

Fine-mapping of an expanded set of type 2 diabetes loci to single-variant resolution using high-density imputation and islet-specific epigenome maps

SUPPLEMENTARY INFORMATION

1. Supplementary Figures and Notes

These tables and notes are provided in this document.

i. Supplementary Figures

- **Supplementary Figure 1** | Sex-differentiated analyses.
- **Supplementary Figure 2** | Distributions of the allele frequency, imputation score, and posterior probability of association.
- **Supplementary Figure 3** | Islet annotation overlap of variant with the highest probability in genetic credible sets.
- **Supplementary Figure 4** | Enrichment of islet epigenetic states in T2D GWAS data.
- **Supplementary Figure 5** | Epigenome landscape of *ST6GAL1* locus.
- **Supplementary Figure 6** | Epigenome landscape of *ANK1* locus.
- **Supplementary Figure 7** | Epigenome landscape of *TCF7L2* locus.
- **Supplementary Figure 8** | Heritability estimates.
- **Supplementary Figure 9** | Polygenic risk score.
- **Supplementary Figure 10** | Genetic correlations between T2D and biomedical-relevant traits estimated by LD score regression implemented in LDHub.
- **Supplementary Figure 11** | Impact of BMI adjustment on genetic correlation estimates between various traits and T2D.

ii. Supplementary Tables

- **Supplementary Table 2** | Summary statistics of distinct T2D association signals.
- **Supplementary Table 3** | Summary of comparison of effect estimates from BMI-adjusted and BMI-unadjusted analyses models obtained from the 28 studies the contributed to both analyses.
- **Supplementary Table 4** | Summary of sex-differentiated analysis.
- **Supplementary Table 5** | Summary of 99% credible sets for 380 distinct T2D association signals.
- **Supplementary Table 6** | Summary of comparison of HRC-based and 1000G-based 99% credible sets.
- **Supplementary Table 7** | Coding variants with posterior probability of association >50%.
- **Supplementary Table 8** | Non-coding credible set variants with >80% posterior probability of association.
- **Supplementary Table 9** | *TCF7L2* credible set variants.
- **Supplementary Table 10** | Summary results of LD Score regression analyses between T2D and various diseases, metabolic, and anthropometric traits.

iii. Supplementary Notes

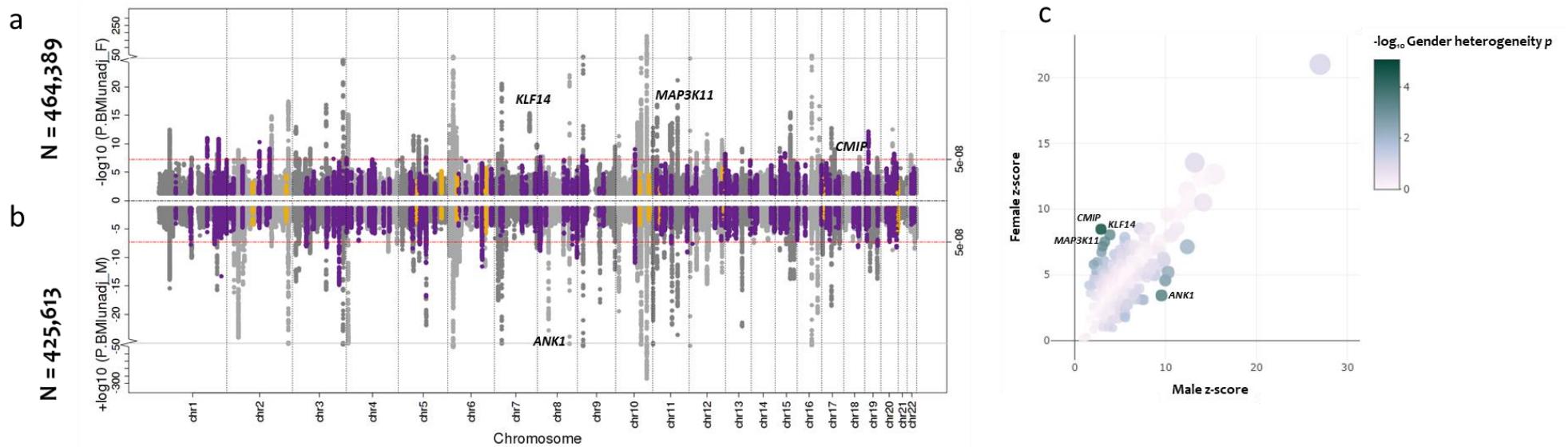
2. Supplementary Tables

This table (excel spreadsheet) is provided in separate document.

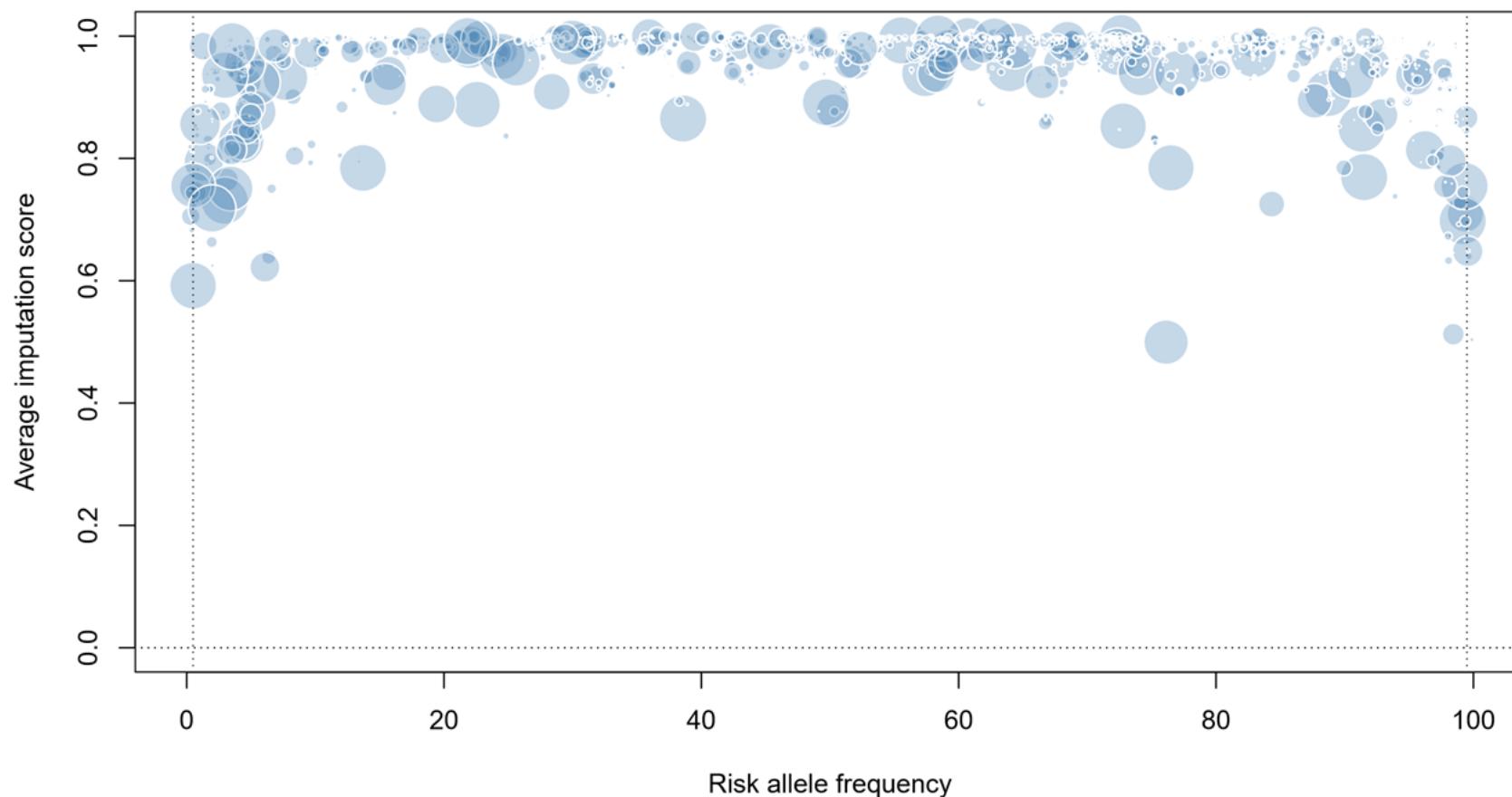
- **Supplementary Table 1** | Study characteristics and analysis details of studies included in the meta-analysis.

i. Supplementary Figures

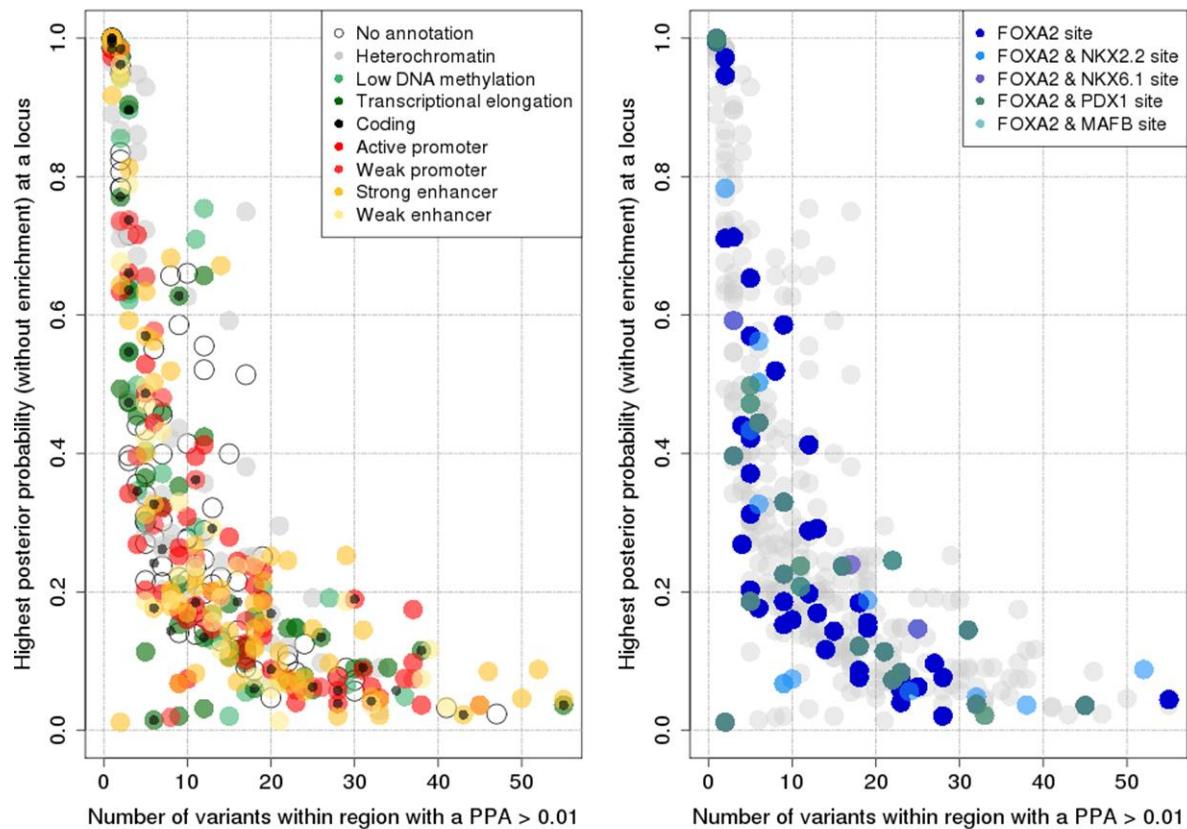
Supplementary Figure 1 | Sex-differentiated analyses. **a**, Manhattan plot (top panel) of genome-wide association results for T2D (without BMI adjustment) from female-specific meta-analysis of up to 30,053 cases and 434,336 controls. The association p -value (on $-\log_{10}$ scale) for each SNP (y-axis) is plotted against the genomic position (NCBI Build 37; x-axis). Association signals that reached genome-wide significance ($p < 5 \times 10^{-8}$) in sex-combined analysis are shown in purple or yellow, if novel. **b**, Manhattan plot (bottom panel) of genome-wide association results for T2D with BMI adjustment from male-specific meta-analysis of up to 41,846 cases and 383,767 controls. **c**, Z-score for each of the 403 distinct signals from male-specific analysis (y-axis) is plotted against the z-score from the female-specific analysis (x-axis). Colour of each point varies with $-\log_{10}$ gender heterogeneity p -value and diameter of the circle is proportional to sex-combined $-\log_{10} p$ -value.



