

SUPPLEMENTARY NOTE

SUMMARY OF ABBREVIATION

I. Affiliation

Stage 1	KNIH	Korea National Institute of Health
	NUS	National University of Singapore
	SERI	Singapore Eye Research Institute
	VU	Vanderbilt University
	SCI	Shanghai Cancer Institute
	NCGM	National Center for Global Health and Medicine
	SINICA	Academia Sinica
Stage 2	UNC	University of North Carolina
	UT	The University of Tokyo
	RIKEN	RIKEN
	KNIH	Korea National Institute of Health
Stage 3	SJTU	Shanghai Jiao Tong University
	NCGM	National Center for Global Health and Medicine
	SJTU	Shanghai Jiao Tong University
	CUHK	Chinese University of Hong Kong
	NTUH	National Taiwan University Hospital
Overall	SNUH	Seoul National University Hospital
	AGEN	Asian Genetic Epidemiology Network

II. Study

Stage 1	KARE	Korea Association Resource Study
	SDCS	Singapore Diabetes Cohort Study
	SP2	Singapore Prospective Study Program
	SIMES	Singapore Malay Eye Study
	CAGE	Cardiometabolic Genome Epidemiology Network
	SDGS	Shanghai Diabetes Genetic Study
	TDS	Taiwan T2D Study
	CLHNS	Cebu Longitudinal Health and Nutrition Survey
Stage 2	BBJ	BioBank Japan
	HT2DS	Health2 T2D Study
	SJTUDS	Shanghai Jiao Tong University Diabetes Study

Stage 3	CAGE	Cardiometabolic Genome Epidemiology Network
	SDIID	Shanghai Diabetes Institute Inpatient Database
	SDS	Shanghai Diabetes Study
	CUHKDS	Chinese University of Hong Kong Diabetes Study
	NTUHDS	National Taiwan University Hospital Diabetes Study
	SNUHDS	Seoul National University Hospital Diabetes Study

STUDY SAMPLES AND T2D DIAGNOSIS

I. Stage 1

Korea Association Resource Study (KARE)

The Korea Association Resource (KARE) study was initiated in 2007 to undertake a large-scale GWA analysis for Type 2 Diabetes (T2D) and numerous complex quantitative traits amongst the 10,038 participants (aged between 40 and 69) of the Ansung (n=5,018) and Ansan (n=5,020) population-based cohorts. About 10,000 subjects from KARE study cohorts were genotyped with Affymetrix Genome-Wide Human SNP array 5.0. Two KARE study cohorts were established as part of the Korean Genome Epidemiology Study (KoGES) in 2001. The sampling base for both cohorts is in KyungGi-Do province, close to Seoul, the capital of the Republic of Korea. Both cohorts were designed to allow longitudinal prospective study and adopted the same investigational strategy. Participants have been examined every two years since baseline (2001). More than 260 traits have been extensively examined through epidemiological surveys, physical examinations, and laboratory tests applied to Ansung and Ansan cohort members. Among a total of 10,038 KARE study participants, 1,042 subjects were included as T2D cases according to the following criteria: (1) treatment of T2D, (2) fasting plasma glucose ≥ 7 mmol/L or plasma glucose 2-h after ingestion of 75gm oral glucose load ≥ 11.1 mmol/L and (3) age of disease onset ≥ 40 years. The inclusion criteria of nondiabetic control subjects (n = 2,943) were as follows: (1) no history of diabetes and (2) fasting plasma glucose < 5.6 mmol/L and plasma glucose 2-h after ingestion of 75gm oral glucose load < 7.8 mmol/L at both baseline and follow up studies.

Singapore Diabetes Cohort Study (SDCS)

The Singapore Diabetes Cohort Study (SDCS) is a research initiative led by the National University of Singapore together with the National Healthcare Group Polyclinics, National University Hospital Singapore and Tan Tock Seng Hospital. Questionnaire data as well as clinical data of consenting patients were obtained together with bio-specimens such as blood and urine archived at -80°C . Patients with an established diagnosis of T2D who were seen at one of these institutions were approached to participate. The participation response rate is excellent at more than 90%. For the purpose of this study, 2,010 Chinese subjects were available for genome wide analysis and they constituted the cases on SDCS/SP2 and SDCS/SP2.

Singapore Prospective Study Program (SP2)

The Singapore Prospective Study Program (SP2) includes 6,968 participants from a random

sample of individuals from the Singapore population, aged 24 to 95 years, with disproportionate sampling stratified by ethnicity to increase the number of minority ethnic groups (Malays and Asian Indians). From 2003–2007, 10,747 participants were invited to participate by linking their unique national identification numbers with national registries, of which 7,742 attended the interview component. Of these 7,742 participants, 5,163 attended the clinical examination. A total of 5,499 Chinese, 1,405 Malays and 1,138 Asian-Indians were available at the time of the study and only the Chinese were used for this study. These Chinese constituted the 1,945 controls on SDCS/SP2 and SDCS/SP2, with no prior history of diabetes and had a fasting glucose level of not more than 6.0 mmol/L.

Singapore Malay Eye Study (SiMES)

The Singapore Malay Eye Study (SiMES) is a population-based, cross-sectional study of Malay adults (N = 3,280), aged 40 – 80 years living in Singapore. Of the 4,168 eligible participants invited, 3,280 participated in the study (78.7% response rate). In brief, age-stratified random sampling of all Malay adults aged from 40–80 years residing in 15 residential districts in the southwestern part of Singapore was performed. Cases and controls were selected from the population based cross sectional study design where diabetic cases were defined as having either a history of diabetes or had HbA1c level more than 6.5%. Controls had no history of diabetes and HbA1c level less than 6%. This yielded 794 Malay diabetic cases with 1,240 controls.

National Center for Global Health and Medicine (NCGM), Japan

T2D cases were enrolled from the clinical practice or the annual medical checkup of university hospitals, medical institutions, and general practitioners that constitute the Study Group of the CAGE (Cardiometabolic Genome Epidemiology) Network according to the 1999 WHO criteria. Exclusion criteria for cases were individuals with diabetes due to: (1) liver dysfunction; (2) monogenic disorder known to cause diabetes; and (3) positive anti-GAD antibody. The inclusion criteria for controls were as follows: (1) no past history of urinary glucose or glucose intolerance (2) HbA1c, <5.6% or a normal glucose (75g) tolerance test and (3) age at examination, ≥ 55 years. Blood samples were drawn after an overnight fast to measure fasting blood glucose and HbA1c when applicable. Sample size is 931 diabetic cases and 1404 non-diabetic controls for T2D case-control study.

Shanghai Diabetes Genetic Study (SDGS)

The SDGS includes genome wide scan data of 1019 diabetes case and 1710 controls. Details of the study design have been described elsewhere¹. Briefly, diabetes cases in the SDGS included 886 incident T2D cases identified in the Shanghai Women's Health Study (SWHS), a population-based cohort study of 75000 women² and 133 prevalent T2D cases identified from female controls of the Shanghai Breast Cancer Study (SBCS), a population-based case-control study³. The 886 diabetes cases identified from the SWHS all met the following criteria: (1) age ≤ 65 with a self-reported diabetes diagnosed after study enrollment; (2) used diabetes medication; (3) had fasting glucose level > 7 mmol/L at least twice, and (4) donated a blood sample. The 133 diabetes cases identified from the controls of SBCS were women who were diagnosed with T2D and were on diabetes medication or had a blood glucose level > 7 mmol/L (measured by study). The 1710 controls used in this GWAS were shared with a GWAS of breast cancer that was recently completed and was based primarily based on the Shanghai Breast Cancer Study (SBCS). Excluded from the control group are women who (1) had a self-reported history of diabetes; or (2) had a blood glucose level between 5.5 and 7

mmol/L and had HbA1C>6.1% or had no HbA1C data.

Taiwan T2D study (TDS)

A total of 2,798 unrelated individuals with T2D, age > 20 years, were recruited from China Medical University Hospital (CMUH), Taichung, Taiwan; Chia-Yi Christian Hospital (CYCH), Chia-Yi, Taiwan; and National Taiwan University Hospital (NTUH), Taipei, Taiwan. All of the T2D cases were diagnosed according to medical records and fasting plasma glucose levels using American Diabetic Association criteria. Subjects with type 1 diabetes, gestational diabetes, and maturity-onset diabetes of the young (MODY) were excluded from this study. The controls were randomly selected from the Taiwan Han Chinese Cell and Genome Bank. The criteria for controls in the association study were (1) no past diagnostic history of T2D, (2) HbA1C ranging from 3.4 to 6, and (3) BMI ≤ 32. The two control groups were comparable with respect to BMI, gender, age at study, and level of HbA1C. All of the participating T2D cases and controls were of Han Chinese origin, which is the origin of 98% of the Taiwan population⁴.

Cebu Longitudinal Health and Nutrition Survey (CLHNS)

The CLHNS is a community-based birth cohort study that originally enrolled 3,327 pregnant women from the Metropolitan Cebu, Philippines area in 1983-4 (3,080 singleton live births), and has since followed them and their offspring. For this study of CLHNS mothers, glucose levels were collected in the year 2005, and the study sample consists of 159 T2D cases and 1624 controls. T2D was defined as overnight fasting plasma glucose level ≥ 7.0 mmol/L or current use of anti-diabetic medication. Controls were defined as overnight fasting plasma glucose level < 7.0 mmol/L and treatment naive. Whole blood glucose levels were converted to plasma glucose level by subtracting 0.97 mmol/L⁵.

II. Stage 2

BioBank Japan (BBJ)

The subjects were recruited from several medical institutes in Japan, including Fukujiji Hospital, Iizuka Hospital, Iwate Medical University School of Medicine, National Hospital Organization Osaka National Hospital, Nihon University, Nippon Medical School, Osaka Medical Center for Cancer and Cardiovascular Diseases, The Cancer Institute Hospital of Japanese Foundation for Cancer Research, Tokushukai Hospitals and Tokyo Metropolitan Geriatric Hospital. We selected T2D cases from individuals registered as having T2D. Diabetes was originally diagnosed according to the World Health Organization (WHO) criteria. Type 2 diabetes is clinically defined as disease with a gradual adult onset. Subjects who tested positive for antibodies to glutamic acid decarboxylase (GAD) and those diagnosed with a mitochondrial disease or MODY were not included in the case group. Controls were individuals registered as individuals not having T2D but with diseases other than T2D, comprised of 13 distinct diseases, or healthy volunteers. Subjects who had been analyzed in the previous report⁶ were excluded and not included in the present study. We first genotyped 4,878 individuals with T2D (case 1, age, 65.8 ± 10.0 years; BMI, 24.1 ± 3.8 kg/m²; (all values are expressed as mean ± s.d.)) and 3,345 controls (control 1, age, 52.5 ± 15.2 years; BMI, 22.5 ± 3.8 kg/m²; (all values are expressed as mean ± s.d.)). We then selected 7,541 subjects belonging to the Hondo cluster (4,470 cases and 3,071 controls).

Health2 T2D Study (H2T2DS)

A total of 1,183 T2D cases were selected from population-based 'Health2 cohort'. Health2 cohort recruited a total of 8,500 participants aged 40-69 years from five regional cities in Korea based on the Korean Genome Epidemiologic Study (KoGES) project. T2D cases were included according to the following criteria: (1) using diabetes medication, (2) fasting plasma glucose ≥ 7 mmol/L or plasma glucose 2-h after ingestion of 75g oral glucose load ≥ 11.1 mmol/L and (3) age of disease onset ≥ 40 years. Among a total of 1,305 T2D controls, 633 subjects were selected from Health2 cohort according to the following inclusion criteria: (1) no past history of diabetes, (2) no anti-diabetic medicine, (3) fasting plasma glucose < 5.6 mmol/L and plasma glucose 2-h after ingestion of 75g oral glucose load < 7.8 mmol/L and (4) subjects older than 60 years of age. Remaining 672 T2D controls were selected from Health Examinee (HEXA) cohort shared control study that was initially designed in 2008 for the Korean cancer and coronary artery disease (CAD) GWA studies by randomly selecting approximately 3,700 of 1,200,000 subjects aged 40-69 from the HEXA cohort. HEXA cohort is another KoGES population-based cohort. Of 3,700 HEXA shared control study subjects, 672 were selected as T2D controls based on the following criteria: (1) (1) no history of diabetes, (2) no anti-diabetic medicine and (3) fasting plasma glucose < 5.6 mmol/L.

Shanghai Jiao Tong University Diabetes Study (SJTUDS)

The cases ($n = 190$) were probands of type 2 diabetic pedigrees with fasting plasma glucose ≥ 7.0 mmol/L and/or 2-h post plasma glucose ≥ 11.1 mmol/L and diagnosed before 40 years old. Type 1 diabetes and mitochondrial diabetes were excluded by clinical, immunological (based on GAD and IAA test) and genetic criteria. The controls ($n = 198$) were subjects with normal glucose regulation with fasting plasma glucose < 6.1 mmol/L and 2-h plasma glucose < 7.8 mmol/L as assessed by standard 75g OGTTs, negative diabetic family history and aged over 50 years old. All subjects were recruited from Shanghai Diabetes Institute of Shanghai Jiao Tong University.

III. Stage 3

Cardiometabolic Genome Epidemiology (CAGE) Network

In the case-control panel of the CAGE Network, T2D cases ($n = 1,110$) were enrolled according to the 1999 WHO criteria, while unaffected controls ($n = 1,014$) were enrolled according to the following criteria: (1) no past history of urinary glucose or glucose intolerance; (2) HbA1c, $< 5.6\%$ or a normal glucose (75g) tolerance test; and (3) age at examination, ≥ 55 years. In addition, population-based subjects (740 cases and 4,889 controls from the CAGE-Fukuoka Study) and the Biobank Japan (<http://biobankjp.org/>) cases ($n = 3,403$, BBJ samples independent of those used for the stage-2 analysis) were included in this case-control study. In the CAGE-Fukuoka Study (12,569 participants in total), diabetes was defined as HbA1c $\geq 7.0\%$ or under treatment of diabetes; the controls were chosen as non-diabetic subjects who met the following conditions: age, ≥ 55 years; HbA1c, $\leq 5.0\%$; no previous and/or current treatment for diabetes; and absence of renal failure (serum creatinine, < 3.0 mg/dL), as previously described⁷.

Shanghai Diabetes Institute Inpatient Database (SHDIID) and Shanghai Diabetes Study (SHDS) at SJTU

The Shanghai Diabetes Inpatient Database (SHDIID) recruited participants from inpatients in

the Department of Endocrinology and Metabolism, Shanghai Jiao Tong University Affiliated Sixth People's Hospital beginning in 2001. Detailed phenotypes of glucose and lipid metabolism, blood pressure, liver and kidney function, and diabetic complications were collected⁸⁻¹⁰. The current study included 5,950 patients of type 1 diabetes, T2D or some genetic syndromes related to diabetes (e.g., Pader Willi, Rabson Manderhall). All participants are Chinese Han ancestry residing in Shanghai area. For stage 3 replication genotyping, 3,410 T2D patients were selected from SHDIID. Shanghai Diabetes Study (SHDS) is a community-based survey of diabetes performed in Shanghai area by SJTU. Baseline surveys were performed for 5,994 participants in 1998–2001 (SHDS I) and for another 5,372 participants in 2007–2008 (SHDS II). The controls of stage 3 analysis were 3,412 subjects from SHDS I and II with normal glucose regulation as assessed by standard 75 g OGTTs. All T2D patients from both studies were negative for glutamic acid decarboxylase (GAD) and insulin autoantibody (IAA) tests.

Seoul National University Hospital (SNUH) Study

The Seoul National University Hospital (SNUH) case-control samples were consisted of 761 T2D patients of the Diabetes Clinic of SNUH and 632 nondiabetic control subjects. T2D was diagnosed using the World Health Organization criteria. Subjects positive for GAD antibodies were excluded. Nondiabetic control subjects were selected according to the following criteria: ≥ 60 years old, no history of diabetes, no first-degree relatives with diabetes, fasting plasma glucose < 6.1 mmol/L, and HbA1c $< 5.8\%$.

National Taiwan University Hospital (NTUH)

The studied population comprised 1520 normoglycemic control subjects and 1520 type 2 diabetic patients from a Han Chinese population in Taiwan. The glucose tolerant control subjects by 75-g oral glucose tolerance tests (OGTT) were recruited from routine health examinations, and type 2 diabetic patients were recruited from the metabolism clinics of the National Taiwan University Hospital. Type 2 diabetic patients were diagnosed based on the criteria of the American Diabetes Association with fasting plasma glucose ≥ 7 mmol/L or 2-hour plasma glucose ≥ 11.1 mmol/L during an OGTT. Patients with ages of onset under 35 years were excluded.

Chinese University of Hong Kong (CUHK)

The subjects in the Hong Kong case-control study were of southern Han Chinese ancestry and resided in Hong Kong, as described previously¹¹. The cases consisted of 1,477 subjects with T2D selected from the Prince of Wales Hospital Diabetes Registry¹². A total of 726 of these individuals had early-onset diabetes (age at diagnosis of < 40 years) and a positive family history; an additional 751 patients were randomly selected from the same registry. Patients with classic type 1 diabetes with acute ketotic presentation or continuous requirement of insulin within 1 year of diagnosis were excluded. The controls consisted of 1584 subjects with normal glucose tolerance (FPG of < 6.1 mmol/L). A total of 608 of these subjects was recruited either from the general population participating in a community-based screening program for cardiovascular risk or from hospital staff; an additional 976 subjects were recruited from a population-based cardiovascular risk-screening program for adolescents¹³. Informed consent was obtained from each participating subject. This study was approved by the Clinical Research Ethics Committee of the Chinese University of Hong Kong.

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Supplementary Table 1. Study cohort information.

Stage	Representative	Study	Ethnic group	Sample size			Study design
				case	control	total	
Stage 1	KNIH	KARE	Korean	1042	2943	3985	Population-based, prospective
		SDCS/SP2(1)	Chinese	1082	1006	2088	Case-control multicenter
	NUS/SERI	SDCS/SP2(2)	Chinese	928	939	1867	Case-control multicenter
		SIMES	Malay	794	1240	2034	Population-based, prospective
	NCGM	CAGE	Japanese	931	1404	2335	Case-control multicenter
	VU/SCI	SDGS	Chinese	1019	1710	2729	Population-based
	SINICA	TDS	Chinese	997	999	1996	Case-control
	UNC	CLHNS	Filipino	159	1624	1783	Population-based
total			6952	11865	18817		
Stage 2	RIKEN/UT	BBJ	Japanese	4470	3071	7541	Case-control multicenter
	KNIH	H2T2DS	Korean	1183	1305	2488	Population-based
	SJTU	SJTUDS	Chinese	190	198	388	Case-control
	total			5843	4574	10417	
Stage 3	NCGM	CAGE	Japanese	5253	5903	11156	Population-based, prospective, part of cases from BBJ
	SJTU	SDIID/SDS	Chinese	3410	3412	6822	SDIID: case-control, SDS: community-based
	CUHK	CUHKDS	Chinese	1477	1584	3061	Case-control
	NTUH	NTUHDS	Chinese	1512	1512	3024	Case-control
	SNUH	SNUHDS	Korean	632	761	1393	Case-control
	total			12284	13172	25456	
Overall	AGEN	AGEN-T2D	East Asian	25079	29611	54690	3 stage design study for genome-wide association meta-analysis

KNIH, Korea National Institute of Health; NUS, National University of Singapore; SERI, Singapore Eye Research Institute; NCGM, National Center for Global Health and Medicine; VU, Vanderbilt University; SCI, Shanghai Cancer Institute; SINICA, Academia Sinica; UNC, University of North Carolina; UT, The University of Tokyo; SJTU, Shanghai Jiao Tong University; NTUH, National Taiwan University Hospital; SNUH, Seoul National University Hospital; KARE, Korea Association Resource Study; SDCS, Singapore Diabetes Cohort Study; SP2, Singapore Prospective Study Program; SIMES, Singapore Malay Eye Study; CAGE, Cardiometabolic Genome Epidemiology Network; SDGS, Shanghai Diabetes Genetic Study; TDS, Taiwan T2D Study; CLHNS, Cebu Longitudinal Health and Nutrition Survey; BBJ, BioBank Japan; H2T2DS, Health2 T2D Study; SJTUDS, Shanghai Jiao Tong University Diabetes Study; SDIID, Shanghai Diabetes Institute Inpatient Database; SDS, Shanghai Diabetes Study; CUHKDS, Chinese University of Hong Kong Diabetes Study; NTUHDS, National Taiwan University Hospital Diabetes Study; SNUHDS, Seoul National University Hospital Diabetes Study; AGEN, Asian Genetic Epidemiology Network

Supplementary Table 2. Information of SNP genotyping and imputation for individual studies participating in one of three stages.

Stage	Representative	Study	Genotyping		SNP imputation			No of SNPs used for Stage 1 meta-analysis	GWAS inflation factor (λ_{GC})
			Platform	QC	Method	Reference panel	QC		
Stage 1	KNIH	KARE	Affy 5.0	Call rate > 0.95	IMPUTE V1.0	HapMap build 36 release 22 (CHB+JPT)	proper_info > 0.5	1419177	1.011
		SDCS/SP2(1)	Illumina610	Call rate > 0.95	IMPUTE V0.5.0	HapMap build 36 release 22 (CHB+JPT)	proper_info > 0.5	1965414	1.054
	NUS/SERI	SDCS/SP2(2)	Illumina1M	Call rate > 0.95	IMPUTE V0.5.0	HapMap build 36 release 22 (CHB+JPT)	proper_info > 0.5	2248003	1.062
		Malay 610	Illumina610	Call rate > 0.95	IMPUTE V0.5.0	HapMap build 36 release 22 (CHB+JPT+CEU+YRI)	proper_info > 0.5	1625733	1.035
	NCGM	CAGE	Illumina Human 550 and 610	Call rate > 0.95	BEAGLE V3.0.4	HapMap build 36 release 24 (CHB+JPT)	Rsqr > 0.3	1988685	1.057
	VU	SDGS	Affy 6.0	Call rate > 0.95	MACH	HapMap build 36 release 23a (CHB+JPT)	Rsqr > 0.3	2241970	1.041
	SINICA	TDS	Illumina Human 550	Call rate > 0.95	MACH	HapMap build 36 release 22 (CHB+JPT)	Rsqr > 0.3	1890143	1.043
	UNC	CLHNS	Affy 5.0	Call rate > 0.95	MACH	HapMap build 36 release 21 (CHB+JPT+CEU)	Rsqr > 0.3	2049920	1.005
Stage 2	RIKEN/Tokyo U	BBJ	Illumina Human 550 and 610	Call rate > 0.95	MACH	HapMap build 36 release 24 (CHB+JPT)	Rsqr > 0.3	na	na
	KNIH	H2T2DS	Affy 6.0	Call rate > 0.95	IMPUTE V1.0	HapMap build 36 release 22 (CHB+JPT)	proper_info > 0.5	na	na
	SJTU	SJTUDS	Illumina Human CNV370DUO	Call rate > 0.95	IMPUTE V1.0	HapMap build 36 release 22 (CHB+JPT)	proper_info > 0.5	na	na
Stage 3	NCGM	CAGE	TaqMan	Call rate > 0.99	na	na	na	na	na
	SJTU	SDIID/SDS	Sequenom MassARRAY and TaqMan	Call rate > 0.99	na	na	na	na	na
	CUHK	CUHK	Sequenom MassARRAY	Call rate > 0.98	na	na	na	na	na
	NTUH	NTUH	TaqMan and Beckman SNP Stream	Call rate > 0.98	na	na	na	na	na
	SNUH	SNUH	TaqMan	Call rate > 0.98	na	na	na	na	na

na, not available; KNIH, Korea National Institute of Health; NUS, National University of Singapore; SERI, Singapore Eye Research Institute; NCGM, National Center for Global Health and Medicine; VU, Vanderbilt University; SCL, Shanghai Cancer Institute; SINICA, Academia Sinica; UNC, University of North Carolina; UT, The University of Tokyo; SJTU, Shanghai Jiao Tong University; NTUH, National Taiwan University Hospital; SNUH, Seoul National University Hospital; KARE, Korea Association Resource Study; SDCS, Singapore Diabetes Cohort Study; SP2, Singapore Prospective Study Program; SIMES, Singapore Malay Eye Study; CAGE, Cardiometabolic Genome Epidemiology Network; SDGS, Shanghai Diabetes Genetic Study; TDS, Taiwan T2D Study; CLHNS, Cebu Longitudinal Health and Nutrition Survey; BBJ, BioBank Japan; H2T2DS, Health2 T2D Study; SJTUDS, Shanghai Jiao Tong University Diabetes Study; SDIID, Shanghai Diabetes Institute Inpatient Database; SDS, Shanghai Diabetes Study; CUHKDS, Chinese University of Hong Kong Diabetes Study; NTUHDS, National Taiwan University Hospital Diabetes Study; SNUHDS, Seoul National University Hospital Diabetes Study; AGEN, Asian Genetic Epidemiology Network

Supplementary Table 3. Association results of stage 1 meta-analysis for established T2D loci

no	Chr	SNP	nearby gene	Risk allele	Other allele	RAF (HapMap JPT/CHB)	Stage1 (GWA meta-analysis)					Reported effect	
							N	OR (CI)	P-value	hetero_p	power [§]	OR (CI)	Reference
1	1	rs10923931	NOTCH2	t	g	0.02	18814	1.05 (0.92-1.20)	4.52E-03	0.31	0.10	1.13 (1.08-1.17)	Zeggini et al., 2008 (14)
2	1	rs340874	PROX1	c	t	0.35	14831	1.08 (1.03-1.14)	2.84E-03	0.39	0.88	1.07 (1.05-1.09)	Dupuis et al., 2010 (15)
3	2	rs780094	GCKR	c	t	0.42	18813	1.05 (1.01-1.10)	2.04E-02	0.26	0.62	1.06 (1.04-1.08)	Dupuis et al., 2010 (15)
4	2	rs7578597	THADA	t	c	1.00	5989	0.93 (0.62-1.40)	7.37E-01	0.36	NA	1.15 (1.10-1.20)	Zeggini et al., 2008 (14)
5	2	rs243021	BCL11A	a	g	0.67	18634	1.05 (1.00-1.10)	4.02E-02	0.79	0.57	1.08 (1.06-1.10)	Voight et al., 2010 (16)
6	2	rs7593730	RBMS1	c	t	0.86	18815	1.03 (0.97-1.09)	3.98E-01	0.05	0.16	1.11 (1.08-1.16)	Qi et al., 2010 (17)
7	2	rs2943641	IRS1	c	t	0.92	18817	1.12 (1.03-1.22)	1.11E-02	0.53	0.80	1.19 (1.13-1.25)	Rung et al., 2009 (18)
8	3	rs1801282	PPARG	c	g	0.96	16770	1.13 (1.01-1.28)	3.93E-02	0.86	0.53	1.14 (1.08-1.20)	Saxena et al., 2007 (19)
9	3	rs6780569	UBE2E2	g	a	0.78	18765	1.13 (1.07-1.20)	2.00E-05	0.50	1.00	1.21 (1.14-1.30)	Yamauchi et al., 2010 (20)
10	3	rs4607103	ADAMTS9	c	t	0.58	18699	0.99 (0.95-1.04)	6.74E-01	0.74	0.08	1.09 (1.06-1.12)	Zeggini et al., 2008 (14)
11	3	rs11708067	ADCY5	a	g	0.99	5869	1.18 (0.80-1.74)	4.04E-01	0.74	0.09	1.12 (1.09-1.15)	Dupuis et al., 2007 (15)
12	3	rs1470579	IGF2BP2	c	a	0.30	18812	1.13 (1.08-1.19)	5.67E-07	0.38	1.00	1.17 (1.11-1.23)	Saxena et al., 2007 (19)
13	4	rs10010131	WFS1	g	a	0.98	18806	1.00 (0.91-1.10)	9.92E-01	0.08	0.05	1.11 (1.05-1.16)	Sandhu et al., 2007 (21)
14	5	rs4457053	ZBED3	g	a	0.03	5887	1.00 (0.85-1.18)	9.77E-01	0.62	0.05	1.08 (1.06-1.11)	Voight et al., 2010 (16)
15	6	rs7754840	CDKAL1	c	g	0.39	18732	1.20 (1.14-1.25)	9.63E-15	0.11	1.00	1.12 (1.08-1.16)	Voight et al., 2010 (22)
16	7	rs2191349	DGKB	t	g	0.72	18645	1.11 (1.05-1.16)	4.31E-05	0.79	0.99	1.06 (1.04-1.08)	Dupuis et al., 2010 (15)
17	7	rs864745	JAZF1	t	c	0.80	18720	1.06 (1.00-1.12)	3.50E-02	0.61	0.59	1.10 (1.07-1.13)	Zeggini et al., 2008 (14)
18	7	rs4607517	GCK	a	g	0.24	16395	1.03 (0.97-1.09)	3.97E-01	0.32	0.20	1.07 (1.05-1.10)	Dupuis et al., 2010 (15)
19	7	rs972283	KLF14	g	a	0.69	10763	0.99 (0.93-1.06)	8.52E-01	0.44	0.06	1.07 (1.05-1.10)	Voight et al., 2010 (16)
20	8	rs896854	TP53INP1	t	c	0.24	18630	1.10 (1.02-1.12)	9.05E-03	0.64	0.78	1.06 (1.04-1.09)	Voight et al., 2010 (16)
21	8	rs13266634	SLC30A8	c	t	0.54	18680	1.11 (1.06-1.16)	9.42E-06	0.61	1.00	1.15 (1.12-1.19)	Sladek et al., 2007 (23)
22	9	rs17584499	PTPRD	t	c	0.12	13042	1.09 (1.01-1.18)	3.56E-02	0.02	0.62	1.57 (1.13-1.83)	Tsai et al., 2010 (4)
23	9	rs10811661	CDKN2A/B	t	c	0.58	12815	1.21 (1.14-1.28)	6.04E-11	0.01	1.00	1.20 (1.14-1.25)	Scott et al., 2007 (22)
24	9	rs13292136	CHCHD9	c	t	0.89	18781	0.99 (0.92-1.07)	8.84E-01	0.41	0.06	1.11 (1.07-1.15)	Voight et al., 2010 (16)
25	10	rs10906115	CDC123/CAMK1D	a	g	0.65	18812	1.09 (1.04-1.14)	2.98E-04	0.00	0.97	1.13 (1.08-1.18)	Shu et al., 2010 (1)
26	10	rs12779790	CDC123/CAMK1D	g	a	0.13	8843	1.12 (1.02-1.23)	1.31E-02	0.47	0.07	1.11 (1.07-1.14)	Zeggini et al., 2008 (14)
27	10	rs1111875	HHEX/IDE	c	t	0.37	14831	1.11 (1.06-1.18)	8.74E-05	0.91	0.99	1.13 (1.09-1.17)	Scott et al., 2007 (22)
28	10	rs7903146	TCF7L2	t	c	0.02	18816	1.16 (1.02-1.31)	2.50E-02	0.77	0.57	1.37 (1.31-1.43)	Scott et al., 2007 (22)
29	11	rs231362	KCNQ1	g	a	0.85	12357	1.10 (1.00-1.20)	3.93E-02	0.10	0.72	1.08 (1.06-1.10)	Voight et al., 2010 (16)
30	11	rs2237892	KCNQ1	c	t	0.62	14830	1.17 (1.11-1.23)	5.41E-09	0.01	1.00	1.40 (1.34-1.47)	Yasuda et al., 2008 (24)
31	11	rs2334499	INS/IGF2 [¶]	t	c	0.81	NC	NC (NC-NC)	NC	NC	NC	1.35 (NA-NA)	Kong et al., 2010 (25)
32	11	rs5215	KCNJ11	c	t	0.34	18750	1.13 (1.08-1.18)	1.52E-07	0.49	1.00	1.14 (1.10-1.19)	Zeggini et al., 2007 (26)
33	11	rs1552224	CENTD2	a	c	0.93	18743	1.16 (1.06-1.27)	1.58E-03	0.22	0.94	1.14 (1.11-1.17)	Voight et al., 2010 (16)
34	11	rs10830963	MTNR1B	g	c	0.48	8710	0.99 (0.93-1.06)	8.64E-01	0.32	0.06	1.09 (1.06-1.12)	Dupuis et al., 2010 (15)
35	12	rs1531343	HMG2	c	g	0.13	18752	1.06 (0.99-1.14)	1.05E-01	0.45	0.47	1.10 (1.07-1.14)	Voight et al., 2010 (16)
36	12	rs7961581	TSPAN8/LGR5	c	t	0.20	18687	1.01 (0.95-1.06)	8.49E-01	0.39	0.07	1.09 (1.06-1.12)	Zeggini et al., 2008 (14)
37	12	rs7957197	HNF1A	t	a	1.00	NA	NA (NA-NA)	NA	NA	NA	1.07 (1.05-1.10)	Voight et al., 2010 (16)
38	13	rs1359790	SPRY2	g	a	0.74	18719	1.02 (0.97-1.08)	3.73E-01	0.00	0.13	1.15 (1.10-1.20)	Shu et al., 2010 (1)
39	15	rs7172432	C2CD4A-C2CD4B	a	g	0.60	18650	1.09 (1.04-1.15)	1.44E-04	0.53	0.98	1.14 (1.09-1.20)	Yamauchi et al., 2010 (20)
40	15	rs11634397	ZFAND6	g	a	0.07	12753	0.99 (0.90-1.09)	8.29E-01	0.35	0.05	1.06 (1.04-1.08)	Voight et al., 2010 (16)
41	15	rs8042680	PRC1	a	c	1.00	5988	1.64 (1.16-2.32)	4.92E-03	0.69	NA	1.07 (1.05-1.09)	Voight et al., 2010 (16)
42	16	rs9939609	FTO	a	t	0.14	18807	1.15 (1.08-1.22)	6.85E-06	0.02	1.00	1.15 (1.09-1.23)	Frayling et al., 2007 (27)
43	17	rs391300	SRR	c	a	0.78	14830	1.03 (0.97-1.09)	3.03E-01	0.01	0.18	1.28 (1.18-1.39)	Tsai et al., 2010 (4)
44	17	rs4430796	HNF1B (TCF2)	g	a	0.28	10317	1.11 (1.05-1.18)	5.71E-04	0.08	0.93	1.14 (1.08-1.20)	Voight et al., 2010 (16)
45	X	rs5945326	DUSP9	g	a	0.61	NA	NA (NA-NA)	NA	NA	NA	1.27 (1.18-1.37)	Voight et al., 2010 (16)

[§]The power was estimated given the stage 1 sample sizes, stage 1 ORs, T2D prevalence of 10% and the risk allele frequency (RAF) in MapMap (JPT/CHB) for $\alpha=0.05$. [¶]A signal near *INS/IGF2* was identified by parental-origin-specific analyses of disease-susceptibility variants. The *PEPD* locus was described previously with suggestive evidence of association ($P=1.4 \times 10^{-5}$). NA, not available; NC, not comparable.

Supplementary Table 4. 297 SNPs and their proxies taken forward to stage 2 in silico replication in three independent GWA studies. A total of 297 SNPs with P-value < 5 × 10⁻⁴ were selected from stage 1 meta-analysis for T2D association.

Locus No	SNP	Chr	BP	Candidate Gene	Allele e1	Allele 2	Stage1 (GWA meta-analysis)				Stage2 (in silico replication (BBJ+KNIH+SJTU))				Combined (Stage1 + Stage2)										
							N	Effect (β)	SE	P-value	Effect direction	hetero_p	N	Effect (β)	SE	P-value	Effect direction	hetero_p	N	Effect (β)	SE	P-value	Effect direction	hetero_p	
1	rs11120894	1	7263431	CAMTA1	a	c	16678	-0.0099	0.0244	1.96E-04	---+---	9.13E-01	10.292	-0.0145	0.0285	6.12E-01	++	5.35E-01	26970	-0.0586	0.0185	1.57E-03	--	4.17E-02	
	rs12409413	1	7264693	CAMTA1	t	c	18765	0.0042	0.0228	5.22E-02	+++----	1.30E-01	10.302	0.014	0.0284	6.22E-01	++	6.33E-01	29067	0.0324	0.0178	6.87E-02	++	4.07E-01	
	rs17036059	1	7265108	CAMTA1	t	c	12408	0.026	0.0261	1.22E-04	+++----	1.12E-01	10.288	0.027	0.0284	6.30E-01	++	6.42E-01	2668	0.0302	0.022	1.31E-01	++	2.13E-01	
	rs9660884	1	7285368	CAMTA1	a	c	14826	-0.0772	0.0256	2.55E-03	---+---	9.75E-01	10.244	-0.0241	0.0286	3.99E-01	++	7.10E-01	25070	-0.0536	0.0191	4.97E-03	--	1.67E-01	
2	rs170139	1	10767145	---	t	c	14832	-0.1178	0.0329	3.36E-04	---+---	0.90E-01	7874	-0.0149	0.0432	7.29E-01	++	6.42E-01	22706	-0.0802	0.0262	2.20E-03	--	5.76E-02	
	rs4373781	1	28162818	XKR8	c	g	13038	0.0936	0.0301	1.87E-03	+++----	8.80E-01	7487	0.015	0.0389	7.00E-01	??	1.00E+00	20525	0.0642	0.0238	7.03E-03	++	1.10E-01	
	rs12565238	1	28171933	EYAS	c	g	14994	0.0991	0.0286	5.41E-04	+++----	8.00E-01	10361	0.0015	0.0335	9.64E-01	++	7.51E-01	25355	0.058	0.0218	7.71E-03	++	2.67E-02	
	rs9661788	1	28184037	EYAS	c	g	17029	-0.0907	0.0269	7.41E-04	---+---	8.43E-01	10360	-0.0012	0.0335	9.71E-01	++	7.43E-01	27389	-0.0566	0.021	8.01E-03	--	3.72E-02	
3	rs10493035	1	28188048	EYAS	t	c	14987	0.0734	0.0286	1.04E-02	+++----	1.63E-01	10360	-0.002	0.0335	9.52E-01	++	7.34E-01	25349	0.0416	0.0218	5.57E-02	++	8.69E-02	
	rs7553480	1	28197067	EYAS	c	g	14998	0.0963	0.0286	7.65E-04	+++----	7.94E-01	10361	-0.0004	0.0335	9.92E-01	++	7.27E-01	25359	0.0555	0.0218	1.07E-02	++	2.81E-02	
	rs11247737	1	28204211	EYAS	c	g	14998	0.096	0.0286	7.87E-04	+++----	7.93E-01	10361	-0.0004	0.0335	9.92E-01	++	7.27E-01	25359	0.0555	0.0218	1.09E-02	++	2.86E-02	
	rs7545441	1	28207154	EYAS	a	g	14998	0.0963	0.0286	7.69E-04	+++----	7.70E-01	10361	-0.0004	0.0335	9.92E-01	++	7.27E-01	25359	0.0555	0.0218	1.07E-02	++	2.81E-02	
	rs12568186	1	28213930	EYAS	t	c	14998	-0.0964	0.0286	7.57E-04	---+---	7.69E-01	10363	-0.0002	0.0335	9.94E-01	++	7.45E-01	25361	-0.0558	0.0218	1.03E-02	++	2.90E-02	
	rs17162928	1	28216105	EYAS	t	c	14997	0.1004	0.0286	4.48E-04	+++----	8.00E-01	10363	0.0002	0.0335	9.94E-01	++	7.43E-01	25360	0.0582	0.0218	7.50E-03	++	2.29E-02	
	rs1578213	1	28222878	EYAS	t	c	16956	0.0951	0.0269	4.07E-04	+++----	7.87E-01	10349	-0.0005	0.0335	9.89E-01	++	7.51E-01	27305	-0.0576	0.021	6.01E-03	++	2.61E-02	
	rs12561882	1	28224417	EYAS	a	g	12662	-0.092	0.0311	3.14E-03	---+---	8.70E-01	10363	-0.0002	0.0335	9.94E-01	++	7.43E-01	23025	-0.0495	0.0228	2.99E-02	++	4.46E-02	
	rs2147025	1	28277281	EYAS	c	g	14913	0.0948	0.0287	9.46E-04	+++----	8.11E-01	10355	-0.0029	0.0334	9.31E-01	++	8.14E-01	25268	0.0533	0.0218	1.43E-02	++	2.65E-02	
	rs2311587	1	28289485	EYAS	a	g	12884	0.0667	0.0314	3.38E-02	+++----	1.11E-01	10352	0.0027	0.0336	9.36E-01	++	8.06E-01	23236	0.0369	0.0209	1.08E-01	++	1.64E-01	
	4	rs12049416	1	47644665	---	t	c	12781	0.0915	0.0431	3.38E-02	+++----	3.11E-02	10329	0.0163	0.0513	7.60E-01	++	8.22E-01	23110	0.0616	0.0335	6.55E-02	++	2.72E-01
		rs12021515	1	47645670	---	a	g	14832	0.1466	0.0417	4.33E-04	+++----	5.39E-01	10320	0.0149	0.0431	7.18E-01	++	8.44E-01	25152	0.0801	0.0293	6.33E-02	++	2.48E-02
5	rs3738317	1	47610448	---	t	g	14529	-0.0682	0.0395	8.39E-02	---+---	1.03E-01	9919	-0.0186	0.0423	6.30E-01	??	7.12E-01	24448	-0.0451	0.0289	1.18E-01	--	3.91E-01	
	rs10749691	1	53741575	GLIS1	g	t	10719	-0.1101	0.0331	8.86E-04	+++----	8.46E-01	7487	-0.02	0.0356	5.32E-01	??	1.00E+00	18197	0.046	0.0236	5.10E-02	++	5.81E-03	
	rs4753322	1	53748204	GLIS1	a	g	12734	-0.1075	0.0299	3.18E-04	---+---	8.96E-01	7838	0.0259	0.0329	4.31E-01	++	3.90E-01	20572	0.0472	0.0221	3.31E-02	++	2.69E-03	
	rs58039	1	53748577	GLIS1	a	g	12791	-0.1072	0.0295	2.73E-04	---+---	8.74E-01	7872	0.0286	0.0328	3.83E-01	++	2.44E-01	20663	0.0465	0.0219	3.41E-02	++	2.08E-03	
	rs2280511	1	53748776	GLIS1	t	c	14831	0.1081	0.0269	5.69E-05	+++----	9.46E-01	7856	-0.0259	0.0329	4.31E-01	??	3.58E-01	22687	-0.0544	0.0208	8.98E-03	++	1.62E-03	
	rs527014	1	53749880	GLIS1	t	c	14712	-0.1147	0.027	2.09E-05	---+---	9.71E-01	7487	0.014	0.0336	6.77E-01	??	1.00E+00	22199	-0.0642	0.021	2.29E-03	++	2.83E-03	
	rs658480	1	53750707	GLIS1	a	g	14827	-0.1123	0.0268	2.85E-05	---+---	9.65E-01	7485	0.012	0.0337	7.22E-01	??	1.00E+00	22312	-0.0641	0.021	2.29E-03	++	3.89E-03	
	6	rs10493411	1	66799712	SGP1	c	g	6847	-0.1372	0.0437	1.70E-03	???-?	7.04E-01	7855	0.0485	0.0563	3.89E-01	??	8.71E-01	14702	-0.0674	0.0345	5.10E-02	++	9.17E-03
		rs7547794	1	66804466	SGP1	a	g	16810	0.0832	0.0262	1.49E-03	+++----	6.42E-02	10363	0.013	0.0416	7.54E-01	++	1.06E-01	21713	0.0633	0.0222	4.32E-03	++	1.53E-01
		rs7547857	1	66804478	SGP1	a	g	16810	0.0832	0.0262	1.49E-03	+++----	6.42E-02	10363	0.0238	0.0416	6.56E-01	++	1.06E-01	21713	0.0662	0.0221	2.78E-03	++	2.25E-01
		rs7537440	1	66804495	SGP1	t	g	12835	0.1145	0.0297	1.15E-04	+++----	1.25E-02	10353	0.0127	0.0416	7.61E-01	++	1.30E-01	23188	0.0801	0.0242	9.16E-04	++	4.64E-02
		rs7535240	1	66804524	SGP1	t	c	18808	-0.0808	0.0244	3.35E-04	---+---	8.96E-01	7838	0.0259	0.0329	4.31E-01	++	1.84E-01	20617	0.0466	0.0207	6.86E-02	++	3.52E-01
rs963140		1	66801614	SGP1	c	g	18812	0.0831	0.0243	6.42E-04	+++----	8.06E-02	10363	0.0083	0.031	7.90E-01	++	1.64E-01	29175	0.0546	0.0191	4.28E-03	++	5.76E-02	
rs11208910		1	66811462	SGP1	t	c	16777	0.0682	0.0258	8.32E-03	+++----	1.07E-02	10363	0.028	0.0415	4.99E-01	++	2.05E-01	27140	0.0507	0.0129	9.29E-03	++	4.11E-01	
rs1373909		1	66813463	SGP1	a	g	18814	0.0861	0.0244	4.08E-04	+++----	5.19E-02	10361	0.021	0.0414	6.13E-01	++	1.43E-01	29175	0.0693	0.021	9.75E-04	++	1.76E-01	
rs12039282		1	66830911	SGP1	a	g	14355	-0.0502	0.0282	7.56E-02	---+---	2.08E-01	10246	-0.0087	0.0317	7.84E-01	++	5.71E-02	24601	-0.0139	0.021	1.30E-01	++	3.28E-01	
rs12089116		1	66832234	SGP1	a	g	16672	0.0642	0.0262	1.41E-02	+++----	1.83E-01	10241	0.0107	0.0316	7.35E-01	++	4.58E-02	26913	0.0424	0.0202	3.55E-02	++	1.93E-01	
rs7530706		1	66835180	SGP1	a	g	16677	-0.0618	0.0261	1.81E-02	---+---	1.91E-01	10242	-0.0104	0.0317	7.44E-01	++	4.52E-02	26919	-0.0421	0.0201	4.17E-02	++	2.11E-01	
rs12023176		1	66836119	SGP1	c	g	16376	0.0689	0.0264	9.11E-03	+++----	1.06E-02	10241	0.0082	0.0316	7.96E-01	++	4.43E-02	26617	0.0359	0.0203	3.01E-01	++	1.40E-01	
7	rs277359	1	75056267	---	a	c	12470	0.0725	0.0333	2.96E-02	+++----	6.56E-03	10313	-0.004	0.0345	9.07E-01	++	3.21E-01	22783	0.056	0.024	1.37E-01	++	1.11E-01	
	rs277355	1	75063372	---	c	g	14828	0.0538	0.0296	6.89E-02	+++----	1.60E-02	10346	0.0159	0.0331	6.31E-01	++	6.43E-01	25174	0.0397	0.0221	9.39E-02	++	3.99E-01	
	rs277354	1	75063661	---	c	g	14830	-0.0535	0.0293	6.94E-02	---+---	1.33E-02	10359	-0.0104	0.0331	7.52E-01	++	5.67E-01	25189	-0.044	0.022	1.18E-01	++	3.31E-01	
	rs1161851	1	75071370	---	a	g	14804	0.0517	0.0293	7.72E-02	+++----	1.12E-02	10349	0.0112	0.0331	7.35E-01	++	5.48E-01	25153	0.0399	0.0219	1.22E-01	++	3.60E-01	
	rs277367	1	75072906	---	a	g	14818	-0.0512	0.0292	7.96E-02	---+---	1.42E-02	10357												

	rs693737	1	93497512	CCDC18	a	g	18779	0.0745	0.0237	1.70E-03	+++++	3.90E-01	10338	0.0476	0.0293	1.04E-01	+++	1.82E-01	29117	0.0639	0.0184	5.29E-04	++	4.75E-01
	rs2926558	1	93509143	CCDC18	t	c	18811	0.0732	0.0237	2.02E-03	+++++	3.79E-01	10362	0.0456	0.0293	1.24E-01	+++	2.41E-01	29173	0.062	0.0184	7.99E-04	++	4.54E-01
	rs6541401	1	93511536	CCDC18	a	g	18815	0.0737	0.0237	1.89E-03	+++++	3.91E-01	10362	0.0436	0.0293	1.36E-01	+++	2.29E-01	29177	0.0618	0.0184	7.99E-04	++	4.25E-01
	rs6541404	1	93516022	CCDC18	a	t	18811	-0.0226	0.0238	3.41E-01	-----	2.33E-02	10283	-0.1198	0.0319	1.71E-01	---	5.71E-18	29094	-0.0574	0.0191	2.64E-03	--	1.46E-02
	rs2932581	1	93520121		t	c	18816	0.0712	0.0245	3.71E-03	+++++	4.41E-01	10362	0.0383	0.031	2.02E-01	+++	9.76E-02	29167	0.058	0.019	2.23E-03	++	4.29E-01
	rs9432681	1	93534844		a	c	18817	-0.0739	0.0237	1.81E-03	-----	4.02E-01	10354	-0.0441	0.0219	1.32E-01	---	2.62E-01	29167	-0.0621	0.0184	7.49E-04	++	4.29E-01
	rs1164897	1	93542960		c	g	18816	-0.0714	0.0245	3.62E-03	-----	4.46E-01	10354	-0.0347	0.0301	2.49E-01	---	8.57E-02	29170	-0.0568	0.019	2.81E-03	--	3.44E-01
	rs6680107	1	93543468		a	g	18814	-0.0745	0.0237	1.67E-03	-----	4.15E-01	10354	-0.0427	0.0293	1.44E-01	---	2.50E-01	29168	-0.0519	0.0184	7.78E-04	++	3.99E-01
	rs9432453	1	93544217		a	g	16434	-0.0829	0.0256	1.20E-03	-----?	3.79E-01	10347	-0.043	0.0293	1.42E-01	---	2.41E-01	26781	-0.0656	0.0193	6.64E-04	++	3.05E-01
	rs1000607	1	93544589		c	g	16482	0.0807	0.0255	1.59E-03	+++++	4.29E-01	10354	0.0441	0.0293	1.32E-01	+++	2.62E-01	26836	0.0649	0.0192	7.37E-04	++	3.46E-01
	rs4240666	1	93547400		a	c	18717	-0.0754	0.0238	1.49E-03	-----	3.47E-01	10311	-0.0432	0.0293	1.40E-01	---	2.25E-01	29028	-0.0626	0.0185	7.02E-04	++	3.84E-01
	rs4000699	1	93552393		t	c	18813	-0.074	0.0237	1.79E-03	-----	4.08E-01	10362	-0.0477	0.0292	1.03E-01	---	1.69E-01	29175	-0.0636	0.0184	5.53E-04	++	4.04E-01
	rs2031225	1	93559302		a	g	18817	-0.0237	0.0237	3.19E-01	-----	2.78E-02	10362	-0.047	0.0292	1.08E-01	---	1.64E-01	29179	-0.0633	0.0184	7.33E-04	--	5.36E-01
	rs2031226	1	93559352		t	g	16361	-0.8002	0.0237	1.82E-03	-----?	3.68E-01	10353	-0.0482	0.0292	9.96E-02	---	1.55E-01	29174	-0.0662	0.0193	5.97E-04	--	4.11E-01
	rs7033310	1	93561328		t	g	18805	0.0741	0.0231	1.79E-03	+++++	4.39E-01	10362	0.0465	0.0292	1.15E-01	+++	3.72E-01	29166	0.0631	0.0184	6.01E-04	++	4.63E-01
	rs1256320	1	93563233		c	g	18812	-0.0718	0.0245	3.40E-03	-----	4.61E-01	10255	-0.0373	0.0301	2.15E-01	---	4.85E-02	29167	-0.0581	0.019	2.25E-03	++	3.74E-01
	rs4847408	1	93564025		c	g	18814	-0.0238	0.0237	3.15E-01	-----	2.48E-02	10362	-0.0477	0.0292	1.03E-01	---	1.69E-01	29178	-0.0383	0.0184	7.04E-04	--	5.25E-01
	rs2391254	1	93567754		a	t	18793	0.0816	0.0238	3.17E-04	+++++	5.26E-01	10360	0.0473	0.0295	1.09E-01	+++	1.20E-01	29153	0.0681	0.0185	2.38E-04	++	3.66E-01
	rs1164908	1	93570979		a	c	18817	0.0716	0.0245	3.53E-03	+++++	4.60E-01	10362	0.0427	0.03	1.54E-01	+++	6.43E-02	29179	0.06	0.019	1.56E-03	++	3.64E-01
	rs942196	1	93574852		a	t	18810	-0.0741	0.0237	1.76E-03	-----	4.29E-01	10356	-0.0487	0.0292	9.56E-02	---	1.66E-01	29166	-0.064	0.0184	5.04E-04	++	4.99E-01
	rs3767964	1	93588078	DR1	t	c	16646	0.0765	0.0253	2.51E-03	+++++	4.29E-01	10351	0.0595	0.029	4.01E-02	+++	1.96E-01	29167	0.0692	0.0191	2.86E-04	++	6.59E-01
	rs1213376	1	93588888	DR1	a	g	16645	0.0152	0.0253	5.49E-01	+++++	9.50E-03	10351	0.059	0.029	4.17E-02	+++	1.86E-01	26996	0.0341	0.0191	7.34E-02	++	2.55E-01
	rs4847240	1	93590534	DR1	a	g	16644	0.0763	0.0253	2.58E-03	+++++	2.21E-01	10351	0.0609	0.029	3.56E-02	+++	2.07E-01	26995	0.0366	0.0191	2.59E-04	++	6.89E-01
	rs12044883	1	93593163	DR1	a	g	12319	-0.0908	0.0304	2.83E-03	-----?	1.53E-01	10330	-0.0623	0.0292	3.80E-02	---	1.29E-01	22649	-0.076	0.0211	3.09E-04	---	4.99E-01
	rs958332	1	93619228	TFE2	a	g	14826	0.1018	0.0293	5.10E-04	+++++	8.09E-01	10352	0.0136	0.0331	1.65E-01	+++	7.33E-01	29151	0.0603	0.0213	4.67E-03	++	3.93E-01
	rs1289671	1	93621459	TFE2	a	g	14828	-0.1025	0.0293	4.65E-04	-----	7.89E-01	10352	-0.0124	0.031	2.90E-01	---	8.65E-01	29180	-0.0601	0.0213	4.81E-03	++	3.50E-01
	rs2801939	1	93623809	TFE2	a	g	14829	0.1007	0.0293	6.01E-04	+++++	8.53E-01	10338	0.009	0.0311	7.72E-01	+++	7.44E-01	29167	0.0576	0.0209	1.83E-03	++	3.19E-01
	rs7540530	1	150057482	RORC	a	g	14830	0.1001	0.0272	2.32E-04	+++++	8.42E-01	7875	0.0463	0.0328	1.59E-01	+++	6.25E-01	27205	0.0782	0.0209	1.19E-03	++	2.07E-01
	rs6693413	1	150058934	RORC	a	g	14627	0.1004	0.0274	2.50E-04	+++++	2.86E-01	7853	-0.0445	0.0486	3.60E-01	-?	9.77E-01	22480	0.0653	0.0239	6.10E-03	++	9.40E-03
	rs7531041	1	150061036	RORC	a	c	630	0.0819	0.0524	1.18E-01	+++++	7.77E-01	7840	-0.062	0.0491	2.07E-01	-?	9.55E-01	14150	0.0535	0.0388	8.83E-01	++	4.51E-02
	rs12036145	1	150062490	RORC	t	c	4512	0.1017	0.0776	1.90E-01	+++++	8.47E-01	7839	-0.0508	0.0489	2.99E-01	-?	9.43E-01	12351	-0.0755	0.0414	8.57E-01	++	9.64E-02
	rs10917835	1	159983917		a	t	18681	0.0907	0.0253	3.40E-04	+++++	7.83E-01	10319	0.0405	0.0377	2.83E-01	+++	5.99E-01	29000	0.0701	0.021	3.50E-04	++	2.69E-01
	rs7519514	1	160005814	ATF6	a	g	18808	0.0999	0.0242	3.68E-05	+++++	7.25E-01	10363	0.0611	0.0292	3.62E-02	+++	9.72E-01	29171	0.0841	0.0186	6.37E-06	+++	3.06E-01
	rs9055959	1	160007898	ATF6	t	c	18814	0.0879	0.0242	2.78E-04	+++++	6.01E-01	10330	0.0615	0.0293	3.56E-02	+++	9.82E-01	29144	0.0772	0.0187	6.37E-06	+++	4.07E-01
	rs1027000	1	160023660	ATF6	t	c	18663	0.1101	0.0266	3.47E-05	+++++	3.64E-01	10350	0.0869	0.0331	9.97E-04	+++	8.95E-01	29120	0.106	0.02	1.15E-07	++	8.77E-01
	rs11576878	1	160042201	ATF6	a	g	16770	0.106	0.0266	6.59E-05	+++++	4.22E-01	10358	0.101	0.0303	8.55E-04	+++	9.04E-01	27128	0.1038	0.02	2.06E-07	++	9.01E-01
	rs1214697	1	160047603	ATF6	a	g	18648	-0.0891	0.0242	2.36E-04	-----	4.20E-01	10287	-0.0815	0.0334	1.46E-02	---	5.89E-01	28935	-0.0865	0.0196	1.02E-05	++	8.54E-01
	rs11581556	1	160060236	ATF6	a	g	18798	-0.0934	0.0253	2.18E-04	-----	3.24E-01	10356	-0.0987	0.0303	1.12E-03	---	9.13E-01	29154	-0.0956	0.0194	8.59E-07	++	8.93E-01
	rs1006310	1	160078179	ATF6	t	c	18810	0.0967	0.0242	6.32E-05	+++++	7.21E-01	10362	0.0144	0.0311	6.44E-01	+++	6.82E-01	29172	0.0519	0.0191	5.86E-04	++	3.68E-02
	rs931778	1	160090242	ATF6	a	g	18793	0.0968	0.0242	6.39E-05	+++++	7.06E-01	10354	0.0087	0.0311	7.79E-01	+++	6.12E-01	29147	0.0636	0.0191	7.83E-04	++	2.54E-02
	rs2340721	1	160116009	ATF6	a	c	18800	0.0968	0.0242	6.30E-05	+++++	7.10E-01	10356	0.01	0.0311	7.48E-01	+++	6.37E-01	29156	0.0641	0.0191	9.76E-04	++	2.76E-02
	rs10918317	1	160129179	ATF6	a	g	18792	-0.0954	0.0243	8.52E-05	-----	7.06E-01	10356	0.0032	0.0306	9.19E-01	---	6.51E-01	29148	-0.0573	0.019	2.62E-03	++	1.16E-02
	rs12124509	1	160134113	ATF6	a	g	18790	-0.0946	0.0243	9.19E-05	-----	7.11E-01	10356	0.0039	0.0306	8.99E-01	---	6.42E-01	29126	-0.0565	0.019	2.98E-03	++	1.17E-02
	rs2247475	1	160145222	ATF6	a	g	16305	0.0991	0.0259	1.29E-04	+++++	2.89E-01	10322	0.0543	0.0446	2.24E-01	+++	9.61E-01	26627	0.0878	0.0224	8.85E-05	+++	3.85E-01
	rs4654121	1	160149707	ATF6	a	g	18784	-0.0938	0.0253	2.05E-04	+++++	3.64E-01	10350	-0.0989	0.0331	1.69E-01	---	9.21E-01	2					

