Supplementary Table 1. Study sample characteristics, genotyping, quality control and association analysis.

					Sample characterist	tics			Sample OC		SNP OC					Associatio	n analysis	
Study	Ethnic group (origin)	Disease status	Sample size	Age (years)	Age at onset (years)	Fasting glucose (mmol/l)	BMI (kg/m2)	Genotyping platform	Callirate	Evolusion criteria	Call rate	HWE	MAF	Imputation	SNDs	Analysis software	Covariates	Genomic
Study	(01,811)	Discuse status	(males) remaies)	mean (30)	illeali (30)	ilicali (3D)	illeali (30)	denotyping platform	Califate	Exclusion circena	Califate	HVVE	IVIAF	imputation	SINFS	Analysis software	Covariates	Control
Stage 1		Cases	775 (416/359)	56.1 (5.6)	50.9 (10.4)	9.3 (3.6)	30 4 (5 4)		1									1.01 (G)
ARIC	European	Controls	7,159 (3,167/3,992)	54.0 (5.7)		5.4 (0.4)	26.4 (4.5)	Affymetrix Human SNP Array 6.0	≥0.95	Relatedness and duplicates	≥0.90	p>10 ⁻⁶	≥0.01	r ² >0.3	2,443,161	ProbABEL	Age, sex, and study centre	1.01 (I)
deCODE	European (Iceland)	Cases Controls	1,465 (868/597) 23,194 (7,316/15,878)	68.4 (10.1) 59.7 (18.1)	55.1 (12.7)	8.5 (2.7) 5.3 (0.7)	30.1 (5.4) 26.8 (5.0)	Illumina HumanHap 300/370	≥0.98	Duplicates	≥0.96	p>10 ⁻⁶	≥0.01	proper-info>0.5	2,338,113	SNPTEST	-	1.31 (G) 1.31 (I)
DGDG	European (France)	Cases Controls	679 (413/266) 697 (281/416)	59.5 (10.1) 53.9 (5.6)	45.1 (8.4)	9.2 (3.1) 5.1 (0.4)	25.9 (2.8) 23.2 (1.8)	Illumina HumanHap 300	≥0.95	Duplicates	≥0.95	p>10 ⁻⁴	≥0.01	proper-info>0.5	2,051,387	SNPTEST	-	1.10 (G) 1.10 (I)
DGI	European (Sweden/Finland)	Cases Controls	1,022 (529/493) 1,075 (540/535)	65.9 (10.0) 58.0 (10.0)	58.0 (10.0)	9.5 (3.1) 5.3 (0.5)	28.1 (4.1) 27.6 (3.7)	Affymetrix GeneChip 500K	≥0.95	Duplicates	≥0.95	p >10 ⁻⁶	≥0.01	proper-info>0.5	2,230,032	PLINK and SNPTEST	Age, sex, BMI, and study centre	e 1.05 (G) 1.06 (I)
EUROSPAN	European	Cases Controls	269 (127/142) 3,710 (1,557/2,153)	62.9 49.9	-	8.0 4.8	29.8 26.5	Illumina HumanHap 300/370	≥0.98	Duplicates	≥0.98	p>10 ⁻⁶	≥0.01	r ² >0.5	2,359,525	GenABEL and SNPTEST	Age and sex	0.97 (G) 0.98 (I)
FHS	European (USA)	Cases Controls	674 (386/288) 7,664 (3,443/4,221)	63.7 (12.4) 52.3 (16.0)	-	8.6 (2.8) 5.3 (0.5)	31.4 (6.5) 27.0 (5.1)	Affymetrix GeneChip 500K & MIPS 50K	≥0.95	Duplicates	≥0.95	p>10 ⁻⁶	≥0.01	r ² >0.3	2,389,929	R (GEE correction for relatedness)	Age, sex, and cohort	1.02 (G) 1.02 (I)
FUSION	European	Cases Controls	1,161 (653/508) 1,174 (574/600)	62.9 (7.6) 63.6 (7.4)	53.7 (9.1)	9.4 (3.1) 5.3 (0.5)	30.2 (4.7) 27.1 (3.9)	Illumina HumanHap 300	≥0.975	Duplicates	≥0.90	p >10 ⁻⁶	≥0.01	r ² >0.3	2,413,085	MACH2DAT	Age, sex, and birth province	1.03 (G) 1.04 (I)