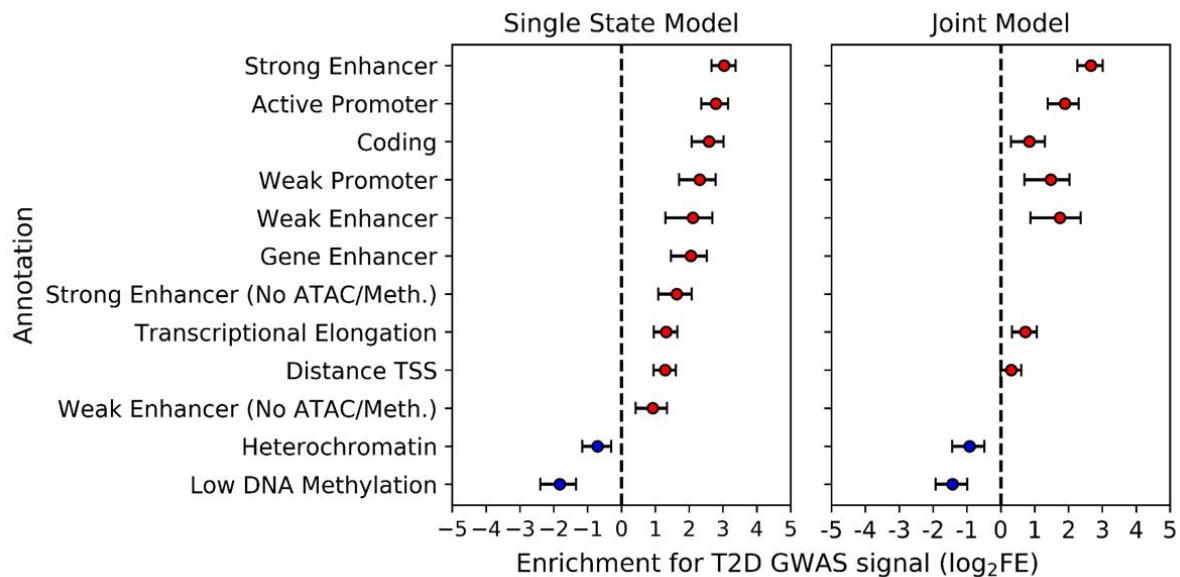
Supplementary Figure 2 | Distributions of the allele frequency, imputation score, and posterior probability of association. Distribution of the risk allele frequencies for all variants having >1% posterior probability of association in genetic credible set (x-axis) plotted against average imputation quality (y-axis). Diameter varies with the posterior probability of association assigned to each variant.



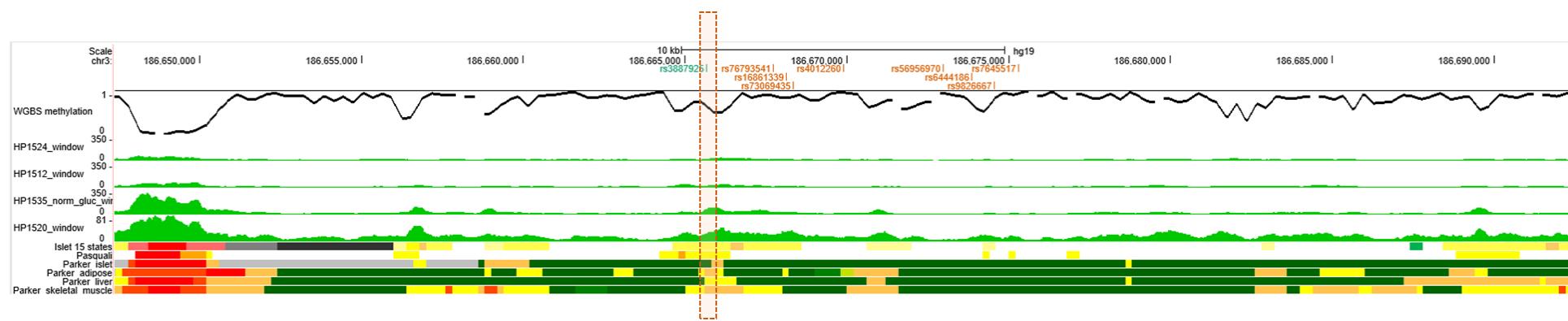
Supplementary Figure 3 | Islet annotation overlap of variant with the highest probability in genetic credible sets. Number of variants with posterior probability of association >1% (x-axis) plotted against the highest posterior probability of association (y-axis) assigned to a variant in the credible set. Points are colour coded according to islet epigenome states.



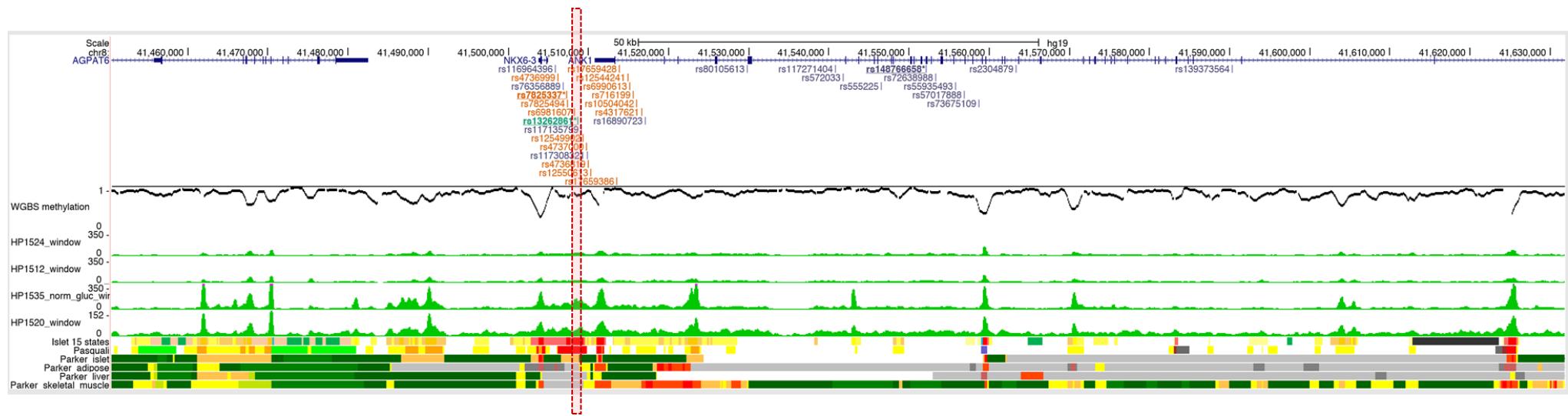
Supplementary Figure 4 | Enrichment of islet epigenetic states in T2D GWAS data. fGWAS log₂ fold enrichment including 95% confidence intervals (x-axis) of all chromatin states (y-axis) genome-wide.



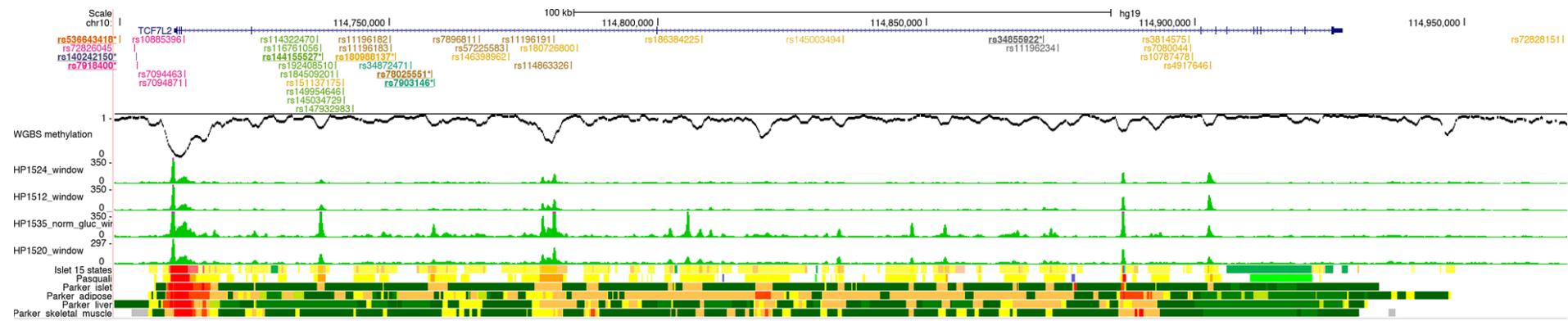
Supplementary Figure 5 | Epigenome landscape of *ST6GAL1* locus. For variants included in 99% credible set (PPA>1%) of each distinct signal at *ST6GAL1* locus, following information is shown: genomic position of each variant (colour coded for each distinct signal; variant with highest PPA in bold); whole genome bisulphite methylation data (black), 4 human islet ATAC-seq tracks (green, middle), islet chromatin states (from Thurner et al, Pasquali et al 2013, and Varshney et al 2017); and adipose, liver and skeletal muscle chromatin states from Varshney et al 2017.