23	rs4665527	22931682	t	c	14832	-0.1992	0.0601	9.23E-04	---?+---	1.83E-01	7872	0.0746	0.0917	4.16E-01	++?	4.37E-01	22704	-0.1169	0.0503	2.00E-02	++	1.25E-02	
	rs738274	22932895	c	g	18012	-0.2128	0.0596	3.57E-04	---?+---	1.10E-01	7872	0.0102	0.1016	9.20E-01	++?	3.49E-01	22684	-0.1557	0.0514	2.46E-03	++	5.83E-02	
	rs4300862	22933902	c	g	18605	-0.1629	0.0605	7.05E-03	---?+---	1.10E-02	7858	0.0388	0.0955	6.85E-01	++?	7.54E-01	22663	-0.1551	0.0511	3.97E-02	++	7.44E-02	
	rs7566322	22941931	t	c	14831	0.1783	0.0624	4.25E-03	+++?+---	2.66E-01	7875	-0.0715	0.0897	4.25E-01	++?	5.60E-01	22706	0.0968	0.0512	5.87E-02	++	2.23E-02	
	rs2370384	22943794	a	g	14830	-0.2073	0.0595	4.96E-04	---?+---	1.54E-01	7875	0.0652	0.2705	4.75E-01	++?	5.50E-01	22705	-0.1259	0.0498	1.15E-02	++	1.25E-02	
	rs2347905	22945195	t	c	12794	-0.1606	0.0712	2.41E-02	---?+---	1.56E-01	7875	0.0321	0.1084	7.61E-01	++?	4.94E-01	22669	-0.1023	0.0595	8.58E-02	++	1.36E-02	
24	rs13411045	36820906	VT	t	c	18636	0.1366	0.0358	1.34E-04	+++++---	7.70E-01	7875	-0.0165	0.0553	7.66E-01	++?	4.56E-01	26111	0.0914	0.0301	2.36E-03	++	2.01E-02
	rs13411239	36820949	VT	a	g	18650	0.134	0.0357	1.74E-04	+++++---	7.50E-01	7874	-0.021	0.0548	7.02E-01	++?	4.69E-01	26524	0.0878	0.0299	3.33E-03	++	1.78E-02
25	rs3796038	39099076	SOS1	c	g	18728	0.0731	0.0281	9.14E-03	+++++---	1.12E-02	7862	0.0531	0.0341	1.95E-01	++?	5.37E-01	26590	0.0667	0.0232	4.00E-03	++	6.87E-01
	rs10204958	39096207	SOS1	c	g	18542	0.0744	0.0279	7.74E-03	+++++---	1.28E-02	7864	0.0615	0.0394	1.18E-01	++?	6.20E-01	26406	0.0701	0.0208	2.08E-03	++	7.89E-01
	rs10184015	39096633	SOS1	a	g	18733	-0.0765	0.0281	6.42E-03	+++++---	1.41E-02	7861	-0.0659	0.0405	1.04E-01	++?	5.15E-01	26594	-0.0731	0.0231	1.55E-03	++	8.30E-01
	rs1470914	39099260	SOS1	t	g	18731	-0.0774	0.0282	6.09E-03	+++++---	1.19E-02	7861	-0.0663	0.0394	9.20E-02	++?	5.12E-01	26592	-0.0736	0.0229	1.33E-03	++	8.19E-01
	rs1377864	39099744	SOS1	t	g	16708	-0.051	0.0297	8.56E-02	+++?+---	1.06E-01	7861	-0.0616	0.0393	1.17E-01	++?	5.29E-01	26459	-0.0499	0.0237	2.06E-02	++	8.30E-01
	rs1212882	39101360	SOS1	t	g	18753	0.0811	0.0282	4.00E-03	+++?+---	1.52E-02	7861	0.052	0.0392	1.84E-01	++?	5.65E-01	26614	0.0712	0.0229	1.88E-03	++	5.47E-01
	rs3033393	39102991	SOS1	a	g	14794	-0.0051	0.0316	8.18E-03	---?+---	3.14E-01	7860	-0.0485	0.0397	2.21E-01	++?	6.88E-01	22654	-0.0525	0.0247	3.36E-02	++	8.97E-01
	rs10190377	39104177	SOS1	a	g	18721	-0.0785	0.0282	5.42E-03	+++++---	2.82E-02	7860	-0.0495	0.0397	2.12E-01	++?	6.85E-01	26581	-0.0689	0.023	2.78E-03	++	5.52E-01
	rs13031871	391106355	SOS1	a	g	16535	-0.0996	0.0303	1.02E-03	+++?+---	4.09E-02	7860	-0.0542	0.0397	1.72E-01	++?	6.65E-01	24395	-0.0289	0.0241	5.79E-04	++	6.35E-01
	rs11674846	39125233	SOS1	t	c	16476	-0.1116	0.0314	3.85E-04	+++?+---	3.42E-02	7849	-0.0492	0.0394	2.11E-01	++?	7.36E-01	24325	-0.0874	0.0246	3.74E-04	++	2.16E-01
26	rs7593324	58919312	a	g	14829	0.0885	0.0329	7.19E-03	+++?+---	4.18E-01	7872	0.0058	0.044	8.95E-01	++?	7.59E-01	22701	0.0588	0.0263	2.55E-02	++	1.32E-01	
	rs7409767	58924706	t	c	14723	0.0817	0.0329	1.31E-02	+++?+---	3.96E-01	7871	0.0079	0.044	8.58E-01	++?	8.39E-01	22594	0.0552	0.0263	3.61E-02	++	1.79E-01	
	rs1400881	58945569	c	g	14760	-0.1144	0.0328	4.79E-04	+++?+---	1.22E-02	7867	0.0006	0.0332	9.85E-01	++?	7.78E-01	22627	-0.0576	0.0263	1.36E-02	++	1.37E-02	
	rs12618463	58949304	a	g	12771	0.0973	0.0363	7.38E-03	+++?+---	2.60E-01	7861	0.0017	0.0332	9.59E-01	++?	5.49E-01	20632	0.0452	0.0245	6.48E-02	++	5.20E-02	
27	rs7589969	61897757	LOC647082	t	g	14829	0.0561	0.0328	8.73E-02	+++?+---	2.76E-02	7859	0.0254	0.0391	5.15E-01	++?	7.55E-02	22688	0.0434	0.0201	8.40E-02	++	5.48E-01
	rs1115893	61900219	LOC647082	a	g	18708	0.076	0.03	1.12E-02	+++++---	2.49E-02	7872	-0.0038	0.0333	9.08E-01	++?	1.13E-01	26580	0.0402	0.0223	7.10E-02	++	7.50E-02
	rs1115894	61900221	FLI3305	a	g	14782	0.0639	0.0332	5.44E-02	+++?+---	2.01E-02	7873	0.0354	0.0405	3.74E-01	++?	5.49E-02	26519	0.0271	0.0259	4.16E-02	++	5.90E-02
	rs1115895	61922862	FLI3305	t	g	18797	-0.0805	0.0297	6.66E-03	+++++---	3.88E-02	7871	-0.0257	0.0409	5.30E-01	++?	4.30E-02	26668	-0.0616	0.024	1.04E-02	++	2.78E-01
	rs1115896	61923032	FLI3305	t	g	18781	0.0819	0.0296	5.76E-03	+++++---	3.46E-02	7873	0.027	0.0409	5.10E-01	++?	6.06E-02	26564	0.063	0.024	8.58E-03	++	2.77E-01
	rs6750080	61925046	FLI3305	t	g	18795	0.0818	0.0296	5.73E-03	+++++---	3.60E-02	7873	0.027	0.0409	5.10E-01	++?	6.06E-02	26668	0.063	0.024	8.65E-03	++	2.78E-01
	rs1115898	61939399	FLI3305	t	g	18786	-0.0825	0.0296	5.34E-03	+++++---	5.18E-02	7872	-0.0266	0.0409	5.16E-01	++?	7.18E-02	26588	-0.0633	0.024	8.31E-03	++	2.68E-01
	rs12463815	61940669	FLI3305	t	g	18785	0.0832	0.0296	4.99E-03	+++++---	5.98E-02	7875	0.0278	0.0409	4.98E-01	++?	7.57E-02	26660	0.0642	0.024	7.46E-03	++	2.73E-01
	rs12997436	61942282	FLI3305	a	g	16446	-0.0756	0.0321	1.85E-02	+++?+---	3.75E-02	7875	-0.0278	0.0409	4.98E-01	++?	7.57E-02	24321	-0.0574	0.0253	2.31E-02	++	3.58E-01
	rs12470090	61947871	CCT4	t	g	18784	-0.0808	0.0296	6.37E-03	+++++---	5.76E-02	7875	-0.0278	0.0409	4.98E-01	++?	7.57E-02	26659	-0.0626	0.024	9.06E-03	++	2.94E-01
	rs2901428	61954848	CCT4	t	g	18785	-0.0776	0.0292	7.85E-03	+++++---	3.53E-02	7875	-0.0319	0.039	4.14E-01	++?	9.76E-02	26660	-0.0612	0.0234	8.86E-03	++	3.48E-01
	rs17573253	61955149	CCT4	a	g	18813	-0.0787	0.0297	7.95E-03	+++++---	4.87E-02	7871	-0.024	0.0409	4.07E-01	++?	6.09E-02	26684	-0.0633	0.024	8.47E-03	++	3.77E-01
	rs12427249	61961136	CCT4	t	g	18785	0.0468	0.029	1.96E-01	+++?+---	1.64E-01	376	0.0745	0.5205	1.54E-01	++?	1.00E+00	19063	0.0157	0.0509	2.69E-03	++	2.56E-01
	rs12466300	61961350	CCT4	a	g	18784	-0.0765	0.029	8.38E-03	+++?+---	3.64E-02	7875	-0.0319	0.039	4.14E-01	++?	9.76E-02	26659	-0.0606	0.0233	9.19E-03	++	3.59E-01
	rs756087	61971889	FLI3305	a	g	16472	0.1186	0.0319	1.99E-04	+++?+---	8.06E-01	7859	0.0363	0.0393	3.55E-01	++?	7.08E-02	24331	0.0859	0.0248	5.23E-04	++	1.04E-01
	rs6755907	61971944	FLI3305	t	g	18782	0.0709	0.0292	1.50E-02	+++++---	2.76E-02	7875	0.0356	0.0392	3.64E-01	++?	9.37E-02	26657	0.0883	0.0234	1.28E-02	++	4.70E-01
28	rs6714614	67303043	a	c	18737	0.0884	0.0264	8.17E-04	+++++---	2.63E-01	7874	-0.0279	0.0382	4.65E-01	++?	9.93E-01	26611	-0.0508	0.0217	1.93E-02	++	1.23E-02	
	rs1384741	67306676	t	c	18775	-0.0851	0.0262	1.18E-03	+++++---	2.89E-01	7873	0.0313	0.0381	4.11E-01	++?	9.46E-01	26648	-0.0477	0.0216	2.71E-02	++	1.18E-02	
	rs875037	67310204	t	c	18780	0.0904	0.0263	5.93E-04	+++++---	2.81E-01	7872	-0.0243	0.0383	5.25E-01	++?	8.92E-01	26652	-0.0366	0.0217	1.34E-02	++	1.36E-02	
	rs1351723	67312811	t	c	16724	0.0804	0.0281	4.16E-03	+++?+---	2.71E-01	7874	-0.0251	0.0382	5.11E-01	++?	9.92E-01	24998	-0.0344	0.0226	5.54E-02	++	2.61E-02	
	rs1483337	67315153	t	g	18658	0.0897	0.0264	6.90E-04	+++++---	3.22E-01	7873	-0.0245	0.0382	5.21E-01	++?	9.59E-01	26531	-0.0528	0.0217	1.51E-02	++	1.99E-02	
	rs1483339	67316231	FLI3305	t	g	18643	0.0928	0.0264	4.42E-04	+++++---	3.49E-01	7871	-0.0262	0.0382	4.92E-01	++?	8.99E-01	26534	-0.0543	0.0217	1.24E-02	++	1.95E-02
	rs1483486	67316486	FLI3305	t	g	16632	0.0599	0.0292	4.04E-02	+++?+---	1.03E-01	7887	0.0309	0.039	4.28E-01	++?	1.00E+00	18311	0.0271	0.0237	2.53E-02	++	5.48E-02
	rs2022204	67326049	FLI3305	a	g	10824	0.0574	0.0364	1.15E-01	+++?+---	8.06E-01	7857	-0.04	0.0411	3.30E-01								

rs13008978	2	215191856	a	t	18739	0.0507	0.0281	7.18E-02	+++++	8.04E-02	7874	-0.0267	0.0366	4.65E-01	-?	4.45E-01	26613	0.022	0.0223	3.24E-01	++	9.35E-02	
rs13009193	2	215191899	t	c	18734	-0.0508	0.0281	7.12E-02	-----	7.96E-02	7874	0.0497	0.0398	2.12E-01	+	5.23E-01	26608	-0.0174	0.023	4.49E-01	++	3.91E-02	
rs10498018	2	2151919481	t	c	18807	0.0856	0.0282	2.36E-03	+++++	5.74E-01	7874	-0.0526	0.04	1.89E-01	-?	5.33E-01	26681	0.0377	0.023	8.49E-01	++	4.75E-03	
rs1040863	2	2151919764	a	g	18816	0.0861	0.0281	2.21E-03	+++++	5.79E-01	7870	-0.0526	0.04	1.89E-01	-?	5.33E-01	26686	0.0403	0.023	7.99E-02	++	4.55E-03	
rs10880806	2	215192000	t	g	18807	0.086	0.0282	2.26E-03	+++++	5.70E-01	7874	-0.0526	0.04	1.89E-01	-?	5.33E-01	26681	0.04	0.023	8.28E-02	++	4.65E-03	
rs9729602	2	215196602	t	g	18807	-0.0861	0.0282	2.22E-03	-----	5.69E-01	7874	0.0526	0.04	1.89E-01	+	5.33E-01	26681	-0.0401	0.023	8.23E-02	++	4.60E-03	
rs972906	2	215197039	t	g	18807	0.0862	0.0282	2.21E-03	+++++	5.68E-01	7874	-0.0526	0.04	1.89E-01	-?	5.33E-01	26681	0.0401	0.023	8.18E-02	++	4.57E-03	
rs222182	2	215197375	t	g	18806	0.0867	0.0282	2.07E-03	+++++	5.66E-01	7874	-0.0526	0.04	1.89E-01	-?	5.33E-01	26680	0.0405	0.023	7.93E-02	++	4.42E-03	
rs9288498	2	215197535	g	18745	-0.0878	0.0282	1.86E-03	-----	5.67E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	26619	-0.042	0.0231	6.88E-02	++	4.73E-03		
rs9646850	2	215197793	a	c	16717	-0.0958	0.03	1.41E-03	-?-----	5.18E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	24591	-0.0432	0.024	7.19E-02	++	3.44E-03	
rs9288499	2	215197923	a	g	18745	-0.0879	0.0282	1.82E-03	-----	5.64E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	26619	-0.042	0.0231	6.84E-02	++	4.70E-03	
rs4513290	2	215198940	c	g	18806	0.0427	0.0281	1.29E-01	+++++	4.26E-02	7874	-0.0507	0.0401	2.06E-01	-?	5.26E-01	26680	0.0119	0.023	6.04E-01	++	5.65E-02	
rs1914515	2	215199321	t	c	18806	0.0889	0.0282	1.59E-03	+++++	5.28E-01	7874	-0.0507	0.0401	2.06E-01	-?	5.26E-01	26680	0.0427	0.0231	6.41E-02	++	4.40E-03	
rs1768996	2	215201034	t	g	18807	-0.0887	0.0282	1.63E-03	-----	5.23E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	26681	-0.0426	0.0231	6.50E-02	++	4.46E-03	
rs5019644	2	215201829	t	g	18806	-0.0872	0.0281	1.94E-03	-----	5.59E-01	7874	0.0501	0.0401	2.06E-01	+	5.26E-01	26680	-0.0418	0.023	6.94E-02	++	4.82E-03	
rs5019643	2	215201886	t	g	18751	0.0846	0.0283	2.75E-03	+++++	6.01E-01	7874	-0.0507	0.0401	2.06E-01	-?	5.26E-01	26625	0.0396	0.0231	8.66E-02	++	5.84E-03	
rs5019642	2	215201905	t	g	18806	-0.09	0.0282	1.40E-03	-----	5.05E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	26680	0.0434	0.0231	5.97E-02	++	4.10E-03	
rs5019641	2	215201929	t	g	16444	0.0879	0.0304	3.87E-03	+++++	4.26E-01	7870	-0.049	0.0401	2.22E-01	-?	6.50E-01	24314	0.0379	0.0242	1.17E-01	++	6.52E-03	
rs10178091	2	215203048	c	g	16775	-0.0983	0.03	1.05E-03	-?-----	4.80E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	24649	-0.0448	0.024	6.20E-02	++	2.93E-03	
rs17764788	2	215203853	a	g	16503	0.097	0.03	1.24E-03	+++++	5.20E-01	7862	-0.0521	0.0401	1.94E-01	-?	4.23E-01	24365	0.0435	0.024	7.02E-02	++	2.91E-03	
rs7574717	2	215204536	a	g	18797	0.0893	0.0282	1.55E-03	+++++	5.56E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24672	0.0428	0.0231	6.33E-02	++	4.18E-03	
rs7601421	2	215204709	t	c	16783	0.0472	0.03	1.15E-01	+++++	3.28E-02	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24658	0.0119	0.024	6.20E-01	++	4.97E-02	
rs7604462	2	215204989	a	g	16781	0.0971	0.03	1.21E-03	+++++	5.19E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24656	0.0439	0.024	6.75E-02	++	3.08E-03	
rs1040217	2	215209400	t	g	18769	0.0851	0.0282	2.56E-03	+++++	6.31E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24644	0.04	0.0231	8.27E-02	++	5.47E-03	
rs7603357	2	215210221	t	g	18817	0.0871	0.0282	2.02E-03	+++++	5.91E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24660	0.0414	0.0231	7.29E-02	++	4.82E-03	
rs4234004	2	215210224	a	g	18792	0.0789	0.0282	2.54E-03	+++++	6.41E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24667	0.0359	0.0231	1.20E-01	++	8.01E-03	
rs1971409	2	215210825	a	g	18781	0.0527	0.0283	2.64E-02	+++++	1.82E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24656	0.0182	0.0231	4.32E-01	++	3.44E-02	
rs10210694	2	215211235	a	g	16410	0.0842	0.0307	6.11E-03	+++++	5.71E-01	7865	-0.0488	0.0402	2.25E-01	-?	5.84E-01	24725	0.0352	0.024	1.49E-01	++	8.55E-03	
rs13406678	2	215211250	c	g	16783	0.0975	0.0301	1.20E-03	+++++	5.58E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24658	0.0439	0.0241	6.79E-02	++	3.04E-03	
rs13410212	2	215212115	t	g	14448	0.0989	0.0328	2.57E-03	+++++	4.22E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	23233	0.0388	0.0254	1.27E-01	++	3.79E-03	
rs6435841	2	215212708	a	g	16759	0.0991	0.0301	1.92E-04	+++++	5.03E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24634	0.045	0.0241	6.18E-02	++	2.74E-03	
rs6435842	2	215212823	t	g	16756	0.099	0.0301	1.02E-03	+++++	4.86E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24631	0.0449	0.0241	6.21E-02	++	2.76E-03	
rs6743885	2	215213267	a	g	16751	0.0994	0.0301	9.53E-04	+++++	4.70E-01	7875	-0.0511	0.0401	2.02E-01	-?	4.95E-01	24626	0.0452	0.0241	6.07E-02	++	2.69E-03	
rs6739136	2	215213460	a	t	16760	0.1089	0.0301	2.91E-04	+++++	4.16E-01	7874	-0.0507	0.0401	2.06E-01	-?	5.26E-01	24574	0.0514	0.0241	3.28E-02	++	1.46E-03	
rs6739865	2	215213675	a	g	16692	0.1066	0.0301	3.98E-04	+++++	4.69E-01	7874	-0.0507	0.0401	2.06E-01	-?	5.26E-01	24566	0.0499	0.0241	3.81E-02	++	1.71E-03	
rs6747271	2	215213903	t	c	16763	-0.1089	0.03	2.87E-04	-?-----	3.00E-01	7874	0.0507	0.0401	2.06E-01	+	5.26E-01	24637	-0.0516	0.024	3.16E-02	++	1.44E-03	
rs10173431	2	215214041	a	g	16700	-0.1091	0.0302	3.11E-04	-?-----	2.67E-01	7853	0.0488	0.0402	2.24E-01	+	6.61E-01	24553	-0.0521	0.0241	3.08E-02	++	1.69E-03	
rs1535322	2	215215323	a	g	16704	0.1072	0.0302	3.79E-04	+++++	2.69E-01	7848	-0.0486	0.0402	2.26E-01	-?	6.79E-01	24552	0.051	0.0241	3.47E-02	++	1.94E-03	
rs16852400	2	215215381	c	g	16704	0.1081	0.0302	3.42E-04	+++++	2.89E-01	7848	-0.0486	0.0402	2.26E-01	-?	6.79E-01	24552	0.0516	0.0241	3.27E-02	++	1.83E-03	
rs4080303	2	215216268	a	g	16703	0.1095	0.0302	2.88E-04	+++++	3.14E-01	7845	-0.0506	0.0402	2.08E-01	-?	5.15E-01	24548	-0.0517	0.0241	3.21E-02	++	1.45E-03	
rs12466043	2	215217141	t	g	14307	-0.1143	0.033	5.36E-04	-?-----	2.45E-01	7844	0.0508	0.0402	2.06E-01	+	4.97E-01	22151	-0.0478	0.0255	6.08E-02	++	1.50E-03	
rs5776443	2	215217785	t	g	14372	0.1103	0.0329	3.18E-04	+++++	2.18E-01	7843	-0.0503	0.0402	2.11E-01	-?	5.34E-01	22215	-0.0459	0.0255	7.13E-02	++	1.99E-03	
rs7606845	2	215218170	a	g	16707	-0.1088	0.0302	3.16E-04	-?-----	3.12E-01	7843	0.0503	0.0402	2.11E-01	+	5.34E-01	24550	-0.0514	0.0241	3.33E-02	++	1.56E-03	
rs7580615	2	215218278	t	c	16667	0.0997	0.0302	9.67E-04	+++++	4.37E-01	7843	-0.0503	0.0402	2.11E-01	-?	5.34E-01	24510	0.0456	0.0241	5.93E-02	++	2.85E-03	
rs7580859	2	215218324	a	g	16687	0.1042	0.0302	5.62E-04	+++++	3.54E-01	7843	-0.0503	0.0402	2.11E-01	-?	5.34E-01	24510	0.0453	0.0241	4.47E-02	++	2.12E-03	
rs10196683	2	215219623	t	c	16665	0.1066	0.0304	4.52E-04	+++++	3.27E-01	7847	-0.045	0.0411	2.74E-01	?	1.00E-01	24152	0.053	0.0244	3.02E-02	++	3.02E-03	
35	rs1959982	2	223956117	ALPP	a	g	16756	-0.1072	0.0296	2.85E-04	-----	9.85E-01	7875	-0.0786	0.0387	4.77E-02	-?	6.00E-01	24631	-0.0959	0.0235	4.52E-05	

rs1613185	23430468	UBE2Z	t	c	18648	0.1106	0.029	1.34e-04	+++++	5.67e-01	10351	0.1742	0.0494	4.24e-04	+++	8.81e-01	28999	0.1269	0.025	3.89e-07	++	2.67e-01	
rs13094957	23432084	UBE2Z	t	c	14822	0.119	0.0324	2.38e-04	+++++	4.36e-01	10348	0.1727	0.0495	4.88e-04	+++	8.74e-01	25170	0.1351	0.0271	6.24e-07	++	3.64e-01	
rs1309303	23438651	UBE2Z	a	g	18747	-0.1336	0.0334	6.46e-05	-----	4.08e-01	2867	-0.1823	0.0701	9.31e-03	--?	6.84e-01	21614	-0.1262	0.0302	2.25e-07	--	5.31e-01	
rs13095736	23439324	UBE2Z	t	c	18756	-0.1322	0.0334	7.59e-05	-----	4.24e-01	2864	-0.1847	0.0701	8.43e-03	--?	6.62e-01	21620	-0.1419	0.0302	2.52e-06	--	4.99e-01	
rs178493	23441259	UBE2Z	t	g	18520	0.0927	0.0252	2.36e-04	+++++	7.03e-01	10286	0.094	0.0421	2.53e-02	+++	2.95e-01	28806	0.093	0.0216	1.68e-05	+++	9.79e-01	
rs1762372	23441259	UBE2Z	a	c	18587	-0.1074	0.029	2.16e-04	+++++	6.20e-01	10342	-0.1801	0.0494	2.69e-04	+++	9.16e-01	28850	-0.126	0.025	4.67e-07	++	2.04e-01	
rs8733229	23442030	UBE2Z	t	c	18791	-0.1362	0.0333	4.33e-05	-----	4.23e-01	10351	-0.1647	0.0393	2.74e-05	---	8.46e-01	29142	-0.1481	0.0254	5.55e-09	---	5.80e-01	
rs2359765	23442777	UBE2Z	a	g	18509	-0.1078	0.0291	2.09e-04	+++++	6.20e-01	10353	-0.1842	0.0494	1.94e-04	---	9.25e-01	28862	-0.1275	0.0251	3.69e-07	--	1.83e-01	
rs17013216	23451012	UBE2Z	t	c	16763	0.1486	0.0352	2.44e-05	+++++	3.78e-01	10352	0.1551	0.065	1.70e-02	+++	5.69e-01	27115	0.1501	0.031	1.24e-06	+++	9.30e-01	
rs17013223	23451411	UBE2Z	a	g	16763	-0.1487	0.0351	2.31e-05	-----	3.77e-01	2865	-0.1818	0.0702	9.62e-03	--?	7.35e-01	29128	-0.1553	0.0314	7.57e-07	--	6.73e-01	
rs17013241	23457114	UBE2Z	t	c	18772	-0.1393	0.0333	2.86e-05	-----	3.71e-01	2858	-0.1851	0.0703	8.44e-03	--?	7.83e-01	21630	-0.1477	0.0301	9.22e-07	--	5.56e-01	
rs17013245	23457379	UBE2Z	a	g	18713	-0.1402	0.0337	3.11e-05	-----	3.50e-01	2836	-0.1987	0.0715	5.47e-03	--?	8.95e-01	21549	-0.1508	0.0305	7.70e-07	--	4.59e-01	
rs17013286	23471138	UBE2Z	t	c	18773	0.1471	0.0353	2.76e-05	+++++	3.68e-01	10341	0.1698	0.0402	2.45e-05	+++	9.80e-01	27114	0.1517	0.0264	2.90e-09	+++	6.72e-01	
rs17013320	23488060	UBE2Z	t	g	18753	-0.1398	0.0333	2.65e-05	-----	4.44e-01	10342	-0.1704	0.0403	2.31e-05	---	9.81e-01	29055	-0.1522	0.0257	3.04e-09	---	5.58e-01	
rs17013322	23492833	UBE2Z	t	c	18720	-0.1476	0.0353	2.93e-05	-----	4.78e-01	10345	-0.183	0.040	1.89e-05	---	9.55e-01	27065	-0.1588	0.0265	2.25e-09	---	6.31e-01	
rs1695192	23497007	UBE2Z	t	g	18729	0.0825	0.0236	4.76e-04	+++++	5.16e-01	10338	0.0839	0.0299	4.94e-03	+++	2.74e-01	29067	0.083	0.0185	7.38e-06	+++	9.71e-01	
rs1692647	23497813	UBE2Z	a	c	18731	-0.0831	0.0236	4.33e-04	-----	5.17e-01	10338	-0.0845	0.0299	4.65e-03	---	2.72e-01	29069	-0.0836	0.0185	6.34e-06	---	9.71e-01	
rs17013333	23498110	UBE2Z	t	g	16724	0.1501	0.0351	1.95e-05	+++++	4.47e-01	10343	0.1752	0.0402	1.33e-05	+++	9.35e-01	27067	0.161	0.0264	1.15e-09	+++	6.38e-01	
rs17013344	23500465	UBE2Z	a	g	16773	-0.1484	0.0351	2.39e-05	-----	4.36e-01	10346	-0.1773	0.0402	1.04e-05	---	9.63e-01	27119	-0.1609	0.0264	1.16e-09	---	5.88e-01	
rs17013348	23501001	UBE2Z	a	g	16777	-0.1473	0.0351	2.74e-05	-----	4.09e-01	10346	-0.1773	0.0402	1.04e-05	---	9.63e-01	27123	-0.1603	0.0264	1.35e-09	---	5.74e-01	
rs801277	23504025	UBE2Z	t	c	18808	-0.0897	0.0237	1.48e-04	-----	5.07e-01	10337	-0.1153	0.03	1.25e-04	---	5.25e-01	29145	-0.0995	0.0186	8.68e-08	---	5.03e-01	
rs9631541	23514783	UBE2Z	a	g	18774	-0.0887	0.0237	1.78e-04	-----	6.12e-01	10351	-0.1186	0.03	7.87e-05	---	6.53e-01	29125	-0.1002	0.0186	7.15e-08	---	4.34e-01	
rs1011754	23516061	UBE2Z	a	g	18805	-0.1382	0.0333	3.34e-05	-----	4.36e-01	10352	-0.1758	0.0402	1.23e-05	---	9.57e-01	29157	-0.1525	0.0256	2.15e-09	---	4.71e-01	
rs17013383	23522375	UBE2Z	a	t	16775	-0.1522	0.0351	1.47e-05	-----	3.65e-01	10354	-0.1742	0.0402	1.48e-05	---	9.53e-01	27129	-0.1617	0.0264	9.57e-10	---	6.80e-01	
rs17013389	23523928	UBE2Z	t	c	16773	-0.1483	0.0354	2.98e-05	-----	4.91e-01	10324	-0.186	0.0494	3.16e-06	---	7.56e-01	27126	-0.1617	0.0266	1.22e-09	---	6.17e-01	
rs17013429	23525558	UBE2Z	a	c	16741	-0.142	0.036	8.07e-05	-----	3.78e-01	10311	-0.165	0.0399	3.47e-05	---	6.94e-01	27052	-0.1523	0.0267	1.21e-08	---	6.69e-01	
rs17013433	23525598	UBE2Z	a	g	16704	0.1394	0.0344	5.10e-05	+++++	7.47e-01	10301	0.186	0.0391	1.99e-06	+++	9.53e-01	27005	0.1597	0.0258	6.22e-10	+++	3.71e-01	
40	rs8957064	24061882	THRB	a	c	12365	0.1001	0.032	1.77e-03	+++++	1.74e-01	7847	0.002	0.068	9.57e-01	??	1.00e+00	29059	0.0241	0.166e-02	++	4.43e-02	
rs2044755	24067930	THRB	a	t	14765	0.1042	0.0288	2.90e-04	+++++	8.41e-01	7848	0.0107	0.0361	7.67e-01	++?	3.95e-01	22613	0.0678	0.0225	2.59e-05	++	4.29e-02	
41	rs1485916	31154343		a	c	16448	0.3563	0.0893	6.61e-05	+++++	7.87e-01	2822	0.0414	0.201	8.43e-01	++?	8.69e-01	19320	0.3077	0.0821	2.19e-04	++	1.66e-01
rs839593	31159203		t	c	16440	-0.2359	0.0952	1.32e-02	-----	7.56e-01	2876	0.0061	0.214	9.77e-01	++?	9.53e-01	19316	-0.1951	0.0868	2.46e-02	++?	2.97e-01	
rs186074	31173029		c	g	14439	-0.218	0.1024	3.33e-02	-----	7.74e-01	2876	-0.1324	0.214	9.50e-01	++?	9.18e-01	17315	-0.1794	0.0922	5.18e-02	++?	3.86e-01	
rs17612607	31176419		c	g	16472	-0.1931	0.0961	4.44e-02	-----	9.54e-01	2876	-0.1324	0.214	9.50e-01	++?	9.18e-01	19348	-0.1626	0.0876	6.34e-02	++?	4.41e-01	
rs1701447	31176976		c	g	16435	-0.1793	0.0959	6.15e-02	-----	9.32e-01	2876	-0.1324	0.214	9.50e-01	++?	9.18e-01	19311	-0.1522	0.0876	8.74e-02	++?	4.77e-01	
rs1687556	31177697		t	g	16421	0.0841	0.0963	8.83e-01	-----	1.62e-01	2875	0.0142	0.1296	0.072	---	9.17e-01	19316	0.072	0.0267	4.11e-01	---	6.17e-01	
42	rs12491561	31524957		c	18808	0.0767	0.0235	1.09e-03	+++++	2.39e-01	10357	0.0145	0.0286	6.13e-01	+++	1.14e-01	29165	0.0516	0.0182	4.46e-03	+++	9.29e-02	
rs7652669	31525512		c	g	16977	-0.0838	0.0239	4.47e-04	-----	2.51e-01	10358	-0.0115	0.0286	6.89e-01	+++	1.20e-01	27335	-0.0518	0.0183	3.20e-03	+++	5.24e-02	
rs4390968	31526784		a	g	18759	-0.0782	0.0235	8.92e-04	-----	1.91e-01	10360	-0.013	0.0286	6.49e-01	+++	1.30e-01	29119	-0.0519	0.0182	4.24e-03	+++	7.82e-02	
rs6791590	31527257		a	t	18752	0.0784	0.0236	8.92e-04	+++++	1.97e-01	10361	0.0128	0.0286	6.55e-01	+++	1.26e-01	29113	0.0518	0.0182	4.41e-03	+++	7.69e-02	
43	rs2168422	31779846	CSBPLO	t	g	18684	0.043	0.0294	2.44e-01	+++++	7.00e-02	10279	0.0216	0.0334	5.18e-01	+++	9.84e-01	28963	0.0337	0.0221	1.27e-01	+++	6.11e-01
rs7613385	31781075	CSBPLO	t	c	18750	0.086	0.0288	2.79e-03	+++++	4.32e-01	10348	0.0282	0.0329	3.91e-01	+++	8.05e-01	29098	0.0609	0.0217	4.93e-01	+++	8.38e-01	
rs17176090	31782734	CSBPLO	a	c	18674	-0.0448	0.0288	1.20e-01	-----	4.23e-02	10328	-0.0275	0.033	4.05e-01	---	7.64e-01	29002	-0.0373	0.0217	8.54e-02	---	6.93e-01	
rs7796221	31782517	CSBPLO	t	g	18531	-0.0507	0.0304	9.52e-02	-----	4.84e-02	10263	-0.025	0.0343	4.06e-01	---	4.97e-01	28794	-0.0394	0.0228	8.34e-02	---	5.75e-01	
rs1902338	31782985	CSBPLO	a	g	16497	-0.0571	0.0309	6.47e-02	-----	4.35e-02	10299	-0.0115	0.0322	7.37e-01	---	8.41e-01	26796	-0.0366	0.0229	1.10e-01	---	3.23e-01	
rs9810972	31785033	CSBPLO	a	g	18642	0.0409	0.0288	1.55e-01	+++++	1.40e-01	10322	0.0165	0.0291	5.82e-01	+++	9.45e-01	28961	0.0343	0.0207	1.22e-01	+++	1.67e-01	
rs12497607	31786629	CSBPLO	a	g	16707	-0.0365	0.0297	2.20e-01	-----	1.98e-02	10350	-0.031	0.0334	3.54e-01	---	6.38e-01	27057	-0.0341	0.0222	1.25e-01	---	9.02e-01	
rs936160	31787589	CSBPLO	t	c	18695	-0.0427	0.0295	1.49e-01	-----	6.31e-02	10354	-0.0245	0.0337	4.67e-01	---	5.21e-01	29049	-0.0348	0.0222	1.17e-01	---	6.85e-01	
rs2219250	31792548	CSBPLO	t	c	18613	0.0828	0.0297	5.33e-03	+++++	3.81e-01	10350	0.0255	0.0337	4.49e-01	+++	5.10e-01	28963	0.0748	0.0223	9.55e-03	+++	2.02e-01	
rs1709187	31795688	CSBPLO	a	g	16572	-0.0505	0.03	9.24e-02	-----	3.56e-02	10276	-0.0278	0.0339	4.12e-01	---	6.21e-01	26848	-0.0405	0.0225	7.12e-02	---	6.16e-01	
rs12171283	31797341	CSBPLO	a	g	18661	-0.0571	0.0299	5.59e-02	-----	1.84e-02	10272	-0.0234	0.0344	4.97e-01	---	4.67e-01	28933	-0.0426	0.0226	5.91e-02	---	4.60e-01	
rs1931078	31801771	CSBPLO	a	g	16884	0.0573	0.0303	5.88e-02	+++++	1.06e-02	10292	0.0243	0.0343	4.80e-01	+++	5.01e-01	27176	0.0428	0.0227	5.93e-02	+++	4.71e-01	
rs4955200	31802319	CSBPLO	t	c	18565	0.0576	0.0289	4.61e-02	+++++	1.47e-02	10264	0.0263	0.034	4.38e-									

rs12638086	60420613	FHIT	t	c	18773	-0.0845	0.0236	3.48E-04	+++++	2.73E-01	10360	-0.0271	0.0289	3.49E-01	+++	4.20E-01	29133	-0.0615	0.0183	7.62E-04	--	1.24E-01
rs12638610	60420846	FHIT	t	c	18777	-0.0848	0.0236	3.29E-04	+++++	1.75E-01	10360	-0.0278	0.0289	3.36E-01	+++	4.24E-01	29137	-0.062	0.0183	6.95E-04	--	1.27E-01
rs12638665	60421098	FHIT	t	c	18817	-0.0405	0.0236	8.60E-02	+++++	2.78E-02	10354	-0.026	0.0289	3.69E-01	+++	4.14E-01	29171	-0.0347	0.0183	5.77E-04	--	1.98E-01
rs1579912	60422380	FHIT	t	g	18763	0.0846	0.0232	2.66E-04	+++++	4.40E-01	10344	0.0238	0.0287	4.07E-01	+++	6.03E-01	29107	0.0606	0.018	7.87E-04	++	9.95E-02
rs15405127	6042301	FHIT	t	c	18815	-0.0812	0.0231	4.53E-04	+++++	4.53E-01	10344	-0.0238	0.0287	4.07E-01	+++	6.03E-01	29109	0.0606	0.018	1.12E-03	++	1.19E-01
rs474675	6042301	FHIT	t	c	18815	-0.0889	0.0232	2.8E-04	+++++	5.11E-01	10338	-0.0234	0.0287	4.15E-01	+++	5.56E-01	29151	-0.063	0.018	4.78E-04	--	7.59E-02
rs378446	60423631	FHIT	a	g	18769	-0.0891	0.0232	1.26E-04	+++++	4.36E-01	10343	-0.0234	0.0287	4.15E-01	+++	5.56E-01	29112	-0.0631	0.018	4.67E-04	--	7.50E-02
rs2594239	60424049	FHIT	c	g	18754	0.081	0.0232	4.81E-04	+++++	3.46E-01	10343	0.0242	0.0287	4.00E-01	+++	5.53E-01	29097	0.0586	0.018	1.17E-03	++	1.24E-01
rs403815	60428440	FHIT	a	g	10636	0.0886	0.0233	6.04E-03	??+?+?	2.67E-01	10234	-0.012	0.0288	6.76E-01	+++	4.48E-01	20870	0.0236	0.0215	1.30E-01	++	2.01E-02
rs12496632	60487512	FHIT	t	c	14983	-0.1387	0.0334	3.33E-05	----?	8.37E-01	10330	0.0174	0.0375	6.41E-01	+++	2.80E-01	25313	-0.0696	0.0249	5.23E-04	++	1.88E-03
rs4679545	60488163	FHIT	a	g	16373	0.0882	0.034	9.45E-03	+++?+?	1.53E-01	10341	-0.0199	0.0376	5.96E-01	+++	3.45E-01	26714	0.0396	0.0252	1.17E-01	++	3.30E-02
rs12485617	60489702	FHIT	t	c	17033	-0.1217	0.0315	1.15E-04	----?	7.31E-01	10232	0.0134	0.0381	7.26E-01	+++	2.01E-01	27265	-0.0668	0.0243	5.90E-03	++	6.28E-03
rs17063484	60496949	FHIT	t	g	14876	-0.1438	0.0341	2.50E-05	----?	8.97E-01	10312	0.0139	0.0371	7.08E-01	+++	5.09E-01	25188	-0.0716	0.0251	4.36E-03	--	1.75E-03
rs9842724	63779801	C3orf99	t	c	18684	-0.0907	0.0235	1.15E-04	----?	6.15E-01	10239	-0.0176	0.0293	6.55E-02	+++	2.52E-01	28923	-0.0354	0.0185	6.36E-03	++	6.16E-01
rs13434089	63923606	ATXN1	t	c	18608	-0.0852	0.0232	2.37E-04	+++++	8.01E-01	10353	0.0631	0.0295	3.26E-02	+++	3.96E-01	29014	0.0768	0.0182	2.57E-05	++	5.56E-01
rs13433909	63923649	ATXN1	t	c	16508	0.0886	0.0249	3.71E-04	+++?+?+	9.04E-01	10211	0.0642	0.0296	3.02E-02	+++	3.21E-01	26819	0.0785	0.0191	3.80E-05	++	5.28E-01
rs33087	63988289	SLC9A9	t	c	15348	0.1036	0.0268	1.07E-04	+++?+?+	5.09E-01	10158	0.0541	0.0299	7.08E-02	+++	2.56E-01	23206	0.081	0.02	4.38E-05	++	2.18E-01
rs35840	63989944	SLC9A9	a	c	14666	-0.1177	0.027	1.31E-05	----?	6.91E-01	7863	-0.0587	0.0346	8.93E-02	-?	3.76E-01	25229	-0.0954	0.0213	7.45E-06	--	1.79E-01
rs831574	64004423	SLC9A9	a	g	14829	-0.0998	0.0278	3.24E-04	----?	8.70E-02	7862	-0.004	0.0337	9.06E-01	-?	8.06E-01	22701	-0.0621	0.014	4.44E-04	++	2.83E-02
rs831571	64023337	SLC9A9	t	c	18783	-0.1078	0.0236	4.85E-06	----?	8.38E-01	10231	-0.0602	0.03	4.46E-02	+++	4.16E-01	29014	-0.0896	0.0185	1.36E-06	--	1.21E-01
rs166230	64028247	SLC9A9	t	g	16176	-0.1088	0.0256	2.20E-05	----?	7.22E-01	10142	-0.044	0.0306	1.50E-01	+++	4.06E-01	26318	-0.0821	0.0196	2.89E-05	++	1.04E-01
rs12636920	65727759	MAGI2	a	t	18657	0.0813	0.0234	2.58E-04	+++++	9.53E-01	10248	0.0159	0.0304	6.01E-01	+++	2.83E-01	28905	0.057	0.0185	2.13E-03	++	8.82E-02
rs12631567	65727807	MAGI2	t	c	18681	-0.0855	0.0232	2.34E-04	+++++	9.14E-01	10223	-0.0126	0.0301	6.76E-01	+++	2.52E-01	29004	-0.0583	0.0184	1.50E-03	++	5.51E-02
rs13078950	65731571	MAGI2	t	c	14192	0.091	0.0272	8.27E-04	+++?+?+	9.48E-01	10181	0.0177	0.0304	5.60E-01	+++	3.95E-01	24773	0.0584	0.0203	3.96E-03	++	7.44E-02
rs9840838	65737292	MAGI2	t	c	18722	0.0567	0.0236	1.44E-02	+++++	4.16E-01	10350	-0.0082	0.0298	8.36E-01	+++	4.41E-01	29072	-0.0376	0.0183	3.59E-02	++	8.81E-01
rs1495453	65737587	MAGI2	a	g	16922	-0.0609	0.0236	9.85E-03	----?	3.61E-01	10336	-0.0113	0.0298	7.06E-01	+++	5.28E-01	27258	-0.0518	0.0185	2.39E-02	++	1.92E-01
rs9852224	65737750	MAGI2	t	c	12548	-0.0905	0.0275	9.91E-04	----?	7.87E-01	10363	-0.0077	0.0298	7.96E-01	+++	4.57E-01	22911	-0.0524	0.0202	9.49E-03	++	4.12E-02
rs9868785	90149113	SLC9A9	a	g	14831	0.1106	0.0287	1.14E-04	+++?+?+	5.35E-01	10337	0.0159	0.0327	6.27E-01	+++	3.39E-01	25168	0.0694	0.0216	3.30E-03	++	2.95E-02
rs9877921	90191181	SLC9A9	t	c	14762	0.1243	0.0355	4.64E-04	+++?+?+	7.76E-01	10341	0.0149	0.0332	6.55E-01	+++	4.33E-01	25103	0.0659	0.0242	6.54E-03	++	2.44E-02
rs6551450	90212018	SLC9A9	t	c	10992	-0.1013	0.033	2.14E-03	----?	3.92E-01	10322	-0.0109	0.0332	7.43E-01	+++	4.48E-01	21314	-0.0564	0.024	1.60E-03	++	5.35E-02
rs9877833	90221002	SLC9A9	a	g	14824	0.1139	0.0294	1.06E-04	+++?+?+	5.06E-01	10309	0.0121	0.0335	7.18E-01	+++	3.07E-01	25133	0.0686	0.0221	1.63E-03	++	2.24E-02
rs9849053	90235192	SLC9A9	a	g	10999	0.103	0.033	1.78E-03	+++?+?+	3.78E-01	10285	-0.008	0.0326	8.06E-01	+++	7.67E-01	21284	0.0692	0.0232	4.35E-02	++	1.67E-02
rs11717504	90261770	SLC9A9	t	c	10889	0.1174	0.0361	1.16E-03	+++?+?+	6.63E-01	10238	-0.0129	0.0314	6.82E-01	+++	5.31E-01	21127	0.0432	0.0237	6.81E-02	++	6.46E-03
rs17653828	90263360	SLC9A9	t	c	10997	-0.1024	0.033	1.90E-03	----?	3.55E-01	10263	0.0148	0.0314	6.38E-01	+++	8.38E-01	21260	-0.0409	0.0227	7.23E-02	++	1.01E-02
rs13072134	90272660	SLC9A9	t	c	10982	-0.1034	0.032	1.73E-03	----?	3.56E-01	10261	0.0092	0.0314	7.70E-01	+++	7.93E-01	21243	-0.0443	0.0227	5.15E-02	++	1.37E-02
rs14729720	90310814	SLC9A9	t	g	13043	0.1182	0.0299	7.57E-05	+++?+?+	2.79E-01	10275	-0.0238	0.0302	4.30E-01	+++	9.88E-01	23318	-0.0479	0.0212	2.42E-02	++	8.34E-04
rs9815264	90359201	SLC9A9	a	g	13044	-0.116	0.0298	9.91E-05	----?	2.96E-01	10244	0.0343	0.0299	2.51E-01	+++	9.49E-01	23388	-0.0411	0.0211	5.15E-02	++	3.70E-04
rs1472932	90366641	SLC9A9	a	g	11014	0.0526	0.0328	1.09E-01	+++?+?+	7.05E-01	10358	0.0366	0.0299	2.22E-01	+++	6.00E-01	21372	0.0439	0.0221	4.72E-02	++	7.19E-01
rs6551461	90369007	SLC9A9	a	g	11007	-0.1107	0.0347	1.42E-03	----?	3.41E-01	10303	-0.0329	0.0303	2.78E-01	+++	5.69E-01	21310	-0.0662	0.028	5.54E-03	++	9.13E-02
rs3772028	90377463	SLC9A9	t	g	10982	-0.096	0.0332	3.78E-03	+++?+?	2.18E-01	10320	-0.0291	0.0301	3.34E-01	+++	6.89E-01	21302	-0.0593	0.0223	7.85E-03	++	1.36E-01
rs7424227	90378260	SLC9A9	a	t	11010	0.099	0.0329	2.63E-03	+++?+?+	3.28E-01	10300	0.0091	0.0302	7.62E-01	+++	6.94E-01	21310	0.0502	0.0222	2.40E-02	++	4.41E-02
rs11718105	124946398	MYLK	t	c	10709	0.1175	0.1148	3.06E-01	+++?+?+	3.84E-01	387	-0.1977	0.6804	7.71E-01	-?	1.00E+00	11096	0.0888	0.1132	3.37E-01	++	6.48E-01
rs11714297	125002269	MYLK	t	c	8540	0.0774	0.0355	5.58E-01	+++?+?+	1.86E-02	382	-0.2919	0.4872	5.49E-01	-?	1.00E+00	8540	0.0798	0.1355	5.58E-01	+	1.00E+00
rs9816400	125006336	MYLK	t	c	14434	-0.0437	0.0393	6.39E-01	---?+?	2.52E-02	382	0.2919	0.4872	5.49E-01	+++	1.00E+00	14016	-0.0524	0.0914	8.56E-01	++	6.17E-01
rs14678086	125017928	MYLK	t	c	16769	0.0285	0.0888	7.48E-01	+++?+?	5.06E-02	382	0.2919	0.4872	5.49E-01	+++	1.00E+00	17151	0.037	0.0874	6.72E-01	++	5.95E-01
rs11720528	125012788	MYLK	t	c	14434	0.0387	0.0926	6.76E-01	+++?+?	3.06E-02	382	0.2919	0.4872	5.49E-01	+++	1.00E+00	14816	0.0475	0.091	6.01E-01	++	6.10E-01
rs13733884	125016973	MYLK	t	c	12497	0.044	0.0827	9.59E-01	+++?+?	6.43E-02	383	0.1664	0.4723	7.25E-01	+++	1.00E+00	12880	0.0476	0.0815	5.59E-01	++	7.99E-01
rs2605417	125032085	MYLK	a	g	14445	-0.2178	0.0712	2.22E-03	----?	1.38E-03	10358	0.0726	0.1045	4.87E-01	+++	3.45E-01	24803	-0.1257	0.0588	3.26E-02	++	2.16E-02
rs2700350	125032860	MYLK	a	g	16452	0.1632	0.0592	5.82E-03	+++?+?	1.20E-02	10345	-0.0483	0.0936	6.06E-01	+++	2.42E-01	26797	0.1028	0.05	4.00E-02	++	5.62E-02
rs2700352	125033153	MYLK	a	g	18813	0.1183	0.0561	3.49E-02	+++++	2.06E-03	10358	-0.0473	0.1133	6.76E-01	+++	2.21E-01	29171	0.0857	0.0503	8.83E-02	++	1.90E-01
rs2700354	125034462	MYLK	a	g	18812	-0.121	0.0567	3.28E-02	+++++	7.73E-04	10358	0.0675	0.0809	4.04E-01	+++	4.38E-01	29170	-0.0589	0.0464	2.05E-01	++	5.64E-02
rs2682																						

