.25E-11	--
rs10732836	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/T	-6.594	4.29E-11	--
rs1137101	<i>LEPR</i>	Missense (Gln223Arg)	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.588	4.45E-11	--
rs2154380	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-6.576	4.83E-11	--
rs10736402	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/T	-6.575	4.86E-11	--
rs10449758	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	A/G	-6.575	4.88E-11	--
rs11208682	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.574	4.91E-11	--
rs12564626	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.571	5.00E-11	--
rs1782763	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	-6.553	5.62E-11	--
rs10749754	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.553	5.65E-11	--
rs10889567	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-6.511	7.49E-11	--
rs10157275	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	6.334	2.39E-10	++
rs10158579	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/C	6.309	2.81E-10	++
rs11808888	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/A	6.306	2.86E-10	++
rs10128072	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/C	6.276	3.48E-10	++
rs1475397	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	6.264	3.76E-10	++
rs17127677	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	G/T	6.258	3.90E-10	++
rs17127673	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	6.256	3.94E-10	++
rs11800275	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	T/A	6.243	4.31E-10	++
rs6694528	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	6.243	4.31E-10	++
rs17127690	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	A/G	6.242	4.33E-10	++
rs9436302	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	G/A	6.239	4.40E-10	++
rs7602	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	G/A	6.189	6.04E-10	++
rs6690625	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/G	-6.118	9.49E-10	++
rs12040007	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.098	1.08E-09	--
rs9436303	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	A/G	6.082	1.19E-09	++
rs4655556	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-6.078	1.22E-09	--
rs1938484	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/A	-6.073	1.26E-09	--
rs12567606	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-6.068	1.30E-09	--
rs6667616	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/A	-5.930	3.03E-09	--
rs6667615	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/A	-5.928	3.07E-09	--

rs6588156	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/A	-5.924	3.13E-09	--
rs4655583	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/A	-5.911	3.40E-09	--
rs6664534	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/T	-5.893	3.80E-09	--
rs4394621	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.891	3.84E-09	--
rs4655795	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/T	-5.883	4.02E-09	--
rs1171279	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	5.881	4.09E-09	++
rs12130476	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/C	-5.862	4.57E-09	--
rs4655792	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	T/G	-5.857	4.70E-09	--
rs2186245	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.856	4.74E-09	--
rs4655555	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/A	-5.855	4.76E-09	--
rs7518632	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/A	-5.846	5.03E-09	--
rs4655793	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/A	-5.845	5.07E-09	--
rs4425959	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.820	5.88E-09	--
rs12043772	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.816	6.03E-09	--
rs12145237	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	T/A	-5.816	6.03E-09	--
rs1892535	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	G/A	-5.810	6.24E-09	--
rs9436300	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	G/A	-5.797	6.74E-09	--
rs12025906	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-5.793	6.90E-09	--
rs11208720	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.789	7.08E-09	--
rs7534511	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	G/A	-5.785	7.23E-09	--
rs11208707	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.785	7.24E-09	--
rs10889576	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.785	7.26E-09	--
rs10736403	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.784	7.30E-09	--
rs11208710	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.783	7.32E-09	--
rs1046011	<i>LEPR</i>	Intron	<i>LEPROT</i>	<i>LOC100128633</i>	C/T	-5.783	7.34E-09	--
rs10749756	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.778	7.58E-09	--
rs12117858	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/T	-5.777	7.61E-09	--
rs12026918	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/G	-5.777	7.62E-09	--
rs4655779	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.775	7.69E-09	--
rs4655782	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/G	-5.774	7.76E-09	--
rs4655780	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	C/A	-5.772	7.84E-09	--
rs11208709	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/A	-5.772	7.85E-09	--
rs4655778	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/A	-5.771	7.87E-09	--
rs12042379	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	G/T	-5.764	8.21E-09	--
rs3828033	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	C/T	-5.690	1.27E-08	--
rs3762274	<i>LEPR</i>	Intron	<i>LOC100128633</i>	<i>PDE4B</i>	T/C	-5.689	1.28E-08	--
rs9436301	<i>LEPR</i>	Intron	<i>DNAJC6</i>	<i>LOC100128633</i>	T/C	5.669	1.44E-08	++
rs10158937	NA	NA	<i>LEPR</i>	<i>PDE4B</i>	A/C	-5.668	1.45E-08	--