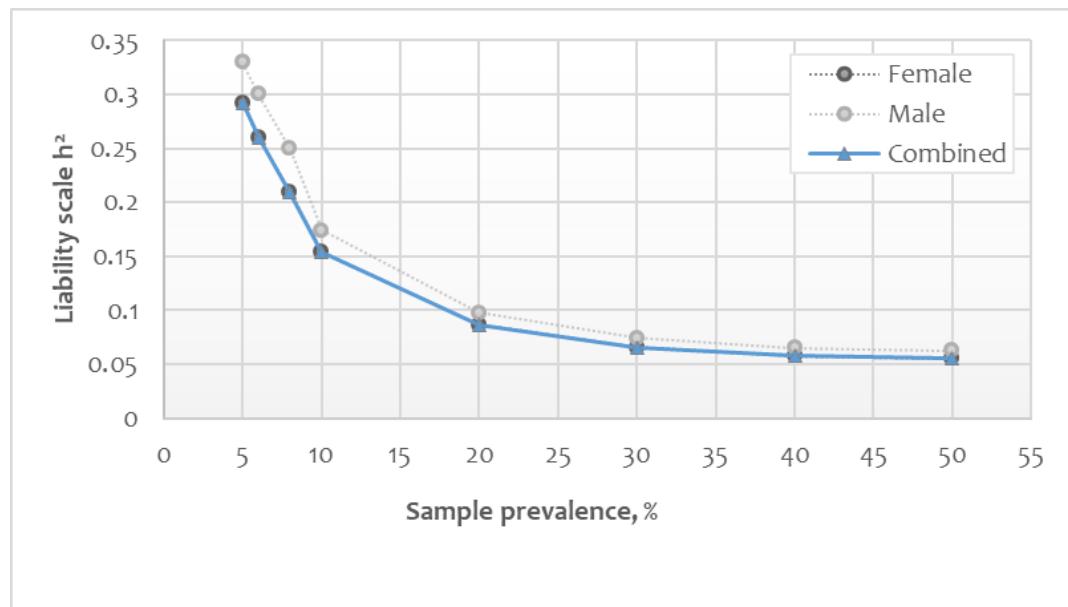
Supplementary Figure 6 | Epigenome landscape of ANK1 locus. For variants included in 99% credible set (PPA>1%) of each distinct signal at ANK1 locus, following information is shown: genomic position of each variant (colour coded for each distinct signal; variant with highest PPA in bold); whole genome bisulphite methylation data (black), 4 human islet ATAC-seq tracks (green, middle), islet chromatin states (from Thurner et al, Pasquali et al 2013, and Varshney et al 2017); and adipose, liver and skeletal muscle chromatin states from Varshney et al 2017.



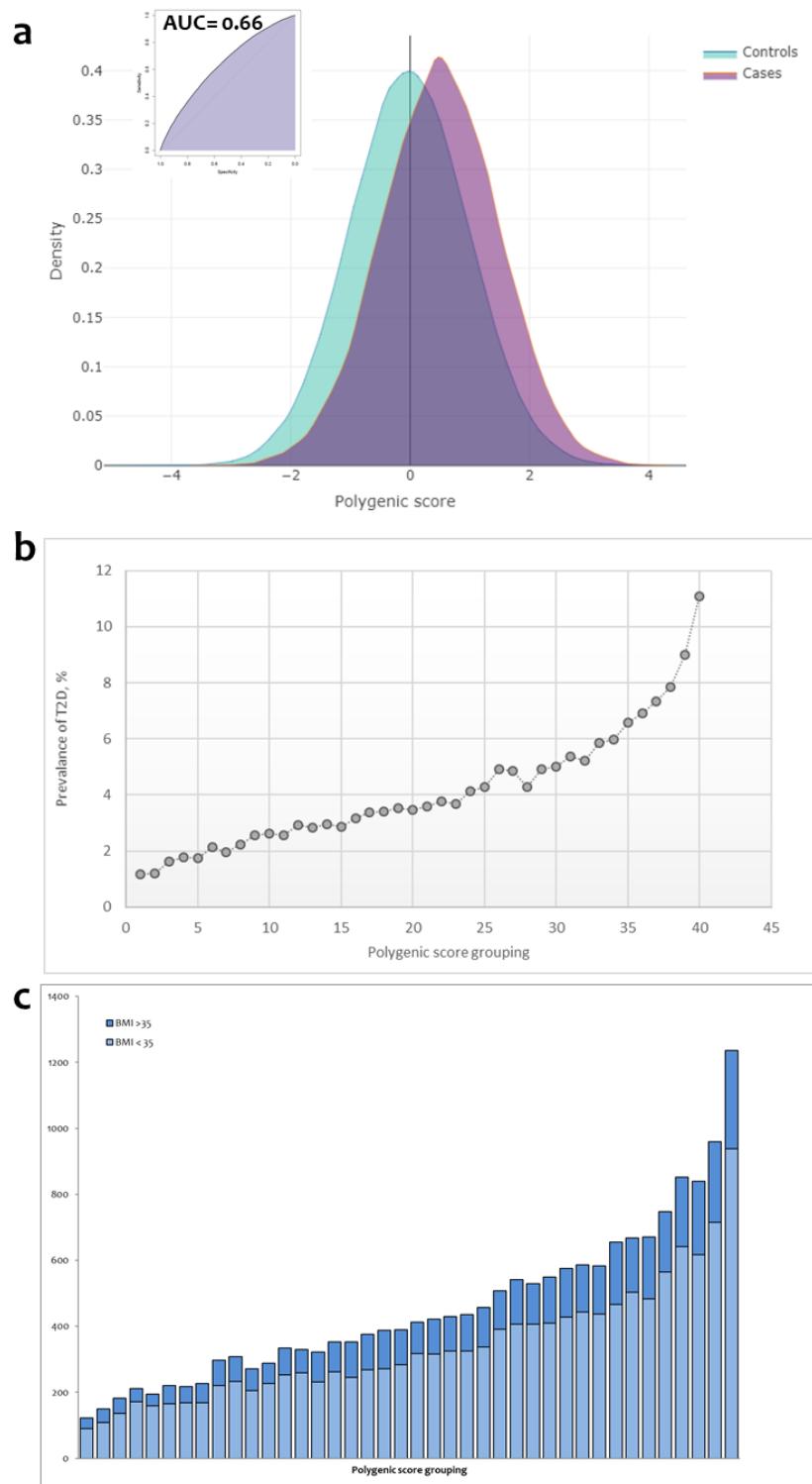
Supplementary Figure 7 | Epigenome landscape of *TCF7L2* locus. For variants included in 99% credible set (PPA>1%) of each distinct signal at *TCF7L2* locus, following information is shown: genomic position of each variant (colour coded for each distinct signal; variant with highest PPA in bold); whole genome bisulphite methylation data (black), 4 human islet ATAC-seq tracks (green, middle), islet chromatin states (from Thurner et al, Pasquali et al 2013, and Varshney et al 2017); and adipose, liver and skeletal muscle chromatin states from Varshney et al 2017.



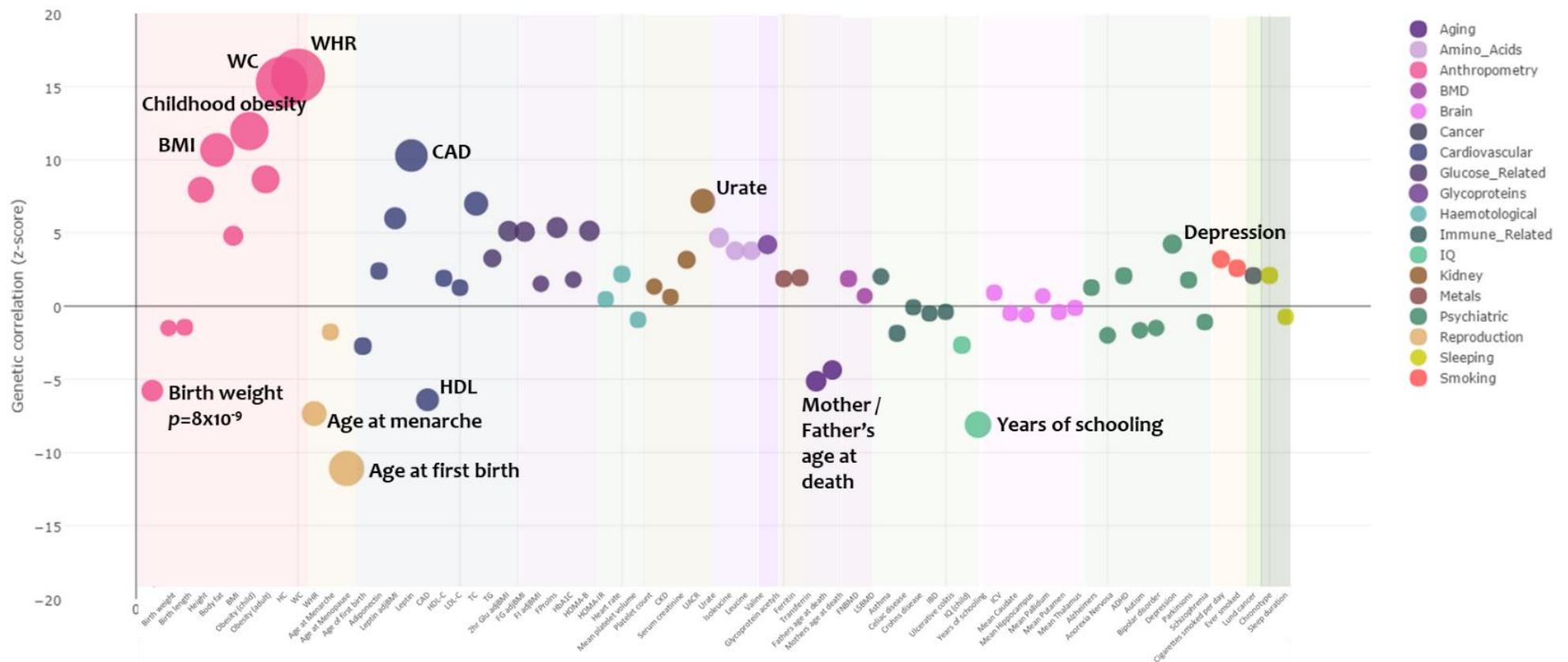
Supplementary Figure 8 | Heritability estimates.



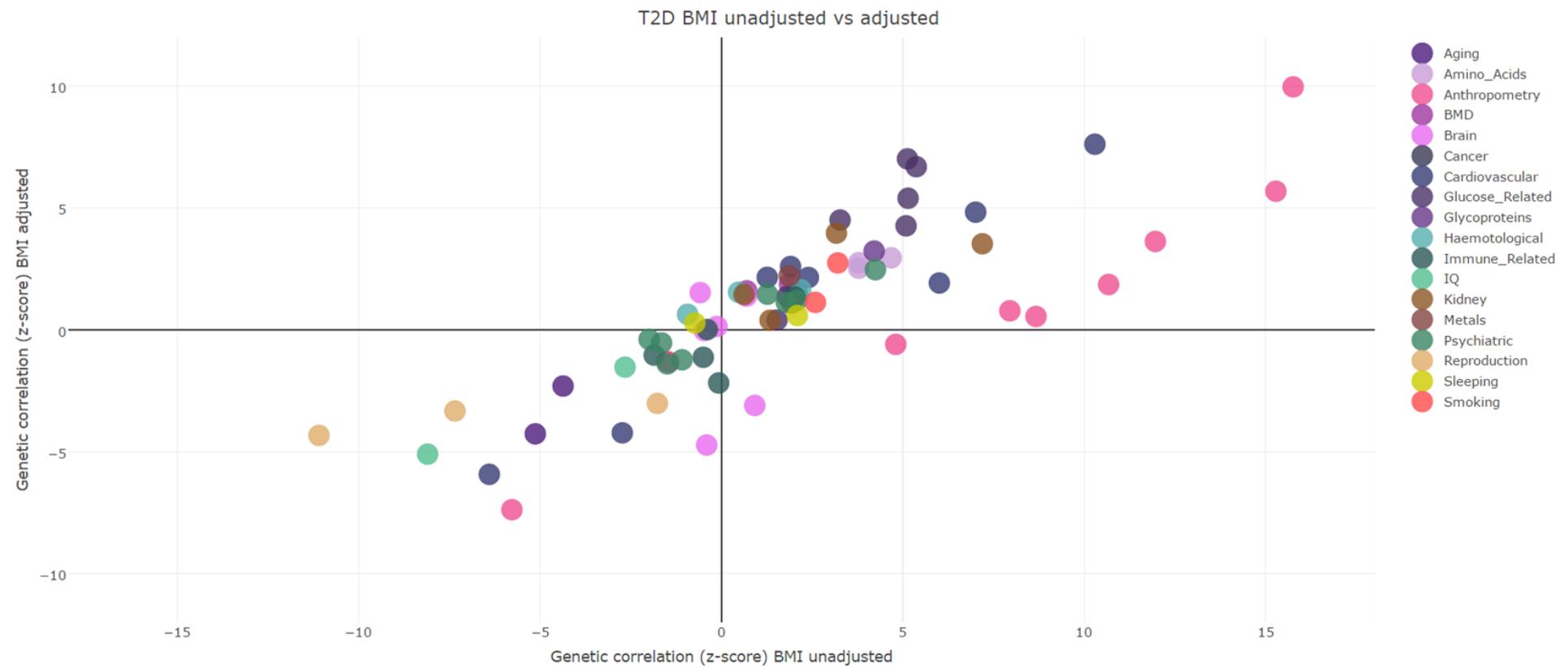
Supplementary Figure 9 | Polygenic risk score. Genome-wide polygenic risk score (PRS) identifies individuals with significantly increased risk of T2D. **a**, PRS in UK Biobank individuals is normally distributed with a shift towards right, observed for T2D cases. PRS is plotted on the x-axis, with values scaled to a mean of 0 and standard deviation of 1. **b**, Individuals were binned into 40 groups based on PRS, with each grouping representing 2.5% of population. **c**, BMI distribution within each bin.