rs344088	3	158044055	FLJ16641	t	c	18812	0.2231	0.0591	1.60E-04	+++++	2.13E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21688	0.205	0.0565	2.89E-04	++	2.92E-01	
rs344091	3	158046801	FLJ16641	t	c	16780	0.2881	0.067	1.68E-05	+++++	5.70E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21656	0.2584	0.0633	4.50E-05	++	1.74E-01	
rs344092	3	158047014	FLJ16641	t	g	18810	-0.2207	0.0585	1.62E-04	+++++	8.05E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21688	-0.2031	0.056	2.88E-04	++	2.97E-01	
rs414883	3	158053397	FLJ16641	a	g	18767	-0.2158	0.0586	2.32E-04	+++++	8.43E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21643	-0.1986	0.0561	4.02E-04	--	3.08E-01	
rs263301	3	158053616	FLJ16641	a	g	18814	-0.2181	0.0585	1.91E-04	+++++	7.85E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21690	-0.2007	0.056	3.40E-04	--	3.05E-01	
rs493423	3	158056770	FLJ16641	a	g	18815	0.2143	0.0581	2.24E-04	+++++	7.00E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21678	0.1811	0.0521	5.01E-04	++	2.09E-01	
rs861359	3	158067040	FLJ16641	c	g	18814	-0.2178	0.0585	1.95E-04	+++++	7.98E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21690	-0.2004	0.056	3.46E-04	--	3.03E-01	
rs16826858	3	158061092	FLJ16641	a	g	18813	-0.2302	0.0584	8.10E-05	+++++	1.85E-02	2876	-0.0246	0.1951	9.00E-01	+	5.71E-01	21688	-0.2133	0.0559	1.38E-04	++	1.13E-01	
rs16826864	3	158062727	FLJ16641	a	g	18815	-0.2185	0.0592	2.22E-04	+++++	1.87E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21691	-0.2007	0.0566	3.94E-04	--	3.02E-01	
rs11927500	3	158063490	FLJ16641	c	g	18814	0.2168	0.0585	2.11E-04	+++++	7.32E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21690	0.1995	0.056	3.68E-04	++	3.06E-01	
rs7630909	3	158064277	FLJ16641	a	t	18813	0.2165	0.0585	2.16E-04	+++++	7.19E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21689	0.1993	0.056	3.75E-04	++	3.06E-01	
rs9878709	3	158064472	FLJ16641	a	g	18814	0.2097	0.0582	3.11E-04	+++++	6.13E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21690	0.1932	0.0558	5.31E-04	++	3.22E-01	
rs641099	3	158066549	FLJ16641	a	g	18814	-0.21	0.0582	3.08E-04	+++++	6.15E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21690	-0.1934	0.0558	5.21E-04	++	3.21E-01	
rs6411100	3	158066825	FLJ16641	t	c	18814	0.2082	0.0584	3.66E-04	+++++	2.08E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21690	0.1917	0.0559	6.10E-04	++	3.26E-01	
rs6786161	3	158073984	FLJ16641	a	g	18815	0.2075	0.0582	3.63E-04	+++++	5.35E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1912	0.0558	6.07E-04	++	3.28E-01	
rs6787535	3	158091660	FLJ16641	a	g	18814	0.2137	0.0585	2.62E-04	+++++	6.02E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21690	0.1967	0.056	4.46E-04	++	3.13E-01	
rs6787952	3	158092069	FLJ16641	a	g	18815	0.2147	0.0592	2.86E-04	+++++	1.46E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1972	0.0566	4.97E-04	++	3.11E-01	
rs16826899	3	158093361	FLJ16641	a	g	18815	0.2274	0.0578	8.47E-05	+++++	1.61E-02	2873	0.0091	0.196	9.63E-01	+	5.89E-01	21688	0.2099	0.0554	1.53E-04	++	2.85E-01	
rs13321802	3	158094022	FLJ16641	a	g	18815	0.2064	0.0582	3.89E-04	+++++	4.94E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1901	0.0558	6.64E-04	++	3.30E-01	
rs6792917	3	158104087	FLJ16641	a	g	18815	0.2067	0.0582	3.83E-04	+++++	4.79E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1904	0.0558	6.37E-04	++	3.29E-01	
rs6795327	3	158104242	FLJ16641	a	g	18815	0.2148	0.0592	2.85E-04	+++++	1.40E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1973	0.0566	4.94E-04	++	3.11E-01	
rs6795516	3	158104373	FLJ16641	t	c	18815	0.2163	0.0592	2.59E-04	+++++	1.29E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1987	0.0566	4.51E-04	++	3.07E-01	
rs6763644	3	158106050	FLJ16641	t	g	17032	-0.1546	0.0617	1.22E-02	+++++	2.93E-02	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	19908	-0.1413	0.0588	1.63E-02	++	4.75E-01	
rs8002042	3	158106159	FLJ16641	t	c	14446	0.2636	0.0693	1.43E-04	+++++	2.60E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	17322	0.2349	0.0653	3.21E-04	++	2.17E-01	
rs16836966	3	158113133	FLJ16641	c	g	18817	-0.2166	0.0583	2.05E-04	+++++	3.58E-03	2874	0.0456	0.1936	3.54E-01	+	5.95E-01	21691	-0.1948	0.0588	6.50E-02	++	1.95E-01	
rs2874490	3	158118207	FLJ16641	a	g	18817	0.2392	0.0592	2.59E-05	+++++	5.37E-03	2875	0.0246	0.1951	9.00E-01	+	5.71E-01	21692	0.2211	0.0566	9.50E-05	++	2.92E-01	
rs7609851	3	158121197	FLJ16641	a	g	18817	-0.2217	0.0592	1.83E-04	+++++	1.10E-02	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21693	-0.2036	0.0566	3.24E-04	--	2.95E-01	
rs7635521	3	158122404	FLJ16641	a	g	18815	0.21	0.0582	3.12E-04	+++++	2.24E-02	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21691	0.1934	0.0558	5.21E-04	++	3.21E-01	
rs7650127	3	158125328	FLJ16641	a	g	18813	-0.2173	0.0586	2.11E-04	+++++	2.92E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21689	-0.1999	0.0561	3.66E-04	--	3.05E-01	
rs6441103	3	158126079	FLJ16641	a	g	18796	-0.2223	0.0587	1.54E-04	+++++	3.62E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21672	-0.2045	0.0562	2.74E-04	--	2.93E-01	
rs2874491	3	158131805	FLJ16641	t	c	18815	-0.2208	0.0593	1.95E-04	+++++	1.07E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21691	-0.2208	0.0567	3.51E-04	--	2.97E-01	
rs6802511	3	158136616	FLJ16641	c	g	18733	0.2308	0.0587	8.47E-05	+++++	2.64E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21609	0.2123	0.0562	1.59E-04	++	2.74E-01	
rs6441104	3	158141335	FLJ16641	a	g	18814	0.224	0.0592	1.55E-04	+++++	1.12E-03	2876	0.0088	0.1944	9.64E-01	+	5.90E-01	21690	0.2057	0.0566	2.80E-04	++	2.90E-01	
rs16826948	3	158164468	FLJ16641	t	c	18814	-0.2243	0.0592	1.52E-04	+++++	1.13E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21690	-0.2056	0.0566	2.75E-04	--	2.89E-01	
rs6743233	3	158165805	FLJ16641	t	c	18814	-0.2229	0.0592	1.56E-04	+++++	1.16E-03	2876	-0.0088	0.1944	9.64E-01	+	5.90E-01	21690	-0.2056	0.0566	2.82E-04	--	2.90E-01	
61	rs488209	3	174602072	NLGN1	a	14829	0.1165	0.0273	1.96E-05	+++++	2.22E-01	10313	0.0402	0.0308	1.92E-01	++	6.28E-01	25142	0.0829	0.0204	4.92E-05	++	6.38E-02	
rs536445	3	174602797	NLGN1	t	c	14728	0.1162	0.0271	1.83E-05	+++++	1.50E-01	10345	0.0313	0.0305	3.05E-01	++	8.36E-01	25073	0.0787	0.0203	1.02E-04	++	3.75E-02	
rs992367	3	175128323	NLGN1	c	g	6847	-0.0469	0.0244	2.68E-01	?????	3.34E-02	9862	-0.0049	0.0306	8.72E-01	??	9.12E-01	16709	-0.0193	0.0248	4.37E-01	++	4.22E-01	
rs963350	3	175164600	NLGN1	t	c	14827	-0.1113	0.0266	2.80E-05	+++++	8.85E-03	7873	0.003	0.034	9.29E-01	??	7.13E-01	22700	-0.0979	0.021	1.93E-01	++	8.10E-03	
rs16833234	3	175293651	NLGN1	t	c	14830	0.0949	0.0262	2.87E-04	+++++	7.98E-01	7875	0.0063	0.0343	8.54E-01	??	8.75E-01	22705	0.0623	0.0208	2.79E-03	++	4.01E-02	
62	rs9842394	3	181109590	PKSL	t	c	18780	0.1158	0.0309	1.79E-04	+++++	2.06E-02	10356	0.0063	0.0435	8.85E-01	++	9.71E-01	19136	0.0791	0.0252	1.70E-03	++	4.02E-02
rs9836757	3	181124915	PKSL	t	c	18638	0.1168	0.0301	1.03E-04	+++++	4.18E-02	10357	-0.0272	0.0431	5.29E-01	++	7.04E-01	28995	0.0696	0.0267	4.80E-03	++	6.16E-03	
rs11748664	3	181132205	PKSL	t	c	18686	0.1002	0.03	8.35E-04	+++++	1.58E-02	10361	-0.0447	0.0462	3.33E-01	++	8.50E-01	29047	0.0592	0.0242	2.30E-02	++	8.53E-03	
rs1416789	3	181167899	PKSL	a	g	18645	0.1099	0.0301	2.60E-04	+++++	2.46E-02	10347	-0.047	0.0461	3.29E-01	++	8.16E-01	28992	0.0636	0.0252	1.16E-02	++	4.90E-03	
rs16830940	3	181140955	PKSL	a	g	12507	0.1113	0.0363	2.19E-04	+++++	1.63E-02	10347	-0.0417	0.0461	3.29E-01	++	8.03E-01	28554	0.0572	0.0285	6.50E-02	++	9.03E-03	
rs7634072	3	181144012	PKSL	a	g	18686	0.1009	0.0301	8.08E-04	+++++	2.62E-02	10357	0.0013	0.0421	9.76E-01	++	5.39E-01	29043	0.0623	0.0245	6.05E-03	++	5.43E-02	
63	rs6784185	3	186955759	IGFBP2	a	g	18804	0.1001	0.0268	1.88E-04	+++++	3.72E-01	10326	0.0694	0.0352	4.84E-02	++	5.21E-03	29130	0.0888	0.0213	3.10E-05	++	4.88E-01
rs7684050	3	186955826	IGFBP2	t	c	18779	0.1011	0.0268	1.62E-04	+++++	3.66E-01	10329	0.0572	0.0346	9.83E-02	++	7.02E-03	29108	0.0846	0.0212	6.48E-05	++	3.16E-01	
rs6799330	3	186961651	IGFBP2	t	g	18751	-0.0995	0.0269	2.10E-04	+++++	3.66E-01	10342	-0.0083	0.0303	7.83E-01	++	2.96E-02	29093	-0.0593	0.0201	3.20E-03	++	2.44E-02	
rs6789741	3	186962427	IGFBP2	t	g	18736	0.0971	0.0269	3.05E-04	+++++	3.11E-													

rs7690771	4	21623733	a	g	16821	-0.0497	0.0285	8.14e-02	---+?--	2.13e-01	10361	-0.0251	0.0348	4.70e-01	+++	8.50e-01	27182	-0.0398	0.022	7.09e-02	--	5.84e-01		
rs1513574	4	21625239	a	g	10662	-0.048	0.0351	1.71e-01	---+?--	4.80e-01	10352	-0.0428	0.0358	2.31e-01	+++	8.03e-01	21014	-0.0455	0.0251	6.98e-02	--	9.17e-01		
rs13125289	4	21626265	a	g	14482	0.0573	0.0311	6.53e-02	---+?--	2.35e-01	10353	0.0382	0.0362	2.91e-01	+++	6.64e-01	24835	-0.0492	0.0376	3.71e-02	--	6.97e-01		
rs2015779	4	21637770	t	c	18605	-0.0748	0.0261	4.12e-03	---+?--	2.30e-02	10265	-0.0519	0.0362	1.51e-01	+++	7.68e-01	28870	-0.067	0.0212	1.56e-03	--	6.08e-01		
rs175778	4	21657874	a	g	18744	0.0752	0.026	3.82e-03	---+?--	2.37e-02	10340	0.0472	0.0361	1.91e-01	+++	7.03e-01	28904	0.056	0.0211	1.86e-03	++	5.29e-01		
rs1652833	4	21658584	a	g	18739	-0.0747	0.026	4.04e-03	---+?--	2.56e-02	10357	-0.0519	0.0322	1.71e-01	+++	9.95e-01	29082	-0.0499	0.0202	1.36e-03	++	1.29e-01		
rs7567801	4	21655334	a	g	12497	-0.0831	0.0315	8.37e-03	---+?--	1.17e-02	10341	-0.0173	0.0325	5.94e-01	---	9.70e-01	22838	-0.0512	0.0226	2.35e-02	--	1.46e-01		
rs13106072	4	21653470	a	g	16404	-0.069	0.028	1.38e-02	---+?--	1.53e-02	10341	-0.0155	0.0326	6.34e-01	---	9.73e-01	26745	-0.0463	0.0212	2.93e-02	--	2.13e-01		
rs6836373	4	21654727	a	g	18739	0.0757	0.026	3.58e-03	---+?--	2.51e-02	10322	0.0324	0.0326	6.51e-01	+++	9.62e-01	29071	0.052	0.0203	1.05e-02	++	1.44e-01		
rs6839252	4	21655010	c	g	16705	-0.1019	0.0281	2.92e-04	---+?--	1.18e-01	10326	-0.0149	0.0326	6.49e-01	---	9.63e-01	27031	-0.0648	0.0213	2.33e-02	++	4.32e-02		
rs12510345	4	21661915	a	g	16730	-0.0448	0.0274	1.02e-01	---+?--	1.87e-01	10305	-0.0296	0.0304	3.30e-01	---	6.90e-01	27035	-0.038	0.0204	6.20e-02	--	7.10e-01		
rs17345664	4	21662382	t	c	16725	-0.0455	0.0274	9.70e-02	---+?--	1.90e-01	10305	-0.0205	0.0298	4.91e-01	---	7.40e-01	27030	-0.0324	0.0202	9.14e-02	---	5.37e-01		
71	rs1626840	4	42533105	t	c	18815	0.0918	0.0249	2.27e-04	-----	6.11e-01	10362	-0.0363	0.0322	2.60e-01	0	1.37e-01	29177	0.0439	0.0197	2.60e-02	++	1.65e-03	
rs1200251	4	42545916	t	c	18795	0.0755	0.0256	3.15e-03	-----	5.99e-01	10363	-0.0249	0.0381	5.14e-01	+++	8.99e-02	29158	0.0403	0.0212	3.72e-02	++	2.87e-02		
rs10938223	4	42549703	a	g	18815	0.0821	0.0229	3.36e-04	-----	1.99e-02	10372	-0.0394	0.0359	5.24e-01	---	4.27e-01	21688	0.0643	0.0211	2.29e-03	++	4.67e-02		
rs10004231	4	42552330	t	c	18813	-0.0798	0.0229	4.80e-04	-----	2.04e-01	10371	0.0371	0.0359	4.92e-01	+++	4.38e-01	21688	-0.0619	0.0211	3.30e-03	++	4.59e-02		
rs11730312	4	42557311	c	g	18803	0.077	0.0255	2.59e-03	-----	7.27e-01	10358	-0.0281	0.0309	3.63e-01	---	1.26e-01	01614	0.0347	0.0197	8.01e-02	++	8.71e-03		
rs6850606	4	42559453	a	g	18794	-0.0759	0.0255	2.96e-03	-----	7.32e-01	10357	0.0164	0.032	6.08e-01	---	7.92e-02	01521	-0.0401	0.0199	4.46e-02	++	2.41e-02		
72	rs1512128	4	46593834	a	c	18743	-0.1351	0.0427	1.56e-03	-----	8.72e-01	10344	-0.0066	0.0457	8.85e-01	---	9.79e-01	29087	-0.0752	0.0312	1.59e-02	---	3.99e-02	
rs7960061	4	46601198	a	t	18797	-0.1342	0.0425	1.58e-03	-----	8.83e-01	10344	-0.0066	0.0457	8.85e-01	---	9.79e-01	01941	-0.075	0.0311	1.59e-02	---	4.09e-02		
rs16859669	4	46602060	t	c	14824	-0.1376	0.0484	4.46e-03	---+?--	8.40e-01	10344	-0.0066	0.0457	8.85e-01	---	9.79e-01	25168	-0.0683	0.0332	3.97e-02	---	4.91e-02		
rs17641147	4	46604292	t	c	16763	-0.1297	0.0451	4.06e-03	---+?--	8.81e-01	10344	-0.0066	0.0457	8.85e-01	---	9.79e-01	27107	-0.069	0.0321	3.17e-02	---	5.52e-02		
rs2351168	4	46601925	t	g	18795	-0.1392	0.0426	1.08e-03	-----	9.03e-01	10344	-0.0066	0.0457	8.85e-01	---	9.79e-01	01939	-0.0775	0.0312	1.28e-02	---	3.38e-02		
rs9952303	4	46614384	GABRA4	t	c	18793	0.139	0.0426	1.10e-03	-----	9.01e-01	10352	0.006	0.0457	8.96e-01	+++	9.84e-01	01945	0.0772	0.0312	1.33e-02	++	3.33e-02	
rs15121347	4	46626376	GABRA4	a	g	16458	-0.1574	0.0474	9.98e-04	-----	8.94e-01	10352	0.006	0.0457	8.96e-01	+++	9.84e-01	01945	0.0772	0.0312	1.33e-02	++	3.33e-02	
rs6815456	4	46627316	GABRA4	a	g	18793	0.1402	0.0426	9.91e-04	-----	8.91e-01	10352	0.006	0.0457	8.96e-01	+++	9.84e-01	01945	0.0772	0.0312	1.25e-02	++	3.17e-02	
rs12508647	4	46628711	GABRA4	c	g	18794	0.1408	0.0426	9.41e-04	-----	8.89e-01	10353	0.0062	0.0457	8.93e-01	+++	9.82e-01	01947	0.0782	0.0312	1.21e-02	++	3.12e-02	
rs10024134	4	46629556	GABRA4	t	c	18799	0.1424	0.0425	7.99e-04	-----	8.86e-01	10357	0.005	0.0457	9.13e-01	+++	9.89e-01	01956	0.0787	0.0311	1.15e-02	++	2.77e-02	
rs1877397	4	46632377	GABRA4	a	g	18799	0.1424	0.0425	8.15e-04	-----	8.65e-01	10357	0.005	0.0457	9.13e-01	+++	9.89e-01	01956	0.0787	0.0311	1.15e-02	++	2.77e-02	
rs10018276	4	46636923	GABRA4	t	c	14824	-0.1417	0.0485	3.47e-03	---+?--	8.08e-01	10357	-0.005	0.0457	9.13e-01	---	9.89e-01	25181	-0.0693	0.0333	3.72e-02	---	4.02e-02	
rs1877403	4	46647212	GABRA4	t	c	18801	-0.1433	0.0425	7.54e-04	-----	8.62e-01	10357	-0.0087	0.0461	8.50e-01	---	9.95e-01	01958	-0.0615	0.0312	9.14e-03	---	3.18e-02	
rs1398175	4	46644635	GABRA4	t	g	18785	-0.1424	0.0425	8.07e-04	-----	9.30e-01	10357	-0.0087	0.0461	8.50e-01	---	9.95e-01	01958	-0.0615	0.0312	9.14e-03	---	3.18e-02	
rs1398176	4	46644483	GABRA4	t	g	18810	-0.1422	0.0425	8.16e-04	-----	8.70e-01	10357	-0.0087	0.0461	8.50e-01	---	9.97e-01	01961	-0.0615	0.0312	9.14e-03	---	3.18e-02	
rs7666667	4	46645470	GABRA4	t	c	18809	0.1416	0.0426	8.75e-04	-----	8.81e-01	10357	0.005	0.0457	9.13e-01	+++	9.97e-01	01961	-0.0615	0.0312	9.14e-03	---	3.18e-02	
rs6896865	4	46646786	GABRA4	a	g	18726	-0.1386	0.0434	4.40e-03	---+?--	8.47e-01	10357	-0.006	0.0457	9.13e-01	---	9.97e-01	01961	-0.0615	0.0312	9.14e-03	---	3.18e-02	
rs7695616	4	46648003	GABRA4	t	c	18813	-0.1503	0.0424	3.94e-04	-----	7.68e-01	10357	-0.0179	0.0457	9.13e-01	---	9.86e-01	01678	-0.126	0.0383	1.01e-03	---	1.81e-01	
rs16851641	4	46647090	GABRA4	a	g	18810	-0.1419	0.0425	8.32e-04	-----	8.43e-01	10357	-0.0241	0.0457	9.13e-01	---	9.93e-01	01679	-0.126	0.0383	1.01e-03	---	1.81e-01	
rs16859751	4	46647816	GABRA4	t	c	18805	-0.1417	0.0427	9.15e-04	-----	8.81e-01	10358	-0.0097	0.0457	9.13e-01	---	9.85e-01	01963	-0.08	0.0312	1.02e-02	---	3.46e-02	
rs7685553	4	46648254	GABRA4	t	c	18805	0.1439	0.0427	7.47e-04	-----	8.61e-01	10357	0.0027	0.0457	9.13e-01	+++	9.97e-01	01676	0.1212	0.0383	1.64e-03	++	2.19e-01	
rs12505347	4	46649834	GABRA4	t	c	16770	0.1331	0.0455	3.46e-03	---+?--	8.63e-01	10357	0.0027	0.0457	9.13e-01	+++	9.97e-01	01961	0.1102	0.0405	6.52e-03	++	2.69e-01	
73	rs12503480	4	73122656	NPPFR2	t	c	16421	0.0928	0.0262	3.93e-04	-----	4.40e-01	9935	-0.0427	0.0386	1.63e-01	?	9.22e-01	26356	0.0359	0.0199	7.44e-02	++	7.69e-04
rs13107558	4	73125670	NPPFR2	c	g	16420	0.0913	0.0263	1.73e-04	-----	3.68e-01	9934	-0.0182	0.0294	5.35e-01	?	7.40e-01	26654	0.0469	0.0187	1.24e-02	++	4.09e-03	
rs7695255	4	73126973	NPPFR2	a	g	16700	0.0941	0.0243	3.53e-04	-----	3.41e-01	9935	-0.0186	0.0294	5.26e-01	?	7.13e-01	26441	0.0404	0.0196	2.48e-02	++	4.28e-03	
rs7670209	4	73127071	NPPFR2	c	g	16744	-0.0901	0.0243	2.07e-04	-----	3.72e-01	9965	0.0166	0.0293	5.71e-01	?	5.90e-01	26709	0.0466	0.0187	1.27e-02	++	5.06e-03	
rs7670216	4	73127071	NPPFR2	c	g	16745	-0.0892	0.0243	2.07e-04	-----	3.72e-01	9965	0.0166	0.0293	5.71e-01	?	5.90e-01	26709	0.0466	0.0187	1.27e-02	++	5.06e-03	
rs13121468	4	73127294	NPPFR2	t	c	16423	-0.0803	0.0228	4.25e-04	-----	3.54e-01	9966	0.0176	0.0293	5.48e-01	?	5.90e-01	26709	0.0434	0.018				

	rs12643619	4	75625891	t	c	14135	-0.0573	0.0314	6.78E-02	-----??	3.80E-03	10250	-0.0036	0.0326	9.12E-01	++	5.10E-01	24385	-0.0315	0.0226	1.64E-01	--	2.36E-01	
	rs6447006	4	75628191	a	g	16772	-0.0465	0.0278	9.40E-02	-----?+	6.61E-03	10340	-0.0163	0.0335	6.26E-01	++	3.50E-01	27112	-0.0342	0.0214	1.10E-01	--	4.88E-01	
76	rs689040	4	87988115	c	g	12714	0.1791	0.0702	1.12E-02	+++++??	6.13E-01	7844	-0.3911	0.1637	1.76E-02	-?	4.69E-01	20558	0.0906	0.0646	1.61E-01	++	1.48E-03	
	rs1497768	4	82954871	a	t	16548	0.16	0.0478	8.14E-04	+++++??	1.34E-01	7868	-0.08	0.0772	3.00E-01	-?	1.38E-02	24416	0.0935	0.0406	2.14E-02	++	8.21E-03	
	rs12643522	4	85731222	g	t	18656	-0.1563	0.0442	4.08E-04	-----??	1.80E-01	7873	0.0653	0.0765	3.94E-01	-?	3.08E-01	26528	-0.1008	0.0383	8.42E-03	++	1.21E-02	
77	rs7659871	4	85702451	t	c	18813	-0.072	0.0239	2.56E-03	-----??	3.94E-01	10263	0.0363	0.0111	0.0287	0.03E-01	++	2.23E-01	9176	0.047	0.0184	1.05E-02	++	1.02E-01
	rs7678268	4	85703738	t	c	18789	0.059	0.0238	1.33E-02	+++++??	2.59E-01	10326	0.0217	0.0289	4.53E-01	++	3.32E-01	29115	0.0439	0.0184	1.68E-02	++	3.19E-01	
	rs4494005	4	85704353	a	g	16452	-0.0811	0.0259	1.76E-03	-----?+	4.34E-01	10340	-0.0223	0.0289	4.40E-01	++	2.30E-01	26792	-0.0549	0.0193	4.42E-03	++	1.30E-01	
	rs4533768	4	85704655	g	t	18787	0.0649	0.0239	6.53E-03	+++++??	2.38E-01	10327	0.0238	0.0289	4.09E-01	++	3.27E-01	29114	0.0482	0.0184	8.86E-03	++	2.73E-01	
	rs6355544	4	85705077	a	c	18794	0.0612	0.0239	1.04E-02	+++++??	1.54E-01	10340	0.0216	0.0289	4.53E-01	++	2.37E-01	29134	0.0451	0.0184	1.83E-02	++	2.91E-01	
	rs6750366	4	85705965	t	c	16470	0.0868	0.0258	7.54E-04	+++++??	7.64E-01	10331	0.0217	0.0289	4.53E-01	++	3.23E-01	26801	-0.0579	0.0192	2.61E-03	++	2.92E-01	
	rs1109652	4	85708656	c	g	14946	-0.0687	0.0256	7.41E-03	-----?+	1.03E-01	10318	-0.0238	0.0289	4.10E-01	++	3.04E-01	25264	-0.0499	0.0192	1.06E-02	++	4.25E-01	
	rs1109653	4	85709521	c	g	18643	0.0712	0.0238	2.82E-03	+++++??	2.77E-01	10324	0.0227	0.0288	4.30E-01	++	1.93E-01	28967	0.0515	0.0183	5.98E-03	++	1.94E-01	
	rs10776519	4	85710291	g	t	16322	-0.1014	0.0238	8.69E-05	-----?+	8.31E-01	10332	-0.0226	0.0288	4.34E-01	++	1.20E-01	26654	-0.0663	0.0192	4.59E-04	++	4.16E-02	
	rs12643523	4	85712722	t	c	18666	0.0761	0.0239	1.42E-03	+++++??	2.43E-01	10274	0.022	0.0288	4.45E-01	++	9.74E-02	29000	0.054	0.0184	3.30E-03	++	1.48E-01	
	rs2868914	4	85715785	t	c	14775	0.0523	0.0268	5.05E-02	+++++??	6.16E-02	10212	0.0154	0.0289	5.94E-01	++	8.84E-02	25087	-0.0352	0.0197	7.29E-02	++	3.49E-01	
	rs2868915	4	85716071	t	c	12764	-0.0693	0.0287	1.56E-02	-----?+	1.41E-01	10355	-0.0211	0.0288	4.63E-01	++	7.45E-02	23119	-0.0453	0.0203	2.59E-02	++	2.36E-01	
	rs2868917	4	85719225	t	c	12982	-0.0632	0.0272	1.99E-02	-----?+	1.16E-01	10332	-0.0218	0.0288	4.50E-01	++	1.20E-01	23314	-0.0437	0.0198	2.72E-02	++	2.96E-01	
	rs11097023	4	85724498	t	c	14832	-0.0705	0.0269	8.85E-03	-----?+	1.95E-01	10262	-0.012	0.0291	6.81E-01	++	1.04E-01	25094	-0.0438	0.0198	2.75E-02	++	1.40E-01	
	rs11725339	4	85728173	a	g	8843	-0.0362	0.0358	3.11E-01	????+??	1.57E-01	10290	-0.0119	0.0292	6.83E-01	++	8.50E-02	19133	-0.0216	0.0226	3.40E-01	++	5.90E-01	
78	rs1514733	4	91883046	t	c	14832	0.0919	0.0255	3.22E-04	+++++??	4.53E-01	10363	0.0151	0.0286	5.99E-01	++	6.00E-01	25195	0.0759	0.0191	2.36E-03	++	4.50E-02	
	rs6843193	4	91884426	a	g	14793	0.0565	0.0255	2.68E-02	+++++??	2.92E-02	10320	0.0152	0.0357	6.70E-01	++	3.09E-01	25113	0.0425	0.0208	4.03E-02	++	3.47E-01	
	rs11932857	4	91934031	a	g	10324	-0.0419	0.0312	1.80E-01	-----?+	3.28E-01	10159	-0.0025	0.029	9.30E-01	++	3.12E-01	20483	-0.0208	0.0212	3.28E-02	++	3.55E-01	
	rs7438052	4	91935637	a	c	4512	-0.0385	0.0555	4.88E-01	????+??	6.16E-01	7487	-0.018	0.0358	5.94E-01	??	1.00E+00	11999	-0.0235	0.0289	4.15E-01	++	7.52E-01	
	rs2852086	4	91936008	a	c	16446	-0.0285	0.0248	2.50E-01	-----?+	1.36E-01	10359	-0.0098	0.0289	7.30E-01	++	3.28E-01	28136	-0.0481	0.0178	6.95E-03	++	4.49E-01	
	rs12168076	4	91939223	t	c	14131	0.0038	0.027	8.88E-01	+++++??	7.32E-02	10156	0.0011	0.029	9.71E-01	++	3.15E-01	24287	0.0205	0.0198	8.98E-01	++	9.46E-01	
	rs1213957	4	91939429	t	c	14512	-0.0081	0.0268	7.62E-01	-----?+	5.71E-02	10156	0.0055	0.029	8.49E-01	++	2.67E-01	24668	-0.0108	0.0197	9.26E-01	++	7.31E-01	
	rs12168077	4	91939662	t	c	16231	0.0199	0.025	4.27E-01	+++++??	1.57E-01	10232	-0.0031	0.0289	9.13E-01	++	2.27E-01	26463	-0.0119	0.0189	5.95E-01	++	5.47E-01	
	rs11097267	4	91940596	a	t	18813	0.0348	0.0229	1.29E-01	+++++??	2.49E-02	10216	-0.0018	0.0289	9.51E-01	++	3.29E-01	29029	-0.0207	0.0179	2.49E-01	++	3.21E-01	
	rs1213960	4	91940938	t	c	18762	-0.0289	0.0229	2.07E-01	-----?+	3.06E-02	10229	0.0044	0.0289	8.79E-01	++	2.31E-01	28991	-0.0161	0.0179	3.71E-01	++	3.67E-01	
	rs1213961	4	91940963	a	g	18784	0.0353	0.0229	1.23E-01	+++++??	4.39E-02	10216	-0.002	0.0289	9.46E-01	++	3.35E-01	29000	-0.0209	0.0179	2.44E-01	++	3.12E-01	
	rs1377918	4	91941262	a	g	17011	0.0402	0.0234	8.61E-02	+++++??	2.96E-02	10228	-0.0053	0.0289	8.55E-01	++	2.48E-01	27239	0.0222	0.0182	2.32E-01	++	2.21E-01	
	rs1377919	4	91941548	a	t	14427	-0.0044	0.0268	8.69E-01	+++++??	3.83E-01	10232	-0.006	0.0289	8.36E-01	++	2.45E-01	24659	-0.0051	0.0197	7.94E-01	++	9.68E-01	
	rs10516878	4	91943890	a	g	18811	-0.0351	0.0229	1.26E-01	+++++??	2.37E-02	10237	0.0007	0.0288	8.08E-01	++	3.27E-01	29048	-0.0188	0.0179	2.94E-01	++	2.53E-01	
79	rs2073017	4	110819267	t	c	18517	0.0691	0.0229	2.60E-02	+++++??	1.57E-01	10330	0.0146	0.0284	6.24E-01	++	6.36E-01	28794	-0.0476	0.0179	7.78E-03	++	1.33E-01	
	rs5234444	4	110780555	t	c	16961	-0.0648	0.0233	5.30E-03	-----?+	1.75E-01	10315	-0.0114	0.0285	6.89E-01	++	7.56E-01	27276	-0.0434	0.018	1.61E-02	++	1.47E-01	
	rs12504784	4	110787207	a	g	16349	-0.0851	0.0246	5.51E-04	-----?+	2.41E-01	10323	-0.0082	0.0285	7.74E-01	++	8.18E-01	26672	-0.0528	0.0186	0.03E-01	++	4.11E-02	
	rs10010159	4	110787247	t	c	18774	-0.0717	0.0228	1.67E-03	-----?+	1.42E-01	10358	-0.0077	0.0285	7.86E-01	++	8.03E-01	29132	-0.0467	0.0178	8.68E-03	++	7.95E-02	
	rs9990765	4	110788160	t	c	18779	0.0706	0.0228	1.97E-03	+++++??	1.36E-01	10359	0.0078	0.0285	7.85E-01	++	8.02E-01	29138	-0.0461	0.0178	9.63E-03	++	8.53E-02	
	rs9993532	4	110788540	t	c	18779	0.071	0.0228	1.87E-03	+++++??	1.40E-01	10359	0.0078	0.0285	7.85E-01	++	8.02E-01	29138	-0.0463	0.0178	9.25E-03	++	8.33E-02	
	rs726585	4	110788996	t	c	18778	-0.0711	0.0228	1.84E-03	-----?+	1.39E-01	10359	-0.0078	0.0285	7.85E-01	++	8.02E-01	29137	-0.0464	0.0178	9.16E-03	++	8.29E-02	
	rs9996730	4	110789544	t	c	18778	0.0712	0.0228	1.81E-03	+++++??	1.38E-01	10359	0.0078	0.0285	7.85E-01	++	8.02E-01	29137	-0.0465	0.0178	9.07E-03	++	8.24E-02	
	rs1029449	4	110789716	t	c	18778	-0.0712	0.0228	1.81E-03	-----?+	1.38E-01	10359	-0.0078	0.0285	7.85E-01	++	8.02E-01	29137	-0.0465	0.0178	9.07E-03	++	8.24E-02	
	rs9825905	4	110793192	t	c	16301	-0.0919	0.0247	2.01E-04	-----?+	1.93E-01	10278	-0.0083	0.0286	7.71E-01	++	7.65E-01	26579	-0.0562	0.0187	2.65E-03	++	2.70E-02	
	rs6851190	4	110794027	t	c	18777	-0.0726	0.0228	1.46E-03	-----?+	1.36E-01	10359	-0.0098	0.0285	7.93E-01	++	7.55E-01	28136	-0.0481	0.0178	6.95E-03	++	4.49E-01	
	rs6523445	4	110794058	a	g	18762	-0.045	0.0228	4.90E-02	-----?+	1.53E-02	10349	-0.											

rs1023962	4	153670806	FBXW7	t	g	18714	0.0781	0.0227	5.94e-04	+++++	2.37e-01	9973	0.0517	0.029	7.45e-02	7++	6.58e-01	28687	0.0681	0.0179	1.40e-04	++	4.74e-01
rs1484879	4	153675078	DKFZ34340714, FBXW7	a	g	18692	0.0779	0.0227	6.15e-04	+++++	2.39e-01	9973	0.0502	0.029	8.32e-02	7++	6.36e-01	28665	0.0674	0.0179	1.64e-04	++	4.52e-01
rs6832998	4	153690116		a	t	18562	0.0847	0.0229	2.16e-04	+++++	1.13e-01	10329	0.0315	0.015	2.68e-01	++	4.79e-01	28891	0.0637	0.0178	3.50e-04	++	4.54e-01
rs469135	4	153730038		t	c	16645	0.0856	0.0243	4.30e-04	+++++	1.42e-01	9883	0.03	0.0292	3.05e-01	7++	7.70e-01	26528	0.0668	0.0187	7.66e-04	++	1.43e-01
rs1680783	4	153730038		a	g	16691	0.0955	0.0243	4.39e-05	+++++	2.68e-01	9877	0.032	0.0292	2.56e-01	7++	9.46e-01	26568	0.07	0.0187	1.78e-04	++	1.30e-01
rs1676833	4	153732819		a	t	16665	-0.0981	0.0244	4.72e-05	+++++	2.68e-01	9870	0.0402	0.0294	1.72e-01	7++	9.93e-01	26535	-0.0745	0.0188	7.28e-05	++	1.30e-01
rs4696327	4	153735030		g	c	16549	0.089	0.0245	2.73e-04	+++++	1.78e-01	9839	0.0398	0.0291	1.72e-01	7++	9.42e-01	26388	0.0686	0.0187	2.52e-04	++	1.96e-01
rs10020796	4	153739231		t	c	16468	0.0936	0.0246	1.40e-04	+++++	1.57e-01	9868	0.036	0.0292	2.18e-01	7++	8.13e-01	26336	0.0697	0.0188	2.12e-04	++	1.31e-01
rs6813195	4	153739925		t	c	18815	-0.0874	0.0227	1.20e-04	-----	2.60e-01	9973	-0.0289	0.0289	3.18e-01	7--	9.95e-01	28788	-0.0651	0.0179	2.67e-04	--	1.11e-01
rs817587	4	15437618		a	g	16420	0.0981	0.0271	2.93e-04	+++++	1.71e-01	10332	0.0061	0.0339	8.56e-01	++	8.54e-01	26752	0.0622	0.0212	3.28e-04	++	3.40e-02
rs6822903	4	15437975		a	g	16640	-0.0007	0.0272	9.81e-01	+++++	3.76e-05	10324	-0.0025	0.0339	9.42e-01	++	8.31e-01	26964	-0.0014	0.0212	9.47e-01	++	9.67e-01
rs6539906	4	15439696		a	g	18732	-0.0607	0.0252	1.62e-02	-----	1.72e-01	10198	-0.0106	0.0342	7.58e-01	---	9.26e-01	28920	-0.0431	0.0203	3.28e-04	---	2.38e-01
rs2711283	4	161172446		t	c	10140	0.1191	0.0416	4.17e-03	+++++	6.84e-01	9851	-0.0089	0.0391	8.20e-01	7--	9.27e-01	19991	0.0711	0.0285	2.72e-04	---	2.50e-02
rs2711274	4	161192589		t	c	14803	0.1163	0.0331	4.49e-04	+++++	7.03e-01	9857	-0.011	0.0447	8.06e-01	7--	1.00e+00	22290	0.0512	0.0266	7.42e-03	---	2.21e-02
rs2712273	4	161319002		t	c	14832	0.1198	0.0352	3.05e-04	+++++	6.36e-01	9865	-0.0095	0.0395	1.00e+00	7?0	1.00e+00	22318	0.071	0.0266	3.82e-03	0	6.79e-02
rs2727448	4	181646514		a	g	18708	0.0452	0.0269	9.32e-02	+++++	2.52e-02	10343	-0.0285	0.0301	3.42e-01	---	3.66e-02	29051	0.0125	0.0201	5.24e-01	---	6.79e-02
rs2727449	4	181647694		a	g	4512	0.0611	0.0632	3.33e-01	+++++	3.79e-02	7487	0.007	0.0338	8.36e-01	7+0	1.00e+00	11999	0.0119	0.0298	5.23e-01	++	4.50e-01
rs2727450	4	181648854		t	c	16734	0.0409	0.0281	1.46e-01	+++++	4.60e-02	10348	-0.0462	0.0294	1.16e-01	---	5.99e-02	27082	-0.0097	0.0203	9.73e-01	---	3.22e-02
rs2546573	4	181649549		t	c	18730	0.0502	0.0269	6.20e-02	+++++	2.70e-02	10348	-0.0399	0.0298	1.80e-01	---	4.06e-02	29078	0.0097	0.02	6.26e-01	---	2.48e-02
rs2546572	4	181651341		c	g	16757	0.0777	0.028	5.45e-03	+++++	4.71e-01	10355	-0.03	0.0303	3.23e-01	---	3.29e-02	27112	0.0281	0.0206	1.72e-01	---	9.04e-03
rs3095962	4	181656252		t	c	12395	-0.076	0.0339	2.50e-02	-----	7.96e-03	10124	0.0287	0.0235	3.77e-01	++	1.57e-03	22519	-0.0214	0.0235	3.61e-01	---	2.58e-02
rs2546570	4	181660102		g	c	18797	-0.0504	0.0268	6.05e-02	-----	3.42e-02	10338	0.0171	0.0327	6.02e-01	---	9.92e-03	29135	-0.0233	0.0207	2.61e-01	---	1.10e-01
rs2546614	4	181663353		g	c	14825	-0.1086	0.0301	3.04e-04	-----	4.60e-01	10338	0.0161	0.0327	6.23e-01	++	1.21e-02	25163	-0.0514	0.0221	2.03e-02	---	5.02e-03
rs2727413	4	181666872		g	c	18635	0.0836	0.0268	1.42e-03	+++++	4.04e-01	10324	-0.0246	0.0328	4.53e-01	---	1.13e-02	28959	0.0415	0.0208	4.56e-02	---	9.28e-03
rs2727414	4	181666872		a	g	16479	-0.0832	0.0253	9.86e-04	-----	8.32e-01	9929	-0.0109	0.0296	9.71e-01	7+	3.29e-01	2451	-0.0487	0.0192	3.12e-01	---	3.08e-02
rs3796649	4	188154024		c	g	18807	0.0843	0.0223	2.90e-04	+++++	9.06e-01	9966	-0.007	0.0298	8.12e-01	7--	2.80e-01	28773	0.0494	0.0183	7.00e-03	---	1.54e-02
rs2964727	5	6219717		a	t	18810	-0.0792	0.0491	1.07e-01	-----	8.49e-03	2875	0.0135	0.0156	8.98e-01	++	2.00e-02	21685	-0.0627	0.0445	1.59e-01	---	4.26e-01
rs1428747	5	6219973		c	g	18805	0.0703	0.0483	1.46e-01	+++++	6.28e-02	2876	-0.0183	0.0156	8.63e-01	++	1.90e-02	21681	-0.055	0.039	2.11e-01	---	4.46e-01
rs2937799	5	6220264		g	c	18814	0.071	0.0483	1.41e-01	+++++	6.66e-03	10362	0.0644	0.0633	3.09e-01	++	4.37e-02	29176	0.0686	0.0384	7.41e-02	---	9.34e-01
rs7445173	5	6223934		t	c	18813	0.0705	0.0483	1.44e-01	+++++	6.66e-03	10362	0.0627	0.0631	3.21e-01	++	4.45e-02	29175	0.0676	0.0384	7.79e-02	---	9.22e-01
rs11745623	5	6229413		c	g	18800	0.0687	0.0484	1.56e-01	+++++	6.98e-03	2874	-0.0143	0.0156	8.92e-01	++	1.98e-02	21674	-0.0543	0.044	2.17e-01	---	4.75e-01
rs2937798	5	6231155		t	c	16461	-0.1662	0.0513	1.20e-03	-----	2.91e-01	2808	0.0355	0.0167	7.39e-01	++	1.57e-02	19269	-0.1283	0.0462	5.51e-03	---	8.84e-02
rs2964722	5	6232771		c	g	18774	-0.17	0.0484	4.50e-04	-----	3.25e-01	10361	-0.0521	0.0632	4.10e-01	++	4.88e-02	29135	-0.1264	0.0384	1.00e-03	---	1.39e-01
rs2964716	5	6234008		a	g	14149	-0.0054	0.0556	9.23e-01	+++++	1.37e-02	7276	-0.0134	0.1108	9.04e-01	++	4.44e-02	16875	-0.007	0.0497	8.88e-01	---	9.49e-01
rs2937797	5	6236277		t	c	18770	0.0512	0.0488	2.94e-01	+++++	6.26e-03	10328	0.0636	0.0639	3.20e-01	++	6.29e-02	29088	0.0589	0.0388	1.51e-01	---	1.51e-01
rs2937789	5	6241709		t	c	18662	0.0383	0.0499	4.43e-01	+++++	8.83e-03	10325	0.0722	0.0643	2.62e-01	++	4.71e-02	28897	0.051	0.0394	1.95e-01	---	6.77e-01
rs2937785	5	6250044		t	c	18816	-0.152	0.0509	7.65e-01	-----	1.03e-01	10362	-0.0754	0.0648	2.45e-01	++	3.91e-02	29178	-0.0382	0.04	3.40e-01	---	4.65e-01
rs16884022	5	10094023		a	c	10618	-0.0716	0.0298	1.64e-02	-----	7.45e-01	10338	0.0043	0.0301	8.86e-01	---	2.88e-01	29556	-0.034	0.0212	1.08e-01	---	7.31e-02
rs11740157	5	10094128		a	g	14778	0.0449	0.0263	8.82e-02	+++++	4.47e-01	10363	0.0018	0.0301	9.52e-01	++	2.45e-01	20141	0.0262	0.0198	1.85e-01	---	2.81e-01
rs11738850	5	10095771		t	c	16777	0.0694	0.0243	4.32e-03	+++++	2.48e-01	10347	-0.0144	0.0304	6.35e-01	---	2.76e-01	27124	0.0367	0.019	5.30e-02	---	3.31e-02
rs7729241	5	10097149		a	g	18803	-0.0894	0.0229	9.08e-05	-----	1.19e-01	10354	-0.0209	0.0292	4.76e-01	++	1.57e-01	29157	-0.0633	0.018	4.42e-04	---	6.19e-02
rs16884037	5	10097397		a	g	16433	0.0755	0.0244	1.97e-03	+++++	4.45e-01	10354	0.0102	0.0305	7.39e-01	++	2.75e-01	26787	0.05	0.0191	8.66e-03	---	9.46e-02
rs2786930	5	27938515		t	c	14633	0.0225	0.0379	5.53e-01	+++++	6.47e-02	9972	-0.0109	0.0364	7.65e-01	7+	3.49e-01	24605	0.0051	0.0263	8.41e-01	++	5.25e-01
rs956306	5	27940408		t	c	14636	-0.0261	0.038	4.92e-01	+++++	6.47e-02	9972	-0.0102	0.0366	7.80e-01	7+	3.79e-01	24611	-0.0073	0.0264	7.83e-01	++	4.91e-01
rs1469439	5	27940555		g	c	12522	-0.0061	0.0414	8.49e-01	+++++	6.47e-02	9971	-0.0109	0.0365	7.71e-01	7+	3.34e-01	24541	-0.0087	0.0275	5.12e-01	---	9.33e-01
rs1711654	5	27944483		g	c	14679	-0.0292	0.0377	4.40e-01	+++++	5.07e-02	9971	0.0266	0.0374	4.77e-01	++	4.57e-01	24650	-0.0011	0.0266	9.68e-01	---	2.93e-01