HPFS	European (USA)	Cases Controls	1,124 (1,124/0) 1,298 (1,298/0)	55.0 (8.6) 55.0 (8.4)	64.0 (8.4)	-	27.8 (4.0) 25.0 (2.9)	Affymetrix Human SNP Array 6.0	≥0.95	Relatedness and duplicates	≥0.95	p>10 ⁻⁶	≥0.01	-	622,575	PLINK	Age and BMI	1.03 (G)
KORAGen	European (Germany)	Cases Controls	433 (255/178) 1,438 (693/745)	65.2 (8.3) 61.9 (7.4)	58.2 (10.3)	= =	30.9 (5.0) 27.7 (4.3)	Affymetrix GeneChip 500K	≥0.93	Duplicates	≥0.95	p>10 ⁻⁶	≥0.01	proper-info>0.5	2,325,232	SNPTEST	Age and sex	1.04 (G) 1.04 (I)
NHS	European (USA)	Cases Controls	1,467 (0/1,467) 1,754 (0/1,754)	43.5 (6.7) 43.1 (6.8)	58.7 (10.6)	-	27.4 (0.1) 23.5 (0.1)	Affymetrix Human SNP Array 6.0	≥0.98	Relatedness and duplicates	≥0.98	p>10 ⁻⁶	≥0.02	-	615,391	PLINK	Age and BMI	0.98 (G)
RS1	European (Netherlands)	Cases Controls	1,178 (488/690) 4.761 (1.928/2.833)	71.7 (8.9) 69.0 (9.1)	71.5 (8.9)	-	27.4 (4.0) 26.0 (3.6)	Illumina HumanHap 550	≥0.975	Duplicates	≥0.98	p>10 ⁻⁶	≥0.01	r ² >0.5	2,439,672	GenABEL and SNPTEST	-	1.01 (G) 1.01 (I)
WTCCC	European (UK)	Cases Controls	1,924 (1,118/806) 2,938 (1,446/1,492)	58.6 (9.2)	50.3 (9.2)	-	30.7 (6.1)	Affymetrix GeneChip 500K	≥0.97	Duplicates	≥0.95 (≥0.99 for MAF<0.05)	p>10 ⁻³	≥0.01	proper-info>0.5	2,308,535	PLINK and SNPTEST	-	1.06 (G) 1.08 (I)
	(UK)	Controls	2,938 (1,446/1,492)			-					TOT MIAP < 0.05)							1.08 (1)
Stage 2 AMC-PAS	European (Netherlands)	Cases Controls	48 (35/13) 442 (333/109)	44.0 (4.7) 43.2 (5.3)	-	8.6 (3.2) 5.3 (0.6)	29.3 (4.4) 26.6 (4.1)	Metabochip	≥0.95	Duplicates and non-European descent	≥0.98	p>10 ⁻⁴	≥0.01	-	109,525	PLINK	Age and sex	1.00 (QT)
BHS	European	Cases	51 (38/13)	72.3 (7.8)		7.8 (4.0)	27.2 (3.9)	Metabochip	≥0.95	Ambiguous sex and relatedness	≥0.95	p>10 ⁻⁶	≥0.01	-	119,893	PLINK	Age and sex	0.96 (QT)
deCODE-Stage2	European	Controls	359 (224/135) 722 (433/289)	70.0 (9.6) 67.3 (13.6)		5.1 (0.9)	26.6 (3.9) 31.2 (6.3)	Metabochip	-	-			≥0.01		125,236	SNPTEST	Sex	0.95 (QT)
DILGOM	(Iceland) European	Controls Cases	10,153 (6,604/3,549) 541 (298/243)	44.3 (26.5) 60.5 (10.3)	-	-	30.1 (5.6)	Metabochip	>0.95	Ambiguous sex and relatedness	>0.95	p>10 ⁻⁶	>0.01		116.634	PLINK	Age, sex, and population	0.93 (QT)
DUNDEE	(Finland) European	Controls Cases	3,357 (1,480/1,877) 3,298 (1,940/1,358)	50.9 (13.5) 63.5 (9.6)	55.9 (8.9)		26.5 (4.5) 31.9 (6.3)	Metabochip	≥0.95	Duplicates, ambiguous sex, and non-	≥0.95 (≥0.99	>5.7x10 ⁻⁷ (p>10 ⁻⁴	≥0.01		121,365	PLINK	structure Population structure	1.07 (QT)