Supplementary Figure 10 | Genetic correlations between T2D and biomedical-relevant traits estimated by LD score regression implemented in LDHub. Genetic correlations (z-score) between T2D (y-axis) and range of metabolic and anthropometric traits (x-axis) as estimated using LD Score regression. The genetic correlation estimates are colour coded according to phenotypic area. Allelic direction of effect is aligned to increased T2D risk. Size of the circle denotes the significance level for the correlation.



Supplementary Figure 11 | Impact of BMI adjustment on genetic correlation estimates between various traits and T2D.



ii. Supplementary Tables

Supplementary Table 2 | Summary statistics of distinct T2D association signals.

| Nearest gene | Index variant | Chromosome | Position (Build 37 bp) | Risk allele | Other allele | RAF (%) | MAF (%) | p-value | OR (95% CI) | BMI adjusted | Novel | Primary | Cases | Controls | Mean imputation quality score |
|---------------------|---------------|------------|------------------------|-------------|--------------|---------|---------|-----------------------|------------------|--------------|-------|---------|--------|----------|-------------------------------|
| <i>MACF1</i> | rs3768321 | 1 | 40,035,928 | T | G | 20.0 | 20.0 | 2.6x10 ⁻²⁶ | 1.09 (1.07-1.10) | N | N | Y | 74,117 | 823,996 | 0.98 |
| <i>FAF1</i> | rs58432198 | 1 | 51,256,091 | C | T | 88.1 | 11.9 | 2.1x10 ⁻¹⁰ | 1.07 (1.05-1.09) | N | N | Y | 74,037 | 819,489 | 0.94 |
| <i>PATJ</i> | rs12140153 | 1 | 62,579,891 | G | T | 90.5 | 9.49 | 1.3x10 ⁻⁸ | 1.07 (1.04-1.09) | N | Y | Y | 74,117 | 823,997 | 0.94 |
| <i>DENND2C</i> | rs184660829 | 1 | 115,144,899 | C | T | 0.0200 | 0.0200 | 2.5x10 ⁻⁸ | 8.05 (3.86-16.8) | N | Y | Y | 33,287 | 471,082 | 0.69 |
| <i>PTGFRN</i> | rs1127215 | 1 | 117,532,790 | C | T | 58.4 | 41.6 | 1.6x10 ⁻¹³ | 1.05 (1.04-1.06) | N | Y | Y | 74,117 | 823,995 | 1.00 |
| <i>NOTCH2</i> | rs1493694 | 1 | 120,526,982 | T | C | 10.9 | 10.9 | 2.7x10 ⁻¹⁶ | 1.09 (1.07-1.11) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>FAM63A</i> | rs10305745 | 1 | 150,786,038 | A | G | 1.45 | 1.45 | 4.4x10 ⁻⁶ | 1.28 (1.15-1.42) | N | N | N | 73,395 | 821,537 | 0.91 |
| <i>FAM63A</i> | rs145904381 | 1 | 151,017,991 | T | C | 98.7 | 1.33 | 2.6x10 ⁻⁸ | 1.19 (1.12-1.26) | N | N | Y | 74,117 | 823,997 | 0.79 |
| <i>SEC16B</i> | rs539515 | 1 | 177,889,025 | C | A | 19.8 | 19.8 | 1.6x10 ⁻¹⁰ | 1.05 (1.04-1.07) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>DSTYK</i> | rs12048743 | 1 | 205,114,873 | G | C | 44.2 | 44.2 | 3.5x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,996 | 1.00 |
| <i>SRGP2</i> | rs9430095 | 1 | 206,593,900 | C | G | 49.4 | 49.4 | 1.9x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 0.97 |
| <i>PROX1</i> | rs79687284 | 1 | 214,150,821 | C | G | 3.48 | 3.48 | 2.6x10 ⁻¹⁶ | 1.16 (1.12-1.21) | N | N | N | 74,117 | 823,997 | 0.81 |
| <i>PROX1</i> | rs340874 | 1 | 214,159,256 | C | T | 55.6 | 44.5 | 1.6x10 ⁻²² | 1.07 (1.05-1.08) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>PROX1</i> | rs114526150 | 1 | 214,175,531 | G | T | 2.25 | 2.25 | 4.2x10 ⁻⁷ | 1.12 (1.07-1.17) | N | N | N | 74,116 | 823,997 | 0.97 |
| <i>LYPLAL1</i> | rs553014999 | 1 | 219,584,164 | C | T | 0.130 | 0.130 | 7.7x10 ⁻⁶ | 1.90 (1.43-2.51) | N | Y | N | 45,408 | 756,103 | 0.67 |
| <i>LYPLAL1</i> | rs2820446 | 1 | 219,748,818 | C | G | 70.6 | 29.5 | 3.3x10 ⁻¹⁶ | 1.06 (1.04-1.07) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>ABCB10</i> | rs348330 | 1 | 229,672,955 | G | A | 36.1 | 36.1 | 2.7x10 ⁻¹⁴ | 1.05 (1.04-1.07) | N | Y | Y | 74,115 | 823,996 | 0.95 |
| <i>GNG4</i> | rs291367 | 1 | 235,690,800 | G | A | 63.2 | 36.8 | 4.7x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 62,669 | 545,620 | 1.00 |
| <i>TMEM18</i> | rs62107261 | 2 | 422,144 | T | C | 95.4 | 4.64 | 3.8x10 ⁻¹² | 1.12 (1.08-1.15) | N | N | Y | 74,117 | 823,997 | 0.85 |
| <i>TMEM18</i> | rs35913461 | 2 | 653,575 | C | T | 82.9 | 17.1 | 1.6x10 ⁻¹¹ | 1.06 (1.04-1.08) | N | N | N | 74,117 | 823,996 | 0.99 |
| <i>FAM49A</i> | rs11680058 | 2 | 16,574,669 | A | G | 86.3 | 13.7 | 1.4x10 ⁻⁸ | 1.06 (1.04-1.08) | N | Y | Y | 74,115 | 823,995 | 0.78 |
| <i>DTNB</i> | rs17802463 | 2 | 25,643,221 | G | T | 73.1 | 26.9 | 2.9x10 ⁻⁸ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>GCKR</i> | rs1260326 | 2 | 27,730,940 | C | T | 60.7 | 39.3 | 6.5x10 ⁻²⁵ | 1.07 (1.06-1.08) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>THADA</i> | rs28525376 | 2 | 43,207,872 | G | T | 42.2 | 42.2 | 2.7x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,116 | 823,995 | 0.99 |
| <i>THADA</i> | rs6708643 | 2 | 43,430,440 | A | G | 50.1 | 49.9 | 3.9x10 ⁻⁸ | 1.04 (1.02-1.05) | N | N | N | 74,116 | 823,995 | 0.97 |
| <i>THADA</i> | rs80147536 | 2 | 43,698,028 | A | T | 90.4 | 9.57 | 2.7x10 ⁻²⁹ | 1.13 (1.11-1.16) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>BNIPL</i> | rs10193538 | 2 | 58,981,064 | T | G | 61.0 | 39.0 | 8.9x10 ⁻⁹ | 1.04 (1.02-1.05) | N | Y | N | 74,116 | 823,996 | 1.00 |
| <i>BNIPL</i> | rs6545714 | 2 | 59,307,725 | G | A | 39.2 | 39.2 | 8.9x10 ⁻⁹ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>BCL11A</i> | rs243024 | 2 | 60,583,665 | A | G | 46.0 | 46.0 | 2.5x10 ⁻²⁰ | 1.06 (1.05-1.07) | N | N | Y | 74,116 | 823,996 | 1.00 |
| <i>CEP68</i> | rs2249105 | 2 | 65,287,896 | A | G | 63.4 | 36.6 | 2.2x10 ⁻¹⁴ | 1.10 (1.08-1.13) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>CEP68</i> | rs2052261 | 2 | 65,355,270 | G | A | 30.4 | 30.4 | 2.5x10 ⁻⁶ | 1.07 (1.04-1.09) | N | N | N | 74,116 | 823,996 | 1.00 |
| <i>CEP68</i> | rs2028150 | 2 | 65,655,012 | C | G | 59.8 | 40.2 | 2.3x10 ⁻¹² | 1.05 (1.03-1.06) | N | N | N | 74,117 | 823,997 | 1.00 |
| <i>TMEM127</i> | rs79046683 | 2 | 96,913,918 | T | G | 0.480 | 0.480 | 3.0x10 ⁻⁸ | 2.34 (1.73-3.16) | Y | Y | Y | 34,775 | 477,429 | 0.91 |
| <i>DDX18</i> | rs562386202 | 2 | 118,071,061 | G | A | 0.0600 | 0.0600 | 4.2x10 ⁻⁸ | 3.20 (2.11-4.86) | N | Y | Y | 26,326 | 457,160 | 0.72 |
| <i>GLI2</i> | rs11688931 | 2 | 121,318,166 | C | G | 84.9 | 15.1 | 8.2x10 ⁻⁶ | 1.04 (1.02-1.06) | N | Y | N | 74,117 | 823,997 | 0.98 |
| <i>GLI2</i> | rs11688682 | 2 | 121,347,612 | G | C | 72.8 | 27.2 | 4.2x10 ⁻⁹ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,996 | 0.85 |
| <i>GLI2</i> | rs66477705 | 2 | 121,378,852 | T | C | 96.7 | 3.30 | 2.6x10 ⁻⁶ | 1.09 (1.05-1.13) | N | Y | N | 74,117 | 823,997 | 0.95 |
| <i>PABPC1P2</i> | rs35999103 | 2 | 147,861,633 | T | C | 15.5 | 15.5 | 9.7x10 ⁻⁹ | 1.05 (1.03-1.07) | N | Y | Y | 74,116 | 823,997 | 1.00 |
| <i>CYTIP</i> | rs13426680 | 2 | 158,339,550 | A | G | 93.7 | 6.27 | 6.7x10 ⁻¹⁰ | 1.09 (1.06-1.11) | N | Y | Y | 74,117 | 823,996 | 0.99 |
| <i>RBMS1</i> | rs3772071 | 2 | 161,135,544 | T | C | 71.4 | 28.7 | 1.2x10 ⁻¹¹ | 1.05 (1.03-1.06) | N | N | Y | 74,115 | 823,995 | 1.00 |
| <i>GRB14/COBLL1</i> | rs10195252 | 2 | 165,513,091 | T | C | 58.6 | 41.4 | 6.0x10 ⁻²⁵ | 1.07 (1.06-1.08) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>GRB14/COBLL1</i> | rs13024606 | 2 | 165,573,194 | T | C | 4.72 | 4.72 | 1.7x10 ⁻⁸ | 1.09 (1.06-1.13) | N | N | N | 74,117 | 823,997 | 0.93 |
| <i>CRYBA2</i> | rs113414093 | 2 | 219,859,171 | A | G | 5.14 | 5.14 | 6.6x10 ⁻⁹ | 1.12 (1.08-1.17) | Y | Y | Y | 50,402 | 523,888 | 0.61 |
| <i>IRS1</i> | rs2972144 | 2 | 227,101,411 | G | A | 63.9 | 36.2 | 2.1x10 ⁻⁴⁶ | 1.10 (1.08-1.11) | N | N | Y | 74,115 | 823,996 | 1.00 |
| <i>PPARG</i> | rs11709077 | 3 | 12,336,507 | G | A | 87.7 | 12.4 | 1.8x10 ⁻³⁶ | 1.14 (1.11-1.16) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>PPARG</i> | rs17819328 | 3 | 12,489,342 | G | T | 42.5 | 42.5 | 4.8x10 ⁻¹⁶ | 1.06 (1.04-1.07) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>UBE2E2</i> | rs35352848 | 3 | 23,455,582 | T | C | 78.8 | 21.2 | 1.3x10 ⁻¹⁷ | 1.07 (1.05-1.09) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>UBE2E2</i> | rs17013314 | 3 | 23,510,044 | G | A | 3.13 | 3.13 | 8.4x10 ⁻⁹ | 1.11 (1.07-1.15) | N | N | N | 74,116 | 823,996 | 0.97 |
| <i>KIF9</i> | rs11926707 | 3 | 46,925,539 | C | T | 62.6 | 37.4 | 2.1x10 ⁻⁸ | 1.27 (1.17-1.38) | N | N | Y | 74,115 | 823,995 | 0.95 |
| <i>KIF9</i> | rs75423501 | 3 | 47,242,923 | G | A | 10.1 | 10.1 | 7.5x10 ⁻⁶ | 1.05 (1.03-1.08) | N | N | N | 74,036 | 819,488 | 0.78 |
| <i>RBM6</i> | rs4688760 | 3 | 49,980,596 | T | C | 68.4 | 31.6 | 3.5x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,996 | 0.93 |
| <i>RFT1</i> | rs2581787 | 3 | 53,127,677 | T | G | 56.3 | 43.7 | 2.4x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>CACNA2D3</i> | rs76263492 | 3 | 54,828,827 | T | G | 4.52 | 4.52 | 6.3x10 ⁻⁹ | 1.09 (1.06-1.13) | N | Y | Y | 74,117 | 823,996 | 0.95 |
| <i>PSMD6</i> | rs3774723 | 3 | 63,962,339 | G | A | 84.4 | 15.6 | 1.6x10 ⁻¹³ | 1.07 (1.05-1.09) | N | N | Y | 74,117 | 823,996 | 0.99 |
| <i>PSMD6</i> | rs74368513 | 3 | 64,460,694 | G | A | 99.6 | 0.440 | 7.0x10 ⁻⁶ | 1.31 (1.16-1.47) | N | N | N | 72,923 | 820,276 | 0.65 |