^asOB-R levels were log-transformed. ^bAverage of *z* statistics corresponding to the *P* value of association for each single SNP in cases and controls, with weights proportional to the square root of the number of individuals in cases and controls, respectively. The *z* statistics were derived from a fixed-effects linear regression model. ^c*P* value for each SNP was based on the average *z* statistics. ^dDirection of association in cases or controls. “++” indicated that each copy of the reference allele was associated with increased sOB-R levels in both cases and controls; “--” indicated that each copy of the reference allele was associated with decreased sOB-R levels in both cases and controls; “+-” or “-+” indicated that the direction of associations was opposite for cases and controls.

Supplementary Table 5. Associations between selected SNPs and diabetes risk factors, as well as type 2 diabetes risk in the Nurses' Health Study

	SNPs										
	rs1137100 ^a		rs1137101 ^b		rs2767485 ^b		rs1751492 ^a		rs4655555 ^b		
N	Beta±s.e.	P	Beta±s.e.	P	Beta±s.e.	P	Beta±s.e.	P	Beta±s.e.	P	
BMI (kg/m ²)	3279	0.04±0.15	0.80	0.04±0.13	0.77	-0.22±0.17	0.20	-0.04±0.14	0.78	-0.18±0.16	0.28
WHR	2236	-0.01±0.004	0.17	-0.002±0.003	0.46	-0.006±0.004	0.18	-0.005±0.004	0.17	0.003±0.004	0.54
Leptin (ng/mL)	1508	-0.01±0.02	0.64	-0.006±0.02	0.74	0.03±0.02	0.21	0.01±0.02	0.59	-0.003±0.02	0.89
HMW adiponectin (μg/mL)	1506	-0.02±0.03	0.42	-0.02±0.03	0.33	0.02±0.03	0.49	-0.01±0.03	0.83	0.02±0.03	0.45
CRP (mg/dL)	830	-0.07±0.06	0.18	-0.05±0.05	0.34	-0.03±0.06	0.61	-0.10±0.05	0.07	0.17±0.06	0.005
TNF-αR2 (pg/mL)	843	-0.03±0.021	0.15	0.001±0.02	0.96	-0.06±0.02	0.02	-0.0±0.02	0.08	0.05±0.02	0.02
IL-6 (ng/mL)	815	-0.09±0.04	0.02	-0.07±0.03	0.04	-0.07±0.04	0.10	-0.07±0.04	0.05	0.08±0.04	0.06
IL-18 (pg/mL)	1505	-0.02±0.02	0.23	-0.01±0.02	0.50	0.02±0.02	0.39	0.001±0.02	0.97	0.04±0.02	0.02
Insulin (μU/mL)	604	0.04±0.05	0.45	0.05±0.04	0.25	-0.14±0.05	0.008	-0.01±0.04	0.75	-0.08±0.05	0.12
Type 2 diabetes ^c	3279	0.07±0.06	0.23	0.007±0.05	0.90	0.02±0.07	0.76	0.08±0.06	0.18	-0.07±0.07	0.30

^aGenotyped SNPs. ^bImputed SNPs. ^cBoth incident and prevalent cases were included. To convert beta ± s.e. to odds ratio (95% CI), exponentiate beta ± 1.96×s.e.

Supplementary Table 6. Genotyped SNPs that only reached nominal significance level ($P < 1 \times 10^{-5}$) for plasma sOB-R levels in the Nurses' Health Study^a