rs10040195	5	113174830	t	c	18541	0.0992	0.0235	2.45E-05	+++++	7.27E-02	10334	0.0232	0.0289	4.21E-01	++	4.66E-01	28875	0.0689	0.0182	1.56E-04	++	4.13E-02
rs2128317	5	113175231	t	c	18622	0.0945	0.0232	4.58E-05	+++++	1.70E-02	10333	0.0242	0.0286	3.98E-01	++	4.26E-01	28955	0.0668	0.018	2.19E-04	++	5.63E-02
rs694762	5	113176166	t	c	18790	-0.0919	0.0231	7.00E-05	-----	9.60E-03	10329	-0.0208	0.0286	4.69E-01	++	3.41E-01	29119	-0.0638	0.018	3.83E-04	--	5.31E-02
rs1384282	5	113177023	t	c	18813	0.0865	0.023	1.74E-04	+++++	1.88E-02	10360	0.019	0.0286	5.05E-01	++	3.56E-01	29173	0.06	0.0179	8.17E-04	++	6.59E-02
rs1213377	5	113177377	t	c	18659	0.0478	0.0237	4.32E-02	+++++	3.16E-04	10353	0	0.0294	1.90E-00	++	5.27E-01	29012	0.029	0.0185	1.16E-01	0	2.06E-01
rs10055433	5	113177785	a	g	18696	-0.045	0.0232	3.8E-01	-----	8.69E-05	10353	-0.0114	0.0287	6.90E-01	++	3.39E-01	29029	-0.0254	0.011	1.67E-01	++	5.31E-01
rs17136028	5	113178099	a	g	10747	-0.1105	0.0327	7.20E-04	??-?-?	4.75E-01	10265	0.0035	0.0292	9.06E-01	++	5.01E-01	21012	-0.0471	0.0218	3.07E-02	++	9.31E-03
rs13355972	5	113178595	a	g	10754	0.1024	0.0327	1.75E-03	??+?+?	6.01E-01	10223	-0.0005	0.0291	9.87E-01	0	4.00E-01	20977	0.045	0.0217	3.86E-02	++	1.87E-02
98																						
rs12518149	5	11320726	a	t	16726	0.0944	0.0265	3.58E-04	+++++?+	7.75E-01	10234	0.0266	0.0315	3.99E-01	++	2.57E-01	26960	0.0663	0.0203	1.08E-03	++	9.96E-02
rs17424081	5	113238402	c	g	16766	0.0415	0.0263	1.14E-01	??+?+?	3.79E-02	10354	0.0278	0.0313	3.75E-01	++	3.43E-01	27120	0.0358	0.0201	7.52E-02	++	7.38E-01
rs975704	5	113238768	c	g	18786	0.0901	0.0246	2.44E-04	+++++	8.08E-01	10357	0.0273	0.0313	3.82E-01	++	3.35E-01	29143	0.0661	0.0193	6.29E-04	++	1.15E-01
rs1482483	5	113239111	a	g	18811	-0.042	0.0245	8.69E-02	-----	4.49E-02	10357	-0.026	0.0313	4.06E-01	++	3.76E-01	29168	-0.0359	0.0193	6.62E-04	--	6.87E-01
rs1743869	5	113240869	a	g	14346	-0.1380	0.0287	1.54E-04	??-?-?	6.26E-01	10353	-0.0302	0.0313	3.34E-01	---	2.70E-01	26499	-0.0727	0.0212	5.85E-04	---	6.52E-02
rs1522099	5	113246774	c	g	18657	-0.0955	0.0248	1.20E-04	-----	7.51E-01	10318	-0.0308	0.0313	3.25E-01	---	2.28E-01	28975	-0.0705	0.0194	2.84E-04	---	1.05E-02
rs1046353	5	113291236	c	g	14597	-0.1033	0.0284	2.77E-04	??-?-?	8.09E-01	10306	-0.0343	0.0291	6.45E-01	---	2.12E-01	24903	-0.0627	0.0209	2.76E-03	---	3.43E-02
99																						
rs1207762	5	12146462			18674	-0.0866	0.0247	4.65E-04	-----	1.00E-01	10243	-0.0741	0.0307	1.56E-02	---	3.68E-02	29017	-0.0817	0.0192	2.19E-05	---	7.51E-01
rs13161508	5	12158355	a	c	18730	0.1487	0.0415	3.40E-04	+++++	6.96E-01	10359	0.0351	0.0698	6.15E-01	++	8.89E-01	29089	0.119	0.0357	8.47E-04	++	1.62E-01
rs1421877	5	121590134	t	c	18671	-0.1488	0.0417	3.56E-04	-----	9.99E-01	10312	-0.0474	0.0713	5.06E-01	++	9.50E-01	28983	-0.123	0.036	6.36E-04	++	2.20E-01
rs1421878	5	121597424	a	g	18655	-0.1481	0.0418	4.02E-04	-----	9.99E-01	10309	-0.0476	0.0713	5.05E-01	++	9.51E-01	28964	-0.1224	0.0361	6.88E-04	++	2.24E-01
rs1421879	5	121597568	c	g	18660	-0.1485	0.0418	3.83E-04	-----	9.99E-01	10309	-0.0515	0.0713	4.70E-01	++	9.72E-01	28969	-0.1237	0.0361	6.03E-04	++	2.41E-01
rs1363163	5	121600149	t	c	18647	0.1494	0.0418	3.45E-04	+++++	9.99E-01	10304	0.0459	0.0713	5.20E-01	++	9.35E-01	28951	0.1229	0.0361	6.52E-04	++	2.11E-01
rs6881951	5	121620497	a	g	18628	-0.1376	0.0399	5.68E-04	-----	8.62E-01	9960	-0.0215	0.0637	7.36E-01	?	9.57E-01	28588	-0.1049	0.0338	1.92E-03	?	1.22E-01
rs20470110	5	121620597	a	g	18727	-0.1576	0.0394	6.23E-05	-----	9.32E-01	9967	-0.033	0.0639	6.06E-01	?	8.83E-01	28694	-0.1233	0.0335	2.37E-04	?	9.70E-02
rs11241634	5	121622952	a	g	18705	0.1374	0.0402	6.23E-04	+++++	9.26E-01	9975	0.0325	0.0641	6.12E-01	?	8.63E-01	28680	0.1078	0.0341	1.55E-03	?	1.66E-01
rs10052044	5	121622991	c	g	16748	-0.1121	0.0449	6.53E-01	-----	9.27E-01	9975	-0.0324	0.0644	6.15E-01	?	8.63E-01	28723	-0.0928	0.0368	1.18E-02	?	2.53E-01
rs2050948	5	121623317	c	g	18782	0.1415	0.0398	3.78E-04	+++++	9.98E-01	9975	0.0324	0.0644	6.15E-01	?	8.63E-01	28757	0.1133	0.0339	1.03E-03	?	1.50E-01
rs2139363	5	121624095	a	g	18774	0.1456	0.0397	2.43E-04	+++++	9.98E-01	9975	0.0393	0.0644	5.42E-01	?	7.99E-01	28749	0.1163	0.0338	5.77E-04	?	1.60E-01
rs9327244	5	121625290	c	g	16443	0.1521	0.0415	2.44E-04	+++++?+	9.38E-01	9975	0.0324	0.0644	6.15E-01	?	8.63E-01	28618	0.117	0.0349	7.99E-04	?	1.18E-01
rs10058354	5	121626139	c	g	16993	-0.1411	0.0405	4.96E-04	-----?	8.70E-01	9975	-0.0324	0.0644	6.15E-01	?	8.63E-01	28668	-0.1103	0.0343	1.30E-03	?	1.53E-01
rs731312	5	121628286	c	g	18669	0.1478	0.0408	2.87E-04	+++++	9.98E-01	9974	0.0398	0.0646	5.38E-01	?	7.32E-01	28643	0.117	0.0345	6.94E-04	?	1.58E-01
rs11241636	5	121629357	c	g	14764	0.1526	0.0443	5.71E-04	??+?+?	7.61E-01	9956	0.0505	0.065	4.37E-01	?	8.96E-01	24720	0.202	0.0366	1.02E-03	?	1.94E-01
rs13328228	5	121641599	a	g	16466	-0.1436	0.0385	1.94E-04	-----?	2.69E-01	9962	0.0323	0.0566	5.69E-01	?	3.30E-01	26428	-0.088	0.0318	5.73E-03	?	1.02E-02
rs11241637	5	121642763	a	t	18769	-0.1312	0.0374	4.45E-04	-----	1.29E-01	9956	0.0312	0.0569	5.84E-01	?	4.24E-01	28725	-0.0822	0.0313	8.53E-03	?	1.71E-02
rs11241638	5	121643310	t	c	16421	0.1419	0.0389	2.66E-04	+++++?+	2.07E-01	9899	-0.0261	0.0581	6.53E-01	?	5.22E-01	26320	-0.0899	0.0323	5.42E-03	?	1.63E-02
rs11733978	5	121643977	a	g	18807	0.1051	0.0304	5.35E-01	-----	4.95E-01	10355	0.0452	0.0557	2.05E-01	++	3.44E-01	29169	0.0799	0.0231	5.54E-04	++	2.01E-01
rs1716003	5	125439556	a	g	18814	0.1045	0.0303	5.68E-04	+++++	8.28E-01	10355	0.0452	0.0557	2.05E-01	++	3.44E-01	29169	0.0797	0.0231	5.63E-04	++	2.05E-01
rs10070545	5	125469352	t	c	18716	0.1109	0.0304	6.1E-04	+++++	8.51E-01	10337	0.0452	0.0557	2.05E-01	++	3.51E-01	29163	0.0833	0.0231	3.20E-04	++	1.61E-01
rs6956669	5	125478687	c	g	18699	0.1116	0.0303	2.33E-04	+++++	8.26E-01	10352	0.0438	0.0556	2.18E-01	++	3.40E-01	29051	0.0831	0.0231	3.16E-04	++	1.47E-01
101																						
rs880770	5	149135028	a	g	18816	0.1231	0.0274	3.71E-06	+++++?+	8.91E-01	10362	0.0142	0.0333	6.70E-01	++	9.78E-01	29178	0.0677	0.0212	1.38E-03	++	1.45E-03
rs1469042	5	149137507	a	g	18613	-0.0775	0.0276	4.98E-03	-----	8.46E-03	10298	0.0152	0.0334	6.49E-01	++	9.82E-01	28911	-0.0399	0.0213	6.08E-02	++	3.24E-02
rs1713988	5	149137819	a	t	14173	0.0581	0.0325	7.36E-02	??+?+?	2.93E-02	10235	-0.0107	0.0336	7.51E-01	++	9.64E-01	24408	-0.0248	0.0234	2.88E-04	++	1.41E-01
rs6829420	5	149138019	a	g	18562	-0.0786	0.0276	4.38E-03	-----	1.01E-02	10296	0.0162	0.0334	6.28E-01	++	9.81E-01	28858	-0.0401	0.0213	5.93E-02	++	2.87E-02
rs405377	5	149143607	a	g	18815	-0.0776	0.0272	4.37E-03	-----	1.22E-02	10344	0.0164	0.0332	6.21E-01	++	9.46E-01	29159	-0.0398	0.021	5.83E-02	++	2.85E-02
rs405378	5	149144856	a	g	18725	-0.0808	0.0274	3.14E-03	-----	1.31E-02	10357	-0.0215	0.0333	5.29E-01	++	9.24E-01	29055	-0.0395	0.0212	6.19E-02	++	1.77E-02
102																						
rs131680	5	159207919	a	g	18647	0.038	0.0254	4.85E-01	??+?+?	2.70E-01	10287	0.0142	0.0346	6.31E-01	?	1.00E-01	14334					

rs9295474	6	20760696	CDKAL1	c	g	18735	-0.1735	0.0231	5.27E-14	-----	1.92E-01	10357	-0.1402	0.0285	8.59E-07	---	8.69E-01	29092	-0.1603	0.0179	4.17E-19	--	3.64E-01
rs9295475	6	20760744	CDKAL1	a	g	18787	-0.1696	0.0231	2.32E-13	-----	4.98E-01	10356	-0.1516	0.0286	1.15E-07	---	9.36E-01	29143	-0.1625	0.018	1.54E-19	--	6.24E-01
rs2328545	6	20761529	CDKAL1	c	g	18776	0.1712	0.0232	1.48E-13	+++++	4.85E-01	10357	0.1525	0.0286	9.61E-08	+++	9.45E-01	29133	0.1638	0.018	9.90E-20	--	6.12E-01
rs9358355	6	20762876	CDKAL1	t	c	18763	-0.1713	0.0232	1.40E-13	-----	4.91E-01	10356	-0.1525	0.0286	9.67E-08	---	9.44E-01	29119	-0.1638	0.018	9.60E-20	--	6.10E-01
rs9362166	6	20763160	CDKAL1	a	g	18760	-0.1706	0.0232	1.85E-13	-----	4.52E-01	10356	-0.1525	0.0286	9.67E-08	---	9.44E-01	29116	-0.1634	0.018	1.19E-19	--	6.23E-01
rs4712522	6	20764779	CDKAL1	a	g	18813	-0.1761	0.023	2.14E-14	-----	1.42E-01	10355	-0.1413	0.0285	6.88E-07	---	8.80E-01	29171	-0.1624	0.0179	1.17E-19	--	8.42E-01
rs4712523	6	20765543	CDKAL1	a	g	18787	-0.1743	0.023	3.47E-14	-----	1.31E-01	10356	-0.1424	0.0285	5.79E-07	---	8.41E-01	29143	-0.1617	0.0179	1.64E-19	--	8.34E-01
rs4710940	6	20765991	CDKAL1	a	g	18813	-0.1771	0.023	1.44E-14	-----	1.57E-01	10358	-0.1413	0.0285	6.88E-07	---	8.80E-01	29171	-0.1617	0.0179	8.57E-20	--	8.28E-01
rs9606327	6	20767438	CDKAL1	a	g	18814	0.1761	0.023	2.04E-14	+++++	1.54E-01	10353	0.1415	0.0284	6.53E-07	+++	8.71E-01	29167	0.1624	0.0179	1.03E-19	++	3.44E-01
rs9456367	6	20767566	CDKAL1	a	t	18813	0.1761	0.023	2.05E-14	+++++	1.49E-01	10363	0.1423	0.0284	5.99E-07	+++	8.54E-01	29176	0.1627	0.0179	8.75E-20	++	3.55E-01
rs9456368	6	20767785	CDKAL1	t	c	18696	-0.1719	0.0231	1.10E-13	-----	1.50E-01	10363	-0.1423	0.0284	5.99E-07	---	8.54E-01	29059	-0.1601	0.0179	4.08E-19	--	4.19E-01
rs9456369	6	20768344	CDKAL1	t	c	18813	-0.122	0.0231	1.22E-07	-----	6.59E-07	10327	-0.1342	0.0286	2.60E-06	---	9.65E-01	29140	-0.1268	0.018	1.70E-19	--	7.40E-01
rs10946398	6	20769013	CDKAL1	a	c	18810	-0.1757	0.023	2.42E-14	-----	1.54E-01	10343	-0.1429	0.0285	5.33E-07	---	8.39E-01	29153	-0.1628	0.0179	9.58E-20	--	3.71E-01
rs1774594	6	20769122	CDKAL1	a	t	18813	0.1761	0.023	2.08E-14	+++++	1.45E-01	10363	0.1423	0.0284	5.99E-07	+++	8.54E-01	29176	0.1627	0.0179	8.75E-20	++	3.55E-01
rs1794840	6	20769222	CDKAL1	c	g	18732	0.1786	0.0231	9.63E-15	+++++	1.09E-07	10457	0.1455	0.0285	3.29E-07	+++	7.64E-01	29079	0.1655	0.0219	2.94E-20	++	3.97E-01
rs9460544	6	20769508	CDKAL1	t	c	18814	0.176	0.023	2.14E-14	+++++	1.45E-01	10363	0.1423	0.0284	5.99E-07	---	8.54E-01	29177	0.1627	0.0179	9.03E-20	++	3.57E-01
rs9460545	6	20769529	CDKAL1	t	c	18814	-0.176	0.023	2.12E-14	+++++	1.44E-01	10363	-0.1423	0.0284	5.99E-07	---	8.54E-01	29177	-0.1627	0.0179	9.03E-20	++	3.57E-01
rs979614	6	20770102	CDKAL1	a	g	14832	0.131	0.0257	3.38E-07	++2++++	2.34E-01	10361	0.1386	0.0297	3.05E-06	+++	6.16E-01	25193	0.1343	0.0194	4.91E-12	++	8.47E-01
rs4712525	6	20770945	CDKAL1	t	c	18814	0.1741	0.023	4.03E-14	+++++	1.73E-01	10363	0.1423	0.0284	5.99E-07	+++	8.54E-01	29177	0.1615	0.0179	1.63E-19	++	3.84E-01
rs4712526	6	20771014	CDKAL1	a	t	18814	0.1742	0.023	3.94E-14	+++++	1.73E-01	10363	0.1423	0.0284	5.99E-07	+++	8.54E-01	29177	0.1616	0.0179	1.58E-19	++	3.83E-01
rs9460546	6	20771611	CDKAL1	t	g	18787	-0.1767	0.0231	2.10E-14	-----	1.62E-01	10274	-0.1461	0.0287	3.46E-07	---	7.37E-01	29061	-0.1647	0.018	5.65E-20	--	4.06E-01
rs742642	6	20773060	CDKAL1	a	g	18814	0.1708	0.0231	1.56E-13	+++++	3.37E-01	10362	0.1539	0.0286	7.19E-08	+++	9.24E-01	29176	0.1641	0.018	6.65E-20	++	4.64E-01
rs17748382	6	20773528	CDKAL1	a	g	18814	0.1738	0.023	4.60E-14	+++++	1.83E-01	10363	0.1409	0.0284	7.22E-07	+++	8.38E-01	29177	0.1608	0.0179	2.37E-19	++	3.68E-01
rs1772603	6	20773925	CDKAL1	c	g	18780	-0.191	0.0245	6.26E-15	-?-----	4.20E-01	10363	-0.1416	0.0284	6.95E-07	---	8.46E-01	27143	-0.1659	0.0186	5.20E-20	--	1.88E-01
rs9358357	6	20774003	CDKAL1	a	g	18814	0.1754	0.023	2.69E-14	+++++	2.17E-01	10338	0.1416	0.0284	6.35E-07	+++	8.46E-01	29007	0.1467	0.0178	1.60E-16	++	6.97E-01
rs1772906	6	20774034	CDKAL1	t	g	16479	0.1617	0.0248	3.76E-11	+++++2+	8.61E-01	10363	0.1416	0.0284	6.35E-07	+++	8.46E-01	26842	0.1517	0.0267	1.60E-16	++	5.94E-01
rs9358356	6	20775361	CDKAL1	t	c	18737	-0.1766	0.0232	2.41E-14	-----	2.26E-01	10271	-0.1446	0.0286	4.17E-07	---	7.88E-01	29008	-0.1639	0.018	9.31E-20	--	3.85E-01
rs9350270	6	20775718	CDKAL1	a	g	14831	-0.0996	0.0254	8.99E-05	-?+-----	5.41E-01	10197	-0.131	0.0286	4.55E-06	---	7.52E-01	25028	-0.1134	0.019	2.32E-20	--	4.12E-01
rs9362129	6	20782670	CDKAL1	t	c	14701	0.1599	0.0261	8.44E-10	+++++2+	4.38E-01	10222	0.1774	0.0288	6.82E-10	+++	7.29E-01	24222	0.1678	0.0193	4.10E-18	++	6.53E-01
rs1569699	6	20787289	CDKAL1	t	g	18636	-0.1416	0.0228	5.41E-10	-----	7.73E-02	10360	-0.1496	0.0283	1.23E-07	---	6.36E-01	28996	-0.1447	0.0178	3.56E-16	--	8.26E-01
rs1756992	6	20787688	CDKAL1	a	g	18649	-0.1116	0.0228	1.01E-06	-----	2.68E-04	10340	-0.1545	0.0284	5.42E-08	---	4.72E-01	28989	-0.1284	0.0178	5.10E-13	--	2.39E-01
rs9350271	6	20791143	CDKAL1	a	g	12447	0.1565	0.0286	4.32E-08	+++++2+	6.51E-01	10349	0.177	0.0286	8.58E-10	+++	7.53E-01	22796	0.1667	0.0202	1.65E-16	++	6.12E-01
rs9356744	6	20793465	CDKAL1	t	c	14870	-0.17	0.0259	5.56E-11	-?-----	4.07E-01	10350	-0.1765	0.0286	6.42E-10	---	7.33E-01	25130	-0.1729	0.0192	2.10E-19	--	8.66E-01
rs1776670	6	20794552	CDKAL1	a	c	14807	0.167	0.0259	1.20E-10	+++++2+	6.72E-01	10350	0.1765	0.0286	6.42E-10	+++	7.33E-01	25157	0.1733	0.0192	4.59E-19	++	8.06E-01
rs9358322	6	20794975	CDKAL1	a	g	14806	0.0892	0.0259	5.78E-04	+++++	2.54E-06	10346	0.1276	0.0286	1.56E-09	---	8.45E-01	29007	0.1362	0.0192	4.01E-11	++	8.51E-01
rs10440833	6	20796100	CDKAL1	t	c	14713	0.1702	0.0263	9.35E-11	+++++2+	4.49E-01	10216	0.1775	0.0288	7.16E-10	+++	7.91E-01	24929	0.1735	0.0194	4.08E-19	++	8.52E-01
rs2206734	6	20802863	CDKAL1	a	t	14832	0.1666	0.026	1.52E-10	+++++2+	4.77E-01	10336	0.1802	0.0286	3.16E-10	+++	6.98E-01	25168	0.1728	0.0192	2.72E-19	++	7.25E-01
rs11753081	6	20813569	CDKAL1	t	g	18655	-0.1367	0.0228	1.93E-09	-----	1.17E-01	10352	-0.1565	0.0284	3.38E-08	---	6.71E-01	29007	-0.1445	0.0178	4.47E-16	--	5.87E-01
rs1040558	6	20821685	CDKAL1	a	g	18718	-0.107	0.0227	2.47E-06	-----	5.16E-04	10359	-0.1564	0.0283	3.44E-08	---	6.74E-01	29077	-0.1263	0.0177	9.69E-13	--	1.73E-01
rs9295478	6	20824232	CDKAL1	a	g	18753	0.1407	0.0227	5.69E-10	+++++	1.49E-01	10269	0.1639	0.0284	8.09E-09	+++	4.26E-01	29022	0.1497	0.0177	3.04E-17	++	5.23E-01
rs2328548	6	20824937	CDKAL1	a	g	18772	0.1394	0.0227	8.35E-10	+++++	1.24E-01	10360	0.1561	0.0283	3.66E-08	+++	6.80E-01	29132	0.1459	0.0177	1.70E-16	++	6.45E-01
rs9359959	6	20825074	CDKAL1	a	g	18777	-0.1398	0.0227	7.36E-10	-----	1.30E-01	10360	-0.1561	0.0283	3.66E-08	---	6.80E-01	29137	-0.1462	0.0177	1.51E-16	--	6.53E-01
rs9456781	6	20825234	CDKAL1	t	c	18578	-0.1138	0.0228	6.02E-07	-----	1.71E-04	10347	-0.1523	0.0283	7.89E-08	---	7.68E-01	28925	-0.1289	0.0178	4.19E-13	--	2.91E-01
rs10946003	6	20825383	CDKAL1	a	g	18765	-0.1379	0.0228	1.37E-09	-----	6.72E-02	10356	-0.1571	0.0283	2.99E-08	---	7.25E-01	29121	-0.1455	0.0178	2.56E-16	--	5.97E-01
rs9358357	6	20827003	CDKAL1	a	g	18662	-0.1411	0.0228	5.59E-10	-----	1.21E-01	10338	-0.1553	0.0284	4.33E-08	---	7.74E-01	29000	-0.1467	0.0178	1.60E-16	--	6.97E-01
rs9362224	6	20827211	CDKAL1	a	t	18661	0.1412	0.0228	5.51E-10	+++++	9.22E-01	10338	0.1553	0.0284	4.33E-08	+++	7.74E-01	28999	0.1467	0.0178	1.55E-16	++	6.99E-01
rs9358358	6	20827372	CDKAL1	a	g	18615	-0.1398	0.0228	8.37E-10	-----	9.90E-02	10338	-0.1553	0.0284	4.33E-08	---	7.74E-01	28953	-0.1459	0.0178	2.31E-16	--	6.70E-01
rs9460550	6	20827540	CDKAL1	a	g	18662	0.1404	0.0228	6.97E-10	+++++	1.10E-01	10338	0.1553	0.0284	4.33E-08	+++	7.74E-01	29000	0.1462	0.0178	1.95E-16	++	6.83E-01
rs9356746	6	20828258	CDKAL1	t	c	18626	-0.14	0.0228	7.70E-10	-----	9.59E-02	10338	-0.1553	0.0284	4.33E-08	---	7.74E-01	28964	-0.1446	0.0178	2.18E-16	--	6.74E-01
rs9368226	6	20831036	CDKAL1	t	c	18570	-0.1468	0.0229	1.34E-10	-----	2.60E-01	9897	-0.1612	0.029	2.77E-08	---	4.06E-01	28467	-0.1523	0.018	2.33E-17	--	6.97

	rs7488392	6	38226733	ZFAND3	a	g	14803	-0.106	0.0275	1.17E-04	---?+..	4.04E-01	13002	-0.0822	0.0351	1.92E-02	---	5.83E-01	25105	-0.0969	0.0216	7.52E-06	---	5.94E-01
	rs786188	6	38228638	ZFAND3	t	c	14811	0.1043	0.0275	1.51E-04	+++?+..	3.02E-01	13002	0.0822	0.0351	1.92E-02	+++	5.83E-01	25113	0.0959	0.0216	9.43E-06	+++	6.20E-01
	rs3800364	6	38229189	ZFAND3	a	g	14811	-0.1036	0.0275	1.68E-04	---?+..	3.02E-01	13003	-0.0804	0.0351	2.18E-02	---	5.72E-01	25114	-0.0948	0.0216	1.20E-05	---	6.03E-01
	rs2395692	6	38232057		a	g	14806	-0.1027	0.0275	1.91E-04	---?+..	3.05E-01	13006	-0.0808	0.0351	2.13E-02	---	5.78E-01	25112	-0.0944	0.0216	1.30E-05	---	6.23E-01
	rs2395693	6	38232058		t	c	14798	0.1028	0.0275	1.89E-04	+++?+..	3.02E-01	13007	0.0785	0.0351	2.51E-02	+++	5.68E-01	25105	0.0931	0.0216	1.53E-05	+++	5.85E-01
118	rs11756091	6	39390784	KCNK16	t	g	13045	0.1171	0.0276	5.54E-06	+++?+..	2.03E-01	7874	0.0695	0.0338	3.99E-02	+++	8.57E-01	20919	0.0991	0.0208	1.78E-05	+++	2.66E-01
	rs1246262	6	39391771	KCNK16	a	g	18764	0.0547	0.0229	1.70E-02	+++++..	2.41E-03	7869	0.0688	0.0339	4.22E-02	+++	7.84E-01	26633	0.0591	0.019	1.84E-03	+++	7.30E-01
	rs1535500	6	39392028	KCNK16	t	g	17010	0.1063	0.0234	3.54E-06	+++++..	3.54E-03	7875	0.072	0.0338	3.33E-02	+++	9.07E-01	24885	0.0952	0.0192	7.52E-07	+++	4.04E-01
	rs3734618	6	39392162	KCNK16	a	g	14829	-0.1165	0.0255	5.08E-06	---?+..	4.30E-01	7873	-0.0717	0.0338	3.41E-02	---	9.72E-01	22702	-0.1003	0.0204	8.45E-07	---	2.90E-01
119	rs782817	6	39409896		t	c	18782	-0.0767	0.0238	1.27E-03	-----	7.35E-01	10356	-0.0119	0.0297	6.89E-01	---	6.78E-01	29138	-0.0514	0.0186	9.56E-07	---	8.86E-02
	rs7740446	6	39410410	KIF6	t	c	18674	-0.0832	0.0238	4.85E-04	-----	7.51E-01	10337	-0.0077	0.0296	7.94E-01	---	3.45E-01	29011	-0.0536	0.0185	3.89E-03	---	4.68E-02
	rs774204	6	39410479	KIF6	t	c	18766	-0.0786	0.0238	9.74E-04	-----	7.59E-01	10355	-0.0168	0.0295	5.68E-01	---	6.18E-01	29121	-0.0542	0.0185	3.41E-03	---	1.03E-01
	rs7754225	6	39410701	KIF6	t	c	18663	0.0806	0.0239	7.34E-04	+++++..	7.52E-01	10353	0.0081	0.0297	7.85E-01	+++	8.19E-01	29016	0.0521	0.0186	5.14E-03	+++	5.72E-02
	rs1887716	6	39410810	KIF6	t	c	18638	0.0621	0.0238	9.12E-04	+++++..	3.02E-01	10352	0.0184	0.0295	5.31E-01	+++	7.23E-01	28990	0.0449	0.0185	1.54E-03	+++	2.49E-01
	rs9462321	6	39421574	KIF6	a	g	18652	0.0769	0.0237	1.20E-03	+++++..	6.90E-01	10360	-0.0088	0.0295	7.66E-01	---	2.83E-01	28955	0.0433	0.0185	1.92E-02	---	2.35E-02
	rs694582	6	39411733	KIF6	a	g	18646	0.0799	0.0239	8.13E-04	+++++..	7.74E-01	10201	-0.0095	0.0299	7.51E-01	---	1.76E-01	28947	0.045	0.0187	1.58E-02	---	1.95E-02
120	rs10456556	6	46052414	C11C5	a	g	18712	0.0771	0.0256	2.56E-03	+++++..	4.20E-02	10347	0.0425	0.0324	1.90E-01	+++	3.60E-01	29059	0.0638	0.0201	1.49E-03	+++	4.02E-01
	rs2039716	6	46053234	C11C5	t	g	16774	0.0513	0.0273	6.06E-02	+?+..	1.93E-02	10351	0.0434	0.0324	1.80E-01	+++	3.11E-01	27125	0.0408	0.0209	2.14E-02	+++	8.52E-01
	rs6930840	6	46059789	C11C5	a	g	18644	0.0683	0.0257	7.98E-03	+?+..	9.91E-03	10258	0.046	0.0326	1.58E-01	+++	1.89E-01	28902	0.0598	0.0202	3.07E-03	+++	5.91E-01
	rs1338467	6	46061231	C11C5	t	g	16486	-0.0435	0.0275	1.15E-01	-?+..	4.10E-03	10240	-0.0405	0.0323	2.10E-01	---	1.75E-01	26726	-0.0422	0.0209	4.37E-02	---	9.44E-01
	rs3776005	6	46065579	C11C5	a	g	18812	0.0666	0.0256	9.44E-03	+++++..	5.92E-03	10363	0.0241	0.0321	4.53E-01	---	1.70E-01	29175	0.0501	0.02	1.24E-02	---	3.01E-01
	rs3776006	6	46065863	C11C5	a	g	18805	0.0678	0.0256	8.13E-03	+++++..	5.09E-03	10363	0.0261	0.032	4.14E-01	---	1.79E-01	29168	0.0514	0.02	9.95E-03	---	3.09E-01
	rs1338469	6	46066913	C11C5	a	g	18749	0.069	0.0256	7.13E-03	+++++..	3.94E-03	10361	0.0265	0.032	4.08E-01	+++	1.87E-01	29110	0.0524	0.02	8.74E-03	+++	3.00E-01
	rs1535109	6	46068107	C11C5	a	g	16220	-0.0574	0.0277	3.85E-02	---?+..	6.35E-03	10337	-0.0273	0.032	3.94E-01	---	2.03E-01	26557	-0.0495	0.0209	3.36E-02	---	4.77E-01
	rs10480486	6	46070430	C11C5	a	g	14766	0.1019	0.0285	2.93E-04	+++?+..	3.70E-01	10311	0.0271	0.0495	3.97E-01	+++	1.90E-01	25143	0.0709	0.0204	1.29E-02	+++	6.55E-02
	rs4077761	6	46073476	C11C5	t	g	18748	0.0711	0.0256	5.46E-03	+++++..	3.10E-03	10361	0.0265	0.0319	4.07E-01	+++	2.01E-01	29109	0.0536	0.02	7.23E-03	+++	2.76E-01
	rs1241463	6	46074782	C11C5	t	c	18760	-0.07	0.0256	6.27E-03	---?+..	2.81E-03	10361	-0.0264	0.0319	4.08E-01	---	1.86E-01	29121	-0.0529	0.02	8.04E-03	---	2.86E-01
	rs1338472	6	46076739	C11C5	a	g	18816	0.0696	0.0256	6.53E-03	+++++..	2.98E-03	10363	0.0261	0.032	4.15E-01	+++	1.77E-01	29179	0.0526	0.02	8.48E-03	+++	2.89E-01
	rs1338474	6	46079747	C11C5	a	g	18702	0.0708	0.0257	5.84E-03	+++++..	3.31E-03	10352	0.0303	0.032	3.45E-01	+++	1.60E-01	29054	0.0549	0.02	6.13E-03	+++	3.24E-01
	rs3776223	6	46084355	C11C5	a	g	18797	0.066	0.0255	9.80E-03	+++++..	3.36E-03	10347	0.0331	0.0321	3.03E-01	+++	1.40E-01	29144	0.0533	0.02	7.63E-03	+++	4.22E-01
	rs10948271	6	46086102	C11C5	t	c	10885	0.0939	0.0234	3.80E-03	+++?+..	2.09E-01	10339	0.0445	0.0323	1.68E-01	+++	3.93E-02	21224	0.0691	0.0229	2.51E-03	+++	2.80E-01
	rs1555216	6	46086137	C11C5	a	g	18662	0.0653	0.0256	1.06E-02	+++++..	5.37E-03	10346	0.0338	0.032	2.92E-01	+++	1.31E-01	29008	0.053	0.02	8.01E-03	+++	4.42E-01
	rs3776225	6	46086193	C11C5	a	g	14822	0.0988	0.0284	5.05E-04	+++?+..	6.12E-02	10348	0.0336	0.032	2.94E-01	+++	1.36E-01	25170	0.0701	0.0212	9.71E-04	+++	1.28E-01
	rs10400472	6	46089124	C11C5	c	g	8843	-0.2281	0.029	4.70E-01	????	4.85E-01	9962	-0.418	0.025	2.88E-01	?	6.89E-02	18805	-0.622	0.025	1.47E-01	?	7.87E-01
121	rs6923157	6	53694474		a	g	12495	0.1271	0.032	3.78E-04	+++++..	3.40E-01	10246	0.0688	0.0407	9.07E-02	+++	1.16E-01	26541	0.1017	0.0268	1.50E-04	+++	3.00E-01
	rs6913301	6	53694508		t	c	12459	-0.1366	0.041	8.57E-04	---?+..	9.00E-01	10235	-0.064	0.0475	1.78E-01	---	1.23E-01	26294	-0.1056	0.031	6.68E-04	---	2.47E-01
122	rs17830060	6	55355234	GRAL1	t	c	8843	-0.1198	0.0358	2.60E-02	????	2.51E-01	7848	-0.0294	0.0604	6.26E-01	?	4.76E-01	16691	-0.0798	0.0402	4.70E-02	?	2.64E-01
	rs783867	6	55344896	GRAL1	t	c	14832	0.1297	0.037	4.64E-04	+++?+..	2.51E-01	7875	0.0265	0.0609	6.63E-01	+++	3.70E-01	22707	0.1019	0.0316	1.27E-03	+++	1.48E-01
	rs16886642	6	55349782	GRAL1	c	g	14719	-0.1198	0.037	1.20E-03	---?+..	2.40E-01	10316	-0.0257	0.0507	6.12E-01	---	5.04E-01	25035	-0.0879	0.0299	3.57E-03	---	1.34E-01
	rs12663965	6	55363750	GRAL1	t	c	14827	-0.1044	0.0368	4.50E-03	---?+..	2.61E-01	10350	-0.0257	0.0503	6.10E-01	---	4.13E-01	25177	-0.0777	0.0297	9.56E-03	---	2.07E-01
	rs12662782	6	55363886	GRAL1	a	g	14830	0.051	0.0368	1.66E-01	+++?+..	5.06E-02	10356	0.0107	0.0513	8.34E-01	---	3.75E-01	25166	0.0373	0.0299	2.12E-01	---	5.23E-01
	rs3846915	6	55365381	GRAL1	a	g	10973	-0.1289	0.0421	2.22E-03	---?+..	4.66E-01	10302	-0.0362	0.0339	2.86E-01	---	4.51E-01	25175	-0.0727	0.0264	5.92E-03	---	8.63E-02
	rs3846916	6	55365741	GRAL1	a	g	14832	-0.0995	0.0365	6.43E-03	---?+..	3.61E-01	10311	0.0097	0.0429	8.20E-01	---	5.48E-01	25143	-0.0536	0.0278	3.56E-03	---	5.25E-02
	rs3846917	6	55365757	GRAL1	t	c	14832	0.0986	0.0365	6.90E-03	+++?+..	3.76E-01	10311	0.027	0.0495	8.85E-01	+++	7.34E-01	25143	0.0734	0.0294	1.25E-02	+++	2.44E-01
	rs1032772	6	55366145	GRAL1	a	g																		

	rs802756	6	128295198	a	c	18772	0.1892	0.0646	3.40E-03	+++++	8.84E-01	10362	-0.1421	0.0549	9.65E-03	+	4.89E-01	29134	-0.0032	0.0418	9.40E-01	+	9.31E-05
	rs802755	6	128296381	a	t	18772	0.1897	0.0646	3.30E-03	+++++	8.83E-01	10362	-0.1421	0.0549	9.65E-03	+	4.89E-01	29134	-0.003	0.0418	9.44E-01	+	9.09E-05
	rs699418	6	128297103	a	t	18772	0.1885	0.0646	3.49E-03	+++++	8.94E-01	10362	-0.1421	0.0549	9.65E-03	+	4.89E-01	29082	-0.0035	0.0418	9.34E-01	+	9.63E-05
	rs699419	6	128298053	c	g	18772	-0.1912	0.0646	3.09E-03	-----	8.79E-01	10362	0.1421	0.0549	9.65E-03	+	4.89E-01	29134	0.0023	0.0418	9.56E-01	+	8.44E-05
	rs1699420	6	145730555	c	g	18755	0.114	0.0276	8.96E-02	-----	4.32E-01	10362	-0.1397	0.0549	1.10E-02	+	4.73E-01	29117	0.0882	0.0425	3.70E-01	+	3.46E-02
129	rs10223879	6	145730555	t	c	18781	0.1132	0.0266	2.11E-05	+++++	4.91E-01	9953	0.0144	0.034	6.71E-01	+	7.73E-01	28734	0.0757	0.021	3.03E-04	+	2.71E-02
	rs6922610	6	145730739	c	g	18784	0.1132	0.0266	2.11E-05	+++++	5.09E-01	9953	0.0148	0.0304	6.27E-01	+	7.76E-01	28737	0.0705	0.02	4.26E-04	+	1.49E-02
	rs976921	6	145733931	a	g	18796	0.118	0.0276	1.97E-05	+++++	5.80E-01	9951	-0.0221	0.0311	4.77E-01	+	7.54E-01	28740	0.0563	0.0206	6.41E-03	+	7.54E-04
	rs976922	6	145734562	c	g	18796	0.1179	0.0276	1.98E-05	+++++	5.80E-01	9957	-0.0048	0.0314	8.79E-01	+	7.35E-01	28753	0.0644	0.0207	1.89E-03	+	3.34E-03
	rs9403694	6	145734571	t	c	18780	-0.118	0.0276	1.96E-05	-----	5.49E-01	9959	0.0004	0.0316	9.91E-01	+	7.65E-01	28719	-0.0668	0.0208	1.32E-03	+	4.77E-03
	rs4896795	6	145735282	c	g	16767	-0.1053	0.0298	4.04E-04	-?-----	5.90E-01	9959	0.0057	0.0316	8.56E-01	+	6.89E-01	26726	-0.0531	0.0217	1.44E-02	+	1.06E-02
	rs9393010	6	145735917	a	g	18799	-0.1173	0.0276	2.18E-05	-----	5.70E-01	9959	0.0128	0.0319	6.88E-01	+	6.08E-01	28758	-0.0166	0.0209	3.16E-03	+	2.04E-03
	rs9479280	6	145736790	c	g	18799	0.1173	0.0276	2.18E-05	+++++	5.70E-01	9959	-0.0023	0.0323	9.44E-01	+	7.28E-01	28758	0.0688	0.021	1.45E-03	+	4.88E-03
	rs6939342	6	145737056	c	g	18799	0.116	0.0276	2.67E-05	+++++	6.20E-01	9959	0.0025	0.0325	9.38E-01	+	7.85E-01	28758	0.0684	0.021	1.14E-03	+	7.77E-03
	rs12661545	6	145738189	a	c	18767	-0.1053	0.0298	4.04E-04	-?-----	5.99E-01	9959	-0.007	0.0331	8.25E-01	+	8.44E-01	26726	-0.0615	0.0221	5.51E-03	+	3.9E-02
	rs9393011	6	145739230	a	g	18799	0.1173	0.0276	2.18E-05	+++++	5.70E-01	9959	-0.0068	0.0345	8.43E-01	+	6.69E-01	28758	0.0689	0.0216	1.40E-03	+	4.97E-03
	rs1970321	6	145740415	a	g	16767	-0.1053	0.0298	4.04E-04	-?-----	5.90E-01	9959	-0.0017	0.0348	9.61E-01	+	7.71E-01	26726	-0.0615	0.0226	6.16E-03	+	2.37E-02
	rs1970320	6	145740538	a	g	14823	0.1388	0.0308	6.74E-06	+++++	7.73E-01	9959	0.0117	0.0359	7.43E-01	+	8.56E-01	24782	0.0849	0.0234	2.81E-04	+	7.21E-03
	rs4370382	6	145741896	t	c	18799	0.1173	0.0276	2.18E-05	+++++	5.70E-01	9959	0.0251	0.037	4.98E-01	+	9.31E-01	28758	0.0843	0.0221	1.38E-04	+	4.58E-02
	rs12181026	6	145742068	a	g	18813	-0.117	0.0276	2.32E-05	-----	5.54E-01	10363	0.0167	0.0364	6.46E-01	+	5.56E-01	29176	-0.0682	0.022	1.93E-03	+	4.32E-03
	rs973455	6	145745918	a	g	16478	0.1245	0.0298	2.98E-05	+++++	4.89E-01	10363	0.0242	0.0343	4.81E-01	+	8.47E-01	26841	0.0814	0.0225	2.99E-04	+	2.73E-02
	rs9403696	6	145746024	c	g	12661	-0.1018	0.0336	2.44E-03	-?-----	5.60E-01	10363	-0.0242	0.0343	4.81E-01	+	8.47E-01	23024	-0.0638	0.024	7.86E-03	+	1.06E-01
	rs9386116	6	145747747	c	g	12644	0.1001	0.0334	2.73E-03	+++++	4.45E-01	10342	0.0297	0.0341	3.84E-01	+	8.38E-01	22986	0.0656	0.0239	5.95E-03	+	1.40E-01
	rs976923	6	145750671	a	g	18816	0.1162	0.0276	2.60E-05	+++++	6.03E-01	10363	0.0242	0.0343	4.81E-01	+	8.47E-01	29179	0.08	0.0215	1.97E-04	+	3.67E-02
	rs12664698	6	145751064	a	g	16471	-0.1146	0.0298	2.98E-05	-?-----	4.75E-01	10362	-0.0249	0.0343	4.57E-01	+	8.49E-01	26833	-0.0817	0.0225	1.49E-04	+	6.32E-01
	rs4896799	6	145754233	c	g	18791	-0.1173	0.0277	2.35E-05	-----	5.61E-01	10366	-0.031	0.0344	3.67E-01	+	7.99E-01	29147	-0.0834	0.0216	1.12E-04	+	5.07E-02
	rs976924	6	145757206	a	c	16476	-0.1256	0.0298	2.55E-05	-----	5.09E-01	10362	-0.0244	0.0343	4.76E-01	+	8.27E-01	26838	-0.0821	0.0225	2.64E-04	+	2.59E-02
	rs2039698	6	145758660	t	c	18804	-0.1183	0.0276	1.89E-05	-----	5.52E-01	10362	-0.0239	0.0343	4.86E-01	+	8.70E-01	29166	-0.0812	0.0215	1.05E-04	+	3.20E-02
	rs1856266	6	145758996	c	g	18802	-0.1196	0.0276	1.52E-05	-----	5.58E-01	10362	-0.0239	0.0343	4.86E-01	+	8.70E-01	29164	-0.082	0.0215	1.37E-04	+	2.97E-02
	rs1856264	6	145759813	c	g	18801	0.1214	0.0277	1.14E-05	+++++	5.42E-01	10362	0.0239	0.0343	4.86E-01	+	8.70E-01	29163	0.0829	0.0216	1.19E-04	+	2.70E-02
	rs976925	6	145763699	a	c	18798	-0.1242	0.0277	7.55E-06	-----	4.87E-01	10361	-0.022	0.0343	5.22E-01	+	8.96E-01	29159	-0.0839	0.0216	9.97E-05	+	2.05E-02
	rs9403698	6	145765403	a	g	18798	-0.1247	0.0277	6.87E-06	-----	4.87E-01	10361	-0.022	0.0343	5.22E-01	+	8.96E-01	29159	-0.0842	0.0216	9.41E-05	+	1.98E-02
	rs9393018	6	145765544	t	c	18798	0.1249	0.0277	6.65E-06	+++++	4.63E-01	10361	0.022	0.0343	5.22E-01	+	8.96E-01	29159	0.0843	0.0216	9.19E-05	+	1.96E-02
	rs4895800	6	145767242	a	g	16485	-0.1334	0.0299	8.27E-06	-?-----	4.11E-01	10361	-0.022	0.0343	5.22E-01	+	8.96E-01	26826	-0.0853	0.0225	1.54E-04	+	1.44E-02
	rs9393045	6	145768187	a	g	16465	0.1437	0.0299	7.86E-06	+++++	4.06E-01	10361	0.022	0.0343	5.22E-01	+	8.96E-01	26826	0.0855	0.0225	1.49E-04	+	1.44E-02
	rs1355145	6	145770275	c	g	16758	0.1182	0.0298	7.38E-05	+++++	4.38E-01	10357	0.02	0.0342	5.60E-01	+	8.86E-01	27115	0.0758	0.0225	7.39E-04	+	3.04E-02
	rs1155417	6	145770647	t	c	18787	-0.1283	0.0277	3.51E-06	-----	4.58E-01	10357	-0.02	0.0342	5.60E-01	+	8.86E-01	29144	-0.0824	0.0215	7.27E-05	+	1.39E-02
	rs976933	6	145780158	a	g	12777	0.1321	0.0339	9.72E-05	+++++	7.83E-01	10337	0.0279	0.0342	4.14E-01	+	9.25E-01	23114	0.0805	0.0241	8.32E-04	+	3.05E-02
	rs4142213	6	145781563	t	c	12783	-0.0499	0.0337	1.39E-01	-?+?+?+?	4.10E-01	10328	-0.0195	0.0343	5.69E-01	+	8.60E-01	23111	-0.035	0.024	1.46E-01	+	5.27E-01
130	rs9383557	6	151504339	t	g	14829	0.1248	0.0296	2.52E-05	+++++	6.22E-02	7457	0.012	0.0374	7.48E-01	??	1.00E+00	22286	0.0814	0.0232	4.56E-04	+	1.80E-02
131	rs2181069	6	155846341	t	c	18771	0.0739	0.032	2.10E-02	+++++	2.18E-02	10341	0.1	0.0372	7.21E-03	+	4.51E-01	29112	0.085	0.0243	4.59E-04	+	5.95E-01
	rs1756131	6	155851382	a	g	18794	0.1044	0.0294	3.78E-04	+++++	3.16E-01	10362	0.0775	0.0345	2.49E-02	+	1.35E-01	29156	0.0931	0.0224	3.19E-05	+	5.53E-01
	rs1156027	6	155852038	a	g	18778	0.0788	0.0312	1.14E-02	+++++	4.12E-02	10323	0.0979	0.036	6.54E-03	+	1.89E-01	29101	0.087	0.0236	2.25E-04	+	6.89E-01
	rs4898956	6	155855586	a	g	18620	0.1145	0.0313	2.59E-04	+++++	4.84E-01	10352	0.0859	0.0357	1.60E-02	+	2.10E-01	28972	0.1021	0.0235	1.45E-05	+	5.47E-01
	rs12665958	6	155861084	a	g	16590	-0.1511	0.0342	7.63E-04	-?-----	2.40E-01	10383	-0.0909	0.0371	1.41E-02	+	2.76E-01	28873	-0.1104	0.0251	3.35E-05	+	6.32E-01
	rs1219778	6	155865681	t	c	14824	0.0744	0.0239	8.25E-02	+++++	2.80E-01	7872	0.0557	0.0399	1.63E-01	-?	2.96E-01	26296	0.0664	0.0258	9.98E-03	+	7.21E-01
132	rs9478656	6	155868148	c	g	8636	-0.1154	0.037	1.83E-03	-?+?+?	4.40E-01	7487	-0.046	0.0342	1.79E-01	??	1.00E+00	16123	-0.078	0.0251	1.90E-03	+	1.68E-01
	rs9397820	6	155879090	t	c	14714	0.0897	0.0266	7.66E-04	+++++	1.20E-01	7842	0.0322	0.0335	3.36E-01	-?	6.48E-02	22556	0.0675	0.0208	1.20E-03	+	1.79E-01
	rs9371892	6	155882462	t	c	14830	-0.0924	0.0265	5.00E-04	-?+?+?	9.26E-02	7842	-0.0291	0.0334	3.83E-01	+	6.26E-02	22672	-0.0679	0.0208	1.06E-03	+	1.38E-01
	rs9478657	6	155900649	a	g	10976	-0.1134	0.0294	1.16E-04	-?+?+?	5.68E-01	7840	-0.024	0.0335	4.73E-01	+	8.31E-02	18816	-0.0745	0.0221	7.47E-04	+	4.49E-02
133	rs6976086	7	15015136	t	g	16955	0.0798	0.0265	2.60E-03	+++++	5.67E-01	10347	0.1148	0.0315	2.75E-04	+	8.04E-01	27302	0.0943	0.0203	3.31E-06	+	3.95E-01