| ADAMTS9 | rs9860730 | 3 | 64,701,146 | A | G | 70.4 | 29.6 | 4.9x10 ⁻¹⁵ | 1.06 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 1.00 |
|-----------------|-------------|---|-------------|---|---|-------|-------|-----------------------|------------------|---|---|---|--------|---------|------|
| <i>SHQ1</i> | rs13085136 | 3 | 72,865,183 | C | T | 92.8 | 7.17 | 1.5x10 ⁻⁸ | 1.08 (1.05-1.10) | N | Y | Y | 74,117 | 823,997 | 0.87 |
| <i>ROBO2</i> | rs2272163 | 3 | 77,671,721 | C | A | 61.8 | 38.2 | 9.6x10 ⁻⁹ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 0.98 |
| <i>ADCY5</i> | rs11708067 | 3 | 123,065,778 | A | G | 77.2 | 22.8 | 5.2x10 ⁻³² | 1.09 (1.08-1.11) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>SLC12A8</i> | rs649961 | 3 | 124,926,637 | T | C | 46.5 | 46.5 | 9.9x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | Y | Y | 74,115 | 823,996 | 0.99 |
| <i>TMCC1</i> | rs9828772 | 3 | 129,333,182 | C | G | 89.8 | 10.2 | 4.2x10 ⁻⁸ | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>TMCC1</i> | rs559138871 | 3 | 129,470,067 | T | C | 0.200 | 0.200 | 7.3x10 ⁻⁶ | 1.49 (1.25-1.77) | N | Y | N | 67,516 | 797,362 | 0.84 |
| <i>TSC22D2</i> | rs62271373 | 3 | 150,066,540 | A | T | 5.53 | 5.53 | 1.0x10 ⁻⁹ | 1.09 (1.06-1.12) | N | Y | Y | 74,117 | 823,997 | 0.92 |
| <i>MBNL1</i> | rs13065698 | 3 | 152,086,533 | A | G | 60.0 | 40.0 | 8.1x10 ⁻¹³ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>MBNL1</i> | rs74653713 | 3 | 152,417,881 | C | A | 95.7 | 4.29 | 1.2x10 ⁻⁸ | 1.10 (1.06-1.13) | N | Y | N | 74,116 | 823,997 | 0.99 |
| <i>MBNL1</i> | rs35497231 | 3 | 152,433,628 | C | T | 31.7 | 31.7 | 7.6x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | N | 74,115 | 823,995 | 1.00 |
| <i>EGFEM1P</i> | rs7629630 | 3 | 168,218,841 | A | T | 85.7 | 14.3 | 2.5x10 ⁻⁸ | 1.05 (1.03-1.07) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>SLC2A2</i> | rs9873618 | 3 | 170,733,076 | G | A | 71.0 | 29.0 | 4.8x10 ⁻²¹ | 1.07 (1.05-1.08) | N | Y | Y | 74,117 | 823,996 | 0.99 |
| <i>ABCC5</i> | rs2872246 | 3 | 183,738,460 | A | C | 45.4 | 45.4 | 1.5x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,117 | 823,996 | 1.00 |
| <i>IGF2BP2</i> | rs6780171 | 3 | 185,503,456 | A | T | 31.4 | 31.4 | 9.0x10 ⁻⁵⁶ | 1.14 (1.12-1.16) | N | N | Y | 74,115 | 823,996 | 1.00 |
| <i>IGF2BP2</i> | rs150111048 | 3 | 185,514,421 | G | A | 23.9 | 23.9 | 2.7x10 ⁻⁷ | 1.12 (1.07-1.16) | N | N | N | 69,270 | 800,165 | 0.50 |
| <i>IGF2BP2</i> | rs11717959 | 3 | 185,541,213 | G | T | 62.1 | 37.9 | 3.0x10 ⁻⁶ | 1.04 (1.02-1.06) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>IGF2BP2</i> | rs1516728 | 3 | 185,829,891 | A | T | 75.9 | 24.1 | 6.3x10 ⁻⁶ | 1.03 (1.02-1.05) | N | N | N | 74,115 | 823,995 | 0.99 |
| <i>ST6GAL1</i> | rs3887925 | 3 | 186,665,645 | T | C | 54.7 | 45.3 | 3.1x10 ⁻²² | 1.07 (1.05-1.08) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>ST6GAL1</i> | rs7645517 | 3 | 186,675,277 | A | G | 5.76 | 5.76 | 2.5x10 ⁻⁸ | 1.08 (1.05-1.11) | N | N | N | 74,117 | 823,996 | 0.96 |
| <i>LPP</i> | rs4686471 | 3 | 187,740,899 | C | T | 61.0 | 39.0 | 1.7x10 ⁻²⁰ | 1.06 (1.05-1.08) | N | N | Y | 74,115 | 823,995 | 1.00 |
| <i>PCGF3</i> | rs111827885 | 4 | 616,608 | C | T | 1.56 | 1.56 | 8.4x10 ⁻⁶ | 1.18 (1.10-1.27) | N | Y | N | 57,868 | 533,395 | 0.51 |
| <i>PCGF3</i> | rs1531583 | 4 | 744,972 | T | G | 4.58 | 4.58 | 3.5x10 ⁻¹⁴ | 1.13 (1.09-1.16) | N | Y | Y | 74,117 | 823,997 | 0.95 |
| <i>PCGF3</i> | rs35654957 | 4 | 1,010,077 | C | T | 36.7 | 36.7 | 4.2x10 ⁻⁷ | 1.03 (1.02-1.05) | N | Y | N | 74,116 | 823,996 | 0.94 |
| <i>MAEA</i> | rs56337234 | 4 | 1,784,403 | C | T | 50.3 | 49.7 | 8.6x10 ⁻¹⁸ | 1.06 (1.04-1.07) | N | N | Y | 74,116 | 823,995 | 0.88 |
| <i>HTT</i> | rs362307 | 4 | 3,241,845 | T | C | 7.68 | 7.68 | 1.1x10 ⁻⁹ | 1.08 (1.05-1.10) | N | Y | Y | 74,117 | 823,997 | 0.93 |
| <i>WFS1</i> | rs1801212 | 4 | 6,302,519 | A | G | 70.9 | 29.1 | 4.5x10 ⁻⁶ | 1.05 (1.03-1.07) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>WFS1</i> | rs10937721 | 4 | 6,306,763 | C | G | 58.8 | 41.2 | 1.5x10 ⁻⁸ | 1.06 (1.04-1.08) | N | N | Y | 74,115 | 823,995 | 0.97 |
| <i>LCORL</i> | rs12640250 | 4 | 17,792,869 | C | A | 71.5 | 28.5 | 3.7x10 ⁻⁸ | 1.04 (1.03-1.05) | N | Y | Y | 74,115 | 823,995 | 0.98 |
| <i>GNPDA2</i> | rs10938398 | 4 | 45,186,139 | A | G | 42.9 | 42.9 | 3.6x10 ⁻¹² | 1.05 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 0.99 |
| <i>USP46</i> | rs2102278 | 4 | 52,818,664 | G | A | 31.9 | 31.9 | 3.7x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,995 | 0.95 |
| <i>USP46</i> | rs114447556 | 4 | 53,207,093 | T | C | 8.39 | 8.39 | 8.4x10 ⁻⁶ | 1.06 (1.03-1.08) | N | Y | N | 74,116 | 823,997 | 0.80 |