SNP	l h r	Position (bp) ^b	Gene	A lles (+/-) s	M A F ^c	Geometric LS mean (95% CI), ng/mL ^d			B eta ^d	s .e.	P value ^d
						0	1	2			
rs120 25906	5 0 3	658739 72	<i>LEPR</i>	T /C	0 .1 9	30.7 (30.2 -31.2)	28.7 (28.0 -29.3)	27.0 (25.0 -29.0)	-0.0 66	.0 12	0 5E-7
rs465 5754	5 0 3	658873 35	NA	T /C	0 .1 9	30.6 (30.1 -31.2)	28.7 (28.1 -29.4)	27.0 (25.0 -29.0)	-0.0 65	.0 12	0 8E-7
rs101 58937	5 0 0	659174 64	NA	C /A	0 .2 0	30.6 (30.1 -31.2)	28.9 (28.2 -29.5)	26.9 (25.1 -28.7)	-0.0 62	.0 12	0 1E-7
rs156 7214	5 0 3	692992 70	NA	G /C	0 .4 0	31.0 (30.3 -31.7)	29.7 (29.1 -30.2)	28.2 (27.3 -29.1)	-0.0 47	.0 10	0 5E-6
rs666 9354 ^e	5 0 4	656979 37	<i>LEPR</i>	T /G	0 .1 3	29.0 (23.6 -35.6)	31.0 (26.4 -37.9)	34.2 (27.8 -42.4)	0.0 0.069	.0 0.14	0 7E-6
rs109 97637 ^e	5 0 3	688636 07	<i>CTNN</i> <i>A3</i>	T /C	0 .0 .4	29.3 (24.0 -36.1)	32.8 (26.9 -40.0)	31.1 (28.7 -70.0)	0.0 116	.0 25	0 2E-6
rs652 7988	5 0	203007 89	NA	N A	0 .2 .6	29.2 (28.7	30.8 (30.1	32.0 (30.5	0.0 0.051	.0 .011	0 1E-6

							-29.7)	-31.5)	-33.7)			
rs100 7033	6 5 0 3	7 7 87	287198 <i>ACCN</i> <i>I</i>	G /A	0 .0 .6 0	29.6 (29.2 -30.0)	32.7 (31.3 -34.2)	31.1 (25.3 -38.8)	0. 093 020	0 0 0	4.5 2E-6	
rs112 08654	5 0 3	657381 54	<i>LEPR</i>	T /C	0 .3 .3 0	30.9 (30.3 -31.5)	29.2 (28.7 -29.8)	28.5 (27.3 -29.8)	0.0 47	0 0 0	4.7 1E-6	
rs943 6747	5 0 4	656841 95	<i>LEPR</i>	T /C	0 .3 .8	31.1 (30.4 -31.8)	29.4 (28.8 -29.9)	28.6 (27.5 -29.7)	0.0 46	0 0 0	4.7 8E-6	
rs311 8667	4 4 9	135280 884	<i>ADAM</i> <i>TS13</i>	T /C	0 .4 .7	31.3 (30.6 -32.1)	29.6 (29.1 -30.2)	28.8 (28.0 -29.6)	0.0 44	0 0 0	5.2 6E-6	
rs595 0304	5 0 4	203033 16	NA	N A	0 .2 .6	29.2 (28.7 -29.7)	30.8 (30.1 -31.5)	32.1 (30.5 -33.7)	0. 050	0 0 0	6.5 6E-6	
rs671 7199	5 0 3	196040 109	NA	T /G	0 .4 .0	31.0 (30.3 -31.8)	29.7 (29.2 -30.3)	28.3 (27.4 -29.2)	0.0 45	0 0 0	8.0 5E-6	
rs202 5804	5 0 2	657187 09	<i>LEPR</i>	T /C	0 .3 .3	30.9 (30.3 -31.5)	29.2 (28.7 -29.8)	28.6 (27.4 -29.8)	0.0 46	0 0 0	8.3 3E-6	
rs171 77643	5 0 5	196083 765	NA	T /C	0 .3 .9	30.9 (30.2 -31.6)	29.8 (29.2 -30.3)	28.2 (27.3 -29.1)	0.0 45	0 0 0	8.9 1E-6	
rs109	5	689446	<i>CTNN</i>	G	0 .0	29.3	32.5	70.0	0. 0	0 0	9.3	

97688 ^e	0 3	0	82	A3	/A	4	(24.0 -36.0)	(26.3 -39.0)	(70.0 -70.0)	108	24	6E-6
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^asOB-R levels were log-transformed. ^bPosition based on NCBI build 36.3. ^cMinor allele frequency among study participants. ^dAdjusted for age at blood draw, diabetes case-control status, fasting status, body mass index, menopausal status, and postmenopausal hormone use. ^eLeast square means could not be obtained due to rare homozygous minor alleles (1.7% for rs6669354, 0.2% for rs10997637, and 0.1% for rs10997688). Median (interquartile range) was provided instead.