FAS	(UK) European	Controls Cases	3,708 (1,918/1,790) 110 (64/46)	59.1 (11.3) 64.0 (5.6)	-	4.9 (0.5)	27.0 (4.5) 25.9 (3.3)	Metabochin	≥0.95	European descent Gender check and ambiguous sex	for MAF<0.05) >0.95	for MAF<0.05) p>10 ⁻⁶	>0.01		119.523	PLINK	Age and sex	1.00 (QT)
EGCUT	European	Controls Cases	641 (301/340) 938 (342/596)	64.5 (5.6) 64.1 (10.5)	-	-	25.5 (3.9) 33.4 (5.4)	Metabochip		Ambiguous sex, relatedness, and non-	20.95		≥0.01	•	120,720	SNPTEST	Age, sex, and population	
EGCUT FMII-III M	(Estonia) European	Controls Cases	915 (326/589) 755 (510/245)	51.7 (10.7) 48.7 (12.0)	45.5 (10.8)		22.3 (2.5) 28.6 (7.1)	Metabochip		European descent		p>10 ⁻⁶	>0.01		120,720	PLINK	structure Age and sex	1.00 (QT)
	European	Controls Cases	1,632 (765/867) 727 (432/295)	45.1 (10.9) 61.8 (8.2)	-		26.7 (5.0) 29.5 (4.4)			imbiguous sex and cryptic relatedness		p>10 ⁻⁶					•	1.14 (QT)
EPIC	(UK) European	Controls Cases	927 (393/534) 1,037 (584/453)	58.8 (9.4) 59.7 (8.4)	-	7.7 (2.3)	26.1 (3.7) 30.8 (5.4)	Metabochip	≥0.95	Gender check and duplicates	≥0.90	p>10 ⁻⁶	≥0.01	-	120,527	PLINK	Age and sex	0.97 (QT)
FUSION-Stage2	(Finland) European	Controls Cases	1,157 (691/466) 454 (269/185)	59.0 (7.6) 63.4 (7.6)	-	5.4 (0.4) 7.7 (1.8)	26.9 (3.9) 30.6 (5.6)	Metabochip	≥0.98	Relatedness and ambiguous sex	≥0.98	p>10 ⁻⁵	≥0.01	-	123,853	PLINK	Age and sex	0.97 (QT)
FUSION-D2D2007	(Finland) European	Controls Cases	1,229 (455/774) 110 (53/57)	58.1 (8.2) 68.4 (5.8)		5.6 (0.3) 6.6 (0.8)	26.2 (4.3) 30.9 (5.6)	Metabochip	≥0.98	Relatedness and ambiguous sex	≥0.98	p>10 ⁵	≥0.01	-	123,461	PLINK	Age and sex	0.96 (QT)
FUSION-DRExtra	(Finland) European	Controls Cases	785 (341/444) 1.239 (628/611)	66.0 (5.3) 64.0 (12.9)	61.9 (11.9)	5.4 (0.3)	26.7 (4.0) 29.2 (4.8)	Metabochip	≥0.98	Relatedness and ambiguous sex	≥0.98	p>10 ⁵	≥0.01	-	120,746	PLINK	Age and sex	0.94 (QT)
FUSION-HUNT	(Norway)	Controls	1,375 (691/684)	63.5 (13.9)			26.5 (4.0)	Metabochip	≥0.98	Relatedness and ambiguous sex	≥0.98	p>10 ⁻⁵	≥0.01	-	125,644	PLINK	Age, sex and collection site	1.10 (QT)
FUSION-METSIM	European (Finland)	Cases Controls	1,169 (1,169/0) 651 (651/0)	60.5 (6.6) 53.8 (5.0)	57.0 (8.0)	7.5 (2.0) 5.5 (0.3)	30.2 (5.2) 25.9 (3.1)	Metabochip	≥0.98	Relatedness and ambiguous sex	≥0.98	p>10 ⁻⁵	≥0.01	-	122,600	PLINK	Age and sex	1.01 (QT)
GMetS	European (France)	Cases Controls	507 (334/173) 2,553 (1,134/1,419)	55.9 (9.9) 47.3 (11.1)	-	-	28.5 (3.9) 24.8 (3.8)	Metabochip	≥0.95	Gender check, duplicates, and ambigous sex	≥0.95	p>10 ⁻⁴	≥0.01	-	123,359	PLINK	Age, sex, and BMI	1.11 (QT)