	rs10257587	7	68177744	AUTS2	t	c	18816	-0.1411	0.0415	6.69E-04	-----	4.74E-01	2875	-0.0255	0.1031	8.05E-01	--?	8.78E-01	21691	-0.125	0.0385	1.17E-03	--	2.98E-01
	rs4305786	6	68780076	AUTS2	a	g	18812	0.1443	0.0415	5.03E-04	+++++	4.91E-01	2875	0.0255	0.1031	8.05E-01	++?	8.78E-01	21687	0.1277	0.0385	9.07E-04	++	2.85E-01
	rs17140817	6	68799931	AUTS2	t	c	18816	-0.0802	0.0414	5.27E-02	-----	4.99E-02	2865	-0.0224	0.1046	8.31E-01	--?	8.68E-01	21681	-0.0774	0.0385	6.01E-02	--	6.07E-01
	rs19375481	6	68800200	AUTS2	a	t	18816	-0.1428	0.0415	5.77E-04	-----	4.96E-01	2875	-0.0255	0.1031	8.05E-01	--?	8.78E-01	21691	-0.1264	0.0385	1.02E-03	--	2.91E-01
	rs102404015	7	68817129	AUTS2	a	g	18816	0.1429	0.0415	5.71E-04	+++++	4.95E-01	2875	0.0255	0.1031	8.05E-01	++?	8.78E-01	21691	0.1265	0.0385	1.01E-03	++	2.91E-01
	rs10242206	6	68809542	AUTS2	t	c	18813	-0.1423	0.0415	6.16E-04	-----	5.04E-01	2875	-0.0255	0.1031	8.05E-01	--?	8.78E-01	21688	-0.1285	0.0385	1.08E-03	--	2.94E-01
	rs10244247	6	68831755	AUTS2	a	g	16507	0.1729	0.0495	4.73E-04	+++++	7.93E-01	2811	0.0225	0.1148	8.45E-01	++?	5.17E-01	19318	0.1493	0.0455	1.02E-03	++	2.29E-01
	rs10269267	6	68857073	AUTS2	a	c	18798	-0.1479	0.0418	3.99E-04	-----	5.34E-01	10362	0.0167	0.0745	4.08E-01	++?	6.14E-01	19160	-0.0977	0.0365	7.35E-03	++	1.41E-02
	rs917725	6	68861926	AUTS2	a	g	18799	0.1485	0.0418	3.77E-04	+++++	5.34E-01	10362	-0.0502	0.0753	5.05E-01	++?	7.00E-01	19161	0.1107	0.0365	5.39E-03	++	1.11E-02
	rs10254086	6	68878420	AUTS2	a	t	18804	-0.1501	0.0416	3.11E-04	-----	5.42E-01	10362	-0.0162	0.0795	8.39E-01	--?	9.61E-01	19166	-0.1213	0.0369	9.97E-04	--	2.36E-01
	rs10274099	6	68882731	AUTS2	a	g	18808	-0.1507	0.0416	2.93E-04	-----	5.47E-01	10362	-0.0153	0.0797	8.48E-01	--?	9.63E-01	19170	-0.1217	0.0369	9.66E-04	--	1.32E-01
	rs11769738	6	68886156	AUTS2	a	c	18807	-0.1498	0.0416	3.20E-04	-----	5.37E-01	10362	-0.017	0.0797	8.32E-01	--?	9.60E-01	19169	-0.1214	0.0369	9.98E-04	--	1.40E-01
	rs10268729	6	68886796	AUTS2	a	c	18808	-0.1509	0.0416	2.90E-04	-----	5.44E-01	10361	-0.0174	0.0798	8.27E-01	--?	9.57E-01	19169	-0.1224	0.0369	9.09E-04	--	1.38E-01
	rs1177151	6	68888242	AUTS2	a	g	18808	-0.1502	0.0416	3.08E-04	-----	5.36E-01	10362	-0.0107	0.0798	8.93E-01	--?	9.57E-01	19170	-0.1205	0.0369	1.09E-03	--	1.22E-01
	rs11789033	7	68888941	AUTS2	c	g	18808	-0.1502	0.0416	3.09E-04	-----	5.38E-01	10362	-0.0082	0.0809	9.39E-01	--?	9.68E-01	19170	-0.1202	0.0369	1.14E-03	--	1.18E-01
	rs10278978	6	68904666	AUTS2	a	g	18806	-0.1498	0.0416	3.22E-04	-----	5.23E-01	10362	0.0052	0.081	9.49E-01	++?	9.55E-01	19168	-0.1174	0.037	1.50E-03	++	8.87E-02
	rs10245690	6	68911178	AUTS2	a	g	18806	-0.1501	0.0416	3.13E-04	-----	5.19E-01	10362	-0.0057	0.0813	9.45E-01	++?	9.62E-01	19168	-0.1201	0.037	1.18E-03	++	1.14E-01
	rs6958450	6	68917806	AUTS2	a	g	18804	0.1517	0.0416	2.69E-04	+++++	5.25E-01	10362	-0.0011	0.0813	8.98E-01	++?	9.61E-01	19166	0.12	0.037	1.20E-03	++	9.43E-02
	rs3800979	6	68920489	AUTS2	a	g	18805	0.1506	0.0417	3.01E-04	+++++	5.12E-01	10362	-0.0035	0.0813	9.65E-01	++?	9.58E-01	19167	0.1185	0.037	1.40E-03	++	9.17E-02
	rs10272781	6	68924823	AUTS2	c	g	18805	0.1496	0.0417	3.30E-04	+++++	5.01E-01	10362	-0.0047	0.0815	9.54E-01	++?	9.56E-01	19167	0.1176	0.0371	1.54E-03	++	9.19E-02
	rs10275721	6	68928332	AUTS2	t	c	18804	-0.1492	0.0417	3.44E-04	-----	4.92E-01	10362	0.0062	0.0817	9.40E-01	++?	9.53E-01	19166	-0.1171	0.0371	1.62E-03	++	9.02E-02
	rs7784950	6	68929094	AUTS2	a	g	18807	0.1491	0.0417	3.44E-04	+++++	4.92E-01	10362	-0.005	0.0818	9.52E-01	++?	9.55E-01	19169	-0.1173	0.0372	1.59E-03	++	9.33E-02
	rs10487943	6	68929369	AUTS2	a	c	18807	-0.1497	0.0417	3.27E-04	-----	4.79E-01	10362	0.014	0.0818	8.64E-01	++?	9.30E-01	19169	-0.1139	0.0372	1.81E-03	++	7.46E-02
	rs10295969	6	68931447	AUTS2	t	c	18807	0.1496	0.0417	3.28E-04	+++++	4.75E-01	10362	-0.0176	0.0822	8.31E-01	++?	9.14E-01	19169	-0.1154	0.0372	1.92E-03	++	6.97E-02
	rs17140938	6	68931833	AUTS2	t	c	18811	-0.1507	0.0418	3.07E-04	-----	4.36E-01	10362	-0.0099	0.0826	9.57E-01	++?	8.84E-01	19167	0.1118	0.0371	1.54E-04	++	8.78E-02
	rs2658504	6	68932444	AUTS2	c	g	18811	0.149	0.0416	3.46E-04	+++++	4.86E-01	10362	0.0034	0.0829	9.74E-01	++?	7.83E-01	14686	0.1292	0.0387	8.34E-04	++	1.97E-01
	rs740946	6	68939549	AUTS2	c	g	16516	0.1879	0.0497	1.55E-04	+++++	8.00E-01	2813	-0.0072	0.1174	9.51E-01	++?	6.69E-01	19329	-0.1582	0.0458	5.45E-04	++	1.26E-01
	rs1000874	6	68945639	AUTS2	a	g	18811	-0.1485	0.0416	3.63E-04	-----	4.90E-01	2875	-0.0034	0.1049	9.74E-01	++?	7.83E-01	12168	-0.1288	0.0387	8.68E-04	++	1.99E-01
	rs10216186	6	69033104	AUTS2	a	c	14803	0.1649	0.0455	2.94E-04	+++++	6.14E-01	10135	-0.0789	0.0693	2.55E-01	++?	6.96E-01	24938	0.0915	0.038	1.62E-02	++	3.27E-03
	rs11770963	6	69425726	AUTS2	a	g	18656	0.146	0.0416	4.46E-04	+++++	8.73E-01	10357	0.0103	0.0697	8.82E-01	++?	9.31E-01	29013	0.1034	0.0357	2.01E-03	++	9.46E-02
141	rs10241548	7	77040630	PTNP2	g	t	18786	0.1714	0.0624	6.00E-03	+++++	1.96E-01	2853	-0.1504	0.1512	3.20E-01	++?	1.07E-02	21639	0.1246	0.0577	3.08E-02	++	4.91E-02
	rs1582304	7	77055410	PTNP2	a	t	18785	0.1709	0.0623	6.10E-03	+++++	1.95E-01	2853	-0.1504	0.1512	3.20E-01	++?	1.07E-02	21638	0.1245	0.0576	3.10E-02	++	4.94E-02
	rs10273942	7	77077392	PTNP2	t	g	18788	-0.1283	0.0626	4.90E-04	-----	5.04E-01	10306	0.0401	0.0775	6.05E-01	--?	4.58E-03	29094	-0.1163	0.0487	1.70E-02	++	9.49E-03
142	rs1779623	7	87229697	RUMDC3B	a	g	1720	0.0833	0.0278	2.78E-03	+++++	6.99E-01	7848	0.0029	0.0299	9.30E-01	++?	1.41E-02	20568	0.0498	0.0212	1.90E-02	++	6.20E-02
	rs1029424	7	87254977	RUMDC3B	t	c	1423	0.0924	0.0256	3.10E-04	+++++	2.79E-02	10287	0.0268	0.0314	9.66E-01	++?	2.27E-02	21689	0.0572	0.0202	4.72E-03	++	2.47E-02
	rs12704370	7	87268605	RUMDC3B	a	g	10710	-0.0449	0.0307	1.44E-01	???	1.07E-01	7841	-0.0075	0.0329	8.20E-01	--?	4.74E-02	18551	-0.0275	0.0224	2.21E-01	++	4.06E-01
	rs2279542	7	87402433	ADAM2	c	g	6847	0.0861	0.0418	3.95E-02	???	1.67E-01	7487	0.021	0.0337	5.33E-01	???	1.00E-04	14334	0.0662	0.0256	7.54E-02	++	2.25E-01
143	rs19781117	6	92922601	CALCR	a	g	14720	-0.0833	0.0283	3.28E-03	+?..?..	1.76E-01	10349	-0.0354	0.0305	2.46E-01	---	6.45E-01	25069	-0.0621	0.0207	3.21E-03	---	2.50E-01
	rs2283006	6	92923658	CALCR	a	g	14720	-0.0832	0.0283	3.31E-03	+?..?..	1.75E-01	10349	-0.0382	0.0305	2.09E-01	---	6.69E-01	25069	-0.0624	0.0207	6.46E-03	---	2.80E-01
	rs6976031	6	92926362	CALCR	c	g	14727	-0.0832	0.0282	3.16E-03	+?..?..	1.62E-01	10303	-0.0349	0.0305	2.52E-01	---	6.34E-01	25030	-0.0609	0.0207	3.25E-03	---	2.45E-01
	rs2074120	6	92945029	CALCR	a	c	18748	0.0861	0.0244	4.27E-04	+++++	3.17E-01	10359	0.0345	0.0304	2.57E-01	+++	6.62E-01	29107	0.0659	0.019	5.36E-04	+++	1.86E-01
	rs2299253	6	92947991	CALCR	t	c	18750	0.0868	0.0244	3.84E-04	+++++	3.32E-01	10360	0.0331	0.0304	2.77E-01	+++	6.78E-01	29110	0.0658	0.019	5.49E-04	+++	1.68E-01
	rs10488542	6	92952300	CALCR	t	g	18749	0.0866	0.0244	3.96E-04	+++++	3.39E-01	10360	0.0345	0.0304	2.57E-01	+++	6.62E-01	29109	0.0662	0.019	5.05E-04	+++	1.81E-01
	rs9690762	6	92953866	CALCR	a	g	18750	0.0863	0.0244	4.18E-04	+++++	3.41E-01	10360	0.0345	0.0304	2.57E-01	+++	6.62E-01	29110	0.066	0.019	5.23E-04	+++	1.75E-01
	rs2301680	6	92954235	CALCR	a	g	18737	0.0858	0.0245	4.50E-04</														

rs11768363	7	147050616	CNTNAP2	t	c	4512	0.1095	0.0706	1.21e-01	??????	1.69e-03	10324	-0.0551	0.0372	1.38e-01	++	1.25e-01	14836	-0.0193	0.0329	5.57e-01	++	3.92e-02
rs1078756	7	156672759	UBE3C	t	c	18775	0.8007	0.0231	4.67e-04	+++++	7.42e-03	10348	0.0454	0.0285	1.12e-01	++	2.17e-01	29123	0.0667	0.0179	2.02e-04	++	3.76e-01
rs10251112	7	156665285	UBE3C	a	g	18781	0.0887	0.0231	1.15e-04	+++++	4.30e-02	10342	0.0463	0.0285	1.05e-01	++	2.29e-01	29123	0.077	0.0179	5.79e-05	++	2.47e-01
rs696244	7	156697773	UBE3C	a	g	18813	0.0793	0.023	5.50e-04	+++++	1.04e-02	10362	0.046	0.0285	1.06e-01	++	2.61e-01	29175	0.0662	0.0179	2.18e-04	++	3.63e-01
rs0979973	7	156700978	UBE3C	a	g	18816	0.0649	0.0233	3.34e-05	+++++	1.09e-01	10354	0.0502	0.0285	7.84e-02	++	6.45e-01	29170	0.059	0.011	1.07e-03	++	6.90e-01
rs10228145	7	156700758	UBE3C	a	g	18824	0.0905	0.023	8.24e-05	+++++	5.37e-02	10362	0.0472	0.0284	9.69e-02	++	2.78e-01	29176	0.0733	0.0179	4.07e-05	++	2.36e-01
rs2051877	7	156701574	UBE3C	a	t	18812	0.0796	0.023	5.31e-04	+++++	1.01e-02	10363	0.0472	0.0284	9.69e-02	++	2.78e-01	29175	0.0668	0.0179	1.87e-04	++	3.75e-01
rs2051876	7	156701798	UBE3C	a	g	18790	-0.0796	0.023	5.38e-04	-----	9.85e-03	10346	-0.045	0.0285	1.14e-01	++	2.32e-01	29136	-0.066	0.0179	2.29e-04	++	3.45e-01
rs933344	7	156704026	UBE3C	a	t	17028	0.0746	0.0233	1.39e-03	+++++	9.58e-03	10363	-0.0551	0.0285	5.31e-02	++	4.32e-01	27391	0.0226	0.018	2.10e-01	++	4.26e-04
rs1182363	7	156711912	UBE3C	a	g	18810	-0.0794	0.023	5.46e-04	-----	-9.63e-03	10363	-0.0464	0.0285	1.04e-01	++	2.84e-01	29173	-0.0664	0.0179	2.08e-04	++	3.68e-01
rs1182360	7	156713311	UBE3C	c	g	18811	0.0917	0.023	6.62e-05	+++++	3.97e-02	10351	0.0495	0.0285	8.27e-02	++	2.95e-01	29162	0.0751	0.0179	2.75e-05	++	2.49e-01
rs6459742	7	156719745	UBE3C	a	g	18770	-0.065	0.0234	5.47e-03	-----	9.81e-02	10356	-0.049	0.0285	8.58e-02	++	7.5e-01	29126	-0.0586	0.0181	1.20e-03	++	6.64e-01
rs6459743	7	156719869	UBE3C	a	g	18799	-0.0755	0.0231	1.07e-03	-----	7.71e-02	10358	-0.0474	0.0285	9.61e-02	++	3.30e-01	29157	-0.064	0.0179	3.35e-04	++	4.44e-01
rs1182438	7	156720199	UBE3C	a	g	12623	0.048	0.0271	7.63e-02	-----	1.84e-03	10358	0.0494	0.0284	8.26e-02	++	3.08e-01	22981	0.0487	0.0196	1.31e-02	++	9.72e-01
rs6948082	7	156721641	UBE3C	a	g	17015	-0.069	0.0235	3.30e-02	-----	1.81e-01	10358	-0.0488	0.0285	8.55e-02	++	3.15e-01	27373	-0.0608	0.0181	7.95e-04	++	5.85e-01
rs3779598	7	156723619	UBE3C	a	g	16766	0.0662	0.0245	6.79e-03	+++++	1.59e-02	10358	0.0488	0.0285	8.55e-02	++	3.15e-01	27124	0.0588	0.0186	1.55e-03	++	6.43e-01
rs6960676	7	156723678	UBE3C	a	g	18729	-0.0721	0.0231	1.83e-03	-----	4.95e-02	10330	-0.0399	0.0285	1.62e-01	++	2.05e-01	29059	-0.0593	0.0179	9.45e-04	++	3.80e-01
rs6459745	7	156723949	UBE3C	c	g	16793	0.0476	0.0247	5.36e-02	+++++	2.57e-01	10354	0.0481	0.0285	9.13e-02	++	3.22e-01	27147	0.0478	0.0187	1.04e-02	++	9.89e-01
rs10266129	7	156725148	UBE3C	a	g	16748	0.0464	0.0247	6.06e-02	+++++	2.60e-01	10354	0.0489	0.0284	8.52e-02	++	3.15e-01	27102	0.0475	0.0186	1.09e-02	++	9.47e-01
rs10266518	7	156725451	UBE3C	a	g	16706	0.0469	0.0248	5.81e-02	+++++	2.64e-01	10353	0.045	0.0284	1.13e-01	++	3.82e-01	27059	0.0469	0.0187	1.36e-02	++	9.60e-01
rs12730254	7	158759008	VPR2	c	g	8713	-0.0775	0.0429	7.08e-02	-----	8.36e-01	10169	-0.0333	0.0332	3.16e-01	++	6.51e-01	18882	-0.0499	0.0263	5.76e-02	++	4.15e-01
rs2540359	7	158529046	VPR2	a	g	6378	0.0632	0.0546	2.47e-01	+++++	7.71e-01	10169	0.0333	0.0332	3.16e-01	++	6.51e-01	16547	0.0414	0.0263	1.45e-01	++	6.40e-01
rs7784586	7	158529140	VPR2	t	c	8713	0.0749	0.0429	8.09e-02	+++++	8.41e-01	10169	0.0325	0.0332	3.27e-01	++	6.50e-01	18882	0.0484	0.0263	5.54e-02	++	4.34e-01
rs7392117	7	158541582	VPR2	a	g	1714	0.0679	0.045	1.31e-01	+++++	7.35e-01	9973	0.0224	0.034	5.10e-01	++	5.99e-01	18687	0.0389	0.0271	1.51e-01	++	4.20e-01
rs3793227	7	158542204	VPR2	a	g	18885	0.1118	0.0208	1.96e-04	+++++	5.37e-02	10350	0.0254	0.0352	4.44e-01	++	2.25e-01	29700	0.0624	0.0224	2.37e-04	++	1.79e-01
rs3793223	7	158556335	VPR2	a	g	18693	-0.1141	0.0305	1.83e-04	-----	5.53e-02	10320	-0.0254	0.0342	4.44e-01	++	4.28e-01	29013	-0.0735	0.0225	1.07e-03	++	4.91e-02
rs3793227	7	158578689	VPR2	a	g	18685	0.1144	0.0301	1.41e-04	+++++	5.46e-01	10329	0.0355	0.0337	2.92e-01	++	3.98e-01	29104	0.0794	0.0224	4.06e-04	++	8.08e-02
rs2270314	7	158589385	VPR2	c	g	18555	0.1189	0.0306	1.01e-04	+++++	6.32e-01	2853	0.0164	0.0657	8.03e-01	++	1.42e-01	21048	0.1006	0.0277	2.86e-04	++	1.57e-01
rs3828962	7	158593367	VPR2	c	g	18575	0.1178	0.0306	1.17e-04	+++++	7.57e-01	2873	0.008	0.0651	9.03e-01	++	1.30e-01	21448	0.0979	0.0277	4.06e-04	++	1.27e-01
rs17837875	7	158595844	VPR2	c	g	16631	0.1243	0.0315	7.84e-05	+++++	6.88e-01	2837	0.0252	0.0657	7.01e-01	++	1.55e-01	19468	0.1058	0.0284	1.96e-04	++	1.74e-01
rs3793237	7	158604794	VPR2	c	g	6574	0.0237	0.0484	6.25e-01	-----	3.91e-02	2467	0.0421	0.0704	5.50e-01	++	1.00e+00	9041	0.0296	0.0399	4.58e-01	++	8.30e-01
rs17837880	7	158607556	VPR2	c	g	10691	-0.0895	0.0393	2.26e-02	-----	4.93e-01	9970	-0.0419	0.0444	3.45e-01	++	7.81e-01	2061	-0.0318	0.0294	2.80e-01	++	2.67e-02
rs6985318	7	322993		a	g	14831	-0.128	0.0364	4.42e-04	-----	6.71e-01	7487	0.048	0.0413	2.45e-01	??	1.00e+00	22318	-0.0511	0.0273	6.15e-02	++	1.39e-03
rs7007915	7	323018		a	g	18730	0.0806	0.0303	7.81e-03	+++++	4.37e-01	10338	0.0287	0.047	1.73e-02	++	3.50e-01	29068	0.01	0.0228	6.63e-01	++	3.93e-04
rs1979081	7	324288		a	g	16335	0.0675	0.0328	4.10e-02	++++	5.31e-01	10348	0.0789	0.0347	2.32e-02	++	3.76e-01	29675	0.0082	0.0239	9.31e-01	++	5.45e-02
rs1550950	7	324709		a	g	18784	-0.0477	0.0301	1.13e-01	-----	1.39e-01	10358	0.0758	0.0347	2.87e-02	++	4.47e-01	29142	0.0053	0.0227	8.15e-01	++	1.78e-03
rs19802017	7	1766104	ARHGAP10	t	c	18789	-0.0863	0.0228	1.58e-04	-----	3.49e-01	9969	-0.0213	0.0289	4.62e-01	++	8.03e-01	29148	-0.0614	0.0179	6.08e-04	++	7.74e-02
rs4291722	7	15830711		a	g	14831	-0.0973	0.0261	1.90e-04	-----	9.60e-01	7874	-0.0451	0.0331	1.72e-01	??	8.11e-02	22705	-0.0773	0.0205	1.63e-04	++	2.16e-01
rs6994058	7	15832852		a	g	14830	-0.0866	0.0258	7.79e-04	-----	8.96e-01	7487	-0.027	0.0339	4.26e-01	??	1.00e+00	22317	-0.0647	0.0205	1.61e-03	++	1.62e-01
rs13282650	7	60465455		a	g	18756	-0.0841	0.0228	2.33e-04	-----	6.17e-02	9973	-0.0294	0.0313	3.48e-01	??	2.61e-01	28729	-0.0651	0.0184	4.09e-04	++	1.58e-01
rs11775907	7	60467881		t	g	18806	0.0827	0.0228	2.95e-04	++++	5.31e-02	9973	0.0373	0.0294	2.05e-01	++	2.09e-01	28779	-0.0656	0.018	2.69e-04	++	2.22e-01
rs6471785	7	60471437		a	g	18765	0.0843	0.0228	2.23e-04	+++++	5.66e-02	9974	0.0333	0.0294	2.57e-01	++	2.55e-01	28739	-0.0651	0.018	2.99e-04	++	1.70e-01
rs3942395	7	60472404		a	g	18787	-0.0846	0.0228	2.08e-04	-----	4.86e-02	9958	-0.0324	0.0294	2.70e-01	++	2.34e-01	28745	-0.0651	0.018	3.09e-04	++	1.61e-01
rs1554246	7	60481367		a	g	18815	-0.0849	0.0228	2.01e-04	-----	5.83e-02	9975	-0.0408	0.031	1.49e-01	++	1.51e-01	28790	-0.0656	0.0184	1.15e-04	++	2.97e-01
rs1949477	7	60481683		a	g	18815	-0.085	0.0228	1.96e-04	-----	5.97e-02	9975	-0.035	0.0294	1.44e-01	++	1.55e-01	28790	-0.0656	0.018	1.27e-04	++	2.97e-01
rs1401423	7	60482787		a	g	18799	0.0847	0.0228	2.08e-														

rs10092477	8	90653170	a	g	18787	-0.0641	0.023	5.39e-03	-----	8.88e-01	10339	0.0218	0.029	4.53e-01	+++	7.45e-01	29126	-0.0309	0.018	8.61e-02	++	2.03e-02	
rs6981677	8	90659633	c	g	18765	-0.0617	0.023	7.29e-03	-----	9.16e-01	10339	0.0218	0.029	4.53e-01	+++	7.45e-01	29104	-0.0295	0.018	1.02e-01	++	2.41e-02	
161	rs454039	8	104172086	t	c	18816	-0.1727	0.0484	3.61e-04	-----	7.24e-01	10241	-0.0248	0.0582	4.62e-01	---	9.44e-01	29057	-0.1196	0.0372	1.31e-01	++	8.62e-02
162	rs1450188	8	106690657	ZFPM2	c	18761	-0.0961	0.036	4.21e-03	-----	4.49e-01	10361	0.0254	0.0408	5.37e-01	+++	1.69e-01	29122	-0.047	0.0259	7.00e-02	++	2.15e-02
	rs1031305	8	106697907	ZFPM2	a	18766	-0.1081	0.035	2.04e-03	-----	4.53e-01	10361	0.0260	0.0467	1.95e-01	+++	7.14e-01	29129	0.0472	0.028	9.12e-03	++	3.81e-03
	rs13281515	8	106697907	ZFPM2	a	18768	-0.1235	0.035	1.19e-04	-----	1.16e-01	10361	0.0803	0.0436	6.37e-02	+++	7.00e-01	29029	-0.0434	0.0271	1.12e-01	++	2.58e-04
	rs17218607	8	106707484	ZFPM2	t	18753	-0.0965	0.0352	6.07e-03	-----	5.98e-01	10362	0.079	0.0466	8.98e-02	+++	7.24e-01	29115	-0.0327	0.0281	2.44e-01	++	2.66e-03
	rs17218886	8	106714133	ZFPM2	t	18742	-0.0382	0.0353	2.79e-01	-----	5.95e-01	10361	0.0931	0.0433	3.16e-02	+++	7.22e-01	29103	-0.0427	0.0241	6.03e-01	++	1.88e-02
	rs10505078	8	106717830	ZFPM2	a	16543	0.0574	0.0381	1.32e-01	-----	8.84e-01	10315	-0.0643	0.0462	1.64e-01	+++	6.54e-01	26588	0.0081	0.0294	7.82e-01	++	4.21e-02
	rs17291297	8	106720186	ZFPM2	a	18712	0.0593	0.0355	9.47e-02	-----	8.88e-01	9953	-0.07	0.0439	1.11e-01	??	4.29e-01	28665	0.0082	0.0276	7.67e-01	++	2.20e-02
	rs13263514	8	106720599	ZFPM2	a	18704	0.0568	0.0355	1.09e-01	-----	9.01e-01	9955	-0.0778	0.0433	7.25e-02	??	4.15e-01	28659	0.0027	0.0275	9.22e-01	++	1.62e-02
	rs17219068	8	106720704	ZFPM2	a	18710	-0.055	0.0355	1.21e-01	-----	9.07e-01	9954	0.0763	0.0433	7.84e-02	??	4.26e-01	28664	-0.0022	0.0275	9.36e-01	++	1.90e-02
	rs13264377	8	106720881	ZFPM2	t	18714	0.0553	0.0354	1.19e-01	-----	9.01e-01	9959	-0.0753	0.0433	8.21e-02	??	4.03e-01	28673	0.003	0.0274	9.13e-01	++	1.95e-02
	rs13254699	8	106721129	ZFPM2	c	18731	-0.0553	0.0354	1.18e-01	-----	9.02e-01	9968	0.0586	0.0469	2.12e-01	??	4.99e-01	28669	-0.014	0.0283	6.21e-01	++	5.26e-02
	rs6989566	8	106721300	ZFPM2	t	14831	-0.0724	0.0399	6.80e-02	-----	9.48e-01	9919	0.0532	0.0477	2.65e-01	??	2.84e-01	24750	-0.0213	0.0302	4.83e-01	++	4.34e-02
163	rs13266634	8	118253964	SLC30A8	t	18880	-0.103	0.0232	9.42e-06	-----	6.10e-01	10279	-0.0945	0.0292	1.22e-03	---	8.94e-01	28959	-0.0997	0.0182	0.40e-05	---	8.20e-02
	rs13820177	8	118254026	SLC30A8	a	18735	-0.0924	0.0231	1.70e-05	-----	6.10e-01	10279	-0.0945	0.0292	1.22e-03	---	2.69e-01	28959	-0.0997	0.0198	5.53e-05	---	1.04e-01
	rs11558471	8	118254914	SLC30A8	a	18716	0.0986	0.023	1.90e-05	-----	5.77e-01	10309	0.0319	0.0387	4.09e-01	+++	3.55e-01	29025	0.0812	0.0128	0.420e-05	++	1.38e-01
	rs11774700	8	118289451	SLC30A8	t	10777	0.0926	0.0231	2.93e-03	-----	4.45e-01	10309	0.0373	0.0404	3.56e-01	+++	3.91e-01	21086	0.072	0.0246	3.47e-03	++	2.78e-01
164	rs4871655	8	127086430	t	g	18806	0.0692	0.0707	3.28e-01	-----	8.62e-01	10356	-0.0155	0.0329	6.38e-01	---	5.68e-01	29162	-0.0004	0.0298	9.89e-01	++	2.77e-01
	rs2038288	8	127088070	a	t	18701	-0.0597	0.0711	4.01e-01	-----	8.40e-01	10300	0.0108	0.0329	7.42e-01	-0+	5.95e-01	29001	-0.0016	0.0299	9.56e-01	++	3.68e-01
	rs13274749	8	127096325	t	g	18699	-0.0871	0.0709	2.19e-01	-----	8.69e-01	10300	-0.0237	0.0329	4.72e-01	-0+	6.43e-01	28999	-0.0349	0.0298	2.42e-01	++	4.77e-01
	rs10094962	8	127099117	t	c	14334	0.1113	0.0812	1.71e-01	-----	6.50e-01	2816	0.0998	0.0605	7.10e-01	+-?	3.61e-01	17150	0.0308	0.0309	0.72e-01	++	1.15e-01
	rs1285598	8	127099435	a	g	18704	0.1434	0.0699	3.97e-02	-----	6.21e-01	10304	0.0068	0.0832	9.35e-01	+++	5.08e-01	29008	0.0871	0.0334	1.03e-01	++	2.08e-01
	rs1016046	8	127101842	a	g	18702	0.1568	0.0692	3.38e-02	-----	6.01e-01	10308	0.0042	0.0839	9.00e-01	+++	5.63e-01	29008	0.0502	0.0433	2.46e-01	++	3.51e-01
	rs6474043	8	127102177	t	c	16669	0.2244	0.0699	1.33e-03	-----	1.09e-01	10309	0.0102	0.03	7.58e-01	+++	6.27e-01	26978	0.0492	0.0298	9.89e-02	++	5.59e-03
	rs1491481	8	127103048	a	g	18699	0.2051	0.0666	2.08e-03	-----	1.22e-01	10310	0.0888	0.033	7.90e-01	+++	6.21e-01	29009	0.0475	0.0296	1.08e-01	++	8.27e-03
	rs1491483	8	127105893	a	g	16667	-0.2267	0.0697	1.15e-03	-----	9.79e-01	10357	-0.0077	0.0328	8.14e-01	+++	5.79e-01	27026	-0.0474	0.0297	1.10e-01	++	4.47e-03
	rs1491485	8	127106120	a	g	18615	0.1156	0.0549	3.58e-02	-----	7.99e-01	10358	-0.0128	0.0808	8.74e-01	+++	5.74e-01	28973	0.075	0.0454	8.94e-02	++	1.89e-01
	rs13281356	8	127109754	ZFPM2	a	16667	0.235	0.0662	3.84e-04	-----	1.65e-01	10342	0.0354	0.0502	4.82e-01	+++	3.80e-01	27009	0.1083	0.04	6.78e-02	++	1.63e-02
	rs4520128	8	127111193	a	g	18615	-0.1155	0.0552	3.64e-02	-----	5.63e-01	10362	0.0248	0.0614	6.86e-01	+++	5.60e-01	28977	-0.0528	0.041	1.99e-01	++	8.93e-02
	rs1552704	8	127113289	a	g	18614	0.0835	0.0551	1.30e-01	-----	1.98e-01	10363	0.0617	0.1	9.07e-01	+++	5.57e-01	28977	0.0613	0.0483	2.04e-01	++	4.04e-01
	rs13279203	8	127114737	a	g	18683	-0.1126	0.0546	3.92e-02	-----	4.42e-01	10339	0.0306	0.0828	7.12e-01	+++	6.68e-01	29022	-0.0492	0.0456	1.29e-01	++	1.49e-01
	rs1602506	8	127115691	a	g	16665	0.2143	0.0701	2.23e-03	-----	8.00e-01	10338	0.0114	0.0785	8.85e-01	+++	7.03e-01	27003	0.2423	0.0523	1.75e-02	++	5.39e-02
	rs1602507	8	127115763	a	g	18651	-0.1014	0.0647	6.37e-02	-----	6.01e-01	10337	0.0048	0.0839	9.00e-01	+++	7.03e-01	29008	0.0502	0.0433	2.46e-01	++	3.47e-01
	rs10956282	8	127117218	a	t	18672	-0.1023	0.0548	6.18e-02	-----	6.90e-01	10337	-0.017	0.0785	8.29e-01	+++	5.78e-01	29009	0.0744	0.0449	9.80e-02	++	3.73e-01
	rs6997268	8	127117450	a	g	16357	-0.2017	0.072	5.07e-03	-----	8.17e-01	10337	-0.01	0.0784	8.98e-01	+++	5.77e-01	26994	-0.114	0.053	3.16e-02	++	7.17e-02
	rs6997036	8	127117547	a	g	18693	0.1974	0.0671	3.28e-03	-----	1.50e-01	10337	0.0121	0.0813	8.77e-01	+++	5.77e-01	29030	0.1189	0.051	1.96e-02	++	7.23e-02
	rs6997697	8	127117898	a	g	18691	-0.1935	0.0672	4.00e-03	-----	1.54e-01	10332	0.0094	0.0812	9.08e-01	+++	5.63e-01	29023	-0.111	0.0518	3.20e-02	++	5.42e-02
	rs7001995	8	127118103	a	g	16612	-0.2007	0.0721	5.40e-03	-----	1.71e-01	10337	0.0102	0.081	9.00e-01	+++	5.62e-01	26949	-0.1075	0.0539	4.60e-02	++	5.18e-02
	rs1032371	8	127119047	a	g	16612	0.2116	0.071	2.87e-03	-----	1.35e-01	10337	-0.0102	0.081	9.00e-01	+++	5.62e-01	26949	-0.1152	0.0534	3.09e-02	++	3.95e-02
	rs1032372	8	127119059	a	t	16612	0.2105	0.0709	3.00e-03	-----	1.59e-01	10337	-0.0091	0.0799	9.09e-01	+++	5.64e-01	26949	-0.1138	0.0534	3.19e-02	++	3.98e-02
	rs1032373	8	127119411	a	g	16608	-0.1052	0.0559	5.81e-02	-----	1.09e-01	10337	-0.0017	0.0799	9.08e-01	+++	5.73e-01	26945	-0.1708	0.0454	1.18e-01	++	2.83e-01
	rs1552706	8	127119597	a	g	16603	-0.0886	0.0505	7.96e-02	-----	2.42e-01	10335	-0.0134	0.0753	5.96e-01	+++	5.77e-01	29028	-0.0653	0.0419	1.20e-01	++	4.07e-01
	rs12220646	8	127121342	a	g	18671	0.0774																

rs10990221	g	104456687	a	c	18796	-0.0865	0.0234	2.21E-04	-----	1.02E-01	2488	0.0113	0.0588	8.48E-01	?	1.00E+00	21284	-0.0731	0.0217	7.70E-04	++	1.22E-01	
rs10990223	g	104457489	a	c	18786	-0.0842	0.0235	3.37E-04	-----	1.02E-01	2487	0.0117	0.0588	8.43E-01	?	1.00E+00	21273	-0.0701	0.0218	1.14E-03	++	1.30E-01	
rs7875761	t	104459820	t	c	18811	-0.092	0.0234	8.58E-05	-----	1.81E-01	9974	0.0235	0.0305	4.40E-01	?	8.08E-01	28785	-0.0492	0.0186	8.04E-03	++	2.66E-03	
rs10817039	g	112349358	SVEP1	a	g	18768	0.057	0.0244	1.93E-02	+++++	1.46E-01	10357	0.0347	0.0293	2.37E-01	++	8.63E-01	29125	0.0479	0.0187	1.07E-02	++	5.59E-01
rs10804077	g	112351799	SVEP1	a	g	18769	0.0277	0.0244	1.45E-04	+++++	9.82E-01	10357	0.0359	0.0293	2.21E-01	++	8.66E-01	29126	0.0694	0.0187	2.13E-04	++	1.36E-01
rs16915123	g	112357483	SVEP1	t	c	18756	-0.0933	0.0246	1.50E-04	-----	9.91E-01	10341	0.0338	0.0294	1.86E-01	++	8.94E-01	28917	-0.0705	0.0189	1.85E-04	++	1.49E-01
rs13294021	g	127108209	GAPV01	a	g	14794	0.0812	0.0315	1.01E-02	+++++	7.40E-02	10286	-0.061	0.0478	2.01E-01	---	7.25E-01	25080	0.0381	0.0263	1.47E-01	++	1.30E-02
rs13297440	c	14772	0.116	0.0316	2.40E-04	+++++	5.98E-01	10286	-0.0568	0.0477	2.34E-01	---	7.70E-01	25058	0.0363	0.0263	1.63E-02	++	2.53E-03				
rs2270746	g	127161413	GAPV01	a	g	18701	-0.0683	0.0297	2.14E-02	-----	2.36E-02	10294	0.0147	0.0398	7.12E-01	++	9.29E-01	28995	-0.0386	0.0238	1.05E-01	++	9.47E-02
rs13291306	g	127164619	GAPV01	a	t	18662	-0.0684	0.0298	2.16E-02	-----	2.68E-02	10299	0.0414	0.0476	3.84E-01	++	7.98E-01	28961	-0.0375	0.0253	1.38E-01	++	5.06E-02
rs7044663	g	127308084	MAPKAP1	t	c	16420	0.0525	0.0259	4.29E-02	+++++	2.73E-01	10355	0.0098	0.0302	7.46E-01	++	7.35E-01	26775	0.0344	0.0197	8.01E-02	++	2.83E-01
rs7875713	g	127308724	MAPKAP1	t	g	16420	0.0526	0.026	4.26E-02	+++++	2.70E-01	10355	0.0091	0.0302	7.64E-01	++	7.27E-01	26775	0.0301	0.0197	8.37E-02	++	2.75E-01
rs7865246	g	127309332	MAPKAP1	t	c	18753	-0.0615	0.0241	1.07E-02	-----	3.00E-01	10356	-0.0088	0.0302	7.70E-01	---	7.32E-01	29109	-0.0401	0.0188	2.95E-02	++	1.73E-01
rs4837018	g	127351812	MAPKAP1	a	g	18755	-0.0866	0.0241	3.27E-04	-----	9.92E-01	10343	-0.01	0.0303	7.42E-01	---	6.98E-01	29098	-0.0569	0.0189	2.55E-02	++	4.79E-02
rs1253637	g	127354985	MAPKAP1	a	g	18755	-0.0608	0.0241	1.11E-02	-----	2.99E-01	10356	-0.0095	0.0302	7.52E-01	---	7.40E-01	29111	-0.0409	0.0188	3.00E-02	++	1.85E-01
rs10739663	g	127318560	MAPKAP1	a	g	18755	-0.0605	0.0241	2.1E-02	-----	2.91E-01	10356	-0.0088	0.0302	7.70E-01	---	7.32E-01	29111	-0.0404	0.0188	3.20E-02	++	1.81E-01
rs12351538	g	127321324	MAPKAP1	a	g	18756	-0.0605	0.0241	2.1E-02	-----	2.86E-01	10353	-0.0095	0.0302	7.53E-01	---	7.12E-01	29109	-0.0407	0.0188	3.09E-02	++	1.87E-01
rs516735	g	127337513	MAPKAP1	t	c	18755	-0.0608	0.0241	1.16E-02	-----	2.82E-01	10352	-0.0112	0.0302	7.12E-01	---	7.20E-01	29107	-0.0415	0.0188	2.76E-02	++	1.99E-01
rs548793	g	127341752	MAPKAP1	t	c	16983	-0.0631	0.0245	9.86E-03	-----	1.85E-01	10350	-0.0065	0.0303	8.29E-01	---	8.01E-01	27333	-0.0401	0.0191	3.26E-02	++	1.46E-01
rs509753	g	127343339	MAPKAP1	a	g	18765	-0.062	0.0241	1.02E-02	-----	2.17E-01	10350	-0.0073	0.0303	8.11E-01	---	8.09E-01	29115	-0.0408	0.0189	3.05E-02	++	1.58E-01
rs490234	g	127343555	MAPKAP1	t	c	16431	0.0525	0.026	4.35E-02	+++++	2.26E-01	10350	0.0087	0.0303	7.74E-01	++	8.24E-01	26781	0.0309	0.0197	8.56E-02	++	2.73E-01
rs569408	g	127344645	MAPKAP1	a	g	18765	-0.0625	0.0241	9.66E-03	-----	2.29E-01	10350	-0.0094	0.0303	7.56E-01	---	8.32E-01	29115	-0.0419	0.0189	2.62E-02	++	1.70E-01
rs536861	g	127353265	MAPKAP1	a	c	18766	0.0618	0.0241	1.05E-02	+++++	2.31E-01	10350	0.0094	0.0303	7.56E-01	++	8.32E-01	29116	0.0415	0.0189	2.78E-02	++	1.76E-01
rs521119	g	127354215	MAPKAP1	t	c	18766	0.0617	0.0241	1.06E-02	+++++	2.28E-01	10350	0.0116	0.0303	7.03E-01	++	8.54E-01	29116	0.0423	0.0189	2.50E-02	++	1.96E-01
rs497531	g	127354885	MAPKAP1	a	g	18766	0.0621	0.0241	1.01E-02	+++++	2.32E-01	10351	0.0116	0.0303	7.04E-01	++	8.89E-01	29116	0.0442	0.0189	2.61E-02	++	1.78E-01
rs506874	g	127358887	MAPKAP1	a	g	18764	-0.0612	0.0241	1.12E-02	-----	2.41E-01	10340	-0.0132	0.0303	6.64E-01	---	8.13E-01	29104	-0.0429	0.0189	2.39E-02	++	2.15E-01
rs562004	g	127361592	MAPKAP1	t	c	18771	0.0666	0.0242	5.88E-03	+++++	2.45E-01	10342	0.0119	0.0303	6.95E-01	++	7.93E-01	29113	0.0453	0.0189	1.66E-02	++	1.58E-01
rs530628	g	127365755	MAPKAP1	c	g	18763	-0.0964	0.0242	6.88E-05	-----	9.96E-01	10354	-0.0157	0.0303	6.04E-01	0	7.06E-01	29117	-0.0618	0.0189	5.91E-04	++	3.74E-02
rs517064	g	127388807	MAPKAP1	t	c	18784	-0.0684	0.0242	4.73E-03	-----	2.16E-01	10360	-0.0181	0.0303	5.51E-01	---	7.00E-01	29144	-0.0488	0.0189	9.84E-03	++	1.95E-01
rs490333	g	127394390	MAPKAP1	t	c	18801	0.0677	0.0242	5.15E-03	+++++	2.17E-01	10361	0.0201	0.0305	5.09E-01	++	7.07E-01	29162	0.0493	0.0191	9.29E-03	++	2.22E-01
rs476819	g	127402558	MAPKAP1	a	g	18717	0.1021	0.0243	2.71E-05	+++++	9.95E-01	10360	0.0174	0.0304	5.67E-01	0	6.60E-01	29077	0.0691	0.019	2.73E-04	++	2.97E-02
rs534234	g	127409045	MAPKAP1	a	g	18772	-0.0995	0.0243	4.24E-05	-----	9.93E-01	10358	-0.0164	0.0304	5.89E-01	0	6.91E-01	29130	-0.0671	0.019	4.07E-04	++	3.25E-02
rs542974	g	127413039	MAPKAP1	a	g	18731	-0.0673	0.0244	5.72E-03	-----	2.30E-01	10356	-0.0165	0.0304	5.86E-01	0	6.88E-01	29087	-0.0474	0.019	1.28E-02	++	1.93E-01
rs2829312	g	127417694	MAPKAP1	t	c	18563	0.0966	0.0244	7.36E-05	+++++	9.95E-01	10352	0.0178	0.0304	5.57E-01	++	7.02E-01	28915	0.0657	0.019	5.52E-04	++	4.32E-02
rs1259356	g	127428812	MAPKAP1	c	g	18728	-0.0673	0.0244	5.78E-03	-----	2.25E-01	10354	-0.0157	0.0304	6.05E-01	0	7.07E-01	29142	-0.0471	0.019	1.24E-02	++	1.94E-01
rs8176748	g	135121110	ABO	t	c	4686	-0.0791	0.0491	1.07E-01	??-???	2.40E-01	7851	-0.0257	0.0683	7.07E-01	?	4.79E-01	12537	-0.0609	0.0399	1.27E-01	++	5.26E-01
rs8176740	g	135121293	ABO	a	t	16875	0.0948	0.0267	3.79E-04	+++++	2.90E-01	10293	0.0096	0.0461	8.35E-01	++	7.69E-01	27168	0.0734	0.0231	1.49E-03	++	1.10E-01
rs8176732	g	135122124	ABO	a	g	16436	0.1088	0.0281	1.07E-04	+++++	6.37E-01	10348	0.0212	0.0519	6.83E-01	++	7.82E-01	26784	0.0808	0.0247	3.19E-03	++	1.38E-01
rs8176728	g	135122382	ABO	a	g	18760	-0.0914	0.0261	4.65E-04	-----	4.48E-01	10348	0.0067	0.0341	8.98E-01	++	8.80E-01	29108	-0.0552	0.0207	7.78E-04	++	2.23E-02
rs8176717	g	135122855	ABO	g	g	16469	-0.1079	0.028	1.16E-04	-----	6.12E-01	10354	0.0137	0.0331	6.78E-01	++	8.75E-01	26823	-0.0572	0.0214	7.48E-03	++	5.04E-03
rs8176714	g	135122999	ABO	a	g	14780	-0.0795	0.0285	5.26E-03	-----	5.27E-01	10249	0.0228	0.0331	4.91E-01	++	9.00E-01	25029	-0.0359	0.0216	9.60E-02	++	1.92E-02
rs512770	g	135123227	ABO	a	g	17028	-0.0923	0.0265	5.02E-04	-----	3.70E-01	10350	0.0147	0.0331	6.57E-01	++	8.60E-01	27378	-0.0505	0.0207	1.46E-02	++	1.16E-02
rs641959	g	135123520	ABO	a	c	18654	0.0828	0.0259	1.40E-03	+++++	3.95E-01	10355	-0.014	0.033	6.71E-01	---	6.79E-01	29009	0.0459	0.0204	2.43E-02	++	2.10E-02
rs641943	g	135123535	ABO	a	g	18746	0.0842	0.0259	1.14E-03	+++++	3.69E-01	10347	0.0134	0.033	6.94E-01	++	8.45E-01	29093	0.047	0.0203	4.41E-02	++	2.15E-02
rs514708	g	135123564	ABO	a	g	18771	-0.0831	0.0258	1.28E-03	-----	3.86E-01	10355	0.0146	0.033	6.94E-01	++	8.89E-01	29126	0.046	0.0203	2.35E-02	++	1.97E-02
rs517414	g	135123855	ABO	a	g	18807	-0.0																

rs12257919	10314043	t	c	18813	0.1739	0.0604	3.99e-03	+++++	8.95e-02	2876	0.0255	0.132	8.47e-01	+	1.18e-01	21689	0.1482	0.0549	6.97e-03	++	3.07e-01
rs10795772	10314212	t	c	16436	0.2451	0.0674	2.74e-04	+++++	1.87e-01	10343	0.0865	0.0985	3.80e-01	+	7.71e-01	26779	0.1945	0.0556	4.70e-04	++	1.84e-01
rs10752191	10314293	a	g	18808	0.1925	0.0604	1.44e-03	+++++	2.59e-01	2876	0.0255	0.132	8.47e-01	+	1.18e-01	21684	0.1364	0.0549	2.90e-03	++	2.50e-01
rs12268979	10314792	t	g	16475	-0.2268	0.0664	6.40e-04	-----	2.27e-01	2876	-0.0255	0.132	8.47e-01	+	1.18e-01	19351	-0.1861	0.0593	1.70e-03	--	1.73e-01
rs15712106	10315126	t	c	18812	-0.1496	0.0616	2.27e-02	-----	1.04e-01	10363	-0.0787	0.0706	2.65e-01	+	2.63e-01	22969	-0.1189	0.0457	8.75e-03	--	4.46e-01
rs15712123	10315940	c	g	18697	-0.1781	0.0616	8.85e-03	-----	2.29e-01	2872	-0.023	0.132	8.62e-01	+	1.19e-01	21569	-0.1504	0.0555	7.07e-03	--	2.87e-01
rs15721222	10315987	c	g	14270	-0.2324	0.0691	7.70e-04	-----	2.79e-01	2807	-0.0133	0.1344	9.21e-01	+	6.32e-02	17077	-0.1866	0.0615	2.40e-03	--	1.47e-01
rs15721241	10316052	c	g	12324	0.1957	0.0811	1.58e-02	+++++	2.18e-01	10294	0.0725	0.0696	2.98e-01	+	1.94e-01	22618	0.1248	0.0528	1.82e-02	++	2.49e-01
rs1333831	10318009	c	g	6508	0.1237	0.1148	2.81e-01	????+	5.39e-01	10241	0.0814	0.071	2.51e-01	+	2.86e-01	16749	0.0931	0.0604	1.23e-01	++	7.54e-01
rs914987	10318853	c	g	18804	-0.187	0.0598	1.77e-03	-----	1.96e-01	10360	-0.0885	0.0707	2.11e-01	+	4.92e-01	29164	-0.1459	0.0457	1.39e-03	--	2.88e-01
rs2895065	10320197	a	g	18811	-0.1859	0.0598	1.87e-03	-----	2.02e-01	2873	-0.0272	0.131	8.36e-01	+	1.22e-01	21684	-0.1585	0.0544	3.57e-03	--	2.70e-01
rs7905941	10320355	c	g	18810	-0.1809	0.0602	2.65e-03	-----	1.51e-01	2875	-0.0263	0.132	8.42e-01	+	1.17e-01	21685	-0.1543	0.0458	4.85e-03	--	2.87e-01
rs10905687	10322200	c	g	18666	-0.1809	0.0602	2.66e-03	-----	1.51e-01	10330	-0.0882	0.071	2.14e-01	+	2.45e-01	28996	-0.1421	0.0459	1.97e-03	--	3.19e-01
rs1333836	10323780	a	c	18810	0.1784	0.0602	3.06e-03	+++++	1.66e-01	2876	0.034	0.1317	7.97e-01	+	1.13e-01	21686	0.1534	0.0548	5.07e-03	++	3.19e-01
rs1333839	10324061	a	c	18811	-0.1809	0.0602	7.52e-03	-----	9.83e-01	10354	-0.0938	0.083	1.84e-01	+	2.46e-01	29165	-0.1327	0.0458	3.79e-03	--	4.70e-01
rs1215195	10324451	c	g	18672	0.1333	0.0633	3.52e-02	+++++	1.67e-01	2844	0.0818	0.1325	5.37e-01	+	8.89e-02	21516	0.1237	0.0571	3.03e-02	++	7.26e-01
rs7908732	10324519	a	g	18814	-0.177	0.0604	3.38e-03	-----	1.48e-01	2876	-0.034	0.1317	7.97e-01	+	1.13e-01	21690	-0.1521	0.0459	5.58e-03	--	3.24e-01
rs118702	10325126	a	g	18805	-0.1808	0.0601	2.63e-03	-----	1.75e-01	2873	-0.0356	0.1317	7.87e-01	+	1.17e-01	21678	-0.1558	0.0547	4.39e-03	--	3.16e-01
rs10795773	10326103	c	g	18809	0.1796	0.0603	2.90e-03	+++++	1.59e-01	10363	0.0865	0.0709	2.23e-01	+	2.55e-01	29172	0.1405	0.0459	2.22e-03	++	3.17e-01
rs10737038	10326509	a	g	18806	-0.1787	0.0604	3.11e-03	-----	1.57e-01	10363	-0.0865	0.0709	2.23e-01	+	2.55e-01	29169	-0.1399	0.0458	3.74e-03	--	3.22e-01
rs10795774	103268873	a	g	18664	-0.1832	0.0602	2.34e-03	-----	1.81e-01	2841	-0.0405	0.1326	7.60e-01	+	1.10e-01	21505	-0.1588	0.0548	3.76e-03	--	3.27e-01
rs10795775	10327100	c	g	18809	-0.1825	0.0604	2.52e-03	-----	1.76e-01	10363	-0.0865	0.0709	2.23e-01	+	2.55e-01	29172	-0.1421	0.046	1.99e-03	--	3.03e-01
rs7070799	10330746	c	g	18809	0.1811	0.0604	2.72e-03	+++++	1.77e-01	10363	0.0865	0.0709	2.23e-01	+	2.55e-01	29172	0.1413	0.046	2.12e-03	++	3.10e-01
rs7094706	10333573	c	g	16355	0.2367	0.0663	3.59e-04	+++++	2.84e-01	10336	0.0835	0.071	2.29e-01	+	2.52e-01	26691	0.1453	0.0485	6.45e-04	++	1.15e-01
rs1413288	10333888	a	t	14711	0.1467	0.0706	3.78e-02	+++++	1.35e-01	10300	0.0832	0.105	2.45e-01	+	9.90e-01	24316	0.1384	0.0498	3.38e-04	++	1.66e-01
rs10795779	10338427	c	g	18759	0.1747	0.0603	3.75e-03	+++++	1.80e-01	10356	0.1116	0.0726	1.24e-01	+	9.32e-01	28712	0.1489	0.0464	1.32e-03	++	5.04e-01
rs12156543	10339399	a	g	18782	0.1904	0.0595	1.37e-03	+++++	1.17e-01	2488	0.1017	0.1412	4.72e-01	+	1.00e+00	21270	0.177	0.0548	1.24e-03	++	5.63e-01
rs7088313	10339586	c	g	16447	0.2352	0.0652	3.11e-04	+++++	1.90e-01	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	26422	0.1774	0.0484	2.94e-04	++	1.86e-01
rs7072070	10339716	a	t	18814	0.1894	0.0594	1.44e-03	+++++	1.15e-01	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	28789	0.1559	0.0459	6.81e-04	++	3.75e-01
rs7087837	10339965	c	g	18814	-0.1889	0.0594	1.48e-03	-----	1.13e-01	2488	-0.1017	0.1412	4.72e-01	+	1.00e+00	21302	-0.1758	0.0548	1.33e-03	--	5.69e-01
rs12415061	10340635	c	g	18814	0.1887	0.0594	1.50e-03	+++++	1.12e-01	2488	0.1017	0.1412	4.72e-01	+	1.00e+00	21302	0.1756	0.0548	1.34e-03	++	5.70e-01
rs9888040	10341486	a	c	18807	-0.1858	0.059	1.64e-03	-----	1.06e-01	2485	-0.0917	0.1415	5.17e-01	+	1.00e+00	21292	-0.1719	0.0545	1.60e-03	--	5.39e-01
rs10795780	10341894	c	g	16479	0.2339	0.0653	3.38e-04	+++++	1.83e-01	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	26454	0.1766	0.0485	2.69e-04	++	1.90e-01
rs10905699	10342670	c	g	18814	-0.1884	0.0594	1.53e-03	-----	1.12e-01	9975	-0.1063	0.0723	1.42e-01	+	9.69e-01	28789	-0.1553	0.0459	7.14e-04	--	3.80e-01
rs2895067	10342872	c	g	18813	0.1764	0.0629	2.89e-03	+++++	1.11e-01	2487	0.1115	0.141	4.29e-01	+	1.00e+00	21300	0.1667	0.0546	2.26e-03	++	6.71e-01
rs2398212	10343086	a	g	18812	-0.1869	0.0592	1.60e-03	-----	1.20e-01	9975	-0.1063	0.0723	1.42e-01	+	9.69e-01	28787	-0.1546	0.0458	7.40e-04	--	3.84e-01
rs2398213	10343219	a	g	18813	0.1878	0.0594	1.58e-03	+++++	1.05e-01	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	28788	0.1545	0.0459	7.35e-04	++	3.88e-01
rs10795782	10343902	c	g	16783	0.1975	0.0617	1.37e-03	+++++	7.27e-02	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	26758	0.1591	0.0469	7.01e-04	++	3.37e-01
rs10795783	10343953	a	c	18762	0.1772	0.0599	3.10e-03	+++++	5.74e-02	2486	0.1196	0.1419	4.00e-01	+	1.00e+00	21248	0.1685	0.0552	2.26e-03	++	7.08e-01
rs2065592	10344157	a	g	16478	-0.2325	0.0652	3.63e-04	-----	1.63e-01	2488	-0.1017	0.1412	4.72e-01	+	1.00e+00	18966	-0.2095	0.0592	4.01e-04	--	4.00e-01
rs2065591	10344203	c	g	16479	-0.2315	0.0652	3.88e-04	-----	1.48e-01	2488	-0.1017	0.1412	4.72e-01	+	1.00e+00	18967	-0.2089	0.0592	4.32e-04	--	4.04e-01
rs10082491	10344424	a	g	18809	0.182	0.0598	2.35e-03	+++++	7.60e-02	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	28784	0.1512	0.0461	1.03e-03	++	4.20e-01
rs1215193	10344583	c	t	14448	0.248	0.0683	2.84e-04	+++++	1.10e-01	9975	0.1063	0.0723	1.42e-01	+	9.69e-01	24423	0.1812	0.0496	2.63e-04	++	1.54e-01
rs7923025	10345106	a	t	18807	-0.1837	0.0591	1.38e-03	-----	8.74e-02	9972	-0.103	0.0722	1.54e-01	+	9.26e-02	28779	-0.1513	0.0457	9.37e-04	--	3.87e-01
rs10085700	10345227	c	g	14369	0.2436	0.0685	8.80e-04	+++++	1.14e-01	9947	0.105	0.0724	1.45e-01	+	9.90e-01	24316	0.1384	0.0498	3.38e-04	++	1.66e-01
rs10950702	10345647	c	g	18777	0.1839	0.0596	2.02e-03	+++++	8.46e-02	2488	0.1017	0.1412	4.72e-01	+	1.00e+00	21265	0.1715	0.0549	1.79e-03	++	5.92e-01
rs1333833	10347524	c	t	14999	0.2001	0.0621	1.27e-03	+++++	6.83e-02	9975	0.1041	0.0723	1.50e-01	+	9.84e-01	24974	0.1593	0.0471	7.18e-04	++	3.14e-01
rs10466299	10349810	a	g	16476	-0.2226	0.0658	7.24e-04	-----	1.31e-01	2488	-0.1017	0.1412	4.72e-01	+	1.00e+00	18964	-0.201	0.0596	7.50e-04	--	4.38e-01
rs10795784	10349980	c	g	18804	0.1776	0.0599	3.01e-03	+++++	7.96e-02	2488	0.1017	0.1412	4.72e-01	+	1.00e+00	21292	0.166	0.0551	2.61e-03	++	6.21e-01
rs10458816	10350267	c	g	18814	0.1808	0.0601	2.62e-03	+++++	7.02e-02	9969	0.1103	0.0726	1.29e-01	+	9.32e-01	28783	0.1521	0.0463	1.02e-03	++	5.44e-01
rs10458804	10350286	a	g	18817	0.1747	0.0601	3.64e-03	+++++	6.60e-02	2487	0.0926	0.1415	5.13e-01	+	1.00e+00	21304	0.1622	0.0553	3.98e-03	++	5.93e-01
rs10458805	10350325	a	g	18817	0.18	0.06	2.70e-03	+++++	7.46e-02	9975	0.1115	0.0726	1.25e-01	+	9.35e-01	28792	0.1522	0.0462	9.99e-04	++	4.67e-01
rs10458817	10350358	c	t	18803	-0.1748	0.0602	3.70e-03	-----	5.20e-02	9960	-0.1138	0.0732	1.20e-01	+	9.86e-01	28763	-0.1502	0.0465	1.27e-03	--	5.20e-01
rs10458818	10350370	c	g	17032	0.1936	0.0603	1.														