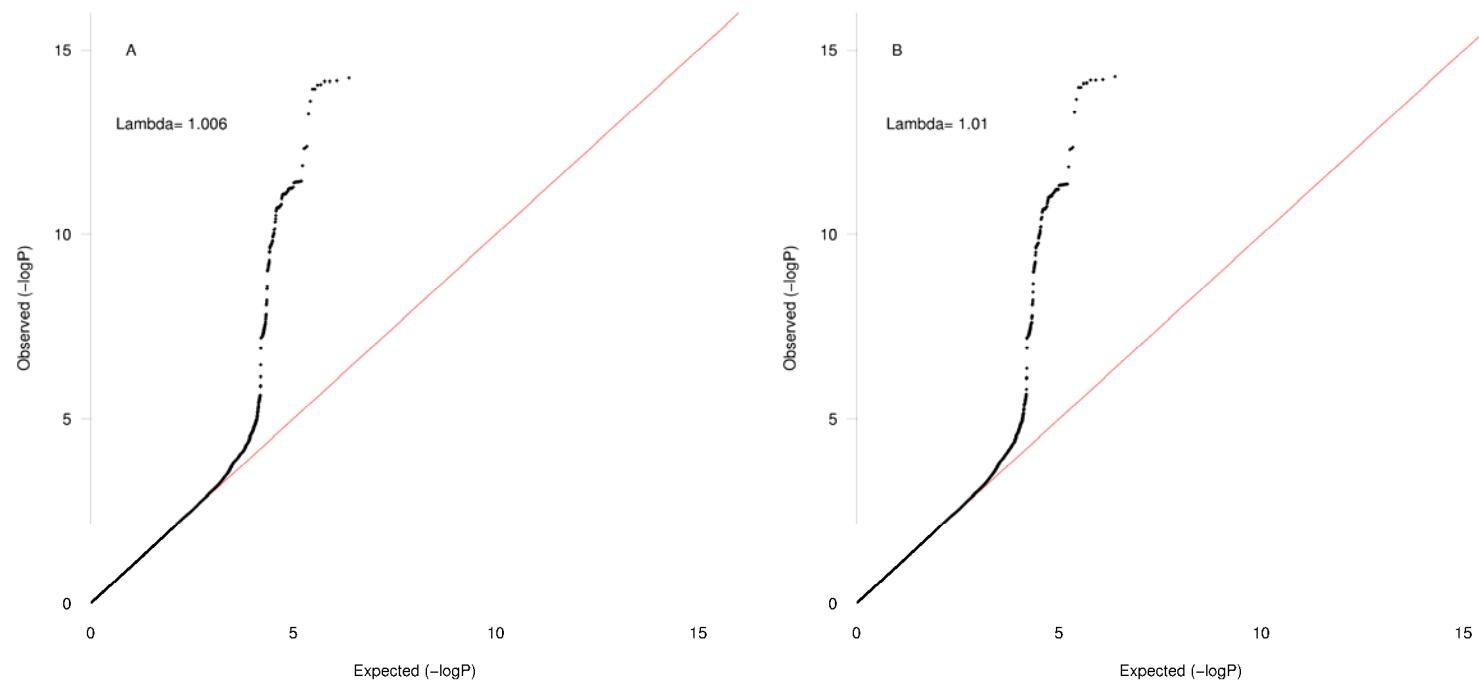
Supplementary Table 7. Genotyped and imputed SNPs that only reached nominal significance level ($P < 1 \times 10^{-5}$) for plasma sOB-R levels^a