| <i>SCD5</i> | rs12642790 | 4 | 83,578,271 | A | G | 33.8 | 33.8 | 4.4x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>FAM13A</i> | rs1903002 | 4 | 89,740,894 | G | C | 50.1 | 50.0 | 2.7x10 ⁻⁸ | 1.04 (1.02-1.05) | N | N | Y | 74,115 | 823,995 | 0.98 |
| <i>FAM13A</i> | rs576406049 | 4 | 89,857,291 | T | C | 0.130 | 0.130 | 1.6x10 ⁻⁶ | 1.65 (1.35-2.03) | N | N | N | 67,914 | 802,131 | 0.63 |
| <i>SMARCAD1</i> | rs6821438 | 4 | 95,091,911 | A | G | 53.4 | 46.6 | 4.0x10 ⁻¹¹ | 1.04 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>SLC9B1</i> | rs1580278 | 4 | 104,140,848 | C | A | 47.3 | 47.3 | 2.2x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | Y | Y | 74,115 | 823,995 | 0.95 |
| <i>PABPC4L</i> | rs1296328 | 4 | 137,083,193 | A | C | 44.6 | 44.6 | 3.5x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,995 | 0.97 |
| <i>TMEM154</i> | rs7669833 | 4 | 153,513,369 | T | A | 70.5 | 29.6 | 1.2x10 ⁻¹⁴ | 1.06 (1.04-1.07) | N | N | Y | 74,116 | 823,997 | 0.99 |
| <i>PDGFC</i> | rs28819812 | 4 | 157,652,753 | C | A | 67.7 | 32.3 | 2.2x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 62,667 | 545,622 | 1.00 |
| <i>ACSL1</i> | rs58730668 | 4 | 185,717,759 | T | C | 85.8 | 14.2 | 1.3x10 ⁻¹³ | 1.07 (1.05-1.09) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>ANKH</i> | rs3845281 | 5 | 14,610,134 | G | A | 90.4 | 9.61 | 2.3x10 ⁻¹¹ | 1.08 (1.06-1.10) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>ANKH</i> | rs78408340 | 5 | 14,751,305 | C | T | 99.4 | 0.620 | 7.8x10 ⁻¹³ | 1.41 (1.28-1.55) | N | N | Y | 71,295 | 813,971 | 0.71 |
| <i>ANKH</i> | rs17250977 | 5 | 14,753,745 | G | A | 3.76 | 3.76 | 2.0x10 ⁻¹¹ | 1.12 (1.09-1.16) | N | N | N | 74,117 | 823,997 | 0.81 |
| <i>ANKH</i> | rs6885132 | 5 | 14,768,092 | C | G | 90.4 | 9.60 | 1.7x10 ⁻⁸ | 1.07 (1.04-1.09) | N | N | N | 74,117 | 823,997 | 0.97 |
| <i>ANKH</i> | rs76549217 | 5 | 14,768,766 | T | C | 2.95 | 2.95 | 3.0x10 ⁻¹⁰ | 1.14 (1.10-1.19) | N | N | N | 74,117 | 823,997 | 0.73 |
| <i>MRPS30</i> | rs62368490 | 5 | 44,534,364 | T | C | 3.13 | 3.13 | 3.4x10 ⁻⁶ | 1.10 (1.06-1.14) | N | Y | N | 74,038 | 819,490 | 0.77 |
| <i>MRPS30</i> | rs6884702 | 5 | 44,682,589 | G | A | 39.3 | 39.3 | 1.5x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 74,037 | 819,490 | 0.99 |
| <i>ITGA1</i> | rs17261179 | 5 | 51,791,225 | T | C | 51.7 | 48.3 | 1.3x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | N | 74,116 | 823,996 | 1.00 |
| <i>ITGA1</i> | rs3811978 | 5 | 52,100,489 | G | A | 16.7 | 16.7 | 7.7x10 ⁻¹¹ | 1.06 (1.04-1.07) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>ITGA1</i> | rs62357230 | 5 | 52,315,682 | A | G | 3.39 | 3.39 | 5.9x10 ⁻⁶ | 1.09 (1.05-1.13) | N | Y | N | 74,117 | 823,997 | 0.99 |
| <i>ARL15</i> | rs62370480 | 5 | 52,774,510 | A | G | 22.0 | 22.0 | 2.0x10 ⁻⁶ | 1.04 (1.02-1.05) | N | N | N | 74,116 | 823,997 | 0.99 |
| <i>ARL15</i> | rs702634 | 5 | 53,271,420 | A | G | 69.0 | 31.0 | 7.7x10 ⁻¹⁴ | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>ARL15</i> | rs279744 | 5 | 53,412,620 | C | A | 69.1 | 30.9 | 3.1x10 ⁻⁸ | 1.04 (1.03-1.05) | N | N | N | 74,117 | 823,997 | 1.00 |
| <i>ANKRD55</i> | rs465002 | 5 | 55,808,475 | T | C | 74.2 | 25.8 | 6.1x10 ⁻³⁸ | 1.11 (1.09-1.12) | N | N | Y | 74,115 | 823,995 | 0.99 |
| <i>ANKRD55</i> | rs2431115 | 5 | 55,848,669 | A | G | 40.2 | 40.2 | 3.9x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | N | N | 74,115 | 823,995 | 0.99 |
| <i>ANKRD55</i> | rs9687832 | 5 | 55,861,595 | A | G | 19.8 | 19.8 | 1.7x10 ⁻²⁰ | 1.08 (1.06-1.10) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>ANKRD55</i> | rs96844 | 5 | 56,196,604 | G | A | 26.2 | 26.2 | 5.4x10 ⁻⁸ | 1.04 (1.03-1.05) | N | N | N | 74,116 | 823,995 | 1.00 |
| <i>PIK3R1</i> | rs4976033 | 5 | 67,714,246 | G | A | 41.1 | 41.1 | 1.0x10 ⁻⁹ | 1.05 (1.03-1.06) | Y | Y | Y | 50,401 | 523,887 | 0.96 |
| <i>POC5</i> | rs2307111 | 5 | 75,003,678 | T | C | 60.5 | 39.5 | 2.1x10 ⁻¹⁶ | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>ZBED3</i> | rs4457053 | 5 | 76,424,949 | G | A | 30.4 | 30.4 | 8.4x10 ⁻¹⁸ | 1.06 (1.05-1.08) | N | N | Y | 74,115 | 823,995 | 0.96 |
| <i>DMGDH</i> | rs1316776 | 5 | 78,430,607 | C | A | 64.8 | 35.2 | 2.6x10 ⁻¹² | 1.05 (1.03-1.06) | N | Y | Y | 74,115 | 823,997 | 1.00 |
| <i>RASA1</i> | rs7719891 | 5 | 86,577,352 | G | A | 25.9 | 25.9 | 2.4x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| <i>SLCO6A1</i> | rs138337556 | 5 | 101,232,944 | G | A | 0.360 | 0.360 | 4.7x10 ⁻⁹ | 1.56 (1.34-1.81) | N | Y | Y | 52,854 | 778,536 | 0.57 |
| <i>PAM</i> | rs78408340 | 5 | 102,338,739 | G | C | 0.830 | 0.830 | 2.1x10 ⁻²⁴ | 1.47 (1.37-1.59) | N | N | N | 73,705 | 823,607 | 0.70 |
| <i>PAM</i> | rs115505614 | 5 | 102,422,968 | T | C | 4.99 | 4.99 | 1.3x10 ⁻³⁰ | 1.19 (1.15-1.22) | N | N | Y | 74,117 | 823,997 | 0.87 |