rs7898493	0.94285296	IDE	c	g	18695	0.1614	0.0406	7.08E-05	+++++	7.34E-01	10342	0.1252	0.0583	3.17E-02	+++	2.56E-01	29037	0.1496	0.0333	7.14E-06	++	6.10E-01
rs1187061	0.94295389	IDE	t	c	18461	0.1659	0.04	3.41E-05	+++++	8.46E-01	10241	0.1201	0.0581	3.86E-02	+++	1.24E-01	28702	0.1512	0.0329	4.47E-06	++	5.16E-01
rs2421943	0.94301795	IDE	a	g	18746	-0.1228	0.024	3.21E-07	+++++	9.71E-01	7870	-0.0262	0.0344	4.46E-01	-?	4.35E-01	26616	-0.0919	0.0192	9.37E-06	++	2.13E-02
rs7908111	0.94304784	IDE	a	g	18795	0.1573	0.0399	8.09E-05	+++++	8.09E-01	10353	0.1302	0.0581	2.52E-02	+++	1.27E-01	29148	0.1486	0.0329	6.23E-06	++	7.01E-01
rs11470442	0.94306045	IDE	c	g	14832	0.1604	0.0442	2.82E-04	+++++	7.16E-01	10361	0.1235	0.058	3.32E-02	+++	1.20E-01	25193	0.1468	0.0352	2.95E-05	++	6.13E-01
rs1187078	0.94306865		t	c	8843	0.0933	0.0377	3.0E-02	????	7.39E-01	7849	0.0235	0.0348	3.50E-01	+	4.05E-01	16687	0.0606	0.0255	1.78E-02	++	2.35E-01
rs8583823	0.94343495		t	c	18522	-0.0946	0.0254	2.01E-04	+++++	9.82E-01	7866	-0.0515	0.0348	1.39E-01	-?	4.95E-01	26388	-0.0796	0.0205	1.04E-04	++	3.17E-01
rs2421941	0.94345889		a	g	16384	-0.1612	0.0437	2.29E-04	+++++	7.94E-01	10329	-0.0004	0.0597	9.95E-01	+	1.99E-03	26713	-0.1051	0.0353	2.88E-03	++	2.98E-02
rs8583826	0.94337810		g	18683	-0.1451	0.0276	1.41E-07	+++++	2.12E-01	10314	-0.0557	0.0381	1.44E-01	+	1.61E-01	28997	-0.1143	0.0224	3.14E-07	++	5.74E-02	
rs1187083	0.94342485	KF11	a	t	18703	-0.1417	0.0366	1.08E-04	+++++	6.68E-01	10360	-0.0858	0.0554	1.22E-01	---	4.31E-01	29063	-0.1247	0.0303	4.03E-05	++	4.00E-01
rs1187084	0.94342998	KF11	t	c	18692	0.1895	0.0494	1.26E-04	+++++	6.89E-01	10347	0.2206	0.0874	1.16E-02	+++	3.03E-01	29039	0.1977	0.043	4.62E-05	++	5.75E-01
rs11818332	0.94343418	KF11	t	c	18810	-0.1446	0.0366	7.90E-05	+++++	6.45E-01	10363	-0.1081	0.0541	4.59E-02	+++	6.14E-01	29173	-0.1331	0.0303	1.12E-05	++	5.57E-01
rs3781236	0.94343843	KF11	a	g	18810	-0.1441	0.0366	8.36E-05	+++++	6.44E-01	10363	-0.1192	0.0543	2.81E-02	+++	7.03E-01	29173	-0.1363	0.0303	7.07E-06	++	7.04E-01
rs3844745	0.94344184	KF11	t	c	18813	0.1367	0.0269	3.71E-07	+++++	9.17E-01	7863	0.0166	0.0342	5.35E-02	++	4.50E-01	22676	0.1097	0.0211	2.13E-07	++	1.04E-01
rs4604797	0.94348292	KF11	a	g	18707	0.1425	0.031	1.25E-04	+++++	5.31E-01	10363	0.0263	0.0581	4.58E-02	+++	6.14E-01	27390	0.1311	0.0306	1.73E-05	++	5.99E-01
rs7107758	0.94357013	KF11	a	g	18810	-0.1437	0.0366	8.82E-05	+++++	6.47E-01	10363	-0.1089	0.0543	4.48E-02	+++	6.20E-01	29173	-0.1328	0.0303	1.21E-05	++	5.95E-01
rs10786500	0.94357210	KF11	a	g	16347	0.1649	0.0433	1.40E-04	+++++	8.30E-01	10353	-0.0015	0.0594	9.80E-01	+++	1.80E-03	26790	0.1072	0.035	2.19E-03	++	2.36E-02
rs12261518	0.94357897	KF11	a	g	18700	-0.1914	0.0493	1.05E-04	+++++	6.73E-01	10348	-0.2228	0.0741	2.63E-03	---	3.87E-01	29048	-0.201	0.041	9.69E-07	++	7.24E-01
rs1187094	0.94358158	KF11	a	g	14821	-0.1125	0.0262	1.74E-05	+++++	9.64E-01	7465	-0.047	0.0341	1.68E-01	??	1.00E+00	22286	-0.0882	0.0208	2.19E-05	++	1.28E-01
rs1187096	0.94359568	KF11	t	c	18635	-0.1377	0.0241	1.17E-08	+++++	6.60E-01	7875	-0.0476	0.0344	1.66E-01	-?	2.87E-01	26510	-0.108	0.0197	4.41E-08	++	3.19E-02
rs10882091	0.94364357	KF11	t	c	18816	-0.1492	0.0413	3.05E-04	+++++	6.53E-01	10362	-0.0005	0.0595	9.94E-01	+	1.42E-03	29178	-0.1009	0.0339	2.95E-03	++	4.01E-02
rs12260688	0.94369694	KF11	t	c	18817	0.1435	0.0366	8.80E-05	+++++	6.43E-01	10361	0.1156	0.0545	3.39E-02	+++	6.73E-01	29178	0.1348	0.0304	9.10E-06	++	6.71E-01
rs93714814	0.94372930	KF11	t	c	18814	0.1493	0.0413	2.97E-04	+++++	6.56E-01	10363	-0.0016	0.0596	9.78E-01	+++	1.43E-03	29180	0.1303	0.0339	3.12E-03	++	3.74E-02
rs78959056	0.94372946	KF11	a	g	18815	0.1445	0.0366	7.69E-05	+++++	6.37E-01	10363	0.0742	0.0529	1.85E-01	+++	3.37E-01	29178	0.1234	0.0306	5.57E-05	++	2.95E-01
rs7073868	0.94373424	KF11	t	c	18700	0.1896	0.0494	1.22E-04	+++++	6.84E-01	10348	0.2118	0.078	1.27E-02	---	2.55E-01	29048	0.1952	0.0426	4.72E-06	++	8.21E-01
rs7087860	0.94376134	KF11	a	g	18815	-0.1444	0.0366	7.78E-05	+++++	6.40E-01	10363	-0.1203	0.0524	2.18E-02	---	7.20E-01	29178	-0.1365	0.03	5.39E-06	++	7.06E-01
rs10882094	0.94376556	KF11	a	g	18817	-0.1488	0.0413	3.17E-04	+++++	6.56E-01	10363	-0.0005	0.0595	9.94E-01	+	1.42E-03	29180	-0.1006	0.0339	3.03E-03	++	4.06E-02
rs10717664	0.94382439	KF11	a	t	18815	-0.1445	0.0366	7.70E-05	+++++	6.40E-01	10363	-0.0948	0.0531	7.43E-02	---	5.12E-01	29178	-0.1285	0.0301	2.01E-05	++	4.41E-01
rs11818705	0.94382708	KF11	t	c	18814	0.1449	0.0366	7.39E-05	+++++	6.45E-01	10358	0.117	0.0519	2.41E-02	+++	6.97E-01	29172	0.1356	0.0299	5.77E-06	++	6.60E-01
rs11818708	0.94382761	KF11	c	18815	-0.1446	0.0366	7.67E-05	+++++	6.39E-01	10363	-0.1236	0.0524	1.84E-02	---	7.45E-01	29178	-0.1377	0.03	4.44E-06	++	7.43E-01	
rs11818750	0.94382810	KF11	t	c	16610	0.152	0.0392	1.06E-04	+++++	7.26E-01	10327	0.1271	0.0517	1.40E-02	+++	6.47E-01	26937	0.1429	0.0312	4.76E-06	++	5.01E-01
rs11819351	0.94384310	KF11	t	c	18518	0.1377	0.0383	3.24E-04	+++++	6.13E-01	10361	0.0952	0.0555	8.65E-02	+++	5.54E-01	28879	0.124	0.0315	8.83E-05	++	7.29E-01
rs10882095	0.94384382	KF11	t	c	16308	0.1489	0.026	1.10E-08	+++++	9.91E-01	7875	0.057	0.0343	9.61E-02	++	3.18E-01	29183	0.1154	0.0207	2.58E-08	++	3.28E-02
rs7079990	0.94384580	KF11	t	c	18781	0.1216	0.0332	2.50E-04	+++++	6.93E-01	10362	0.0459	0.0483	3.43E-01	+++	7.71E-01	29143	0.0973	0.0274	3.76E-04	++	1.97E-02
rs10736069	0.94385373	KF11	t	c	18513	0.1467	0.0414	3.91E-04	+++++	6.84E-01	10348	0.2118	0.078	1.27E-02	---	4.36E-01	29048	0.1952	0.0426	4.72E-06	++	8.21E-01
rs7906089	0.94385728	KF11	a	g	18813	0.1496	0.0413	2.95E-04	+++++	6.47E-01	10356	0.0013	0.0594	9.83E-01	+++	1.56E-03	29169	0.1013	0.0339	2.82E-03	++	4.04E-02
rs7096980	0.94387741	KF11	a	t	16367	0.2001	0.0517	1.09E-04	+++++	5.61E-01	10348	0.2124	0.0836	1.11E-02	+++	2.86E-01	26715	0.2035	0.044	3.69E-06	++	9.00E-01
rs6583830	0.94388098	KF11	a	g	18812	0.1512	0.0413	2.52E-04	+++++	7.06E-01	10356	0.0013	0.0594	9.83E-01	+++	1.56E-03	29168	0.1024	0.0339	2.54E-03	++	3.83E-02
rs7079583	0.94391873	KF11	c	18782	-0.1194	0.0336	3.72E-04	+++++	2.47E-01	10362	-0.0459	0.0484	3.43E-01	---	7.71E-01	29144	-0.0955	0.0276	5.40E-03	++	1.21E-01	
rs7079602	0.94391950	KF11	a	g	18702	-0.1905	0.0494	1.13E-04	+++++	6.19E-01	10348	-0.2091	0.0847	1.35E-02	---	2.65E-01	29050	-0.1955	0.0427	4.77E-06	++	8.50E-01
rs7088685	0.94393578	KF11	a	t	18814	-0.1436	0.0366	8.85E-05	+++++	6.23E-01	10322	-0.1041	0.0546	5.65E-02	---	2.60E-01	29136	-0.1314	0.0304	1.56E-05	++	5.48E-01
rs7902436	0.94394677	KF11	a	t	17034	0.1443	0.0371	9.84E-05	+++++	5.21E-01	10363	0.0859	0.0529	1.05E-01	+++	4.80E-01	27397	0.125	0.0304	3.84E-05	++	3.66E-01
rs7917359	0.94394804	KF11	t	c	18816	0.1449	0.0366	7.51E-05	+++++	5.96E-01	10265	0.1307	0.0531	1.38E-02	+++	4.16E-01	29081	0.1403	0.0301	3.22E-06	++	8.26E-01
rs1118714	0.94395021	KF11	t	c	16301	0.1488	0.0261	1.15E-08	+++++	9.93E-01	7875	0.0712	0.0342	3.72E-02	++	3.67E-01	29176	0.1202	0.0207	6.83E-09	++	4.52E-02
rs12264712	0.94395382	KF11	t	c	18703	0.1878	0.0494	1.45E-04	+++++	6.61E-01	10347	-0.2105	0.078	9.86E-03	---	4.36E-01	29048	-0.1952	0.0426	4.72E-06	++	8.21E-01
rs6583833	0.94399780	KF11	c	18637	-0.1056	0.0283	1.94E-04	+++++	6.12E-01	7875	-0.0436	0.0375	2.45E-01	-?	1.44E-01	26512	-0.0831	0.0226	2.34E-04	++	1.87E-01	
rs7078243	0.94404243	KF11	a	g	18636	0.105	0.0284	2.13E-04	+++++	6.11E-01	7873	0.0474	0.0375	2.06E-01	++	1.50E-01	26509	0.084	0.0226	2.07E-04	++	2.21E-01
rs4933734	0.94404547	KF11	a	t	18810	0.151	0.0414	2.64E-04	+++++	6.34E-01	10356	-0.0015	0.0595	9.80E-01	+++	1.58E-03	29166	0.1013	0.034	2.89E-03	++	3.54E-02
rs6583834	0.94409276	KF11	a	g	18741	0.1368	0.0371	2.29E-04	+++++	5.65E-01	10344	0.0972	0.0559	8.24E-02	+++	5.11E-01	29085	0.1247	0.0309	5.49E-05	++	5.55E-01
rs7914504	0.94411998	KF11	a	t	18743	-0.1401	0.0372	1.64E-04	+++++	5.69E-01	10341	-0.1074	0.0553	5.20E-02	---	6.30E-01	29084	-0.1299	0.0309	2.57E-05	++	6.24E-01
rs7096187	0.94413406	KF11	a	g	18742	-0.1412	0.0372	1.44E-04	+++++	5.62E-01	10339	-0.095	0.0557	8.79E-02	---	5.74E-01	29081	-0.1269	0.0309	4.07E-05	++	4.39E-01
rs2488087	0.944166																					

208	rs4757880	11	20006435	NAV2	a	g	14732	0.0517	0.0256	4.33E-02	+++?++	3.26E-01	10336	0.0017	0.0337	9.61E-01	---	6.09E-01	25068	0.0334	0.0204	1.01E-01	++	2.37E-01
	rs920675	11	20006846	NAV2	a	g	18691	0.0651	0.0229	4.48E-03	++++++	4.69E-01	10288	0.0128	0.0291	6.60E-01	---	3.19E-01	28979	0.0451	0.018	1.22E-02	++	1.58E-01
	rs1101166	11	20007755	NAV2	t	c	12480	-0.0846	0.0279	2.39E-03	---?+?	1.55E-01	7487	-0.037	0.0341	2.78E-01	??	1.00E+00	19967	-0.0655	0.0216	0.41E-03	---	2.80E-01
	rs1230601	11	20008788	NAV2	t	c	12497	-0.0878	0.0281	1.77E-03	---?+?	2.38E-01	7487	-0.036	0.034	2.90E-01	??	1.00E+00	19964	-0.0668	0.0217	2.05E-03	---	2.40E-01
	rs1033224	11	20012324	NAV2	a	g	18643	0.0903	0.023	8.67E-04	++++++	2.29E-01	9972	0.011	0.0295	7.09E-01	??	1.09E+00	28615	0.0603	0.0181	8.83E-04	++	3.40E-02
	rs12276043	11	20012329	NAV2	a	g	14648	-0.0797	0.0263	2.60E-03	---?+?	1.14E-01	9972	-0.002	0.0294	9.46E-01	??	1.86E+00	19489	-0.0469	0.0197	2.62E-03	---	4.66E-02
	rs1731550	11	20012849	NAV2	a	g	14654	-0.0769	0.0264	3.57E-03	---?+?	1.92E-01	9948	-0.0008	0.0295	9.79E-01	??	1.62E+00	24602	-0.0431	0.0197	2.86E-02	---	5.46E-02
	rs1945337	11	21378572		a	g	12698	0.0449	0.0287	1.18E-01	+++?+?	1.59E-02	10330	-0.0349	0.0297	2.40E-01	---	9.37E-01	23028	0.0604	0.0206	7.58E-01	++	5.33E-02
	rs11026216	11	2142439		a	g	16750	-0.0911	0.0246	2.18E-04	-----	8.57E-01	10347	0.0498	0.0291	8.78E-02	+++	9.35E-01	27097	-0.0324	0.0188	8.48E-02	++	2.18E-04
	rs12289507	11	21752855		t	c	16691	-0.0959	0.0245	9.22E-05	-----	8.00E-01	10353	0.0469	0.0291	1.07E-01	+++	9.77E-01	27044	-0.0367	0.0187	5.04E-02	++	1.74E-04
rs1725896	11	21785869		a	g	8702	-0.0913	0.0251	9.22E-03	-----	8.60E-01	10169	0.0438	0.0295	1.38E-01	+++	8.99E-01	18871	-0.0121	0.0226	5.91E-01	++	3.21E-03	
rs10766882	11	21790876		t	c	16461	0.055	0.0246	2.56E-02	+++++?	4.11E-03	10363	-0.0501	0.0291	8.55E-02	---	9.72E-01	26824	0.0112	0.0188	5.51E-01	++	5.81E-03	
rs1026257	11	21791089		a	g	18795	0.0143	0.023	5.33E-01	+++++	9.26E-04	10360	-0.0487	0.0291	9.45E-02	---	9.82E-01	29155	-0.0099	0.018	8.28E-01	++	8.94E-02	
rs1083638	11	21791344		a	g	18794	0.0576	0.023	1.25E-02	++++++	9.04E-03	10362	-0.0502	0.0292	8.49E-02	---	9.75E-01	29156	-0.0163	0.0181	3.66E-01	++	3.73E-03	
rs10333641	11	21796969		a	g	18798	0.0569	0.023	1.36E-02	++++++	6.74E-03	10346	-0.0513	0.0291	7.32E-02	---	9.57E-01	29134	0.0151	0.018	3.97E-01	++	3.53E-03	
rs1948121	11	21809061		a	g	16177	0.0488	0.0251	5.17E-02	+++++?	2.94E-03	10305	-0.0591	0.0293	4.37E-02	---	8.52E-01	26482	0.0031	0.0191	8.70E-01	++	5.16E-03	
rs1032229	11	21809745		t	c	18775	-0.0542	0.023	1.85E-02	+++++?	5.14E-03	10356	-0.0533	0.0292	6.76E-02	---	9.52E-01	29131	0.0118	0.0171	4.71E-01	++	3.83E-03	
210	rs1451327	11	57747669		a	g	14831	-0.1536	0.0369	3.11E-05	-----	6.26E-01	7874	0.0162	0.0416	6.98E-01	++	9.85E-01	22705	-0.0788	0.0276	4.30E-03	++	2.26E-03
	rs12802604	11	57748268		t	c	14828	0.1519	0.0369	3.88E-05	+++++?	6.03E-01	387	0.0161	0.2	9.36E-01	??	1.00E+00	15215	0.1474	0.0363	4.85E-05	++	5.04E-01
	rs1031358	11	57750500	OR10Q1	a	g	14784	-0.1422	0.037	1.22E-04	-----	6.16E-01	385	0.004	0.2005	9.84E-01	??	1.00E+00	15169	-0.1374	0.0364	1.60E-04	++	4.73E-01
	rs11229304	11	57754915		t	c	16391	-0.1255	0.035	3.41E-04	-----	5.28E-01	2841	-0.099	0.2007	1.40E-01	++	5.53E-01	19232	-0.1198	0.031	1.13E-04	++	7.26E-01
	rs11229305	11	57754937		t	c	14426	-0.1175	0.038	2.00E-03	---?+?	4.76E-01	2843	-0.0812	0.0666	2.23E-01	++	6.52E-01	17269	-0.1086	0.033	1.00E-03	++	6.36E-01
	rs11229307	11	57756724		t	c	18812	-0.1244	0.032	1.00E-04	-----	6.93E-01	2871	-0.0853	0.0664	1.99E-01	++	6.37E-01	21683	-0.117	0.0288	4.93E-01	++	5.96E-01
	rs1938659	11	57765556		a	g	18813	-0.1231	0.032	1.18E-04	-----	6.97E-01	2871	-0.0853	0.0664	1.99E-01	++	6.37E-01	21684	-0.116	0.0288	5.74E-05	++	6.08E-01
	rs1938656	11	57767287		a	g	18855	0.0956	0.0324	1.38E-01	+++++	3.89E-02	2855	0.0515	0.068	4.99E-01	++	7.69E-01	23510	0.0678	0.0241	4.95E-03	++	9.91E-01
	rs1938668	11	57772714		a	g	18810	0.0478	0.032	1.35E-01	+++++	1.16E-02	2872	0.0918	0.0662	1.66E-01	++	6.13E-01	21682	0.0681	0.0288	5.14E-02	++	5.50E-01
	rs1938667	11	57774878		a	g	18812	0.0449	0.032	1.60E-01	+++++	6.87E-03	10359	0.023	0.0362	5.25E-01	++	4.16E-01	29171	0.0353	0.024	1.41E-01	++	6.50E-01
rs1938665	11	57775170		t	c	18738	-0.1257	0.032	8.74E-05	-----	5.12E-01	2872	-0.0903	0.0662	1.73E-01	++	5.96E-01	21610	-0.119	0.0288	3.62E-05	++	6.30E-01	
rs7934043	11	57778966		t	c	18815	0.1214	0.032	1.46E-04	+++++	6.33E-01	10362	0.0741	0.0615	2.29E-01	++	6.98E-01	29177	0.1113	0.0284	8.80E-05	++	4.95E-01	
rs7935090	11	57779198		t	c	18815	0.1212	0.032	1.50E-04	+++++	6.38E-01	2875	0.0877	0.0662	1.85E-01	++	5.23E-01	21690	0.1149	0.0288	6.70E-05	++	6.49E-01	
rs1938640	11	57779992		t	c	18815	-0.1221	0.032	1.51E-04	-----	6.35E-01	2875	-0.0877	0.0662	1.85E-01	++	5.23E-01	21690	-0.1148	0.0288	6.78E-05	++	6.50E-01	
rs12223901	11	57781920		t	c	18815	-0.1211	0.032	1.54E-04	-----	6.34E-01	2875	-0.0877	0.0662	1.85E-01	++	5.23E-01	21690	-0.1147	0.0288	6.87E-05	++	6.51E-01	
rs10750892	11	57784885		a	g	18815	0.1172	0.0319	2.44E-04	+++++	5.98E-01	2875	0.0877	0.0662	1.85E-01	++	5.23E-01	21690	0.1116	0.0287	1.02E-04	++	6.88E-01	
rs7173339	11	57787530		a	g	18815	-0.1224	0.032	1.28E-04	-----	6.22E-01	2875	-0.0877	0.0662	1.85E-01	++	5.23E-01	21690	-0.1158	0.0288	5.81E-05	++	6.37E-01	
rs9680182	11	57788287		a	g	18763	0.1214	0.0324	1.07E-04	+++++	6.27E-01	10363	0.0877	0.0662	1.85E-01	++	5.23E-01	21688	0.1174	0.0289	4.80E-05	++	6.16E-01	
rs7930778	11	57792932		t	c	18810	-0.1194	0.032	1.92E-04	-----	6.20E-01	10360	-0.0215	0.0363	5.54E-01	++	3.99E-01	29170	-0.0766	0.024	1.42E-03	++	4.31E-02	
rs4127346	11	57798978		a	t	18752	-0.1183	0.0322	2.36E-04	-----	5.53E-01	2849	-0.1115	0.0667	9.46E-02	++	7.83E-01	21601	-0.117	0.029	5.45E-05	++	9.27E-01	
rs10792159	11	57803648		a	t	18750	-0.1148	0.0321	3.46E-04	-----	5.88E-01	2850	-0.1122	0.0667	9.25E-02	++	8.08E-01	21600	-0.1143	0.0289	7.75E-05	++	9.72E-01	
rs7101864	11	57809764		a	g	18514	-0.1168	0.0322	2.84E-04	-----	6.91E-01	10328	-0.1044	0.0357	5.19E-02	++	7.80E-01	28842	-0.1135	0.0276	3.94E-05	++	8.43E-01	
rs10897472	11	57815997		a	g	18707	-0.1192	0.0326	2.60E-04	-----	7.14E-01	10269	-0.1027	0.0369	7.30E-01	++	5.17E-01	28976	-0.0725	0.0244	3.00E-03	++	3.05E-02	
rs7109083	11	57816658		a	g	18778	-0.1145	0.0322	3.69E-04	-----	7.03E-01	10338	-0.1033	0.0362	7.13E-01	++	3.01E-01	29116	-0.0698	0.0241	3.72E-03	++	3.67E-02	
rs4287351	11	57820690		a	t	18870	0.1132	0.0321	4.27E-04	+++++	7.70E-01	10338	0.0333	0.0362	7.13E-01	++	3.01E-01	29118	0.0692	0.024	3.95E-03	++	3.89E-02	
rs4938893	11	57851580		a	g	18810	0.1119	0.0309	2.94E-04	+++++	8.40E-01	10359	0.0065	0.0355	8.54E-01	++	2.33E-01	29169	0.0665	0.0233	4.35E-03	++	5.21E-02	
rs10896754	11	57861630		t	c	14800	-0.1336	0.0355	1.70E-04	-----	9.42E-01	10355	-0.0069	0.0355	8.47E-01	++	2.24E-01	29155	-0.0702	0.0251	5.13E-03	++	1.16E-02	
rs1928028	11	57861784		t	c	17920	-0.1131	0.0323	4.61E-04	-----	6.70E-01	10363	-0.0108	0.0362	7.65E-01	++	2.69E-01	27383	-0.0678	0.0241	4.95E-03	++	3.50E-02	
rs65921475	11	57950127	OR582	a	g	18792	0.1252	0.0312	6.7E-05	+++++	3.02E-01	10356	0.007	0.03										

rs1144222	11	133668611	GLB1L3	t	c	18639	-0.0507	0.0279	6.89E-02	-----	1.34E-02	9769	-0.0088	0.0356	8.04E-01	7+	3.77E-01	28408	-0.0348	0.022	1.14E-01	--	3.54E-01
rs2129504	12	18765617	PLCZ1	a	g	8843	0.0077	0.0375	1.88E-01	????-??	8.32E-02	7487	-0.01	0.036	7.81E-01	??	1.00E+00	16330	-0.0015	0.026	9.54E-01	++	7.34E-01
rs1021767	12	18767268	PLCZ1	a	g	14830	0.0914	0.0282	1.18E-03	+++++?++	1.25E-01	10332	0.033	0.0308	2.84E-01	++	3.54E-01	25162	0.0648	0.0208	1.85E-03	++	1.62E-01
rs12307562	12	18768845	PLCZ1	a	g	8464	0.0025	0.0399	9.50E-01	????-??	2.13E-02	9789	0.0364	0.0318	2.53E-01	7+	8.33E-02	18253	0.0322	0.0249	3.50E-01	++	5.06E-01
rs12305012	12	18769012	PLCZ1	a	g	4512	-0.1307	0.0637	4.03E-02	????-??	3.89E-01	7487	0.008	0.0361	8.25E-01	??	1.00E+00	19999	-0.0257	0.0314	4.13E-01	++	5.82E-02
rs10144267	12	18769915	PLCZ1	c	g	18587	0.0409	0.0291	1.03E-01	+++++?++	8.32E-02	10295	0.033	0.0308	2.83E-01	++	3.25E-01	28882	0.0377	0.0195	5.24E-02	++	8.24E-01
rs1487064	12	18770818	PLCZ1	c	g	16615	-0.0329	0.0268	2.19E-01	-?+----	1.81E-01	10314	-0.0382	0.0308	2.14E-01	---	2.77E-01	26929	-0.0352	0.0202	8.18E-02	---	8.97E-01
rs10144269	12	18771904	PLCZ1	a	t	16370	0.0378	0.0274	1.69E-01	+++++?+	1.26E-02	10317	0.035	0.0308	2.56E-01	---	2.35E-01	26687	0.0366	0.0205	7.41E-02	---	9.46E-01
rs1027032	12	18772921	PLCZ1	a	g	16930	0.0468	0.0255	6.69E-02	+++++?+	6.34E-02	10315	0.0363	0.0308	2.40E-01	---	2.51E-01	27245	0.0425	0.0196	3.04E-02	---	7.93E-01
rs10144270	12	18773964	PLCZ1	t	g	18629	0.0507	0.0253	4.55E-02	+++++?+	9.24E-02	10313	0.0381	0.0308	2.15E-01	---	2.74E-01	28942	0.0456	0.0195	1.96E-02	---	7.52E-01
rs9764296	12	18774030	PLCZ1	t	g	18629	0.0507	0.0253	4.54E-02	+++++?+	9.22E-02	10312	0.0416	0.0308	1.76E-01	---	3.06E-01	28941	0.047	0.0195	1.61E-02	---	8.19E-01
rs10144272	12	18775683	PLCZ1	a	g	18587	0.0525	0.0253	3.82E-02	+++++?+	7.87E-02	10305	0.0386	0.0308	2.06E-01	---	2.73E-01	28892	0.0468	0.0195	1.62E-02	---	7.26E-01
rs10770395	12	18775923	PLCZ1	a	g	16640	-0.0312	0.0264	2.36E-01	-?+----	9.65E-02	10316	-0.0394	0.0304	1.94E-01	---	4.32E-01	26956	-0.0427	0.0199	8.15E-02	---	8.39E-01
rs9792408	12	18776163	PLCZ1	a	g	16371	0.0394	0.0268	1.53E-01	+++++?+	4.51E-02	10314	0.0468	0.0303	1.23E-01	---	5.01E-01	26685	0.0401	0.0201	3.60E-02	---	8.36E-01
rs10144273	12	18777098	PLCZ1	a	g	10808	0.0308	0.0336	3.59E-01	????-??	1.81E-01	10325	0.0465	0.0307	1.90E-01	---	2.34E-01	21033	0.0394	0.0227	8.25E-02	---	7.30E-01
rs7485517	12	18779723	PLCZ1	a	g	18806	-0.0901	0.0247	2.58E-04	-----	7.99E-01	10253	-0.0354	0.0302	2.42E-01	---	3.32E-01	29159	-0.0662	0.0191	3.62E-04	---	1.61E-01
rs10144275	12	18779950	PLCZ1	t	c	18722	-0.0485	0.0247	4.98E-02	-----	5.33E-02	10342	-0.0395	0.0304	1.94E-01	---	3.66E-01	29064	-0.0449	0.0192	1.91E-02	---	8.18E-01
rs12300257	12	18780416	PLCZ1, CAP2A3	c	g	18727	-0.0444	0.0248	7.37E-02	-----	4.70E-02	10360	-0.042	0.0303	1.66E-01	---	3.54E-01	29087	-0.0438	0.0192	2.36E-02	---	9.51E-01
rs10841080	12	18780814	PLCZ1, CAP2A3	a	c	16325	0.0454	0.0272	9.47E-02	+++++?+	4.90E-02	10302	0.0418	0.0305	1.71E-01	---	2.61E-01	26287	0.0434	0.0203	3.09E-02	---	9.30E-01
rs12306752	12	18780832	PLCZ1, CAP2A3	t	c	18715	-0.0489	0.0248	4.88E-02	-----	5.35E-02	10343	-0.0409	0.0304	1.79E-01	---	3.24E-01	29058	-0.0457	0.0192	1.74E-02	---	8.38E-01
rs1075421	12	18782346	PLCZ1, CAP2A3	t	c	18511	-0.0499	0.0253	4.82E-02	-----	5.60E-02	10293	-0.0429	0.0305	1.59E-01	---	2.58E-01	28804	-0.047	0.0195	1.57E-02	---	8.60E-01
rs7131551	12	18786111		a	g	8443	0.0067	0.0379	8.60E-01	????-??	4.79E-02	10103	0.0363	0.0308	2.39E-01	---	4.41E-01	18946	0.0245	0.0239	3.05E-01	++	5.44E-01
rs16914583	12	18787049		c	g	14108	-0.0295	0.0297	3.20E-01	-?+----	7.76E-02	10134	-0.0367	0.0308	2.34E-01	---	3.97E-01	24422	-0.033	0.0214	1.23E-01	---	8.66E-01
rs10770397	12	18789552	PLCZ1	a	g	10374	0.0005	0.0331	9.88E-01	+++++?+	1.12E-01	7855	0.0147	0.0348	6.73E-01	-?	8.61E-01	18169	0.0077	0.024	7.83E-01	++	7.68E-01
rs10144279	12	18793297	PLCZ1	a	g	8843	0.0094	0.0378	8.25E-01	????-??	1.98E-02	10356	0.0344	0.0307	1.89E-01	---	6.28E-02	25247	-0.0351	0.0256	6.65E-02	---	8.60E-04
rs10841134	12	19831174		a	g	14371	-0.0871	0.0289	2.57E-03	-?+----	2.13E-01	10353	-0.0396	0.0308	1.98E-01	---	1.26E-01	25124	-0.0778	0.0221	1.88E-01	---	2.70E-03
rs7298390	12	19832243		c	g	16804	0.0807	0.0267	2.56E-03	+++++?+	2.77E-01	10350	-0.0409	0.0308	1.84E-01	---	1.42E-01	27154	-0.0285	0.0202	1.57E-01	---	2.85E-03
rs9762547	12	19834638		a	g	14667	0.0864	0.0289	2.80E-03	+++++?+	2.81E-01	10347	-0.035	0.0308	2.55E-01	---	9.45E-02	25014	0.0296	0.0211	1.61E-01	---	4.05E-03
rs9762553	12	19834650		t	g	16810	0.0801	0.0267	2.74E-03	+++++?+	2.87E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27173	0.0287	0.0202	1.56E-01	---	3.27E-03
rs9776559	12	19834849		t	c	16810	-0.08	0.0267	2.75E-03	+++++?+	2.87E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27173	-0.0286	0.0202	1.56E-01	---	3.29E-03
rs10144855	12	19835818		a	g	16808	-0.03	0.0267	2.61E-01	-?+----	1.89E-02	10363	0.0392	0.0308	2.02E-01	---	1.21E-01	27171	-0.0063	0.0202	9.88E-01	---	8.96E-02
rs10144856	12	19835914		a	g	14398	-0.1069	0.0296	3.03E-04	-----	7.78E-01	10343	0.0405	0.0308	1.88E-01	---	1.38E-01	24741	-0.0301	0.0213	9.05E-02	---	5.59E-04
rs10144857	12	19836199		t	c	16811	0.0804	0.0267	2.63E-03	+++++?+	2.81E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27174	0.0288	0.0202	1.53E-01	---	3.19E-03
rs1044858	12	19836311		a	g	16811	-0.0804	0.0267	2.62E-03	+++++?+	2.81E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27174	-0.0288	0.0202	1.53E-01	---	3.19E-03
rs2731621	12	19836487		a	g	16640	-0.0735	0.0271	3.35E-04	-----	4.32E-01	10348	-0.0386	0.0308	2.10E-01	---	1.03E-01	26888	-0.0408	0.0202	1.69E-01	---	3.19E-03
rs10841360	12	19836833		a	g	16811	-0.0804	0.0267	2.64E-03	+++++?+	2.81E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27174	-0.0288	0.0202	1.53E-01	---	3.19E-03
rs1527262	12	19837209		a	c	16810	-0.0804	0.0267	2.63E-03	+++++?+	2.81E-01	10363	-0.0398	0.0308	1.96E-01	---	1.28E-01	27175	-0.0288	0.0202	1.53E-01	---	3.19E-03
rs10841362	12	19838993		t	g	16433	-0.1042	0.0273	1.35E-04	-----	8.46E-01	10316	0.0394	0.0308	2.01E-01	---	1.22E-01	26749	-0.0401	0.0204	4.47E-02	---	4.85E-04
rs10505848	12	19839175		a	g	18813	-0.0828	0.0251	9.54E-04	-----	3.84E-01	10362	0.0383	0.0308	2.13E-01	---	1.36E-01	29175	-0.0345	0.0195	7.65E-02	---	2.30E-03
rs1465089	12	19840118		a	g	18800	0.082	0.0251	1.09E-03	-----	4.34E-01	10346	-0.0445	0.0307	1.46E-01	---	9.55E-02	29146	-0.0313	0.0194	1.07E-01	---	1.42E-03
rs9758669	12	19841180		a	g	10308	-0.1223	0.0359	6.51E-04	????-??	9.35E-01	10238	0.0399	0.0308	1.96E-01	---	1.66E-01	20546	-0.0289	0.0234	2.17E-01	---	6.06E-04
rs9737387	12	19841195		a	t	18710	0.0824	0.0251	1.05E-03	+++++?+	3.89E-01	10343	-0.0414	0.0308	1.79E-01	---	1.22E-01	29053	-0.033	0.0195	8.99E-02	---	1.83E-03
rs2694973	12	19843747		t	c	14846	-0.0959	0.0275	4.90E-04	-?+----	1.99E-01	10322	0.0485	0.0309	1.16E-01	---	8.95E-02	25168	-0.0321	0.0205	1.10E-01	---	4.81E-04
rs9736176	12	19844460		a	g	14829	0.0671	0.0279	1.62E-02	+++++?+	3.60E-01	10323	0.0395	0.0308	2.00E-01	---	4.77E-02	25152	0.0191	0.0207	4.57E-01	++	1.03E-02
rs1094728	12	19845487		a	g	14891	-0.0957	0.0278	5.89E-04	-?+----	1.49E-01	10356	0.0404	0.0307	1.89E-01	---	6.28E-02	25247	-0.0351	0.0256	6.65E-02	---	8.60E-04
rs10144913	12	19911210																					

	rs17674423	66439715	t	c	16693	0.0902	0.0241	1.76E-04	++*+*	3.98E-01	10351	-0.03	0.0288	2.97E-01	++	2.25E-01	27044	0.0407	0.0185	2.77E-02	++	1.37E-03
	rs12370196	66440777	a	g	18738	0.0884	0.0226	9.51E-05	+++++	4.95E-01	10335	-0.0291	0.0288	3.12E-01	++	2.20E-01	29073	0.0436	0.0178	1.42E-02	++	1.33E-03
	rs4913255	66443318	t	c	18784	-0.0874	0.0226	1.11E-04	-----	4.85E-01	10330	0.0283	0.0288	3.26E-01	++	2.15E-01	29114	0.0430	0.0178	1.49E-02	++	1.58E-03
	rs7957455	66443673	t	c	16441	0.0842	0.0243	5.16E-04	+++++?	3.79E-01	10329	-0.0291	0.0288	3.12E-01	++	2.13E-01	26770	0.0371	0.0186	4.59E-02	++	2.64E-03
	rs10402207	66443857	t	c	18647	-0.0865	0.0227	1.41E-04	-----	3.77E-01	10328	0.0151	0.0291	6.05E-01	++	2.20E-01	28985	0.0481	0.0179	7.25E-03	++	5.51E-03
	rs2014130	66454320	a	g	18693	-0.0474	0.0227	3.70E-02	-----	1.16E-02	10361	0.0127	0.0291	6.61E-01	++	1.75E-01	29054	0.0407	0.0179	1.68E-02	++	1.03E-01
	rs10492026	66454597	a	g	18672	0.0869	0.0227	1.31E-04	+++++	3.54E-01	10360	-0.0111	0.0291	7.05E-01	++	1.76E-01	29032	0.0499	0.0179	5.34E-03	++	7.99E-03
	rs1176866	66458793	a	t	16630	-0.1061	0.0244	1.32E-05	-----?	8.00E-01	10367	0.0184	0.0291	5.27E-01	++	2.00E-01	26927	0.0547	0.0187	3.44E-02	++	1.04E-03
231	rs1176917	66553869	a	c	14691	-0.0348	0.0287	2.26E-01	?-+*	7.51E-03	9975	0.0037	0.0322	9.09E-01	?-	3.35E-01	24666	0.0178	0.0214	4.07E-01	++	3.72E-01
	rs1905446	66559766	c	c	14679	0.1066	0.0283	1.63E-04	+++++	6.44E-01	10270	0.0004	0.0316	8.89E-01	++	5.91E-01	24949	0.0593	0.0211	4.89E-02	++	1.23E-02
	rs10878683	66560646	c	g	14776	0.0985	0.0283	5.15E-04	+++++	5.66E-01	10323	0.0071	0.0321	8.26E-01	++	4.95E-01	25099	0.0585	0.0212	5.83E-03	++	3.27E-02
	rs4103248	66561584	a	g	14830	-0.0949	0.0283	7.99E-04	-+*	5.60E-01	10362	0.0062	0.032	8.47E-01	-	5.17E-01	25192	0.0586	0.0212	8.88E-02	++	3.79E-02
	rs10878684	66561761	a	t	14823	0.0301	0.0283	2.87E-01	+++++	4.78E-02	10363	0.0066	0.032	8.37E-01	++	5.02E-01	25186	0.0198	0.0212	3.51E-01	++	5.82E-01
	rs1796311	66689423	a	g	14642	-0.069	0.052	1.85E-01	-+*-?	1.65E-02	10359	-0.0801	0.053	1.30E-01	++	1.02E-01	26821	0.0144	0.0371	4.49E-02	++	8.81E-01
	rs17963209	66703853	a	g	18395	-0.1003	0.0509	4.85E-02	-----	6.62E-02	10259	-0.121	0.0613	4.83E-02	++	3.21E-01	28844	-0.1087	0.0392	5.49E-03	++	9.95E-01
	rs2250248	66702637	a	g	18794	-0.0813	0.0472	8.49E-02	-----	6.36E-01	10359	-0.0801	0.053	1.30E-01	++	1.02E-01	29153	0.0808	0.0352	2.19E-02	++	9.87E-01
	rs1675575	66702905	t	c	18812	0.0801	0.0471	8.86E-02	+++++	6.73E-01	10361	0.1108	0.0571	5.21E-02	++	5.94E-02	29173	0.0925	0.0363	1.09E-02	++	6.78E-01
	rs1614694	66703012	a	g	18812	-0.0801	0.0471	8.86E-02	-----	6.74E-01	10361	-0.0811	0.053	1.26E-01	++	1.02E-01	29173	-0.0805	0.0352	2.22E-02	++	9.89E-01
	rs1796325	66703620	a	g	18812	-0.0801	0.0471	8.86E-02	-----	6.74E-01	10361	-0.0811	0.053	1.26E-01	++	1.02E-01	29173	-0.0805	0.0352	2.22E-02	++	9.89E-01
	rs1675576	66703708	c	g	18814	-0.0803	0.0471	8.80E-02	-----	6.71E-01	10361	-0.0811	0.053	1.26E-01	++	1.02E-01	29175	-0.0807	0.0352	2.20E-02	++	9.91E-01
	rs1675577	66703721	a	g	16469	0.0612	0.052	2.39E-01	+++++?	6.79E-01	10361	0.1109	0.057	5.16E-02	++	5.94E-02	26830	0.0838	0.0384	2.92E-02	++	5.20E-01
	rs2708024	66703778	a	g	18812	-0.0801	0.0471	8.86E-02	-----	6.74E-01	10361	-0.1122	0.0571	4.93E-02	++	5.79E-02	29173	-0.0931	0.0363	1.04E-02	++	6.65E-01
	rs2251464	66703903	a	t	18814	-0.0794	0.0471	9.15E-02	-----	6.80E-01	10361	-0.0804	0.053	1.29E-01	++	1.03E-01	29175	-0.0798	0.0352	2.33E-02	++	9.89E-01
	rs1796266	66704187	t	g	14777	-0.0618	0.0519	2.34E-01	-+*-?	6.50E-01	10361	-0.1143	0.057	4.50E-02	++	5.57E-02	26838	-0.0856	0.0384	2.57E-02	++	4.96E-01
	rs1658806	66704242	c	g	18812	-0.0804	0.0472	8.49E-02	-----	6.59E-01	10359	-0.0811	0.053	1.35E-01	++	1.05E-01	29129	-0.0813	0.0352	2.15E-02	++	9.71E-01
	rs1656807	66704513	c	g	14997	-0.0581	0.0501	2.46E-01	-+*-?	7.70E-01	10361	0.0811	0.057	1.26E-01	++	1.02E-01	25358	0.069	0.0362	5.82E-02	++	7.53E-01
	rs1675578	66704854	c	g	18812	-0.0826	0.0471	7.93E-02	-----	6.53E-01	10361	-0.0789	0.053	1.36E-01	++	1.05E-01	29173	-0.081	0.0352	2.15E-02	++	9.58E-01
	rs1284717	66707309	t	c	18812	-0.0833	0.0471	7.66E-02	-----	6.51E-01	10361	-0.1156	0.057	4.26E-02	++	5.42E-02	29173	-0.0964	0.0363	7.93E-03	++	6.62E-01
	rs1284715	66708061	a	g	18812	-0.0843	0.047	7.29E-02	-----	6.46E-01	10361	-0.1123	0.057	4.89E-02	++	5.79E-02	29174	-0.0956	0.0363	8.36E-03	++	7.05E-01
	rs2253566	66710963	a	c	18813	0.0842	0.0469	7.26E-02	+++++	6.48E-01	10361	0.1073	0.0568	5.91E-02	++	6.38E-02	29174	0.0936	0.0362	9.68E-03	++	7.54E-01
	rs1796305	66711605	a	g	18813	-0.0844	0.0469	7.22E-02	-----	6.48E-01	10361	-0.077	0.0528	1.45E-01	++	1.08E-01	29174	-0.0811	0.0351	2.07E-02	++	9.17E-01
	rs1796304	66711814	t	c	16477	0.0648	0.0518	2.11E-01	+++++?	6.34E-01	10361	0.1065	0.0568	6.12E-02	++	6.46E-02	26838	0.0837	0.0383	2.89E-02	++	5.88E-01
	rs19304445	66712574	t	c	18813	-0.0844	0.0469	7.20E-02	-----	6.49E-01	10361	-0.0783	0.0529	1.39E-01	++	1.06E-01	29174	-0.0817	0.0351	1.99E-02	++	9.30E-01
	rs2253279	66713117	t	c	18813	0.0844	0.0469	7.20E-02	+++++	6.49E-01	10361	0.0755	0.0528	1.53E-01	++	1.10E-01	29174	0.0805	0.0351	2.17E-02	++	9.00E-01
	rs2252375	66713244	t	c	18813	0.0857	0.0469	6.78E-02	+++++	6.59E-01	10361	0.0783	0.0529	1.39E-01	++	1.06E-01	29174	0.0834	0.0351	1.89E-02	++	8.17E-01
	rs2254885	66713750	a	g	18813	-0.0855	0.0469	6.84E-02	-----	6.37E-01	10361	-0.0783	0.0529	1.39E-01	++	1.06E-01	29174	-0.0823	0.0351	1.90E-02	++	9.19E-01
	rs1461284	66714037	a	g	18781	0.084	0.0471	7.49E-02	+++++	6.41E-01	10361	0.1073	0.0568	5.91E-02	++	6.38E-02	29142	0.0935	0.0363	9.92E-03	++	7.52E-01
	rs1922751	66716029	a	t	18813	-0.0857	0.0469	6.80E-02	-----	6.47E-01	10358	-0.076	0.0531	1.52E-01	++	1.09E-01	29171	-0.0814	0.0352	2.05E-02	++	8.91E-01
	rs2261431	66716492	a	g	18813	-0.0842	0.0469	7.27E-02	-----	6.42E-01	10361	-0.1073	0.0568	5.91E-02	++	6.38E-02	29174	-0.0936	0.0362	9.68E-03	++	7.54E-01
	rs2261433	66729643	a	g	16337	-0.0392	0.0516	4.48E-01	-+*-?	5.65E-01	10309	-0.0717	0.0576	2.13E-01	++	5.88E-02	26646	-0.0537	0.0384	1.63E-01	++	6.74E-01
	rs1796308	66731631	a	g	18808	-0.0865	0.0468	6.50E-02	-----	7.01E-01	10362	-0.074	0.0531	1.63E-01	++	1.59E-01	29170	-0.081	0.0351	2.10E-02	++	8.60E-01
	rs2708019	66732175	t	c	18811	-0.0644	0.0469	1.70E-01	-----	5.21E-01	10362	-0.0767	0.0532	1.50E-02	++	1.53E-01	29173	-0.0698	0.0352	4.73E-02	++	8.62E-01
	rs1796310	66733093	c	t	18774	-0.0867	0.0468	6.40E-02	-----	7.06E-01	10363	-0.1091	0.0563	5.26E-02	++	8.35E-02	29137	-0.0959	0.036	7.74E-03	++	7.60E-01
	rs2611437	66733650	t	c	16446	-0.0771	0.052	1.38E-01	-----?	7.40E-01	10361	-0.0709	0.052	1.82E-01	++	1.66E-01	26807	-0.0741	0.0372	4.64E-02	++	9.94E-01
	rs1658810	66733747	t	c	18770	0.1004	0.0472	3.46E-02	+++++	6.30E-01	10359	0.0747	0.0529	1.58E-01	++	1.71E-01	29129	0.089	0.0351	1.15E-02	++	9.17E-01
	rs1658811	66737247	a	g	18678	0.109	0.0473	2.11E-02	+++++	6.18E-01	10298	0.0817	0.0526	1.21E-01	++	1.85E-01	28976	0.0968	0.0352	5.92E-03	++	7.00E-01
	rs1675585	66738018	c	t	18767	-0.1101	0.0474	2.02E-02	-----	6.83E-01	10346	-0.0693	0.0533	1.94E-01	++	1.12E-01	29113	-0.0921	0.0354	9.33E-03	++	5.67E-01
	rs165																					