SNP	Chr.	Position (bp) ^b	Gene	Gene region	Alleles (+/-)	MAF ^c	r ²	Beta ^d	s.e.	P ^d
rs11208720	1	65931643	NA	NA	A/G	0.20	0.9603	-0.067	0.012	5.0E-08
rs4425959	1	65940407	NA	NA	A/G	0.21	0.8548	-0.070	0.013	5.2E-08
rs3828033	1	65837298	<i>LEPR</i>	Intron	C/T	0.37	0.9137	-0.056	0.010	5.2E-08
rs12025906	1	65873972	<i>LEPR</i>	Intron	T/C	0.19	1	-0.067	0.012	5.3E-08
rs11208707	1	65912101	NA	NA	C/T	0.20	0.9960	-0.065	0.012	5.7E-08
rs11208710	1	65918627	NA	NA	A/G	0.20	0.9968	-0.065	0.012	5.8E-08
rs10736403	1	65912984	NA	NA	A/G	0.20	0.9961	-0.065	0.012	5.8E-08
rs12042379	1	65931052	NA	NA	G/T	0.20	0.9833	-0.066	0.012	5.8E-08
rs10749756	1	65913099	NA	NA	C/T	0.20	0.9962	-0.065	0.012	6.0E-08
rs12026918	1	65918834	NA	NA	C/G	0.20	0.9969	-0.065	0.012	6.0E-08
rs12117858	1	65914682	NA	NA	C/T	0.20	0.9962	-0.065	0.012	6.0E-08
rs4655779	1	65915118	NA	NA	A/G	0.20	0.9963	-0.065	0.012	6.1E-08
rs4655782	1	65915560	NA	NA	A/G	0.20	0.9965	-0.065	0.012	6.1E-08
rs11208709	1	65918605	NA	NA	G/A	0.20	0.9971	-0.065	0.012	6.1E-08
rs4655780	1	65915365	NA	NA	C/A	0.20	0.9966	-0.065	0.012	6.2E-08
rs4655778	1	65914924	NA	NA	G/A	0.20	0.9961	-0.065	0.012	6.2E-08
rs10889576	1	65941154	NA	NA	C/T	0.21	0.8306	-0.070	0.013	6.3E-08
rs10158937	1	65917464	NA	NA	A/C	0.20	0.9999	-0.064	0.012	1.2E-07
rs4655811	1	65695745	<i>LEPR</i>	Intron	G/C	0.32	0.9870	-0.053	0.010	3.4E-07
rs11800909	1	65734368	<i>LEPR</i>	Intron	A/G	0.08	0.9778	0.088	0.018	7.1E-07
rs1567214	1	69299270	NA	NA	G/C	0.40	1	-0.048	0.010	1.2E-06
rs12131840	1	66004359	NA	NA	C/T	0.15	0.9466	-0.070	0.015	1.3E-06
rs11208700	1	65906031	NA	NA	T/A	0.46	0.9013	0.048	0.010	2.2E-06
rs970468	1	65679078	<i>LEPR</i>	Intron	A/C	0.37	0.9350	-0.049	0.010	2.3E-06
rs6657868	1	65686295	<i>LEPR</i>	Intron	G/A	0.37	0.9956	-0.048	0.010	2.3E-06
rs4655783	1	65915632	NA	NA	C/T	0.46	0.9012	0.048	0.010	2.4E-06
rs11804091	1	65676467	<i>LEPR</i>	Intron	A/G	0.15	0.9893	0.063	0.013	2.4E-06
rs9436747	1	65684195	<i>LEPR</i>	Intron	C/T	0.37	1	-0.047	0.010	2.6E-06
rs12022410	1	65926521	NA	NA	G/A	0.46	0.9017	0.047	0.010	2.6E-06
rs6669354	1	65697937	<i>LEPR</i>	Intron	T/G	0.13	1	0.067	0.014	2.8E-06
rs1007033	17	28719887	<i>ACCN1</i>	Intron	C/T	0.06	0.9994	0.095	0.020	3.0E-06
rs1327121	1	65729925	<i>LEPR</i>	Intron	A/G	0.33	0.9997	-0.048	0.010	3.1E-06
rs11208654	1	65738154	<i>LEPR</i>	Intron	T/C	0.33	1	-0.048	0.010	3.1E-06
rs2025804	1	65718709	<i>LEPR</i>	Intron	A/G	0.33	1	-0.048	0.010	3.1E-06
rs10997637	10	68863607	<i>CTNNA3</i>	Intron	C/T	0.04	0.9995	0.115	0.025	3.3E-06
rs3790433	1	65666930	<i>LEPR</i>	Intron	C/T	0.24	0.7239	0.060	0.013	3.4E-06
rs13438451	7	14581455	<i>DGKB</i>	Intron	C/T	0.09	0.7836	0.083	0.018	3.5E-06