| | | | | | | | | | | | | | | | |
|------------|-------------|---|-------------|---|---|--------|--------|-----------------------|------------------|---|---|---|--------|---------|------|
| PHF15 | rs244665 | 5 | 133,414,622 | A | G | 70.3 | 29.7 | 9.9x10 ⁻⁶ | 1.03 (1.02-1.05) | N | N | N | 74,115 | 823,995 | 0.98 |
| PHF15 | rs329122 | 5 | 133,864,599 | A | G | 42.9 | 42.9 | 3.6x10 ⁻⁹ | 1.04 (1.03-1.05) | N | N | Y | 74,117 | 823,997 | 0.99 |
| EBF1 | rs3934712 | 5 | 157,928,196 | C | T | 20.6 | 20.6 | 3.2x10 ⁻⁸ | 1.05 (1.03-1.07) | Y | Y | Y | 50,402 | 523,888 | 1.00 |
| RREB1 | rs112498319 | 6 | 7,035,734 | C | A | 40.9 | 40.9 | 4.2x10 ⁻⁷ | 1.03 (1.02-1.05) | N | N | N | 74,116 | 823,996 | 1.00 |
| RREB1 | rs9379084 | 6 | 7,231,843 | G | A | 88.7 | 11.3 | 3.3x10 ⁻²¹ | 1.11 (1.08-1.13) | N | N | Y | 74,117 | 823,997 | 0.91 |
| RREB1 | rs9505097 | 6 | 7,255,650 | C | T | 79.9 | 20.1 | 8.6x10 ⁻¹⁰ | 1.05 (1.03-1.07) | N | N | N | 74,117 | 823,997 | 0.99 |
| CDKAL1 | rs7756992 | 6 | 20,679,709 | G | A | 27.4 | 27.4 | 2.4x10 ⁻⁸⁸ | 1.15 (1.13-1.17) | N | N | Y | 74,117 | 823,997 | 1.00 |
| MHC | rs601945 | 6 | 32,573,415 | G | A | 17.8 | 17.8 | 4.7x10 ⁻⁸ | 1.06 (1.04-1.08) | N | N | Y | 74,038 | 819,490 | 0.88 |
| HMGAA1 | rs77136196 | 6 | 34,247,047 | T | C | 4.20 | 4.20 | 1.6x10 ⁻⁸ | 1.11 (1.07-1.16) | Y | Y | Y | 50,402 | 523,888 | 0.93 |
| HMGAA1 | rs2233632 | 6 | 34,524,698 | T | C | 68.8 | 31.2 | 5.3x10 ⁻⁷ | 1.04 (1.03-1.06) | Y | Y | N | 50,400 | 523,886 | 0.99 |
| LRFN2 | rs34298980 | 6 | 40,409,243 | T | C | 49.7 | 49.7 | 9.3x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | Y | Y | 74,115 | 823,995 | 0.89 |
| VEGFA | rs11967262 | 6 | 43,760,327 | G | C | 48.6 | 48.6 | 8.8x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | N | N | 74,116 | 823,995 | 0.96 |
| VEGFA | rs6458354 | 6 | 43,814,190 | C | T | 28.9 | 28.9 | 2.1x10 ⁻¹² | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,995 | 0.99 |
| TFAP2B | rs3798519 | 6 | 50,788,778 | C | A | 18.4 | 18.4 | 2.6x10 ⁻¹² | 1.06 (1.04-1.08) | N | N | Y | 74,117 | 823,997 | 1.00 |
| TFAP2B | rs2465043 | 6 | 51,180,765 | G | A | 64.4 | 35.6 | 2.9x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,116 | 823,997 | 0.99 |
| SLC25A51P1 | rs555402748 | 6 | 67,387,490 | T | C | 0.0400 | 0.0400 | 4.6x10 ⁻⁸ | 3.67 (2.30-5.86) | N | Y | Y | 45,657 | 492,696 | 0.55 |
| BEND3 | rs4946812 | 6 | 107,431,688 | G | A | 67.4 | 32.6 | 8.2x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,997 | 0.95 |
| CENPW | rs11759026 | 6 | 126,792,095 | G | A | 23.2 | 23.2 | 2.4x10 ⁻¹⁸ | 1.07 (1.05-1.08) | N | N | Y | 74,117 | 823,997 | 0.97 |
| SOGA3 | rs2800733 | 6 | 127,416,930 | A | G | 71.7 | 28.4 | 6.0x10 ⁻¹¹ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| SLC35D3 | rs9494624 | 6 | 137,300,960 | A | G | 29.0 | 29.0 | 6.1x10 ⁻⁹ | 1.04 (1.03-1.06) | N | N | Y | 74,116 | 823,995 | 0.99 |
| MIR3668 | rs2982521 | 6 | 139,835,329 | A | T | 38.0 | 38.0 | 1.3x10 ⁻⁹ | 1.05 (1.03-1.06) | Y | Y | Y | 50,402 | 523,887 | 1.00 |
| MIR3668 | rs616279 | 6 | 140,249,466 | A | G | 73.8 | 26.2 | 6.7x10 ⁻⁷ | 1.04 (1.03-1.06) | Y | Y | N | 50,321 | 519,379 | 0.99 |
| SLC22A3 | rs474513 | 6 | 160,770,312 | A | G | 51.7 | 48.3 | 8.1x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | N | Y | 74,117 | 823,997 | 1.00 |
| QKI | rs4709746 | 6 | 164,133,001 | C | T | 86.8 | 13.2 | 5.8x10 ⁻⁹ | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,997 | 0.96 |
| DGKB | rs17168486 | 7 | 14,898,282 | T | C | 18.1 | 18.1 | 2.3x10 ⁻¹⁷ | 1.07 (1.06-1.09) | N | N | N | 74,117 | 823,996 | 0.99 |
| DGKB | rs10228066 | 7 | 15,063,569 | T | C | 53.7 | 46.3 | 1.1x10 ⁻²⁸ | 1.07 (1.06-1.09) | N | N | Y | 74,116 | 823,995 | 1.00 |
| DGKB | rs2908334 | 7 | 15,206,239 | T | C | 63.1 | 36.9 | 5.9x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,117 | 823,996 | 1.00 |
| IGF2BP3 | rs78840640 | 7 | 23,434,606 | G | C | 2.20 | 2.20 | 2.8x10 ⁻⁶ | 1.11 (1.06-1.16) | N | Y | N | 74,117 | 823,997 | 0.94 |
| IGF2BP3 | rs4279506 | 7 | 23,512,896 | G | C | 61.0 | 39.0 | 4.8x10 ⁻⁸ | 1.06 (1.04-1.08) | N | Y | Y | 74,116 | 823,997 | 0.96 |
| JAZF1 | rs1708302 | 7 | 28,198,677 | C | T | 51.2 | 48.8 | 1.1x10 ⁻⁴⁸ | 1.10 (1.08-1.11) | N | N | Y | 74,116 | 823,996 | 1.00 |
| CRHR2 | rs917195 | 7 | 30,728,452 | C | T | 77.0 | 23.0 | 4.2x10 ⁻¹¹ | 1.05 (1.04-1.07) | N | Y | Y | 74,117 | 823,997 | 0.94 |
| GCK | rs878521 | 7 | 44,255,643 | A | G | 24.5 | 24.5 | 1.9x10 ⁻¹³ | 1.06 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.97 |
| GCK | rs116913033 | 7 | 44,365,549 | C | T | 83.0 | 17.0 | 7.1x10 ⁻⁶ | 1.04 (1.02-1.06) | N | N | N | 74,117 | 823,997 | 0.93 |
| FBXL13 | rs56376556 | 7 | 102,038,318 | T | C | 5.33 | 5.33 | 1.7x10 ⁻⁶ | 1.08 (1.04-1.11) | N | Y | N | 74,117 | 823,997 | 0.84 |
| FBXL13 | rs11496066 | 7 | 102,486,254 | T | C | 81.8 | 18.2 | 1.1x10 ⁻⁸ | 1.08 (1.05-1.11) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| RELN | rs62482405 | 7 | 102,987,583 | G | T | 8.19 | 8.19 | 6.9x10 ⁻⁶ | 1.05 (1.03-1.08) | N | Y | N | 74,117 | 823,996 | 0.99 |
| RELN | rs39328 | 7 | 103,444,978 | T | C | 43.3 | 43.3 | 3.7x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| CTTNBP2 | rs6976111 | 7 | 117,495,667 | A | C | 31.3 | 31.3 | 1.2x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 62,668 | 545,622 | 0.98 |
| KLF14 | rs2268382 | 7 | 130,027,037 | C | A | 32.7 | 32.7 | 7.4x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,117 | 823,997 | 1.00 |
| KLF14 | rs1562396 | 7 | 130,457,914 | G | A | 31.9 | 31.9 | 9.9x10 ⁻¹⁰ | 1.06 (1.05-1.08) | N | N | Y | 74,116 | 823,997 | 0.92 |
| AOC1 | rs62492368 | 7 | 150,537,635 | A | G | 30.8 | 30.8 | 1.1x10 ⁻¹⁰ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| MNX1 | rs6459733 | 7 | 156,930,550 | G | C | 67.3 | 32.7 | 2.4x10 ⁻⁷ | 1.06 (1.05-1.07) | N | N | Y | 74,117 | 823,997 | 0.97 |
| MSRA | rs17689007 | 8 | 9,974,824 | G | A | 53.3 | 46.7 | 2.5x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,037 | 819,490 | 1.00 |
| XKR6 | rs57327348 | 8 | 10,808,687 | A | T | 78.2 | 21.8 | 4.5x10 ⁻⁸ | 1.04 (1.02-1.06) | N | Y | Y | 74,038 | 819,490 | 0.95 |
| LPL | rs10096633 | 8 | 19,830,921 | C | T | 87.7 | 12.3 | 1.1x10 ⁻¹² | 1.07 (1.05-1.09) | N | N | Y | 74,117 | 823,997 | 1.00 |
| PURG | rs10954772 | 8 | 30,863,938 | T | C | 31.4 | 31.4 | 1.8x10 ⁻⁹ | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,995 | 0.99 |
| ANK1 | rs13262861 | 8 | 41,508,577 | C | A | 82.9 | 17.1 | 4.0x10 ⁻¹² | 1.07 (1.05-1.09) | N | N | Y | 74,117 | 823,997 | 0.97 |
| ANK1 | rs4736819 | 8 | 41,509,915 | T | C | 55.4 | 44.6 | 5.4x10 ⁻⁷ | 1.04 (1.02-1.05) | N | N | N | 74,115 | 823,995 | 1.00 |
| ANK1 | rs148766658 | 8 | 41,552,046 | C | T | 3.78 | 3.78 | 5.7x10 ⁻⁷ | 1.09 (1.05-1.13) | N | N | N | 74,117 | 823,997 | 0.97 |
| TP53INP1 | rs11786992 | 8 | 95,685,147 | A | C | 64.4 | 35.6 | 5.3x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,117 | 823,997 | 1.00 |
| TP53INP1 | rs10097617 | 8 | 95,961,626 | T | C | 48.5 | 48.5 | 3.3x10 ⁻¹¹ | 1.04 (1.03-1.06) | N | N | Y | 74,117 | 823,996 | 1.00 |
| TP53INP1 | rs187936726 | 8 | 96,092,422 | G | A | 2.39 | 2.39 | 5.4x10 ⁻⁶ | 1.11 (1.06-1.16) | N | N | N | 74,117 | 823,997 | 0.95 |
| CPQ | rs149364428 | 8 | 97,737,741 | A | G | 1.04 | 1.04 | 1.8x10 ⁻¹² | 1.27 (1.19-1.36) | N | Y | Y | 73,967 | 822,104 | 0.86 |
| TRHR | rs12680028 | 8 | 110,123,183 | C | G | 53.4 | 46.6 | 2.5x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| SLC30A8 | rs3802177 | 8 | 118,185,025 | G | A | 68.5 | 31.5 | 1.1x10 ⁻⁵⁵ | 1.11 (1.10-1.13) | N | N | Y | 74,116 | 823,996 | 0.98 |
| SLC30A8 | rs80244329 | 8 | 118,404,672 | G | A | 97.8 | 2.19 | 6.9x10 ⁻⁶ | 1.11 (1.06-1.17) | N | N | N | 74,117 | 823,997 | 0.75 |
| CASC11 | rs17772814 | 8 | 128,711,742 | G | A | 91.5 | 8.49 | 5.4x10 ⁻¹⁰ | 1.08 (1.05-1.11) | N | Y | Y | 74,117 | 823,997 | 0.77 |
| PVT1 | rs1561927 | 8 | 129,568,078 | C | T | 26.9 | 26.9 | 1.5x10 ⁻⁹ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| BOP1 | rs4977213 | 8 | 145,507,304 | C | T | 37.5 | 37.5 | 9.1x10 ⁻¹⁴ | 1.05 (1.04-1.07) | N | Y | Y | 74,115 | 823,995 | 0.97 |
| BOP1 | rs12719778 | 8 | 145,879,883 | T | C | 53.8 | 46.2 | 5.0x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | N | 74,117 | 823,996 | 0.97 |
| GLIS3 | rs510807 | 9 | 3,965,689 | A | C | 49.1 | 49.1 | 1.4x10 ⁻⁶ | 1.03 (1.02-1.04) | N | N | N | 74,115 | 823,995 | 0.98 |
| GLIS3 | rs79103584 | 9 | 4,243,045 | T | A | 98.6 | 1.38 | 4.4x10 ⁻⁶ | 1.14 (1.08-1.21) | N | N | N | 74,117 | 823,996 | 0.98 |
| GLIS3 | rs10974438 | 9 | 4,291,928 | C | A | 35.7 | 35.7 | 1.5x10 ⁻¹⁴ | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,997 | 0.98 |
| HAUS6 | rs7022807 | 9 | 19,067,833 | G | A | 40.1 | 40.1 | 2.7x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | Y | Y | 74,116 | 823,997 | 1.00 |
| FOCAD | rs7867635 | 9 | 20,241,069 | C | T | 41.2 | 41.2 | 4.0x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,115 | 823,995 | 0.95 |
| FOCAD | rs7847880 | 9 | 20,662,703 | C | T | 84.3 | 15.7 | 2.1x10 ⁻⁶ | 1.04 (1.03-1.06) | N | Y | N | 74,115 | 823,995 | 0.98 |