	rs17068014	13	77064315	SCCL	a	t	16783	0.0633	0.0365	8.30E-02	+?+?+?	9.37E-03	10344	-0.0919	0.046	4.58E-02	---	7.31E-01	27127	0.0033	0.0286	9.07E-01	++	8.22E-03
	rs17068019	13	77064319	SCCL	a	c	16783	0.0633	0.0365	8.27E-02	+?+?+?	9.36E-03	10344	-0.0919	0.046	4.58E-02	---	7.31E-01	27127	0.0033	0.0286	9.07E-01	++	8.22E-03
	rs17068033	13	77065536	SCCL	t	c	18816	0.0548	0.0345	1.12E-01	+?+?+?	1.47E-02	10341	-0.0929	0.0459	4.30E-02	---	7.46E-01	29157	0.0015	0.0276	9.57E-01	++	1.01E-02
	rs1472583	13	77092268	SCCL	t	c	16649	0.0647	0.0366	7.73E-02	+?+?+?	8.44E-03	10328	-0.092	0.0459	4.51E-02	---	8.28E-01	26977	0.0038	0.0286	8.95E-01	++	7.60E-03
	rs17068013	13	77108013	SCCL	t	c	16611	0.073	0.0367	4.68E-02	+?+?+?	4.94E-03	10324	-0.0942	0.046	4.05E-02	---	8.64E-01	26977	0.0058	0.0287	7.81E-01	++	4.49E-03
	rs17068137	13	77109029	SCCL	a	g	16723	0.1546	0.0631	1.80E-02	+?+?+?	5.51E-03	10355	-0.1026	0.0456	2.44E-02	---	9.22E-01	27078	0.0058	0.0383	4.99E-02	++	9.77E-06
	rs2274014	13	77116246	SCCL	t	g	16616	0.0665	0.0363	6.67E-02	+?+?+?	3.02E-03	10342	-0.0912	0.0454	4.47E-02	---	9.97E-01	26958	0.005	0.0284	8.60E-01	++	6.67E-03
238	rs131616380	13	79603316		t	g	14655	-0.1193	0.0296	5.70E-05	-?+?+?	8.02E-02	9927	-0.0669	0.0325	3.98E-02	?	7.09E-01	24582	-0.0955	0.0219	1.27E-05	++	2.33E-01
	rs17072370	13	79603731		a	t	14688	-0.1299	0.0292	8.49E-06	-?+?+?	2.45E-01	9971	-0.066	0.0323	4.11E-02	?	9.58E-01	24659	-0.1012	0.0217	3.01E-06	++	1.42E-01
	rs1215468	13	79605430	SPRY2	a	g	14695	0.1282	0.0289	9.54E-06	-?+?+?	2.57E-01	9974	-0.0688	0.0324	3.36E-02	?	9.90E-01	24669	-0.1019	0.0216	2.31E-06	++	1.71E-01
	rs1327315	13	79606733		t	c	16679	-0.1038	0.027	1.23E-04	-?+?+?	6.29E-02	10340	-0.0874	0.0319	6.19E-03	---	8.71E-03	27019	-0.097	0.0206	2.55E-06	++	9.92E-01
	rs1215451	13	79613894		a	g	18793	-0.0909	0.026	4.87E-04	-?+?+?	7.48E-02	10337	-0.096	0.032	2.74E-03	---	7.48E-03	29130	-0.0929	0.0202	4.12E-06	++	6.05E-01
	rs1359790	13	79615157		a	g	18719	-0.0232	0.026	3.73E-01	+?+?+?	1.04E-03	10337	-0.0952	0.032	2.95E-03	---	7.41E-03	29056	-0.0718	0.0202	1.02E-02	++	8.08E-02
239	rs9301717	13	90672066		t	c	18615	-0.0987	0.0349	4.62E-03	-?+?+?	9.87E-03	10353	-0.0456	0.0414	2.71E-01	---	8.87E-01	28968	-0.0566	0.0267	4.08E-03	++	3.27E-01
	rs9397769	13	90676691		t	c	18725	-0.0953	0.0345	5.72E-03	-?+?+?	8.36E-03	10360	-0.0456	0.0413	2.69E-01	---	9.31E-01	29085	-0.0749	0.0265	4.69E-03	++	3.56E-01
	rs7320223	13	90676691		a	g	16676	0.1219	0.0382	1.44E-03	+?+?+?	4.97E-01	10254	0.0466	0.0418	6.65E-01	---	8.70E-01	27030	0.0876	0.0282	1.89E-03	++	1.84E-01
	rs1330068	13	90683486		t	c	18680	-0.0792	0.0346	2.20E-02	-?+?+?	9.95E-01	10356	-0.0451	0.0415	2.77E-01	---	9.15E-01	29386	-0.0652	0.0266	1.14E-02	++	5.28E-01
	rs9523274	13	90684065		t	c	16683	-0.1161	0.0401	3.83E-03	-?+?+?	9.82E-01	10340	-0.0393	0.0425	3.55E-01	---	9.47E-01	27023	-0.0799	0.0292	6.14E-03	++	1.89E-01
	rs9583901	13	90684716		t	c	18767	-0.1252	0.0359	4.80E-04	-?+?+?	9.99E-01	10323	-0.0518	0.0419	2.17E-01	---	8.71E-01	29090	-0.0941	0.0273	5.55E-04	++	1.83E-01
	rs9523275	13	90686041		a	g	16737	0.1279	0.0382	8.11E-04	+?+?+?	9.96E-01	10322	0.0471	0.0419	6.26E-01	---	8.36E-01	29059	0.0912	0.0282	1.23E-03	++	1.54E-01
	rs9515895	13	90687732		t	c	18691	-0.1181	0.0378	1.76E-03	-?+?+?	9.89E-01	10329	-0.0404	0.0425	3.43E-01	---	9.52E-01	29020	-0.0838	0.0282	3.01E-03	++	1.72E-01
	rs9301719	13	90688458		a	g	18692	0.1089	0.0348	1.73E-03	+?+?+?	8.81E-01	10299	0.0478	0.0418	2.52E-01	---	8.28E-01	28991	0.0389	0.0267	1.71E-03	++	2.61E-01
	rs9523276	13	90690675		t	c	16721	0.1216	0.0383	1.49E-03	+?+?+?	9.87E-01	10338	0.0514	0.0419	2.19E-01	---	7.39E-01	27059	0.0896	0.0283	1.52E-03	++	2.16E-01
240	rs2025452	13	100964561	ITGBL1	a	g	14825	-0.0559	0.0253	2.70E-02	-?+?+?	4.71E-02	10348	0.0296	0.0284	2.98E-01	---	4.73E-01	25173	-0.0181	0.0189	3.39E-01	++	2.46E-02
	rs1414313	13	100965284	ITGBL1	a	g	13933	-0.0469	0.026	2.79E-02	-?+?+?	2.96E-02	10325	0.034	0.0284	3.31E-01	---	5.85E-01	25157	-0.0361	0.0192	6.02E-02	++	3.56E-02
	rs9549792	13	100969172	ITGBL1	a	g	14743	0.0893	0.0256	4.71E-04	+?+?+?	9.09E-01	10321	0.0285	0.0285	9.71E-01	---	9.09E-01	25080	0.0602	0.019	1.58E-03	++	8.88E-02
	rs12586000	13	100970426	ITGBL1	t	c	14738	0.053	0.0255	3.77E-02	+?+?+?	4.48E-01	10313	-0.0348	0.0284	2.20E-01	---	4.66E-01	25051	0.038	0.019	4.67E-01	++	2.14E-02
	rs12586014	13	100970657	ITGBL1	t	g	14780	0.0615	0.0254	1.56E-02	+?+?+?	3.22E-01	10343	0.0281	0.0284	3.23E-01	---	4.58E-01	25123	0.0467	0.0189	1.37E-02	++	3.81E-01
	rs9300677	13	100981929	ITGBL1	a	g	14826	0.0475	0.0254	6.17E-02	+?+?+?	3.61E-01	10272	-0.0348	0.0285	2.22E-01	---	7.28E-01	25098	-0.0111	0.019	5.99E-01	++	3.11E-02
	rs954795	13	100985996	ITGBL1	a	g	18702	-0.0484	0.0227	3.32E-02	-?+?+?	2.18E-01	10290	0.0323	0.0285	2.57E-01	---	7.22E-01	28992	-0.0717	0.0178	3.36E-02	++	2.68E-02
	rs1414312	13	100995842	ITGBL1	a	g	18550	-0.0595	0.0229	9.33E-03	-?+?+?	2.57E-01	10312	0.0337	0.0285	2.38E-01	---	5.76E-01	28862	-0.0229	0.0179	1.99E-01	++	1.08E-02
	rs885304	13	101002835	ITGBL1	t	c	16491	0.0686	0.0243	4.78E-03	+?+?+?	8.09E-01	10338	-0.0281	0.0284	3.23E-01	---	8.81E-01	26829	-0.0277	0.0183	1.33E-02	++	9.68E-03
	rs13069448	13	101019787	ITGBL1	t	c	18774	0.0522	0.0228	2.19E-02	+?+?+?	4.35E-01	10353	0.0215	0.0284	4.49E-01	---	4.04E-01	29127	-0.0402	0.0178	2.39E-02	++	3.99E-01
	rs9557704	13	101025496	ITGBL1	a	t	18737	-0.0528	0.0228	2.03E-02	-?+?+?	4.68E-01	10326	-0.0663	0.0284	8.24E-01	---	4.01E-01	28977	-0.0437	0.0182	5.18E-02	++	2.02E-01
	rs10139195	13	101032322	ITGBL1	t	g	12711	-0.1618	0.035	2.51E-02	-?+?+?	8.31E-01	10269	-0.0712	0.0259	7.71E-01	---	5.17E-01	22880	-0.0301	0.0178	4.09E-02	++	2.18E-02
241	rs2213246	13	21873203		t	c	18680	-0.1515	0.043	4.18E-04	-?+?+?	7.63E-01	10356	0.0229	0.0572	6.89E-01	---	8.06E-01	29036	-0.0885	0.0344	1.00E-02	++	1.48E-02
	rs2213247	13	21873224		a	g	16651	-0.1778	0.0483	2.34E-04	-?+?+?	8.40E-01	10356	0.0229	0.0572	6.89E-01	---	8.06E-01	27007	-0.0943	0.0369	1.06E-02	++	7.34E-03
	rs981423	13	21878594		a	g	18699	-0.1534	0.0427	3.31E-04	-?+?+?	8.32E-01	10359	0.0263	0.0571	6.45E-01	---	8.57E-01	29058	-0.0889	0.0342	9.29E-03	++	1.17E-02
	rs8014927	13	21908470		a	g	18710	0.1099	0.0428	1.01E-02	+?+?+?	1.44E-01	10350	-0.0356	0.0575	5.36E-01	---	5.60E-01	29060	-0.0589	0.0343	9.10E-02	++	4.24E-02
	rs2051536	13	21911637		t	c	18704	-0.1534	0.0426	3.23E-04	-?+?+?	8.13E-01	10357	0.0205	0.0571	7.20E-01	---	8.24E-01	29061	-0.0912	0.0341	7.55E-03	++	1.47E-02
	rs2877454	13	21919601		a	c	18683	-0.1451	0.0426	6.64E-04	-?+?+?	7.45E-01	10355	0.0208	0.0571	7.15E-01	---	8.52E-01	29038	-0.0858	0.0341	1.20E-02	++	1.99E-02
	rs2204983	13	21919803		t	c	14815	-0.1482	0.0479	1.96E-03	-?+?+?	6.47E-01	10336	0.0147	0.0555	7.91E-01	---	6.78E-01	25151	-0.0787	0.0363	3.01E-02	++	2.63E-02
	rs1243473	13	21926789		t	c	16316	0.1659	0.0451	2.31E-04	+?+?+?	9.03E-01	10352	-0.0231	0.0577	6.89E-01	---	7.91E-01	26688	-0.0492	0.0355	8.01E-03	++	9.68E-03
	rs9826224	13	21927364		a	g	14778	-0.1441	0.0448	1.28E-03	-?+?+?	8.32E-02	2852	0.1197	0.117	2.63E-01	---	9.71E-01	27630	-0.1048	0.0413	1.13E-02	++	2.30E-02
	rs13811244	13	21929522</																					

	rs4775248	-0.0526	0.0229	2.16E-04	-----	2.45E-02	10342	-0.0086	0.0412	8.35E-01	++	7.10E-04	29026	-0.0422	0.02	3.49E-02	--	3.51E-01
	rs4775253	0.0525	0.0229	2.16E-04	+++++	3.10E-02	10335	-0.0044	0.0392	9.11E-01	++	7.30E-04	29078	0.038	0.0198	5.45E-02	++	2.10E-01
	rs4775256	0.0525	0.0229	2.16E-04	+++++	3.09E-02	10335	-0.0207	0.0387	5.94E-01	++	1.66E-04	29078	0.0335	0.0197	8.90E-02	++	1.04E-01
	rs21258529	0.0606	0.0229	8.04E-03	+++++	4.59E-02	10347	0.0189	0.0389	6.29E-01	++	2.90E-02	29110	-0.0402	0.0197	4.18E-02	++	7.86E-02
	rs21258715	0.0601	0.0229	8.04E-03	+++++	4.66E-02	10347	-0.0001	0.0386	9.99E-01	++	2.04E-03	29110	0.0440	0.0197	2.41E-02	++	1.80E-01
	rs8036882	0.0779	0.0246	1.51E-03	+++++	3.45E-01	10335	-0.0275	0.0376	4.65E-01	++	1.27E-04	26744	-0.0463	0.0206	2.45E-02	++	1.90E-02
	rs8037063	-0.0601	0.0229	8.64E-03	-----	4.70E-02	10347	0.0193	0.0379	6.10E-01	++	4.17E-04	29110	-0.0389	0.0196	4.74E-02	++	7.30E-02
	rs8038522	-0.0883	0.0246	3.26E-04	-----	6.35E-01	10347	0.0245	0.0376	5.14E-01	++	2.70E-04	26775	-0.0545	0.0206	8.12E-03	++	1.21E-02
	rs1716125	-0.0578	0.0228	1.14E-02	-----	6.91E-02	10352	-0.0006	0.029	9.83E-01	++	1.31E-02	29148	-0.0359	0.0179	4.49E-02	++	1.21E-01
	rs2470655	-0.0604	0.023	8.49E-03	-----	5.98E-02	10338	0.024	0.0368	5.14E-01	++	8.08E-04	29018	-0.0367	0.0195	5.99E-02	++	5.18E-02
	rs11070505	-0.0592	0.0229	9.57E-03	-----	5.89E-02	10346	0.012	0.0326	7.13E-01	++	3.31E-03	29119	-0.0357	0.0187	5.69E-02	++	7.39E-02
	rs355396	0.0582	0.0229	1.09E-02	+++++	7.22E-02	10349	0.0032	0.029	9.38E-01	++	6.04E-03	29161	0.0371	0.018	3.91E-02	++	1.37E-01
	rs355399	-0.0138	0.0415	7.39E-01	?????	4.82E-02	9794	-0.0184	0.0293	5.13E-01	??	1.41E-02	16641	-0.0168	0.0243	4.88E-01	??	9.28E-01
253	rs492365	0.0573	0.0244	1.31E-04	+++++	1.87E-01	10349	0.012	0.0353	1.09E-01	??	5.73E-01	22706	-0.046	0.0203	3.10E-03	??	7.95E-01
	rs575231	0.0867	0.0244	3.17E-04	+++++	5.00E-01	7850	0.0609	0.0334	6.84E-02	??	4.13E-01	24473	-0.0777	0.0197	7.99E-05	??	5.33E-01
	rs9920076	-0.0815	0.0256	1.43E-03	-----	3.09E-01	7863	-0.0555	0.0331	9.35E-02	??	3.71E-01	22509	-0.0718	0.0203	3.94E-04	??	5.34E-01
	rs9920246	0.1012	0.0278	2.67E-04	+++++	6.04E-01	7865	0.054	0.0331	1.03E-01	??	4.12E-01	20577	0.0817	0.0213	1.25E-04	??	2.75E-01
	rs17171318	0.0822	0.0257	1.38E-03	+++++	3.88E-01	7861	0.0555	0.0331	9.34E-02	??	3.73E-01	22510	0.0722	0.0203	3.79E-04	??	5.24E-01
	rs1718379	-0.0816	0.0256	1.43E-03	-----	3.63E-01	10181	-0.0529	0.029	6.83E-02	??	7.10E-01	24855	-0.0609	0.0192	3.22E-04	??	4.58E-01
	rs10851589	-0.0828	0.0255	1.16E-03	-----	3.53E-01	10274	-0.0521	0.0289	7.12E-02	??	8.08E-01	25047	-0.0694	0.0191	2.86E-04	??	4.26E-01
	rs4774800	-0.0762	0.0255	2.81E-03	-----	2.95E-01	10333	-0.0495	0.0288	8.55E-02	??	8.04E-01	25163	-0.0645	0.0191	7.34E-04	??	4.88E-01
	rs12438177	0.05	0.0282	7.69E-02	+++++	2.92E-01	10333	0.0547	0.0288	5.76E-02	??	8.20E-01	25141	0.0623	0.0201	9.44E-03	??	9.07E-01
254	rs24320402	-0.1009	0.0278	2.75E-04	-----	3.88E-01	7870	-0.0091	0.0334	7.85E-01	??	9.03E-01	22696	-0.0633	0.0214	3.04E-03	??	3.46E-02
255	rs4775203	-0.0941	0.0251	7.93E-02	-----	2.90E-02	10349	0.012	0.0351	6.91E-01	++	6.47E-01	26852	-0.0211	0.0193	2.24E-04	++	1.52E-01
	rs17601517	-0.0991	0.025	1.18E-01	-----	2.16E-02	9971	0.0274	0.0306	6.64E-01	??	6.51E-01	26561	-0.0145	0.0194	5.45E-01	??	1.20E-01
	rs17162645	0.0448	0.0251	7.40E-02	+++++	2.21E-02	10353	-0.0119	0.0301	6.92E-01	++	8.08E-01	26954	-0.0215	0.0193	2.64E-01	++	1.48E-01
	rs6494140	-0.0789	0.0287	6.00E-03	-----	2.06E-01	7868	0.0196	0.0353	5.79E-01	??	9.44E-01	20594	-0.0397	0.0223	7.46E-02	??	3.04E-02
	rs12439645	0.0407	0.0235	8.32E-02	+++++	3.67E-02	10354	-0.0121	0.0301	6.87E-01	++	8.25E-01	28984	-0.0207	0.0185	2.64E-01	++	1.67E-01
	rs12439707	-0.0922	0.0261	4.07E-04	-----	6.23E-01	10295	0.01	0.0303	7.41E-01	++	5.75E-01	25107	-0.0487	0.0198	1.39E-02	++	1.06E-02
	rs8026404	-0.0443	0.0235	6.46E-02	-----	3.45E-02	10355	0.0339	0.0301	6.45E-01	++	7.92E-01	29008	-0.0216	0.0185	2.43E-01	++	1.34E-01
256	rs2241492	-0.1074	0.026	3.53E-05	-----	4.47E-01	10270	-0.0445	0.032	1.65E-01	??	8.25E-01	28848	-0.0824	0.0202	4.45E-05	??	1.27E-01
	rs8037603	-0.0913	0.0254	3.34E-04	-----	4.44E-01	10173	-0.0321	0.0286	2.62E-01	??	1.81E-01	24892	-0.0652	0.019	5.97E-04	??	1.22E-01
	rs17364010	-0.0971	0.0253	1.24E-04	-----	3.52E-01	10292	-0.0343	0.0284	2.28E-01	??	1.23E-01	25080	-0.0693	0.0189	2.43E-04	??	1.87E-01
	rs17618164	-0.0937	0.0253	2.07E-04	-----	4.61E-01	10281	-0.0353	0.0284	2.42E-01	??	1.43E-01	25111	-0.0617	0.0189	3.92E-04	??	1.76E-01
	rs17163757	-0.1024	0.0231	1.43E-03	-----	2.47E-01	10267	-0.0721	0.0294	1.43E-02	??	1.17E-01	20869	-0.0859	0.0217	7.40E-05	??	4.86E-01
	rs6494307	0.0804	0.024	7.92E-04	+++++	3.65E-01	10343	0.0931	0.0294	1.56E-03	??	3.05E-01	28885	0.0585	0.0186	2.72E-06	??	7.38E-01
	rs17167878	-0.0485	0.0436	2.66E-01	?????	2.75E-01	9769	-0.1023	0.0305	8.04E-04	??	3.09E-02	16283	-0.0846	0.025	7.09E-04	??	3.12E-01
	rs17172432	0.0895	0.0236	1.45E-04	+++++	5.31E-01	10299	0.1189	0.0288	3.62E-05	??	7.62E-02	28949	0.1013	0.0183	2.86E-04	??	4.30E-01
	rs17173964	-0.079	0.0256	2.00E-03	-----	5.24E-01	10256	-0.121	0.0298	4.99E-05	??	5.31E-02	26532	-0.0668	0.0194	6.14E-07	??	2.85E-01
	rs920178	-0.0677	0.0269	1.17E-02	-----	3.44E-01	10361	-0.1151	0.0333	5.53E-04	??	1.71E-01	25168	-0.0864	0.0209	3.63E-05	??	2.68E-01
	rs1436953	-0.0683	0.0268	1.09E-02	-----	3.42E-01	10355	-0.1261	0.029	1.37E-05	??	9.06E-02	25186	-0.0949	0.0197	1.42E-06	??	1.43E-01
	rs4774658	0.0364	0.0269	1.77E-01	+++++	9.45E-02	10349	0.1133	0.0334	7.00E-04	??	1.47E-01	25114	0.0667	0.0201	1.46E-03	??	7.30E-02
	rs6494309	-0.0564	0.03	6.00E-02	-----	4.27E-01	10314	-0.1142	0.035	6.48E-04	??	1.56E-01	22744	-0.0821	0.0223	2.38E-04	??	1.79E-01
	rs6494310	-0.0533	0.0301	1.68E-02	-----	4.13E-01	10314	-0.1142	0.035	6.48E-04	??	1.56E-01	22744	-0.0821	0.0224	2.38E-04	??	1.79E-01
	rs17205526	-0.0789	0.0235	1.55E-02	-----	3.98E-01	10210	-0.1097	0.0337	1.15E-03	??	9.67E-02	20861	-0.0938	0.0234	6.28E-05	??	5.11E-01
257	rs17166293	-0.1127	0.0314	3.34E-04	-----	5.05E-01	10357	-0.0297	0.0383	4.39E-01	++	5.62E-01	29106	-0.0793	0.0243	1.09E-03	++	9.38E-02
	rs11637488	-0.1131	0.0314	1.16E-04	-----	5.06E-01	10357	-0.0297	0.0383	4.39E-01	++	5.62E-01	29106	-0.0793	0.0243	1.05E-03	++	9.22E-02
	rs12913233	0.0842	0.0234	3.25E-04	+++++	5.90E-01	10359	0.0657	0.0289	2.30E-02	++	7.87E-01	29156	0.0769	0.0182	2.37E-05	++	6.19E-01
	rs17388406	-0.103	0.0351	3.36E-03	-----	7.69E-01	10363	-0.0277	0.0417	5.06E-01	++	5.85E-01	29175	-0.0718	0.0269	7.52E-03	++	1.67E-01
	rs11638043	-0.1047	0.0351	2.88E-03	-----	7.51E-01	10362	-0.027	0.0417	5.17E-01	++	5.91E-01	29174	-0.0725	0.0269	6.95E-03	++	1.54E-01
	rs11633318	-0.1218	0.0312	9.50E-05	-----	5.14E-01	10361	-0.027	0.0383	4.81E-01	++	5.42E-01	29167	-0.0824	0.0242	5.17E-03	++	5.05E-02
	rs11636554	0.087	0.0234	2.00E-04	+++++	6.04E-01	10339	0.0614	0.0289	3.36E-02	++	7.31E-01	29142	0.0769	0.0182	2.37E-05	++	4.91E-01
	rs10519164	-0.1002	0.0352	4.42E-03	-----	3.89E-01	10362	-0.0279	0.0417	5.02E-01	++	5.91E-01	29168	-0.0701	0.0269	9.14E-03	++	1.85E-01
	rs1544737	0.0822	0.0234	2.10E-04	+++++	5.36E-01	10363	0.0657	0.0289	2.31E-02	++	7.73E-01	29179	0.0757	0.0182	3.13E-05	++	6.57E-01
	rs17162362	0.119	0.0313	1.45E-04	+++++	5.74E-01	10357	0.0295	0.0383	4.41E-01	++	5.35E-01	29106	0.0832	0.0242	6.01E-04		

rs10514454	16	77966519	a	g	18735	0.1039	0.0293	4.00E-04	+++++	5.67E-01	9922	0.0199	0.0355	4.30E-02	7++	4.03E-01	28657	0.0909	0.0226	5.72E-05	++	4.87E-01		
rs16950314	16	77966683	t	c	14717	0.1192	0.0326	2.57E-04	+++++	5.62E-01	10274	0.0711	0.0341	3.74E-02	+++	5.88E-01	24991	0.0967	0.0236	4.43E-05	++	3.08E-01		
rs17724843	16	77968611	c	g	18643	0.1133	0.0303	1.84E-04	+++++	5.84E-01	9974	0.0568	0.0358	1.13E-01	7++	4.06E-01	28617	0.0892	0.0231	1.05E-04	++	2.28E-01		
rs13353724	16	77940412	c	g	14751	0.1143	0.0327	4.73E-04	+++++	6.23E-01	10280	0.0734	0.0342	3.20E-02	+++	5.99E-01	25031	0.0948	0.0236	6.08E-05	++	3.87E-01		
rs13346466	16	77966519	a	t	12452	0.1239	0.0366	7.03E-04	+++++	5.84E-01	10276	0.0878	0.0346	1.12E-02	+++	5.82E-01	27268	0.1048	0.0251	3.05E-05	++	4.74E-01		
rs4889555	16	77955373	c	t	14755	0.057	0.0326	8.01E-02	-7-+-	5.06E-02	10227	0.0647	0.034	5.66E-02	+++	6.96E-01	25052	0.0607	0.0235	9.91E-03	++	8.70E-01		
rs17724884	16	77946971	t	c	18536	0.116	0.03	1.09E-04	+++++	6.89E-01	10274	0.0802	0.0349	2.16E-02	+++	7.37E-01	28810	0.1008	0.0228	9.41E-06	++	4.37E-01		
rs4606729	16	77959067	c	g	12668	0.1239	0.0346	3.41E-04	+++++	5.91E-01	10236	0.0107	0.0303	7.25E-01	+++	9.14E-01	22904	0.0988	0.0228	8.67E-03	++	1.38E-02		
rs4128452	16	77959638	t	c	12638	0.1268	0.0351	2.98E-04	+++++	6.71E-01	10159	-0.0175	0.0301	5.61E-01	---	7.69E-01	22797	0.0436	0.0228	5.61E-02	++	1.80E-03		
rs17797882	16	77964419	t	c	18771	-0.1113	0.0297	1.76E-04	+++++	3.72E-01	10355	-0.0818	0.0326	1.21E-02	---	6.09E-01	29126	-0.0979	0.022	8.19E-06	++	5.04E-01		
rs3812994	16	77966170	a	t	18611	-0.101	0.0297	6.86E-04	+++++	4.80E-01	10243	-0.0912	0.034	7.42E-03	---	4.91E-01	28854	-0.0968	0.0224	1.52E-05	++	8.28E-01		
rs3812993	16	77966327	c	g	16550	-0.104	0.0301	5.58E-04	+++++	-7-+-	2.42E-01	10259	-0.0827	0.0343	1.58E-02	---	5.51E-01	26809	-0.0947	0.0226	2.82E-05	++	6.41E-01	
rs4390610	16	77971170	t	c	18728	-0.0732	0.0298	1.40E-02	+++++	1.17E-01	10318	-0.0745	0.0347	3.17E-02	---	4.53E-01	29046	-0.0738	0.0226	1.11E-02	++	9.77E-01		
rs1059548	16	77978861	a	g	8945	-0.1298	0.0429	2.50E-03	????	3.49E-01	9921	-0.0236	0.037	5.23E-01	?	8.08E-01	18764	-0.0689	0.0228	1.39E-03	++	6.09E-02		
rs8044853	16	80051360	t	c	18651	0.0803	0.0227	4.10E-04	+++++	5.77E-01	7875	0.0306	0.0331	3.55E-01	++	6.39E-01	28256	0.0644	0.0187	5.82E-04	++	2.16E-01		
rs8051009	16	80016916	t	g	16427	-0.0884	0.0244	2.86E-04	-3-+-	4.65E-01	7863	-0.0278	0.0331	4.01E-01	?	5.22E-01	24290	-0.0671	0.0196	6.69E-04	++	1.41E-01		
267	rs13333580	16	80045873	CMP1	a	t	14755	0.0564	0.0298	8.95E-02	+++++	2.15E-01	0334	0.322E-02	+++	4.22E-01	25069	0.0635	0.0224	4.62E-03	++	7.18E-01		
rs16955379	16	80046874	CMP1	c	14786	0.1263	0.0298	2.20E-05	+++++	7.96E-01	10280	0.0908	0.0334	6.59E-03	+++	4.74E-01	25066	0.1106	0.0222	6.61E-07	++	4.28E-01		
rs9930117	16	80047458	CMP1	a	12452	-0.1159	0.0327	4.02E-04	-7-+-	7.62E-01	10255	-0.0916	0.0334	6.14E-03	---	4.43E-01	22707	-0.104	0.0224	8.54E-06	++	6.03E-01		
rs12596103	16	80048054	CMP1	c	12654	0.1449	0.0318	5.25E-06	+++++	8.37E-01	9798	0.0836	0.0342	1.46E-02	7++	3.58E-01	22452	0.1165	0.0233	5.69E-07	++	1.89E-01		
rs12596138	16	80048250	CMP1	c	14826	0.1244	0.0298	2.97E-05	+++++	8.39E-01	10259	0.0809	0.0339	1.70E-02	+++	2.73E-01	22585	0.1054	0.0224	2.74E-07	++	3.35E-01		
rs12599890	16	80054403	CMP1	g	12738	0.137	0.0309	9.43E-06	+++++	9.06E-01	10349	0.0725	0.0321	2.40E-02	+++	1.29E-01	23807	0.106	0.0223	1.93E-06	++	1.48E-01		
rs12596548	16	80061378	CMP1	c	14787	-0.0941	0.0266	3.99E-04	-7-+-	5.30E-01	10318	-0.0649	0.0296	2.85E-02	---	5.30E-01	25105	-0.0811	0.0198	4.19E-05	++	4.63E-01		
rs2961221	16	80061378	CMP1	a	14829	0.0897	0.0252	3.73E-04	+++++	8.50E-01	9920	0.0107	0.0291	7.13E-01	??	9.19E-01	24749	0.0588	0.019	3.27E-03	++	4.02E-02		
rs1210275	16	80064455	ATP2C2	g	14829	0.1216	0.0329	2.39E-04	+++++	3.82E-01	7486	0.014	0.0404	7.29E-01	??	1.00E-01	22865	0.0387	0.0255	4.99E-03	++	8.89E-02		
rs17814267	16	83004576	ATP2C2	a	14830	0.1491	0.0374	6.74E-05	+++++	3.21E-01	7486	-0.011	0.0434	8.00E-01	??	1.00E-01	22316	0.0809	0.0283	4.31E-03	++	5.20E-03		
rs17741450	16	83006024	ATP2C2	c	14731	-0.1197	0.033	2.87E-04	-7-+-	1.32E-01	7487	-0.019	0.0404	6.38E-01	??	1.00E-01	22318	-0.0794	0.0256	1.89E-03	++	5.36E-02		
269	rs8082455	17	14240061	DCC	a	g	18797	0.1039	0.0244	2.04E-05	+++++	6.09E-01	7875	0.0406	0.037	2.73E-01	++	9.63E-01	26672	0.084	0.0204	3.20E-05	++	1.53E-01
270	rs9910148	17	16109734	PGL1	a	g	8843	-0.1591	0.0584	6.48E-03	????	4.22E-01	10210	-0.0615	0.0498	2.16E-01	---	9.28E-01	19553	-0.1026	0.0379	6.78E-03	++	2.04E-01
rs2170990	17	16111489	PGL1	c	14812	-0.1314	0.0415	1.26E-03	-7-+-	7.00E-01	10352	-0.0938	0.048	5.08E-02	---	7.24E-01	25164	-0.1168	0.0314	1.99E-04	++	5.26E-01		
rs2014604	17	16117608	PGL1	c	18787	0.1049	0.0311	7.46E-04	+++++	7.36E-01	10334	0.0985	0.0426	2.08E-02	+++	9.78E-01	29121	0.1027	0.0251	4.36E-05	++	9.03E-01		
rs2192687	17	16118112	PGL1	c	18812	0.1082	0.031	4.82E-04	+++++	7.20E-01	10336	0.0977	0.0426	2.18E-02	+++	9.77E-01	29148	0.1046	0.0251	3.02E-05	++	8.42E-01		
rs12948244	17	16118825	PGL1	c	12602	0.1026	0.0371	5.72E-03	+7+7+	5.11E-01	10275	0.0829	0.0427	5.21E-02	+++	8.82E-01	22877	0.0941	0.028	7.77E-04	++	7.28E-01		
rs2192622	17	16123534	PGL1	t	14831	0.1365	0.0415	9.90E-04	+++++	5.94E-01	10361	0.0885	0.0479	6.48E-02	+++	8.19E-01	25192	0.1159	0.0314	2.19E-04	++	4.49E-01		
rs13785626	17	16129373	PGL1	g	14832	0.136	0.0415	1.03E-02	+++++	5.96E-01	10348	0.0856	0.0479	5.95E-02	+++	7.89E-01	25194	0.1164	0.0314	2.05E-04	++	6.88E-02		
rs11651578	17	16128317	PGL1	a	14830	-0.1338	0.0415	1.26E-03	-7-+-	5.65E-01	10363	-0.0885	0.0479	6.48E-02	---	8.19E-01	25193	-0.1144	0.0314	2.66E-04	++	4.75E-01		
rs13785255	17	16131143	PGL1	c	14830	-0.1326	0.0414	1.38E-03	-7-+-	5.93E-01	10363	-0.0885	0.0479	6.48E-02	---	8.19E-01	25193	-0.1137	0.0312	2.82E-04	++	4.86E-01		
rs4791665	17	16136781	PGL1	a	g	18784	0.1017	0.0311	1.07E-03	+++++	7.71E-01	10328	0.0985	0.0422	1.97E-02	+++	9.76E-01	29112	0.1006	0.025	5.89E-05	++	9.51E-01	
rs11868284	17	16137000	PGL1	c	12494	0.1324	0.0448	3.12E-03	+++++	4.64E-01	10361	0.0905	0.0479	5.92E-02	+++	7.86E-01	22855	0.1128	0.0237	5.63E-04	++	5.23E-01		
rs4792731	17	16142666	PGL1	c	14831	0.136	0.0415	1.04E-03	+++++	5.96E-01	10360	0.0907	0.0479	5.86E-02	+++	7.82E-01	25191	0.1166	0.0314	2.02E-04	++	4.75E-01		
rs6502499	17	16156930	PGL1	a	18693	0.1004	0.0313	1.32E-03	+++++	8.87E-01	10329	0.1014	0.0434	1.94E-02	+++	9.74E-01	29022	0.1007	0.0254	7.24E-05	++	9.85E-01		
rs3785621	17	16159571	PGL1	c	16893	0.095	0.0319	2.88E-03	+++++	8.02E-01	10329	0.0994	0.0435	2.21E-02	+++	9.75E-01	27222	0.0968	0.0257	1.75E-04	++	9.35E-01		
271	rs13744797	17	74311430	USP36	a	c	14738	0.1040	0.0391	3.32E-04	+++++	9.25E-01	10279	0.0017	0.0444	9.69E-01	+++	3.35E-01	25017	0.0965	0.0293	6.53E-03	++	1.91E-02
rs2171960	17	74313000	USP36	c	18727	0.0961	0.0303	1.50E-03	+++++	7.54E-01	10301	-0.017	0.0442	7.01E-01	---	1.95E-01	29028	0.0599	0.025	1.65E-02	++	3.46E-02		
rs2172231	17	74317794	USP36	a	g	18637	0.1071	0.0322	8.69E-04	+++++	5.06E-01	10348	0.0056	0.0458	9.93E-01	---	1.91E-01	29056	0.0735	0.0263	5.29E-03	++	6.88E-02	
rs374792	17	74334822	USP36	g	16707	-0.1468	0.0384	1.34E-04	-7-+-	8.19E-01	2835	0.0807	0.0808	9.14E-01	++	1.48E-01	19542	-0.1181	0.0347	6.58E-04	++	8.22E-02		
rs3744791	17</																							

rs2051312	18	55987695	t	c	16778	-0.1	0.0281	3.74E-04	---+---	1.24E-01	10356	-0.0448	0.0324	1.66E-01	--	3.30E-01	27134	-0.0763	0.0212	3.25E-04	--	1.98E-01	
rs474112	18	55987812	t	c	16778	-0.1	0.0281	3.72E-04	---+---	1.24E-01	10356	-0.0448	0.0324	1.67E-01	--	3.30E-01	27134	-0.0763	0.0212	3.25E-04	--	1.98E-01	
rs2051311	18	55987860	t	c	18808	-0.0987	0.0275	3.27E-04	---+---	1.41E-01	10349	-0.0411	0.0324	2.05E-01	---	4.01E-01	19157	-0.0746	0.021	3.75E-04	--	1.75E-01	
rs1619975	18	55988204	t	c	18816	-0.0989	0.0275	3.18E-04	---+---	1.47E-01	10349	-0.0483	0.0327	1.99E-01	---	4.00E-01	19165	-0.0779	0.021	2.13E-04	--	2.36E-01	
rs1673539	18	55988205	a	t	16389	-0.1257	0.0269	2.60E-05	---+---	9.62E-01	10349	-0.049	0.0327	1.34E-01	---	4.00E-01	19168	-0.0908	0.0221	3.89E-05	--	8.35E-02	
rs666181	18	55988728	a	t	18801	-0.0968	0.0267	2.92E-04	+++++	1.66E-01	10362	-0.0487	0.0324	1.33E-01	---	2.92E-01	19169	-0.0773	0.0206	1.74E-04	--	2.52E-01	
rs663129	18	55988881	a	c	18803	0.1124	0.0286	8.76E-05	+++++	1.01E-01	10349	0.0551	0.0329	9.41E-02	---	3.07E-01	19152	-0.0877	0.0216	4.81E-05	--	1.89E-01	
rs571312	18	55990749	a	c	18800	0.1126	0.0287	8.70E-05	+++++	9.09E-02	10263	0.0313	0.034	3.58E-01	---	2.30E-01	19063	0.0788	0.0219	3.28E-04	--	6.77E-02	
rs545708	18	55991199	t	c	18800	0.0998	0.0276	2.95E-04	+++++	3.78E-01	10352	0.0402	0.0326	2.17E-01	---	4.47E-01	19152	-0.0749	0.0211	3.76E-04	--	1.63E-01	
rs591166	18	55992569	a	t	16766	0.1037	0.0291	3.64E-04	+++++	2.92E-01	10351	0.0399	0.0324	2.19E-01	---	4.45E-01	19177	-0.0752	0.0216	5.33E-04	--	1.43E-01	
rs589580	18	55992871	a	c	18799	0.097	0.0268	2.94E-04	+++++	1.48E-01	10361	0.0436	0.0318	1.70E-01	---	2.93E-01	19160	-0.0748	0.0205	2.61E-04	--	1.99E-01	
rs975918	18	55993388	t	c	18750	0.1009	0.0276	2.55E-04	+++++	3.87E-01	10350	0.0498	0.032	1.20E-01	---	4.55E-01	19100	-0.0791	0.0209	1.54E-04	--	2.27E-01	
rs1350341	18	55993531	a	c	18796	0.0971	0.0268	2.90E-04	+++++	1.56E-01	10351	0.0444	0.0318	1.62E-01	---	4.29E-01	19147	-0.0752	0.0205	2.42E-04	--	2.05E-01	
rs611428	18	55993865	a	c	16764	0.1035	0.0269	3.56E-04	+++++	3.55E-01	10347	0.0435	0.0319	1.73E-01	---	3.96E-01	19171	-0.0764	0.0215	3.74E-04	--	1.64E-01	
rs553622	18	55994476	a	t	18796	-0.0966	0.0266	2.87E-04	---+---	1.79E-01	10342	-0.0535	0.0319	8.76E-02	---	2.79E-01	19148	-0.0785	0.0203	1.07E-04	--	4.57E-01	
rs559623	18	55994855	a	c	16641	-0.1256	0.0289	1.39E-05	---+---	7.49E-01	10349	-0.025	0.0293	3.92E-01	---	3.20E-01	16810	-0.076	0.0206	2.21E-04	--	1.45E-02	
rs6567161	18	55995915	a	c	18746	0.1008	0.0276	2.68E-04	+++++	5.31E-01	10343	0.0198	0.0299	5.07E-01	---	3.89E-01	19089	-0.0303	0.0203	1.73E-03	--	4.65E-02	
rs523288	18	55999349	a	t	18788	-0.1083	0.0286	1.52E-04	---+---	1.28E-01	10346	-0.0174	0.0315	5.81E-01	---	2.07E-01	19134	-0.0677	0.0212	1.50E-03	--	3.26E-02	
rs2168711	18	55999511	t	c	18791	-0.1088	0.0286	1.42E-04	---+---	1.29E-01	10346	-0.0278	0.0327	3.96E-01	---	2.29E-01	19137	-0.0732	0.0215	6.19E-04	--	6.23E-02	
rs12967135	18	56000003	a	c	16457	0.1378	0.031	8.73E-06	+++++	5.01E-01	10345	0.0438	0.0345	2.04E-01	---	2.70E-01	16802	-0.0958	0.0231	3.25E-05	--	4.17E-02	
rs538656	18	56001402	t	c	18800	0.1081	0.0286	1.57E-04	+++++	1.02E-01	10344	0.0376	0.0309	2.67E-01	---	3.08E-01	19144	-0.0788	0.0219	3.13E-04	--	1.22E-01	
rs1782313	18	56002077	t	c	18808	-0.1064	0.0286	1.95E-04	---+---	1.43E-01	10335	-0.0333	0.0337	3.22E-01	---	2.66E-01	19143	-0.0758	0.0218	5.09E-04	--	9.82E-02	
rs1087177	18	56002743	a	c	18810	-0.1088	0.0286	1.40E-04	---+---	1.26E-01	10340	-0.0393	0.0334	2.39E-01	---	2.48E-01	19150	-0.0794	0.0217	2.57E-04	--	1.14E-01	
rs475134	18	56003734	t	c	18752	0.0944	0.0267	4.03E-04	+++++	1.68E-01	10338	0.0366	0.0323	2.58E-01	---	2.18E-01	19070	-0.0709	0.0206	5.67E-04	--	1.68E-01	
rs1152213	18	56003928	a	c	18807	-0.1082	0.0286	1.54E-04	---+---	1.20E-01	10340	-0.036	0.0335	2.82E-01	---	2.32E-01	19147	-0.0748	0.0218	3.50E-04	--	1.93E-01	
rs1808964	18	56009809	t	c	14731	-0.0782	0.0334	1.92E-02	---+---	6.31E-02	10337	-0.0359	0.0347	3.02E-01	---	1.71E-01	15068	-0.0579	0.0241	1.62E-02	--	3.80E-01	
rs21969709	18	56010543	a	c	14790	0.0773	0.0333	2.03E-02	---+---	6.46E-02	10337	0.0359	0.0347	3.02E-01	---	1.71E-01	15127	-0.0575	0.024	1.68E-02	--	3.89E-01	
rs921971	18	56012643	t	c	14830	-0.0757	0.0332	2.27E-02	---+---	6.91E-02	10337	-0.0366	0.0348	2.92E-01	---	1.72E-01	15167	-0.0571	0.024	1.75E-02	--	4.16E-01	
rs12955983	18	56023969	a	c	16348	-0.1171	0.0319	2.40E-04	---+---	3.46E-01	10339	-0.0609	0.0358	8.48E-01	---	1.03E-01	16687	-0.0683	0.0238	4.12E-03	--	2.16E-02	
rs12970134	18	56035730	a	c	18803	0.1078	0.0297	8.05E-04	+++++	1.16E-01	10327	0.0089	0.0374	8.13E-01	---	9.90E-02	19130	-0.0696	0.0233	2.79E-03	--	3.84E-02	
277	rs5954832	18	65474134	DK6	a	16655	0.1438	0.0366	8.70E-05	+++++	9.02E-01	7860	0.0275	0.0512	5.91E-01	-?	8.27E-01	22515	-0.1045	0.0298	4.50E-04	--	6.46E-02
rs11660034	18	65475733	DK6	a	14633	-0.1498	0.0368	4.71E-05	---+---	8.90E-01	7863	-0.0251	0.0512	6.23E-01	-?	7.88E-01	22496	-0.1073	0.0299	3.29E-04	--	4.80E-02	
rs4495886	18	65477133	DK6	t	14831	0.1549	0.0383	5.19E-05	+++++	9.13E-01	7872	0.0233	0.0521	6.55E-01	-?	8.82E-01	22703	-0.1087	0.0309	4.26E-04	--	4.18E-02	
278	rs2628125	18	70722593	ZNF407	a	16738	-0.0496	0.0243	4.16E-02	---+---	1.13E-01	10330	0.0247	0.0285	3.95E-01	---	8.20E-01	27068	-0.0185	0.0185	3.16E-01	--	4.88E-02
rs2554162	18	70722734	ZNF407	a	16999	0.0821	0.0223	4.44E-04	+++++	9.90E-01	10331	0.0223	0.0285	4.33E-01	---	8.42E-01	27139	-0.0403	0.018	2.56E-02	--	4.57E-01	
rs8098151	18	70727962	ZNF407	t	14699	-0.0811	0.0226	1.53E-03	---+---	9.67E-01	10345	0.0251	0.0284	3.78E-01	---	6.99E-01	25044	-0.0385	0.019	7.82E-02	--	5.48E-03	
rs2628123	18	70728913	ZNF407	a	18692	-0.0809	0.0228	3.91E-04	---+---	9.88E-01	10323	-0.0287	0.0285	3.14E-01	---	6.64E-01	29015	-0.0381	0.0178	3.72E-02	--	2.67E-03	
rs2554130	18	70744915	ZNF407	a	18646	0.0803	0.023	4.71E-04	+++++	8.67E-01	10321	0.0242	0.0338	4.74E-01	---	1.28E-01	28967	-0.0625	0.019	1.01E-03	--	1.70E-01	
rs2628117	18	70746183	ZNF407	a	8843	0.088	0.0342	1.01E-02	+++++	6.94E-01	10192	0.0288	0.0341	3.99E-01	---	1.68E-01	19035	-0.0583	0.0241	1.57E-02	--	2.20E-01	
279	rs12966258	18	74899539		a	12704	0.096	0.0316	2.41E-03	+++++	9.58E-01	10253	-0.005	0.0313	8.73E-01	---	8.69E-02	22957	-0.0545	0.0222	4.29E-02	--	2.32E-02
rs4247367	18	74903836		a	12703	0.0968	0.0316	2.22E-03	+++++	9.58E-01	10255	-0.0052	0.0313	8.67E-01	---	8.70E-02	22958	-0.0543	0.0222	4.16E-02	--	2.28E-02	
rs1893571	18	74906352		t	14740	-0.0942	0.0288	1.08E-03	---+---	9.46E-01	10317	-0.0327	0.0313	9.18E-01	---	7.12E-02	25057	-0.0495	0.0228	1.94E-02	--	2.20E-02	
rs12966045	18	74907827		a	8638	0.1184	0.0368	1.29E-03	+++++	9.13E-01	10317	-0.0032	0.0313	9.18E-01	---	7.12E-02	25057	-0.0495	0.0228	1.94E-02	--	2.20E-02	
rs506774	18	74908769		a	16689	0.0922	0.0256	3.10E-04	+++++	9.80E-01	10315	-0.0028	0.0311	9.29E-01	---	1.19E-01	27204	-0.0538	0.0198	6.46E-03	--	1.34E-02	
rs19790001	18	74913438		t	14830	-0.0831	0.0271	2.75E-03	---+---	9.70E-01	10309	0.007	0.0309	8.28E-01	---	1.09E-01	25189	-0.0458	0.0206	5.67E-02	--	4.35E-02	
rs12957405	18	74915134		c	16338	0.094	0.028	7.74E-04	+++++	9.78E-01	10347	0.0023	0.0313	9.41E-01	---	3.28E-02	26685	-0.0532	0.0209	1.07E-02	--	2.90E-02	
rs4567807	18	74919034	ATP9B	c	18677	0.0913	0.0258	3.94E-04	+++++	9.87E-01	10343	0.0006	0.0313	9.85E-01	---	3.22E-02	29020	-0.0546					