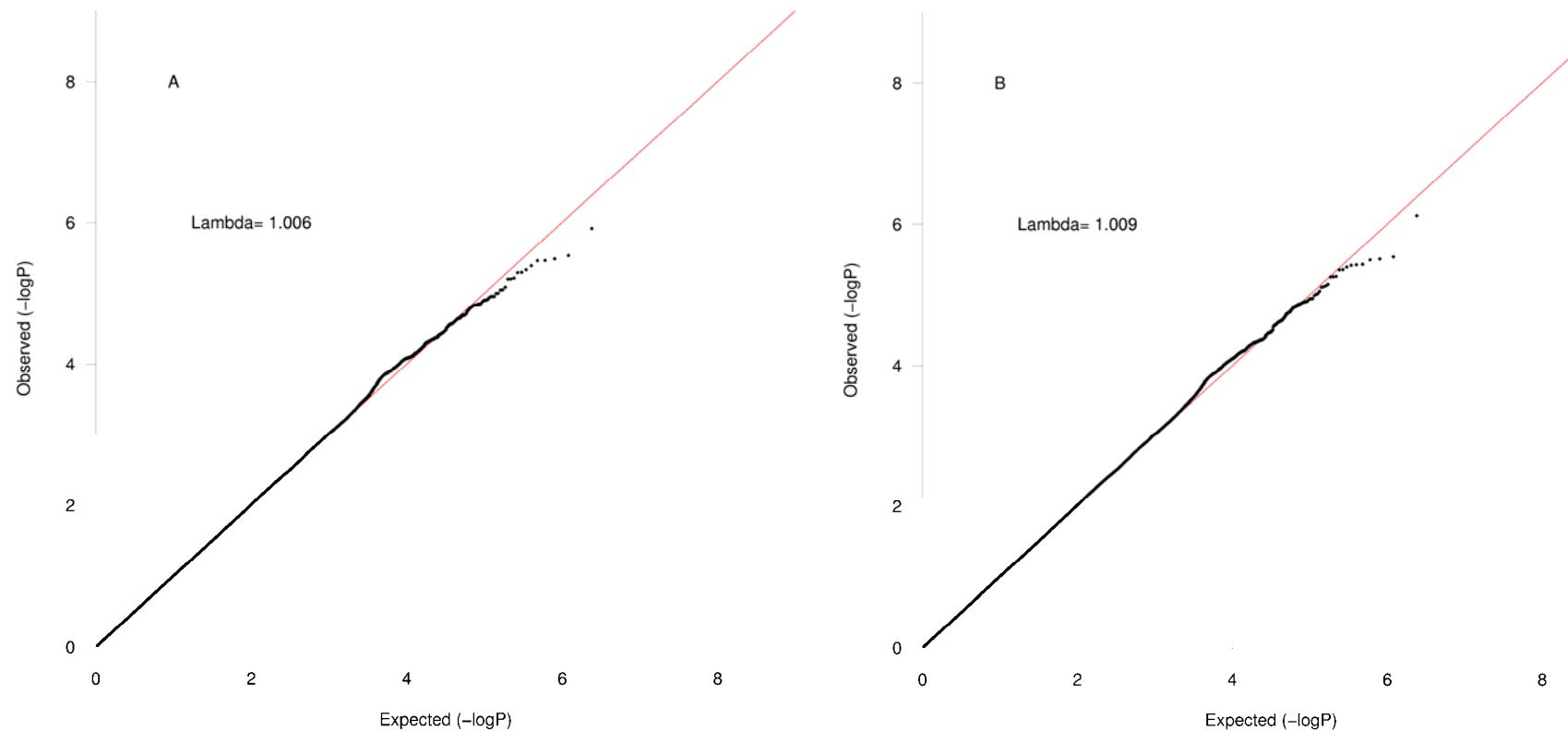
rs17557153	17	35890351	<i>TNS4</i>	Intron	C/A	0.48	0.9171	-0.046	0.010	3.5E-06
rs4655794	1	65931889	NA	NA	C/T	0.47	0.8791	0.047	0.010	3.6E-06
rs10019936	4	161650948	NA	NA	T/C	0.25	0.8978	-0.053	0.012	4.1E-06
rs7907340	10	68864243	<i>CTNNA3</i>	Intron	C/G	0.04	0.9933	0.116	0.025	4.7E-06
rs7524581	1	65939087	NA	NA	C/T	0.36	0.8656	-0.049	0.011	5.1E-06
rs7907588	10	68864374	<i>CTNNA3</i>	Intron	C/G	0.04	0.9888	0.115	0.025	5.1E-06
rs7915595	10	68866036	<i>CTNNA3</i>	Intron	T/A	0.04	0.9845	0.115	0.025	5.2E-06
rs12753193	1	65942267	NA	NA	A/G	0.36	0.8366	-0.050	0.011	5.5E-06
rs12131089	1	69301953	NA	NA	C/A	0.39	0.9919	-0.045	0.010	6.1E-06
rs7900877	10	68874863	<i>CTNNA3</i>	Intron	T/G	0.04	0.9763	0.114	0.025	6.3E-06
rs7900696	10	68875017	<i>CTNNA3</i>	Intron	C/A	0.04	0.9757	0.114	0.025	6.4E-06
rs7541434	1	65941087	NA	NA	C/A	0.36	0.8427	-0.049	0.011	6.5E-06
rs1327115	1	65739427	<i>LEPR</i>	Intron	G/T	0.38	0.9732	-0.046	0.010	6.8E-06
rs6678033	1	65850212	<i>LEPR</i>	Intron	G/A	0.36	0.9718	-0.046	0.010	7.7E-06
rs743516	22	20851498	NA	NA	G/A	0.38	0.9334	-0.046	0.010	8.3E-06
rs4655557	1	65853375	<i>LEPR</i>	Intron	T/C	0.36	0.9720	-0.046	0.010	8.7E-06
rs10997688	10	68944682	<i>CTNNA3</i>	Intron	G/A	0.04	1	0.109	0.024	9.0E-06
rs1507860	1	69293184	NA	NA	G/A	0.39	1	-0.043	0.010	9.0E-06