HNR	European	Cases Controls	520 (315/205) 3,932 (1,911/2,021)	62.6 (7.2) 59.2 (7.8)	-	-	30.4 (5.2) 27.5 (4.4)	Metabochip	≥0.97	Ambiguous sex, relatedness, and non- European descent	≥0.95	p>10 ⁻⁶	≥0.01	-	126,675	PLINK	Age and sex	1.00 (QT)
IMPROVE	European	Cases Controls	898 (513/385) 2,521 (1,134/1,387)	64.3 (5.6) 64.2 (5.3)	-	7.7 (2.2) 5.3 (0.7)	29.2 (4.6) 26.5 (3.9)	Metabochip	≥0.95	Ambiguous sex, cryptic relatedness, non-European descent	≥0.95	p>10 ⁻⁶	≥0.01	-	122,320	PLINK	Age, sex and population structure	1.13 (QT)
KORAGen-Stage2	European (Germany)	Cases Controls	940 (504/436) 4,209 (2,000/2,209)	61.6 (10.0) 53.8 (13.2)	-		30.8 (5.4) 27.2 (4.6)	Metabochip	≥0.95	Ambiguous sex and relatedness	≥0.95	p>10 ⁻⁶	≥0.01	-	120,547	PLINK	Age and sex	1.02 (QT)
PIVUS	European (Sweden)	Cases Controls	113 (70/43) 864 (419/445)	70.2 (0.2) 70.2 (0.2)	-		29.5 (5.1) 26.8 (4.2)	Metabochip	≥0.95	Relatedness and ambiguous sex	≥0.95	p>10 ⁻⁶	≥0.01	-	120,892	PLINK	Age and sex	0.97 (QT)
РМВ	European (Sweden/Finland)	Cases Controls	4,976 (2,911/2,065) 3,500 (1,700/1,800)	58.8 (12.5) 58.3 (8.4)	:	-	28.2 (8.7) 26.4 (4.4)	Metabochip	≥0.95	Outlying heterozygosity, relatedness and non-European descent	≥0.95	p>10 ⁻⁶	≥0.01	-	119,674	PLINK	Population structure	1.05 (QT)
PROMIS	South Asian (Pakistan)	Cases Controls	1,178 (904/274) 2,472 (2,088/384)	53.5 (9.1) 52.1 (10.2)	:	-		Metabochip	≥0.95	Ambiguous sex and relatedness	≥0.95	p>10 ⁻⁶	≥0.01	-	121,792	PLINK	Age, sex, population structure and MI status	1.00 (QT)
SCARFSHEEP	European (Sweden)	Cases Controls	341 (246/95) 3,073 (2,211/862)	59.6 (7.0) 57.9 (7.3)	-	8.12(5.4) 4.0 (3.8)	27.4 (8.4) 25.4 (6.1)	Metabochip	≥0.95	Ambiguous sex and relatedness	≥0.95	p>10 ⁻⁶	≥0.01	-	119,375	PLINK	Age, sex, population structure and CAD status	0.98 (QT)
STR	European (Sweden)	Cases Controls	320 (141/179) 1,318 (612/706)	71.5 (9.3) 74.3 (10.5)	*		27.0 (4.1) 24.9 (3.8)	Metabochip	≥0.95	-	≥0.95	p >10 ⁻⁶	≥0.01	-	120,509	PLINK	Age and sex	0.99 (QT)
THISEAS	European (Greece)	Cases	327 (229/98) 1,180 (643/537)	63.6 (10.6) 58.2 (13.6)	1	8.0 (2.6) 5.3 (0.6)	29.4 (4.8) 28.3 (5.0)	Metabochip	≥0.95	Duplicates, ambiguous sex, and non- European descent	≥0.98	p>10 ⁻⁴	≥0.01	-	108,868	PLINK	Age and sex	1.00 (QT)
ULSAM	European (Sweden)	Cases Controls	233 (233/0) 942 (942/0)	71.0 (0.6) 71.0 (0.6)			27.7 (3.8) 25.9 (3.2)	Metabochip	≥0.95	Relatedness	≥0.95	p>10 ⁻⁶	≥0.01	-	119,018	PLINK	Age and sex	0.95 (QT)
WARREN2	European	Cases	1,117 (647/470)		45.5 (11.0)	-	32.2 (6.6)	Metabochip	≥0.95	Duplicates, ambiguous sex, and non-	≥0.95 (≥0.99	p>10 ⁻⁶	≥0.01	-	120,521	PLINK	-	1.07 (QT)
	(UK)	Controls	4,224 (2,394/1,830)	-	-	-	-	1		European descent	for MAF<0.05)							