

Study	Ethnic group (origin)	Disease status	Phenotypic characteristics					Genotyping platform	Sample QC			SNP QC			Association analysis			Genomic control
			Sample size (males/females)	Age (years) mean (SD)	Age at onset (years) mean (SD)	Fasting glucose (mmol/l) mean (SD)	BMI (kg/m ²) mean (SD)		Call rate	Exclusion criteria	Call rate	HWE	MAF	Imputation	SNPs	Analysis software	Covariates	
Stage 1																		
ARIC	European	Cases Controls	775 (416/359) 7,159 (3,167/3,992)	56.1 (5.6) 54.0 (5.7)	50.9 (10.4) -	9.3 (3.6) 5.4 (0.4)	30.4 (5.4) 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 (G) 1.01 (I)
deCODE	European (Iceland)	Cases Controls	1,465 (868/597) 5,330 (2,132/3,197)	68.4 (10.1) 53.0 (7.7)	55.1 (12.7) -	8.5 (2.7) 5.0 (0.7)	30.1 (5.4) 26.8 (5.0)	Illumina HumanMap 300/370	≥0.98	Duplicates	≥0.96	$p > 10^{-6}$	≥0.01	proper-info=0.5	2,338,113	SNPTTEST	-	1.31 (G) 1.31 (I)
DGI	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 HumanMap 300	≥0.95	Duplicates	≥0.95	$p > 10^{-6}$	≥0.01	proper-info=0.5	2,051,387	SNPTTEST	-	1.10 (G) 1.10 (I)
VDG	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 SNPTTEST	Age, sex, BMI, and study centre	1.05 (G) 1.06 (I)
EUROSPAN	European	Cases Controls	267 (127/140) 3,710 (1,557/2,153)	63.8 49.9	-	8.0 4.8	29.8 26.5	Illumina HumanMap 300/370	≥0.98	Duplicates	≥0.98	$p > 10^{-6}$	≥0.01	$r^2 > 0.5$	2,359,525	GenABEL and SNPTTEST	Age and sex	0.98 (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 & MIP5 50K	≥0.95	Duplicates	≥0.95	$p > 10^{-6}$	≥0.01	$r^2 > 0.3$	2,389,929	R (IGE 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 (1.1) 5.3 (0.5)	30.2 (4.7) 27.1 (7.4)	Illumina HumanMap 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) -	- 25.0 (2.9)	27.8 (4.0) 20.9 (5.0)	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) 1.03 (I)
KORA Gen	European (Germany)	Cases Controls	433 (255/178) 1,438 (693/745)	60.5 (8.3) 61.0 (8.3)	58.2 (10.3) -	- 27.7 (4.3)	30.9 (5.9) 27.7 (4.3)	Affymetrix GeneChip 500K	≥0.93	Duplicates	≥0.95	$p > 10^{-6}$	≥0.01	proper-info=0.5	2,325,232	SNPTTEST	Age and sex	1.04 (G) 1.04 (I)
NHS	European (USA)	Cases Controls	1,467 (971,467) 1,754 (901,754)	43.5 (6.7) 43.1 (6.8)	58.7 (10.6) -	- 23.5 (0.1)	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) 0.98 (I)
RS1	European (Netherlands)	Cases Controls	1,178 (488/690) 4,761 (1,928/2,833)	71.7 (8.9) 69.0 (8.1)	71.5 (8.9) -	- 26.0 (3.6)	27.4 (4.0) 26.0 (3.6)	Illumina HumanMap 550	≥0.975	Duplicates	≥0.98	$p > 10^{-6}$	≥0.01	$r^2 > 0.5$	2,439,672	GenABEL and SNPTTEST	-	1.01 (G) 1.01 (I)
WTCCC	European (UK)	Cases Controls	1,178 (488/690) 2,538 (1,446/1,492)	58.6 (9.2) 50.3 (9.2)	-	- -	26.7 (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 SNPTTEST	-	1.06 (G) 1.08 (I)
Stage 2																		
AMC-PAS	European (Netherlands)	Cases Controls	48 (35/13) 442 (335/109)	44.0 (4.7) 42.3 (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	$p > 10^{-6}$	≥0.01	-	109,525	PLINK	Age and sex	1.00 (QT) 1.00 (QT)
BHS	European	Cases Controls	51 (38/13) 359 (224/135)	72.3 (7.8) 70.0 (9.6)	-	7.8 (4.0) 5.1 (0.9)	27.2 (3.9) 26.6 (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)<