^an = 1,504; sOB-R levels were transformed on a natural logarithm scale. ^bPosition based on NCBI build 36.3. ^cMinor allele frequency among study participants. ^dRegression coefficients and P values for every one copy of minor allele were estimated from linear regression models adjusted for age at blood draw, diabetes case-control status, fasting status, body mass index, menopausal status, and postmenopausal hormone use.

Supplementary Figure 1. Log Quantile-Quantile (QQ) P value plot for genome-wide scan for log-transformed plasma sOB-R levels. (A) Age at blood draw, fasting and diabetes case-control status, body mass index, menopausal status, and postmenopausal hormone use were adjusted for. (B) Top three eigenvectors for European ancestry were further adjusted for.



Supplementary Figure 2. Log Quantile-Quantile (QQ) P value plot for genome-wide scan for log-transformed plasma sOB-R levels after excluding SNPs in and adjacent to *LEPR* region on chromosome 1 (65000 kb and 67000 kb on chromosome 1). (A) Age at blood draw, fasting and diabetes case-control status, body mass index, menopausal status, and postmenopausal hormone use were adjusted for. (B) Top three eigenvectors for European ancestry were further adjusted for.



Supplementary Figure 3. Eigenvectors 1 and 2 from the principal components analysis of all unduplicated NHS samples ($n = 3,369$) and all unduplicated and unrelated HapMap controls ($n = 209$: 59 CEU, 60 YRI, 45 JPT, 45 CHB). Color- and symbol-coding is by self-identified ethnic group.