	rs6036127	22393714	t	c	18766	-0.0839	0.0236	3.84e-04	++++	1.56e-01	9938	-0.0351	0.0294	2.32e-01	?	5.50e-01	28704	-0.0648	0.0184	4.32e-04	--	1.96e-01
	rs3952534	22395954	c	g	16709	-0.0206	0.0252	4.15e-01	+?+?	5.33e-02	9974	-0.0444	0.0292	8.79e-01	?	8.83e-01	26683	-0.0137	0.0191	4.73e-01	--	6.75e-01
288	rs6073383	42302978	a	t	12577	-0.0934	0.0292	1.39e-03	-?+?	4.11e-02	10291	-0.0481	0.0288	9.44e-02	?	4.94e-01	22688	-0.0704	0.0205	5.92e-04	--	2.69e-01
	rs2867800	42307923	a	c	14532	0.0832	0.0267	1.86e-03	+?+?+	3.98e-02	10327	0.0462	0.0287	1.08e-01	+++	5.92e-01	24859	0.0666	0.0195	7.30e-04	++	3.45e-01
	rs4810415	42307923	a	c	18759	-0.085	0.0229	2.04e-04	+?+?+	1.17e-02	10356	-0.0483	0.0289	9.50e-02	---	7.09e-01	29115	-0.0708	0.0179	7.91e-05	--	3.20e-01
	rs7273969	42328183	a	c	18720	0.0856	0.0229	1.86e-04	+?+?+	1.26e-02	10356	0.0483	0.0289	9.50e-02	+++	7.09e-01	29076	0.0712	0.0179	7.26e-05	++	3.12e-01
	rs12481690	42329874	a	g	18759	-0.086	0.0229	1.72e-04	-----	1.37e-02	10356	-0.0483	0.0289	9.50e-02	---	7.09e-01	29115	-0.0715	0.0179	6.85e-05	--	3.07e-01
	rs4812810	42329946	a	g	16426	-0.1083	0.0246	1.05e-05	-----?	1.73e-02	10356	-0.049	0.0289	9.02e-02	---	7.18e-01	26782	-0.0834	0.0187	8.53e-06	--	1.18e-01
	rs6017310	42329852	a	g	18747	-0.085	0.0229	2.06e-04	-----	1.05e-02	10362	-0.0498	0.0289	8.50e-02	---	7.02e-01	29109	-0.0714	0.0179	6.91e-05	--	3.40e-01
	rs1884607	42340956	a	g	18788	-0.0831	0.0229	2.82e-04	-----	9.28e-03	10361	-0.0512	0.0289	7.64e-02	---	7.20e-01	29149	-0.0708	0.0179	8.00e-05	--	3.87e-01
	rs2208064	42344736	a	g	18790	-0.0832	0.0229	2.78e-04	-----	9.37e-03	10362	-0.0512	0.0289	7.64e-02	---	7.20e-01	29152	-0.0709	0.0179	7.89e-05	--	3.86e-01
	rs2268090	42351181	a	t	18754	-0.0835	0.0229	2.67e-04	-----	8.76e-03	10349	-0.0552	0.0289	5.67e-02	---	5.90e-01	29103	-0.0729	0.0179	5.25e-05	--	4.43e-01
	rs7265147	42352171	a	g	18684	0.093	0.0231	5.53e-05	+++++	7.62e-02	10347	0.043	0.0286	1.33e-01	+++	5.98e-01	29031	0.0723	0.018	4.57e-05	++	1.74e-01
	rs6031503	42358335	a	g	18706	-0.0924	0.0231	5.98e-05	+++++	6.56e-02	10347	-0.0407	0.0286	1.53e-01	+++	5.62e-01	29053	-0.0731	0.0179	5.76e-05	++	1.59e-01
	rs4812816	42362866	a	c	18697	0.0933	0.023	5.22e-05	+++++	7.13e-02	10354	0.0437	0.0286	1.27e-01	+++	6.13e-01	29051	0.0738	0.0179	3.81e-05	++	1.77e-01
	rs13045189	42367314	a	g	18726	-0.0839	0.0229	2.55e-04	-----	1.13e-02	10349	-0.0496	0.0289	8.68e-02	---	7.59e-01	29075	-0.0707	0.0179	8.24e-05	--	3.52e-01
	rs4812817	42367435	a	g	18680	-0.093	0.0231	5.49e-05	+++++	8.02e-02	10360	-0.0395	0.0286	1.68e-01	---	6.59e-01	29040	-0.0719	0.018	6.34e-05	--	1.46e-01
	rs13037113	42370171	a	c	18555	-0.0949	0.0231	4.04e-05	-----	8.91e-02	10361	-0.0405	0.0289	1.61e-01	---	6.67e-01	28916	-0.0737	0.018	4.43e-05	--	1.42e-01
	rs6031507	42375042	a	c	18688	-0.0952	0.0231	3.64e-05	-----	1.10e-01	10372	-0.0193	0.0302	5.23e-01	---	2.26e-01	28960	-0.0672	0.0183	2.51e-02	++	4.59e-02
	rs6017317	42380380	a	g	18595	-0.0981	0.0232	2.43e-05	-----	1.17e-01	10237	-0.0637	0.0369	8.42e-02	---	5.09e-01	28832	-0.0884	0.0196	6.84e-06	--	4.30e-01
	rs11907772	42401240	a	g	12699	0.0921	0.0279	9.60e-04	+?+?+	2.39e-01	7858	0.0186	0.0368	6.14e-01	+?	2.46e-01	20557	0.0653	0.0222	3.33e-03	++	1.12e-01
	rs4812822	42404268	a	c	8843	0.0864	0.0346	1.26e-02	????+	1.56e-01	7487	-0.012	0.0346	7.29e-01	??	1.00e+00	16330	-0.0372	0.0245	1.28e-01	++	4.43e-02
	rs4812823	42404514	a	g	14823	-0.0738	0.0256	3.90e-03	-?+?+	6.02e-01	7481	-0.03	0.0339	3.76e-01	??	1.00e+00	22304	-0.0579	0.0204	6.06e-03	++	3.03e-01
	rs4812826	42405689	a	g	8843	0.074	0.0344	1.17e-02	????+	1.53e-01	9965	0.0439	0.0297	1.40e-01	??	8.11e-04	18808	0.0568	0.0225	1.16e-02	++	5.08e-01
289	rs1126631	50688817	a	c	18693	0.0943	0.0265	3.65e-04	+++++	8.99e-01	10305	-0.021	0.0388	4.95e-01	+++	7.04e-02	28998	0.0453	0.0201	2.43e-02	++	4.54e-03
	rs1916648	50691425	a	c	18671	0.0911	0.0265	5.78e-04	+++++	9.84e-01	10305	-0.0148	0.0307	6.30e-01	+++	5.71e-02	28976	0.0509	0.0201	2.22e-02	++	9.02e-03
	rs1916647	50691463	t	c	18671	-0.0911	0.0265	5.78e-04	-----	9.84e-01	10306	0.0149	0.0307	6.26e-01	---	6.23e-02	28977	-0.0458	0.0201	2.23e-02	++	8.96e-03
	rs8124744	50695302	a	g	18511	0.086	0.0261	9.82e-04	+++++	8.32e-01	10343	-0.0144	0.0304	6.37e-01	+++	4.29e-02	28854	-0.0434	0.0198	2.84e-02	++	1.22e-02
	rs8126232	50695583	a	g	18586	-0.0863	0.0261	9.25e-04	-----	9.30e-01	10349	0.0107	0.0304	7.25e-01	---	6.16e-02	28935	-0.0451	0.0198	2.26e-02	++	1.55e-02
	rs6126634	50696649	a	c	18694	-0.0842	0.0259	1.15e-03	-----	8.59e-01	10354	-0.0114	0.0304	7.09e-01	---	7.48e-02	29048	-0.0401	0.0197	2.57e-02	++	1.67e-02
	rs13041661	50697111	a	g	18695	0.0805	0.0258	1.81e-03	+++++	8.55e-01	10354	-0.0104	0.0303	7.30e-01	+++	1.00e-01	29049	0.0423	0.0196	3.13e-02	++	2.24e-02
	rs6202217	50698164	t	c	18650	0.0818	0.0257	1.46e-03	+++++	8.41e-01	10325	-0.0122	0.0303	6.89e-01	+++	1.08e-01	28975	0.0426	0.0196	2.99e-02	++	1.82e-02
	rs6062829	50700220	a	g	18636	0.0812	0.0257	1.55e-03	+++++	8.00e-01	10308	-0.0118	0.0302	6.97e-01	+++	9.35e-02	28944	0.0421	0.0196	3.13e-02	++	1.90e-02
	rs6062920	50701972	a	g	18657	0.0827	0.0257	1.30e-03	+++++	7.84e-01	10297	-0.0109	0.0307	7.22e-01	+++	7.37e-02	28954	0.0401	0.0197	2.51e-02	++	1.94e-02
290	rs1276430	50930012	a	g	14831	-0.1746	0.0464	1.85e-04	-?+?+	2.72e-01	10324	-0.0121	0.0509	8.12e-01	---	9.98e-01	23155	-0.1003	0.0343	3.44e-03	++	1.90e-02
	rs11699823	50938819	a	g	16679	-0.1529	0.0476	1.31e-03	-?+?+	6.45e-01	10333	0.0051	0.0515	9.22e-01	---	7.16e-01	27012	-0.0801	0.035	2.19e-02	++	2.43e-02
	rs13039710	50944179	a	c	16706	0.1528	0.0474	1.27e-03	+?+?+	6.83e-01	10343	-0.0013	0.0516	9.79e-01	+++	6.16e-01	27049	0.0823	0.0349	1.84e-02	++	2.79e-02
291	rs6069971	55073362	c	g	14821	-0.0663	0.0636	2.97e-01	-?+?+	1.55e-03	10346	0.0354	0.0666	5.95e-01	---	3.91e-01	25167	-0.0178	0.046	6.99e-01	++	2.69e-01
	rs6069972	55073413	c	g	14822	-0.0663	0.0636	2.97e-01	-?+?+	1.55e-03	10346	0.001	0.098	9.92e-01	---	3.60e-01	25168	-0.0464	0.0533	3.85e-01	++	5.65e-01
	rs6069973	55073780	c	g	14823	0.2355	0.0636	2.11e-04	+?+?+	1.70e-01	10337	-0.0528	0.0669	4.30e-01	---	3.08e-01	25160	0.0986	0.0461	3.24e-02	++	1.79e-03
	rs6069974	55074698	t	c	14829	0.0659	0.0634	2.99e-01	+?+?+	1.24e-01	10342	-0.0252	0.0663	7.04e-01	---	3.90e-01	25171	0.0224	0.0458	6.25e-01	++	3.21e-01
	rs6069975	55074752	a	g	12795	-0.0928	0.0672	1.68e-01	-?+?+	1.17e-01	10346	0.0125	0.0978	8.98e-01	---	3.78e-01	23141	-0.059	0.0554	2.87e-01	++	3.75e-01
	rs6025344	55075405	a	g	14822	-0.0661	0.0636	2.98e-01	-?+?+	1.42e-03	10345	0.0086	0.0979	9.30e-01	---	3.87e-01	25167	-0.0439	0.0533	4.10e-01	++	5.22e-01
	rs6025345	55075676	a	g	14819	-0.0659	0.0636	3.00e-01	-?+?+	1.40e-03	10345	0.0336	0.0666	6.14e-01	+++	3.96e-01	25164	-0.0184	0.046	6.99e-01	++	2.80e-01
	rs5099432	55093252	a	t	14779	-0.0801	0.0582	8.90e-01	-?+?+	3.56e-03	10302	0.022	0.0623	7.25e-01	---	5.87e-01	25081	0.0505	0.0425	8.89e-01	++	7.24e-01
	rs259984	57200438	a	g	16632	0.0924	0.0263	4.52e-04	+?+?+	3.69e-01	10328	0.0097	0.032	7.63e-01	---	9.88e-01	26960	0.0591	0.0203	3.65e-03	++	4.59e-02
	rs260008	57204799	a	c	16678	0.0927	0.0264	4.48e-04	+?+?+	3.93e-01	10350	0.0123	0.0362	7.35e-01	---	1.00e+00	27028	0.0648	0.0213	2.39e-03	++	7.27e-02
	rs4603838																					

Supplementary Table 5. Nineteen SNPs taken forward to stage 3 for *de novo* replication and their DIAGRAM+ lookup results

SNP	Chr	Nearby gene	Risk allele	Other allele	RAF (HapMap JPT/CHB)	Stage 1 (discovery)	Stage 2 (in silico replication)	Stage 3 (de novo replication)	Combined (stage 1+2+3)	DIAGRAM+			
						P-value	P-value	P-value	P-value	RAF (HapMap CEU)	OR (CI)	P-value	power [§]
						up to 6,952 cases and 11,865 controls	up to 5,843 cases and 4,574 controls	up to 12,284 cases and 13,172 controls	up to 25,079 cases and 29,611 controls	up to 8,130 cases and 38,987 controls			
Replicated in Stage 3													
rs831571	3	<i>PSMD6</i>	c	t	0.61	4.85E-06	4.46E-02	1.41E-05	8.41E-11	0.76	1.02 (0.97-1.07)	4.45E-01	0.16
rs6815464 [†]	4	<i>MAEA</i>	c	g	0.58	8.21E-04	3.67E-05	4.15E-15	1.57E-20	0.97	1.19 (1.05-1.36)	6.56E-03	0.91
rs9470794	6	<i>ZFAND3</i>	c	t	0.27	1.45E-04	1.48E-02	3.20E-06	2.06E-10	0.14	0.97 (0.90-1.04)	4.00E-01	0.23
rs1535500 [‡]	6	<i>KCNK16</i>	t	g	0.42	5.34E-06	3.33E-02	3.50E-03	2.30E-08	0.48	1.00 (0.96-1.04)	9.21E-01	0.05
rs6467136	7	<i>GCC1-PAX4</i>	g	a	0.79	6.47E-05	2.09E-03	2.31E-05	4.96E-11	0.56	0.99 (0.95-1.03)	4.89E-01	0.09
rs7041847	9	<i>GLIS3</i>	a	g	0.41	1.29E-04	2.20E-03	2.89E-09	1.99E-14	0.54	1.04 (1.00-1.08)	6.43E-02	0.62
rs17797882	16	<i>WWOX</i>	t	c	0.32	1.76E-04	1.21E-02	1.61E-02	9.49E-07	0.02	1.20 (0.95-1.52)	1.22E-01	0.87
rs16955379*	16	<i>CMIP</i>	c	t	0.8	2.20E-05	6.59E-03	2.19E-02	2.84E-07	0.99	1.20 (1.01-1.42)	3.33E-02	0.52
rs3786897	19	<i>PEPD</i>	a	g	0.56	3.74E-06	1.28E-01	5.46E-04	1.30E-08	0.6	1.02 (0.98-1.06)	3.61E-01	0.2
rs6017317	20	<i>FITM2-R3HDML-HNF4A</i>	g	t	0.48	2.43E-05	8.42E-02	3.96E-07	1.12E-11	0.18	1.07 (1.01-1.13)	1.47E-02	0.86
Not replicated in Stage 3													
rs1027700	1	<i>ATF6</i>	a	t	0.32	3.47E-05	8.97E-04	7.01E-01	6.16E-04	0.25	1.01 (0.97-1.05)	6.36E-01	0.08
rs6441806	3	<i>ABHD5</i>	t	c	0.64	1.22E-06	7.26E-02	9.16E-01	3.57E-04	0.05	0.99 (0.95-1.04)	7.74E-01	0.06
rs6807248	3	<i>TIPARP</i>	c	t	0.17	7.56E-06	6.61E-02	6.81E-01	1.55E-04	0.23	1.02 (0.97-1.06)	4.95E-01	0.16
rs4927868	3	<i>TFRC</i>	t	c	0.38	2.67E-04	1.13E-02	5.80E-01	6.44E-05	0.6	0.95 (0.84-1.08)	4.40E-01	0.83
rs1265099	6	<i>PSORS1C1</i>	g	a	0.49	2.82E-04	1.15E-02	3.09E-01	9.71E-04	0.61	1.03 (0.99-1.08)	1.19E-01	0.39
rs2191651	6	<i>PRIM2</i>	g	a	0.52	3.65E-05	5.23E-03	6.87E-01	1.52E-04	0.33	1.01 (0.96-1.06)	7.66E-01	0.08
rs17715285	7	<i>GCC1</i>	t	c	0.79	2.14E-04	8.38E-03	2.88E-01	5.93E-06	0.47	0.98 (0.94-1.02)	3.95E-01	0.22
rs10282790	8	<i>CNBD1</i>	g	t	0.19	1.36E-04	1.55E-02	5.43E-02	4.25E-03	0.17	1.01 (0.97-1.06)	5.64E-01	0.07
rs4993062	16	<i>ARHGAP17-LCMT1</i>	a	g	0.13	3.32E-05	1.31E-02	5.44E-01	2.93E-04	0.23	0.99 (0.95-1.04)	7.14E-01	0.08

*Proxy SNP, rs9930117 ($r^2=1$), was genotyped in the stage3 CAGE study. Proxy SNPs used for DIAGRAM+ lookup are [†]rs11247991 ($r^2=0.96$) and [‡]rs3734618 BBJ ($r^2=1$). [§]The power was estimated given the 8,130 cases/38,987 controls, DIAGRAM+ ORs, T2D prevalency of 10% and RAF in HapMap (CEU) for $\alpha=0.05$. RAF, risk allele frequency. NA, not available.

Supplementary Table 6. Results of association analyses for T2D with and without the adjustment for BMI. The primary association of each T2D locus detected in this study was adjusted with BMI.

SNP	Chr	Position (bp)	Nearby gene	Risk allele	Other allele	Combined (stage 1+2+3) before BMI adjustment		Combined (stage 1+2+3) after BMI adjustment	
						OR (CI)	P-value	OR (CI)	P-value
rs6815464	4	1,299,901	<i>MAEA</i>	c	g	1.13 (1.10-1.16)	1.57E-20	1.14 (1.11-1.18)	8.70E-22
rs7041847	9	4,277,466	<i>GLIS3</i>	a	g	1.10 (1.07-1.13)	1.99E-14	1.11 (1.08-1.14)	6.27E-14
rs6017317	20	42,380,380	<i>FITM2-R3HDML-HNF4A</i>	g	t	1.09 (1.07-1.12)	1.12E-11	1.09 (1.06-1.12)	2.27E-10
rs6467136	7	126,952,194	<i>GCC1-PAX4</i>	g	a	1.11 (1.07-1.14)	4.96E-11	1.11 (1.07-1.14)	3.00E-10
rs831571	3	64,023,337	<i>PSMD6</i>	c	t	1.09 (1.06-1.12)	8.41E-11	1.08 (1.06-1.11)	4.15E-09
rs9470794	6	38,087,158	<i>ZFAND3</i>	c	t	1.12 (1.08-1.16)	2.06E-10	1.12 (1.08-1.16)	2.22E-09
rs3786897	19	38,584,848	<i>PEPD</i>	a	g	1.10 (1.07-1.14)	1.30E-08	1.12 (1.08-1.16)	1.28E-09
rs1535500	6	39,392,028	<i>KCNK16</i>	t	g	1.08 (1.05-1.11)	2.30E-08	1.09 (1.06-1.12)	3.17E-09

*Proxy SNP, rs9930117 ($r^2=1$), was genotyped in the stage3 CAGE study. Sample size is up to 25,079 cases and 29,611 controls.

Supplementary Table 7. List of regions contained in the 38 loci identified by genome-wide association meta-analysis (using published PubMed abstracts prior to December, 2006) with the gene identified by GRAIL analysis. GRAIL P_{Region} -value column presents the best significant P_{Gene} -value adjusted for the multiple comparisons within a locus with at least one gene. The right panel presents the keywords over-described in PubMed abstracts featuring the most connected genes functionally.

REGION	GRAIL P_{Region} -value	CANDIDATE GENE	Keywords Describing Functional Connections
region_18	1.52E-06	<i>HNF1B</i>	'insulin'
region_7	8.54E-06	<i>CDKN2A</i>	'diabetes'
region_20	2.05E-05	<i>IDE</i>	'pancreatic'
region_15	2.07E-05	<i>GCKR</i>	'nuclear'
region_19	2.17E-05	<i>HNF4A</i>	'glucose'
region_8	2.96E-05	<i>CDKN2B</i>	'beta'
region_22	0.001055	<i>IRS1</i>	'pancreas'
region_24	0.001124	<i>KCNJ11</i>	'hepatocyte'
region_2	0.003171	<i>FAM148A</i>	'ppargamma'
region_16	0.003893	<i>GLIS3</i>	'channel'
region_36	0.005116	<i>TCF7L2</i>	'talk'
region_29	0.006847	<i>PPARG</i>	'kappab'
region_39	0.009422	<i>WWOX</i>	'catenin'
region_37	0.021412	<i>TP53INP1</i>	'endothelial'
region_17	0.031825	<i>HHEX</i>	'acute'
region_25	0.03307	<i>KCNK16</i>	'factor'
region_26	0.033255	<i>KCNQ1</i>	'promoter'
region_3	0.059807	<i>FAM148B</i>	'zinc'
region_35	0.076044	<i>SLC30A8</i>	'tumor'
region_33	0.111309	<i>PTPRD</i>	'mutations'
region_1	0.209535	<i>BCL11A</i>	
region_11	0.33995	<i>DGKB</i>	
region_31	0.399287	<i>PROX1</i>	
region_21	0.410371	<i>IGF2BP2</i>	
region_30	0.419303	<i>PRC1</i>	
region_38	0.573858	<i>UBE2E2</i>	
region_23	0.630808	<i>JAZF1</i>	
region_14	0.694944	<i>GCC1</i>	
region_5	0.734563	<i>CDC123</i>	
region_4	0.744804	<i>CAMK1D</i>	
region_27	0.74898	<i>MAEA</i>	
region_32	0.823088	<i>PSMD6</i>	
region_28	0.84532	<i>PEPD</i>	
region_40	0.936323	<i>ZFAND3</i>	
region_10	0.958475	<i>CMIP</i>	
region_6	1	<i>CDKAL1</i>	
region_9	1	<i>CENTD2</i>	
region_13	1	<i>FTO</i>	
region_12	N/A		
region_34	N/A		

Supplementary Table 8. List of connection results from the GRAIL analysis for selected genes ($P_{\text{Gene}} < 0.05$) in Supplementary Table 7. GRAIL P_{Gene} -values are the results for the test of the gene itself, and thus are not adjusted for the number of genes tested within a locus. Analysis results present connections between previously known diabetes associated genes based on their prior literature.

GENE	GRAIL P_{Gene} value	SELECTED SIMILAR GENES (Rank in parantheses)
<i>HNF1B</i>	1.52E-06	HNF4A(3), GCKR(98), TCF7L2(111), GLIS3(224), PPARG(340), PROX1(355), PEPD(389), FAM148B(412), WWOX(433), HHEX(474), CDKN2A(608), FAM148A(652), IRS1(1463), KCNJ11(1465), CDKN2B(1732), CAMK1D(1993)
<i>CDKN2A</i>	8.54E-06	CDKN2B(1), WWOX(18), TP53INP1(64), CENTD2(320), PEPD(349), HNF1B(460), TCF7L2(606), PPARG(1059), PSMD6(1245), PTPRD(1299), BCL11A(1414), PROX1(1479), CAMK1D(1482), HNF4A(1663)
<i>IDE</i>	2.05E-05	IRS1(5), PEPD(50), PPARG(160), SLC30A8(278), KCNJ11(367), HNF4A(378), HNF1B(407), CAMK1D(670), GCKR(681), TCF7L2(702), WWOX(1202), PSMD6(1437)
<i>GCKR</i>	2.07E-05	HNF4A(18), HNF1B(24), PEPD(138), IRS1(147), KCNJ11(177), PPARG(196), IDE(581), WWOX(834), TCF7L2(896), CAMK1D(1021), GLIS3(1512), PROX1(1733)
<i>HNF4A</i>	2.17E-05	HNF1B(3), GCKR(85), TCF7L2(121), PPARG(127), PROX1(348), HHEX(394), PEPD(575), WWOX(822), IRS1(833), CAMK1D(1138), GLIS3(1161), BCL11A(1497), KCNJ11(1717)
<i>CDKN2B</i>	2.96E-05	CDKN2A(1), WWOX(29), TP53INP1(273), BCL11A(401), PEPD(407), JAZF1(462), CAMK1D(652), TCF7L2(751), HNF1B(863), PROX1(1039), HNF4A(1521), PTPRD(1623), PSMD6(1807), PPARG(1984)
<i>IRS1</i>	0.001055	IDE(90), PPARG(102), KCNJ11(354), HNF4A(472), SLC30A8(609), TCF7L2(623), HNF1B(748), CAMK1D(807), WWOX(830), PEPD(888), IGF2BP2(902), GCKR(922)
<i>KCNJ11</i>	0.0011244	KCNQ1(103), IRS1(229), KCNK16(267), PPARG(365), HNF1B(376), HNF4A(395), GCKR(491), IDE(655), PEPD(782), TCF7L2(828), SLC30A8(1790), WWOX(1994)
<i>FAM148A</i>	0.0031708	FAM148B(1), HNF1B(64), HNF4A(492), CAMK1D(747), TP53INP1(1040), PPARG(1327)
<i>GLIS3</i>	0.003893	HNF1B(174), ZFAND3(196), JAZF1(383), SLC30A8(403), HNF4A(500), BCL11A(611), PROX1(694), CAMK1D(802), HHEX(952), WWOX(1174), PEPD(1286), TCF7L2(1646)
<i>TCF7L2</i>	0.0051165	HNF4A(185), HNF1B(192), WWOX(344), PPARG(505), HHEX(524), PROX1(634), CAMK1D(756), CDKN2A(810), PEPD(1009), IRS1(1058), CDKN2B(1378), IGF2BP2(1877), BCL11A(1921)
<i>PPARG</i>	0.0068465	IRS1(71), HNF4A(142), HNF1B(371), TCF7L2(451), WWOX(531), PEPD(683), KCNJ11(1125), CDKN2A(1187), CAMK1D(1326), IDE(1536), GCKR(1878)
<i>WWOX</i>	0.0094217	CDKN2A(50), TP53INP1(112), CDKN2B(167), PTPRD(514), PEPD(650), PROX1(1311), CAMK1D(1440), TCF7L2(1454), BCL11A(1847)
<i>TP53INP1</i>	0.0214116	WWOX(39), CDKN2A(82), CDKN2B(425), CAMK1D(662), PEPD(1009), PROX1(1243), CDC123(1946)
<i>HHEX</i>	0.0318255	PROX1(97), HNF4A(420), HNF1B(492), TCF7L2(496), BCL11A(614), CAMK1D(1140), WWOX(1253), PEPD(1295), GLIS3(1474)
<i>KCNK16</i>	0.0330698	KCNJ11(74), KCNQ1(130), CENTD2(514), GLIS3(686), CAMK1D(889), PEPD(1010), SLC30A8(1657)
<i>KCNQ1</i>	0.0332554	KCNJ11(78), PEPD(350), KCNK16(496), HNF1B(597), CAMK1D(757), CDKN2A(1101), WWOX(1109), GLIS3(1652), PROX1(1662), HNF4A(1673)

Supplementary Table 9. eQTL analysis results showing the suggestive evidence of cis-association ($P = 5 \times 10^{-2}$) between each lead SNP and genes within 1Mb of the lead SNP. The A1 allele listed in the table is the reference/effect allele. eQTL data in 776 adipose tissues, 667 skin tissues and 778 LCLs were available from the MuTHER consortium. The cis-association was examined by the linear regression analysis for the additive effect of SNP. All data were analyzed in genABEL and probABEL. Genome-wide FDR of 1% for a multiple-testing correction corresponds to a P-value threshold of 5.06×10^{-5} in adipose, 3.81×10^{-5} in skin and 7.80×10^{-5} in LCL.

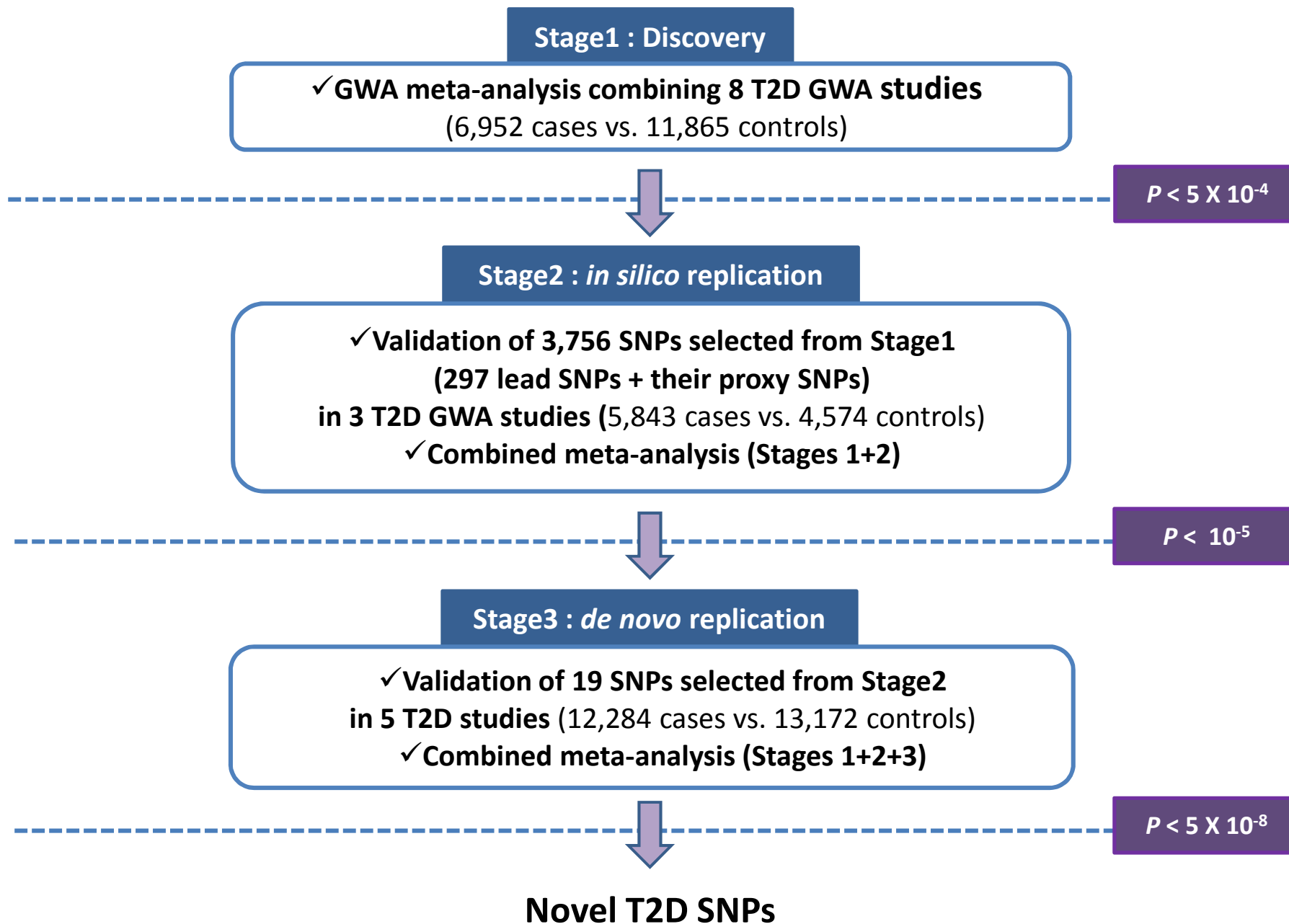
Locus	Lead SNP	Chr	PROBE_Illumina_V3	Gene	TSS	TSE	INFO	SNP_Coor	A1	Freq1	Adipose			Skin			LCL		
											beta	se	P_{eQTL}	beta	se	P_{eQTL}	beta	se	P_{eQTL}
<i>PSMD6</i>	rs831571	3	ILMN_1674559	<i>FLJ44379</i>	63624416	63624465	1	64023337	T	0.197	0.0079	0.0068	2.46E-01	0.0027	0.0084	7.49E-01	-0.0132	0.0062	3.40E-02
<i>ZFAND3</i>	rs9470794	6	ILMN_1763775	<i>DNAH8</i>	38798529	39106545	0.97	38214822	T	0.906	-0.0030	0.0084	7.21E-01	-0.0250	0.0097	1.00E-02	0.0053	0.0097	5.85E-01
		6	ILMN_1651535	<i>TMEM217</i>	37287932	37333391	0.97	38214822	T	0.906	0.0169	0.0079	3.35E-02	0.0031	0.0089	7.28E-01	0.0181	0.0135	1.80E-01
<i>KCNK16</i>	rs1535500	6	ILMN_1732271	<i>BTBD9</i>	38244204	38715902	0.97	38214822	T	0.906	-0.0252	0.0097	9.15E-03	-0.0026	0.0112	8.13E-01	0.0044	0.0099	6.59E-01
		6	ILMN_1766918	<i>KCNK5</i>	39264724	39305229	0.99	39392028	T	0.486	0.0051	0.0060	3.96E-01	0.0628	0.0256	1.42E-02	-0.0091	0.0102	3.70E-01
		6	ILMN_1717702	<i>KCNK17</i>	39374754	39390214	0.99	39392028	T	0.486	-0.0028	0.0059	6.33E-01	-0.0137	0.0060	2.28E-02	-0.0060	0.0058	3.03E-01
		6	ILMN_1702177	<i>GLO1</i>	38751679	38778930	0.99	39392028	T	0.486	-0.0043	0.0166	7.97E-01	-0.0546	0.0249	2.83E-02	0.0159	0.0115	1.65E-01
		6	ILMN_1716728	<i>C6orf64</i>	39179817	39190843	0.99	39392028	T	0.486	-0.0125	0.0162	4.39E-01	0.0334	0.0164	4.15E-02	0.0176	0.0108	1.01E-01
<i>GCC1</i>	rs6467136	6	ILMN_1763775	<i>DNAH8</i>	38798529	39106545	0.99	39392028	T	0.486	0.0109	0.0051	3.18E-02	-0.0025	0.0058	6.66E-01	0.0024	0.0059	6.85E-01
		7	ILMN_1689635	<i>FSCN3</i>	127020924	127029079	0.98	126952194	G	0.525	0.0056	0.0046	2.23E-01	0.0126	0.0048	8.64E-03	0.0072	0.0046	1.17E-01
		7	ILMN_1794846	<i>LRRC4</i>	127454359	127458238	0.98	126952194	G	0.525	-0.0068	0.0057	2.30E-01	-0.0116	0.0056	3.74E-02	-0.0021	0.0064	7.44E-01
		7	ILMN_1729212	<i>GRM8</i>	125865887	126670805	0.98	126952194	G	0.525	-0.0019	0.0043	6.65E-01	0.0086	0.0049	7.88E-02	0.0102	0.0177	5.64E-01
		7	ILMN_1690343	<i>IMPDH1</i>	127819566	127837272	0.98	126952194	G	0.525	-0.0136	0.0056	1.58E-02	-0.0057	0.0060	3.44E-01	-0.0020	0.0082	8.09E-01
<i>GLIS3</i>	rs7041847	7	ILMN_1659990	<i>C7orf68</i>	127883119	127885708	0.98	126952194	G	0.525	0.0509	0.0187	6.51E-03	-0.0083	0.0158	6.00E-01	-0.0013	0.0270	9.61E-01
		9	ILMN_1693427	<i>GLIS3</i>	3814127	4290035	1	4277466	G	0.502	-0.0001	0.0059	9.83E-01	-0.0042	0.0062	4.99E-01	-0.0145	0.0058	1.18E-02
		9	ILMN_2238640	<i>RFX3</i>	3214646	3515983	1	4277466	G	0.502	0.0108	0.0047	2.29E-02	-0.0020	0.0048	6.83E-01	0.0072	0.0049	1.46E-01
<i>CMIP</i>	rs16955379	9	ILMN_2402600	<i>GLIS3</i>	3814127	4142183	1	4277466	G	0.502	0.0083	0.0078	2.88E-01	0.0002	0.0067	9.81E-01	-0.0304	0.0098	1.87E-03
		16	ILMN_1708103	<i>GAN</i>	79906071	79971302	0.8	80046874	T	0.015	-0.0235	0.0227	3.02E-01	-0.0084	0.0258	7.45E-01	-0.0769	0.0270	4.35E-03
		16	ILMN_1772540	<i>ATMIN</i>	79626958	79638452	0.8	80046874	T	0.015	0.1322	0.0471	5.02E-03	-0.0174	0.0656	7.91E-01	-0.0925	0.0568	1.04E-01
<i>PEPD</i>	rs3786897	16	ILMN_1697317	<i>DYNLRB2</i>	79132354	79142042	0.8	80046874	T	0.015	-0.0911	0.0404	2.43E-02	-0.0084	0.0581	8.85E-01	0.0137	0.0356	7.01E-01
		19	ILMN_1675172	<i>RGS9BP</i>	37858152	37861046	0.94	38584848	G	0.438	-0.0064	0.0045	1.55E-01	-0.0134	0.0073	6.75E-02	-0.0140	0.0046	2.37E-03
		19	ILMN_1794063	<i>ANKRD27</i>	37779746	37857942	0.94	38584848	G	0.438	0.0143	0.0060	1.71E-02	0.0073	0.0064	2.53E-01	0.0067	0.0144	6.41E-01
<i>HNF4A</i>	rs6017317	19	ILMN_1717674	<i>PEPD</i>	38569694	38704639	0.94	38584848	G	0.438	-0.1168	0.0209	2.14E-08	-0.0182	0.0204	3.74E-01	0.0194	0.0169	2.51E-01
		20	ILMN_1713752	<i>SERINC3</i>	42561314	42584140	0.99	42380380	T	0.843	0.0053	0.0074	4.76E-01	0.0187	0.0076	1.41E-02	0.0028	0.0086	7.43E-01
		20	ILMN_1779974	<i>C20orf62</i>	42514038	42527398	0.99	42380380	T	0.843	0.0027	0.0075	7.22E-01	-0.0109	0.0073	1.33E-01	-0.0189	0.0083	2.27E-02
		20	ILMN_2277099	<i>YWHAB</i>	42947757	42970575	0.99	42380380	T	0.843	-0.0159	0.0151	2.94E-01	0.0212	0.0157	1.76E-01	0.0633	0.0301	3.52E-02
		20	ILMN_1788421	<i>KCNK15</i>	42807901	42813082	0.99	42380380	T	0.843	-0.0197	0.0090	2.92E-02	0.0126	0.0100	2.08E-01	0.0068	0.0131	6.03E-01
		20	ILMN_1803686	<i>ADA</i>	42681576	42713790	0.99	42380380	T	0.843	-0.0190	0.0174	2.75E-01	-0.0121	0.0208	5.60E-01	-0.0714	0.0255	5.05E-03
		20	ILMN_1744442	<i>TTPAL</i>	42537960	42556658	0.99	42380380	T	0.843	-0.0146	0.0153	3.38E-01	0.0082	0.0255	7.49E-01	0.0510	0.0191	7.69E-03
		20	ILMN_1812548	<i>GTSF1L</i>	41788214	41789056	0.99	42380380	T	0.843	0.0128	0.0060	3.30E-02	-0.0020	0.0070	7.70E-01	0.0022	0.0069	7.48E-01
		20	ILMN_2079042	<i>WFDC5</i>	43171506	43177217	0.99	42380380	T	0.843	0.0158	0.0071	2.59E-02	-0.0068	0.0352	8.47E-01	-0.0163	0.0110	1.41E-01
		20	ILMN_1760208	<i>GDAP1L1</i>	42309321	42342427	0.99	42380380	T	0.843	-0.0004	0.0067	9.48E-01	-0.0001	0.0077	9.88E-01	-0.0210	0.0078	7.48E-03

Supplementary Table 10. Association results of known T1D loci for T2D.

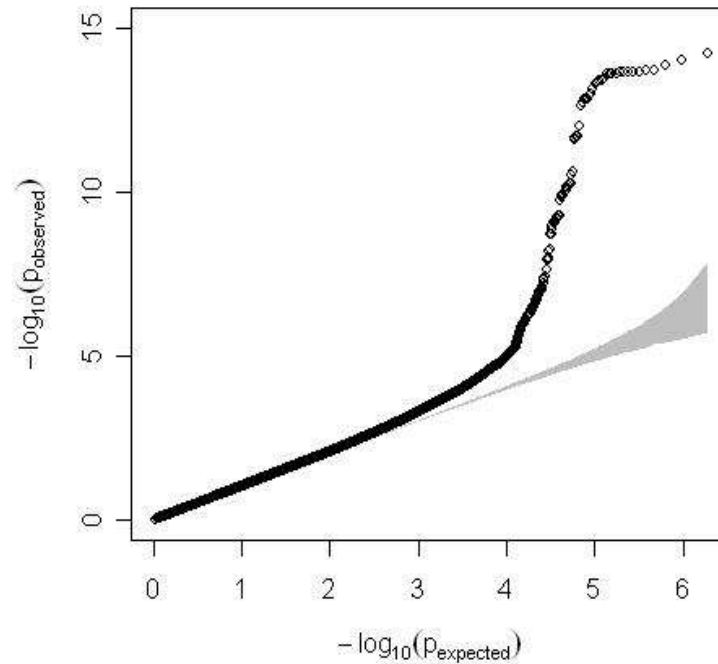
Locus	SNP	Risk allele	N	OR (CI)	P-value	Reference
<i>HLA</i>	rs9267992	na	na	na	na	33
<i>HLA</i>	rs660895	A	14832	1.02 (0.97-1.08)	0.411	29-32
<i>HLA</i>	rs3129941	A	18812	1.05 (0.97-1.13)	0.211	34
<i>HLA</i>	rs926070	G	18649	1.00 (0.95-1.06)	0.957	33
<i>HLA</i>	rs2187668	T	14829	1.02 (0.92-1.13)	0.718	29-32
<i>HLA</i>	rs1048709	G	10725	1.04 (0.97-1.11)	0.226	29-32
<i>HLA</i>	rs630379	C	13034	1.06 (1.00-1.13)	0.045	29-32
<i>INS</i>	rs3842748	G	10891	1.02 (0.85-1.22)	0.845	29-32
<i>PTPN22</i>	rs2476601	na	na	na	na	35
<i>CD25</i>	rs2104286	C	18807	1.05 (0.98-1.12)	0.141	21
<i>CTLA4</i>	rs1427676	T	18796	1.03 (0.98-1.08)	0.220	35
Intergenic region	rs2165738	C	10832	1.07 (1.00-1.14)	0.035	36
Gene desert	rs6887079	C	12699	1.01 (0.95-1.07)	0.747	36
<i>BACH2</i>	rs11755527	C	16758	1.03 (0.98-1.08)	0.221	36
<i>PRKCQ</i>	rs947474	G	18733	1.03 (0.96-1.10)	0.436	36
<i>CTSH</i>	rs3825932	T	12486	1.10 (1.01-1.19)	0.023	36
<i>C16orf75, PRM3, TNP2</i>	rs416603	T	18744	1.06 (1.00-1.13)	0.051	36
<i>C1QTNF6</i>	rs229541	G	18731	1.02 (0.97-1.07)	0.369	36
<i>RGS1</i>	rs2816316	C	18795	1.05 (1.00-1.11)	0.072	33
<i>IL18RAP</i>	rs917997	C	18802	1.04 (0.99-1.08)	0.118	33
<i>IFIH1</i>	rs1990760	C	18761	1.08 (1.02-1.15)	0.007	33
<i>CTLA4</i>	rs3087243	G	18696	1.03 (0.97-1.08)	0.312	33
<i>CCR5</i>	rs11711054	G	18802	1.02 (0.97-1.07)	0.526	33
<i>IL2</i>	rs4505848	G	14831	1.02 (0.96-1.07)	0.560	33
<i>IL7R</i>	rs6897932	C	18816	1.03 (0.97-1.09)	0.315	33
<i>MHC</i>	rs9268645	G	16665	1.00 (0.95-1.05)	0.977	33
<i>BACH2</i>	rs11755527	G	16758	1.03 (0.98-1.08)	0.221	33
<i>TNFAIP3</i>	rs2327832	G	5983	1.42 (0.84-2.38)	0.191	33
<i>TAGAP</i>	rs1738074	C	14831	1.02 (0.97-1.07)	0.509	33
<i>IL2RA</i>	rs12251307	C	18815	1.08 (1.02-1.14)	0.005	33
<i>PRKCQ</i>	rs11258747	G	18815	1.03 (0.91-1.16)	0.666	33
<i>INS</i>	rs7111341	C	13049	1.03 (0.94-1.13)	0.501	33
<i>ERBB3</i>	rs2292239	G	12690	1.01 (0.94-1.08)	0.809	33
Intergenic region	rs1678536	G	4512	1.05 (0.92-1.20)	0.431	33
<i>SH2B3</i>	rs3184504	C	5986	1.15 (0.48-2.76)	0.750	33
<i>CLEC16A</i>	rs12708716	G	16562	1.02 (0.96-1.08)	0.624	33
<i>PTPN2</i>	rs1893217	G	18805	1.01 (0.96-1.08)	0.620	33
<i>CD226</i>	rs763361	C	13047	1.02 (0.97-1.07)	0.499	33
<i>UBASH3A</i>	rs11203203	G	18815	1.01 (0.90-1.13)	0.892	55
<i>IL10</i>	rs3024505	G	18806	1.02 (0.89-1.17)	0.735	33
Gene desert	rs10517086	G	5987	1.18 (0.71-1.96)	0.534	33

<i>C6orf173</i>	rs9388489	G	16481	1.02 (0.90-1.16)	0.718	33
Intergenic region	rs7804356	C	18812	1.03 (0.96-1.11)	0.431	33
<i>COBL</i>	rs4948088	na	na	na	na	33
<i>GLIS3</i>	rs7020673	G	18631	1.08 (1.03-1.13)	0.0008	33
<i>C10orf59</i>	rs10509540	C	14718	1.01 (0.95-1.08)	0.676	33
<i>CD69</i>	rs4763879	G	16459	1.03 (0.98-1.08)	0.193	33
Intergenic region	rs1465788	C	18816	1.02 (0.97-1.07)	0.494	33
Gene desert	rs4900384	G	18695	1.02 (0.98-1.07)	0.317	33
<i>IL27</i>	rs4788084	C	18810	1.04 (0.99-1.09)	0.138	33
Intergenic region	rs7202877	G	14823	1.01 (0.95-1.08)	0.732	33
<i>ORMDL3</i>	rs2290400	C	14832	1.01 (0.96-1.07)	0.662	33
Intergenic region	rs7221109	C	18798	1.01 (0.96-1.06)	0.746	33
Intergenic region	rs425105	C	14829	1.01 (0.95-1.07)	0.794	33
Intergenic region	rs2281808	C	14673	1.01 (0.94-1.08)	0.761	33
Intergenic region	rs5753037	C	18751	1.01 (0.97-1.06)	0.624	33
Intergenic region	rs2664170	na	na	na	na	33
<i>PGM1</i>	rs2269241	C	18792	1.05 (0.99-1.11)	0.086	33
Gene desert	rs1534422	G	14831	1.02 (0.96-1.08)	0.588	33
Intergenic region	rs12444268	T	5677	1.37 (0.83-2.29)	0.220	33
Intergenic region	rs16956936	na	na	na	na	33

na, not available

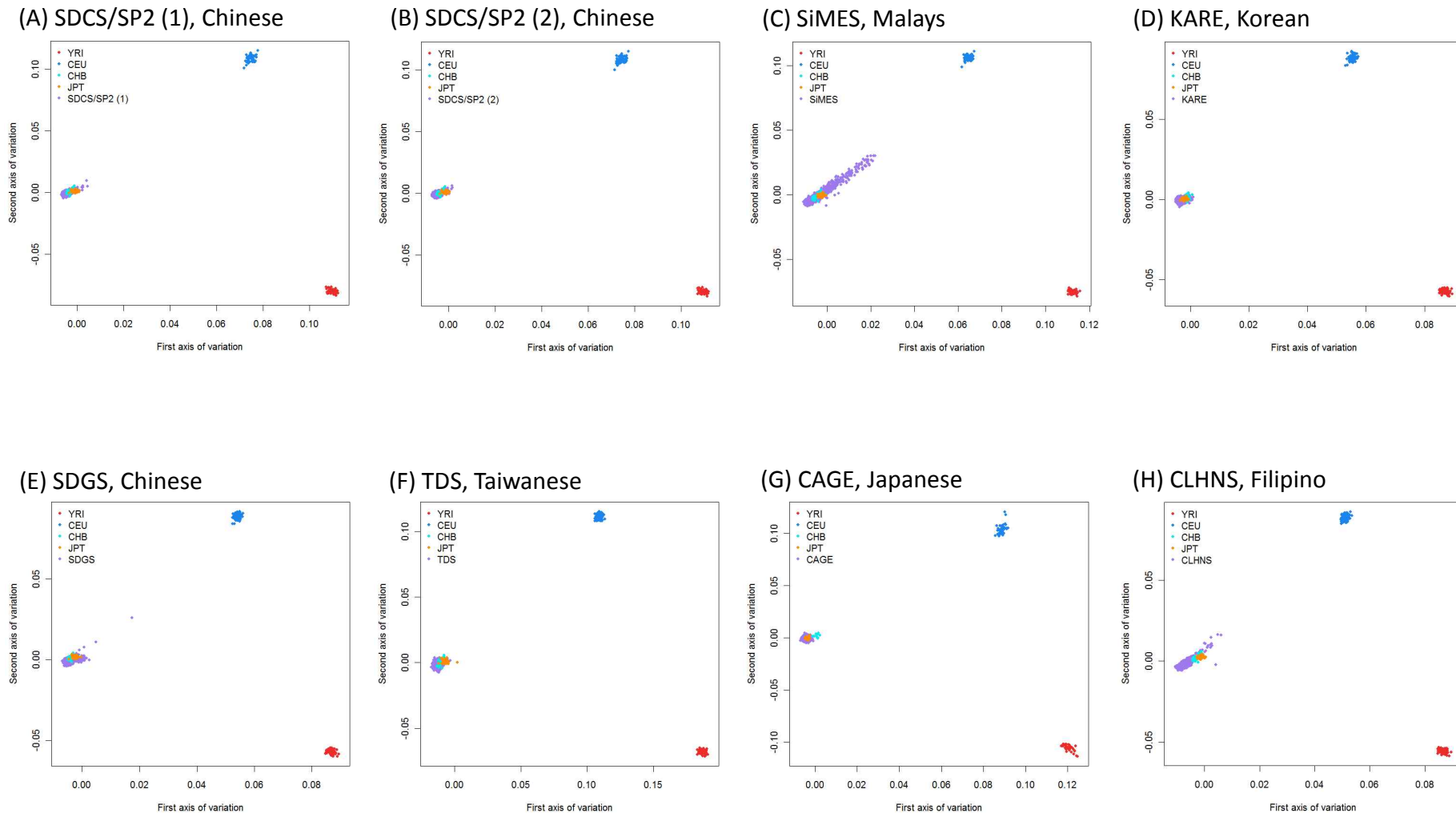


Supplementary Fig. 1. Overall study scheme of East Asian T2D GWA meta-analysis.



Supplementary Figure 2. Quantile-quantile plot for the EA T2D stage 1 meta-analysis.

Observed P -values of given SNPs against the theoretical distribution of expected P -values were plotted from the stage 1 meta-analysis for T2D. The shaded region in the QQ plot represents the 95% pointwise confidence interval. A total of 1,934,619 SNPs that were present in ≥ 5 stage 1 studies were used to generate a plot.



Supplementary Figure 3. Principal Component Analysis (PCA) in individuals from each component study and HapMap data. Individuals from 8 each component study - (A) SDCS/SP2 (1) (Chinese), (B) SDCS/SP2 (2) (Chinese), (C) SiMES (Malays), (D) KARE (Korean), (E) SDGS (Chinese), (F) TDS (Taiwanese), (G) CAGE (Japanese), and (H) CLHNS (Filipino) - and 270 HapMap individuals were plotted based on the first two eigenvectors obtained by PCA. Individuals in each component study, HapMap-JPT, HapMap-CHB, HapMap-CEU and HapMap-YRI are represented by violet, orange, cyan, blue and red colored-dots, respectively.