| | | | | | | | | | | | | | | | |
|----------------------|-----------------|----|-------------|---|---|--------|--------|------------------------|------------------|---|---|---|--------|---------|------|
| <i>CDKN2A/B</i> | rs1412830 | 9 | 22,043,612 | C | T | 62.8 | 37.2 | 9.1x10 ⁻⁸ | 1.04 (1.02-1.05) | N | N | N | 74,117 | 823,995 | 0.97 |
| <i>CDKN2A/B</i> | rs76011118 | 9 | 22,133,773 | A | G | 3.41 | 3.41 | 1.4x10 ⁻⁷ | 1.11 (1.07-1.15) | N | N | N | 74,117 | 823,997 | 0.75 |
| <i>CDKN2A/B</i> | rs10811660 | 9 | 22,134,068 | G | A | 82.8 | 17.2 | 1.4x10 ⁻¹¹⁵ | 1.27 (1.24-1.29) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>CDKN2A/B</i> | rs10757283 | 9 | 22,134,172 | T | C | 43.0 | 43.0 | 1.7x10 ⁻⁴¹ | 1.11 (1.09-1.13) | N | N | N | 74,116 | 823,996 | 0.98 |
| <i>CDKN2A/B</i> | rs1333052 | 9 | 22,157,908 | A | C | 66.0 | 34.0 | 6.3x10 ⁻⁷ | 1.03 (1.02-1.05) | N | N | N | 74,116 | 823,997 | 0.99 |
| <i>CDKN2A/B</i> | rs1575972 | 9 | 22,301,092 | T | A | 96.7 | 3.29 | 3.1x10 ⁻⁷ | 1.10 (1.06-1.14) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>LINGO2</i> | rs1412234 | 9 | 28,410,683 | C | T | 32.3 | 32.3 | 1.9x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 74,116 | 823,997 | 0.99 |
| <i>UBAP2</i> | rs12001437 | 9 | 34,074,476 | C | T | 37.2 | 37.2 | 2.8x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,997 | 0.99 |
| <i>MTND2P8</i> | rs11137820 | 9 | 81,359,113 | C | G | 57.5 | 42.5 | 2.9x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 0.99 |
| <i>TLE4</i> | rs17791513 | 9 | 81,905,590 | A | G | 93.2 | 6.83 | 3.1x10 ⁻¹⁴ | 1.10 (1.08-1.13) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>TLE4</i> | rs2796441 | 9 | 84,308,948 | G | A | 59.2 | 40.8 | 4.4x10 ⁻²⁴ | 1.07 (1.05-1.08) | N | N | Y | 74,116 | 823,997 | 0.98 |
| <i>ZNF169</i> | rs55653563 | 9 | 97,001,682 | A | C | 73.2 | 26.8 | 2.2x10 ⁻⁹ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>ZNF169</i> | rs12236906 | 9 | 97,497,494 | T | C | 98.7 | 1.28 | 3.4x10 ⁻⁶ | 1.15 (1.08-1.22) | N | Y | N | 74,116 | 823,996 | 0.98 |
| <i>ABO</i> | rs505922 | 9 | 136,149,229 | C | T | 33.2 | 33.2 | 3.9x10 ⁻¹² | 1.05 (1.03-1.06) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>GPSM1</i> | rs78403475 | 9 | 139,235,606 | G | C | 89.6 | 10.4 | 1.2x10 ⁻⁶ | 1.06 (1.03-1.08) | N | N | N | 74,117 | 823,997 | 0.85 |
| <i>GPSM1</i> | rs28505901 | 9 | 139,241,030 | G | A | 75.2 | 24.8 | 6.7x10 ⁻²⁶ | 1.09 (1.07-1.11) | N | N | Y | 67,154 | 810,074 | 0.83 |
| <i>GPSM1</i> | rs11793035 | 9 | 139,507,212 | C | T | 33.1 | 33.1 | 2.2x10 ⁻⁷ | 1.04 (1.02-1.05) | N | N | N | 67,154 | 810,073 | 0.86 |
| <i>GPSM1</i> | 9:139737088:G:A | 9 | 139,737,088 | A | G | 0.0700 | 0.0700 | 7.9x10 ⁻⁶ | 2.74 (1.76-4.25) | N | N | N | 31,447 | 464,429 | 0.62 |
| <i>CDC123/CAMK1D</i> | rs11257655 | 10 | 12,307,894 | T | C | 21.8 | 21.8 | 1.5x10 ⁻³² | 1.09 (1.08-1.11) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>NEUROG3</i> | rs177045 | 10 | 71,321,279 | G | A | 31.6 | 31.6 | 6.6x10 ⁻¹⁸ | 1.07 (1.05-1.08) | N | Y | N | 74,117 | 823,997 | 0.99 |
| <i>NEUROG3</i> | rs61850200 | 10 | 71,321,658 | C | G | 27.7 | 27.7 | 7.3x10 ⁻⁶ | 1.04 (1.02-1.05) | N | Y | N | 74,117 | 823,997 | 0.95 |
| <i>NEUROG3</i> | rs41277236 | 10 | 71,332,301 | T | C | 4.31 | 4.31 | 1.5x10 ⁻⁶ | 1.09 (1.05-1.12) | N | Y | N | 74,117 | 823,997 | 0.83 |
| <i>NEUROG3</i> | rs549498088 | 10 | 71,347,311 | T | C | 0.600 | 0.600 | 4.7x10 ⁻⁷ | 1.56 (1.31-1.86) | N | Y | N | 68,385 | 805,130 | 0.89 |
| <i>NEUROG3</i> | rs2642588 | 10 | 71,466,578 | G | T | 70.2 | 29.8 | 2.2x10 ⁻¹⁴ | 1.05 (1.04-1.07) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| <i>ZMIZ1</i> | rs703972 | 10 | 80,952,826 | G | C | 53.3 | 46.7 | 1.7x10 ⁻²⁹ | 1.07 (1.06-1.09) | N | N | Y | 74,117 | 823,996 | 0.99 |
| <i>ZMIZ1</i> | rs1317617 | 10 | 81,096,589 | G | A | 79.8 | 20.2 | 1.8x10 ⁻⁶ | 1.04 (1.02-1.06) | N | N | N | 74,117 | 823,997 | 0.97 |
| <i>PTEN</i> | rs11202627 | 10 | 89,769,340 | T | C | 15.2 | 15.2 | 4.7x10 ⁻⁸ | 1.06 (1.04-1.08) | Y | Y | Y | 50,402 | 523,888 | 1.00 |
| <i>HHEX/IDE</i> | rs7078559 | 10 | 93,924,663 | T | C | 57.8 | 42.2 | 4.1x10 ⁻⁷ | 1.03 (1.02-1.05) | N | N | N | 74,116 | 823,997 | 1.00 |
| <i>HHEX/IDE</i> | rs10882101 | 10 | 94,462,427 | T | C | 58.7 | 41.3 | 1.4x10 ⁻⁸ | 1.06 (1.04-1.08) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>HHEX/IDE</i> | rs1112718 | 10 | 94,479,107 | A | G | 59.8 | 40.2 | 5.0x10 ⁻⁷ | 1.06 (1.03-1.08) | N | N | N | 74,117 | 823,996 | 1.00 |
| <i>TCF7L2</i> | rs536643418 | 10 | 114,699,835 | G | C | 0.520 | 0.520 | 2.6x10 ⁻⁸ | 1.50 (1.30-1.73) | N | N | N | 70,881 | 810,159 | 0.66 |
| <i>TCF7L2</i> | rs140242150 | 10 | 114,702,962 | A | G | 0.500 | 0.500 | 2.2x10 ⁻⁸ | 1.36 (1.22-1.52) | N | N | N | 70,758 | 813,597 | 0.59 |
| <i>TCF7L2</i> | rs7918400 | 10 | 114,703,136 | T | C | 47.6 | 47.6 | 2.0x10 ⁻¹⁵ | 1.06 (1.04-1.07) | N | N | N | 74,117 | 823,995 | 0.98 |
| <i>TCF7L2</i> | rs184509201 | 10 | 114,740,337 | C | G | 98.2 | 1.82 | 1.2x10 ⁻¹³ | 1.21 (1.15-1.27) | N | N | N | 74,117 | 823,997 | 0.91 |
| <i>TCF7L2</i> | rs180988137 | 10 | 114,751,173 | G | A | 1.04 | 1.04 | 6.1x10 ⁻⁶ | 1.17 (1.09-1.25) | N | N | N | 74,117 | 823,997 | 0.73 |
| <i>TCF7L2</i> | rs7903146 | 10 | 114,758,349 | C | T | 70.6 | 29.5 | 5.8x10 ⁻⁴⁴⁷ | 1.37 (1.35-1.39) | N | N | Y | 74,116 | 823,996 | 0.99 |
| <i>TCF7L2</i> | rs78025551 | 10 | 114,757,956 | C | G | 85.1 | 14.9 | 1.6x10 ⁻⁷ | 1.05 (1.03-1.07) | N | N | N | 74,117 | 823,997 | 0.98 |
| <i>TCF7L2</i> | rs34855922 | 10 | 114,871,594 | A | G | 71.6 | 28.4 | 5.5x10 ⁻¹² | 1.05 (1.04-1.07) | N | N | N | 74,117 | 823,997 | 0.91 |
| <i>WDR11</i> | rs72631105 | 10 | 122,915,345 | A | G | 19.0 | 19.0 | 3.7x10 ⁻⁹ | 1.06 (1.04-1.08) | Y | Y | Y | 50,402 | 523,888 | 0.88 |
| <i>PLEKHA1</i> | rs2280141 | 10 | 124,193,181 | T | G | 51.6 | 48.4 | 1.4x10 ⁻¹³ | 1.05 (1.03-1.06) | N | N | Y | 74,116 | 823,996 | 1.00 |
| <i>INS/IGF2</i> | rs12802972 | 11 | 1,704,596 | A | G | 42.8 | 42.8 | 1.5x10 ⁻⁶ | 1.03 (1.02-1.05) | N | N | N | 74,115 | 823,997 | 0.96 |
| <i>INS/IGF2</i> | rs11042596 | 11 | 2,118,860 | G | T | 66.5 | 33.5 | 2.0x10 ⁻⁸ | 1.04 (1.03-1.05) | N | N | N | 74,115 | 823,995 | 0.93 |
| <i>INS/IGF2</i> | rs555759341 | 11 | 2,151,761 | C | G | 0.490 | 0.490 | 3.6x10 ⁻⁸ | 1.38 (1.23-1.55) | N | N | N | 69,523 | 804,881 | 0.76 |
| <i>INS/IGF2</i> | rs571342427 | 11 | 2,182,519 | C | T | 0.150 | 0.150 | 1.0x10 ⁻⁶ | 1.68 (1.36-2.07) | N | N | N | 64,480 | 793,265 | 0.69 |
| <i>INS/IGF2</i> | rs4929965 | 11 | 2,197,286 | A | G | 38.3 | 38.3 | 4.0x10 ⁻²⁶ | 1.07 (1.06-1.09) | N | N | Y | 74,115 | 823,996 | 0.89 |
| <i>KCNQ1</i> | rs4930091 | 11 | 2,372,356 | C | T | 75.9 | 24.1 | 3.7x10 ⁻⁶ | 1.04 (1.02-1.05) | N | N | N | 74,115 | 823,995 | 0.97 |
| <i>KCNQ1</i> | rs2283164 | 11 | 2,579,163 | A | G | 94.7 | 5.32 | 1.2x10 ⁻⁷ | 1.08 (1.05-1.12) | N | N | N | 74,117 | 823,997 | 0.88 |
| <i>KCNQ1</i> | rs80102379 | 11 | 2,634,177 | G | T | 98.2 | 1.78 | 9.3x10 ⁻⁸ | 1.15 (1.09-1.21) | N | N | N | 74,117 | 823,997 | 0.80 |
| <i>KCNQ1</i> | rs231349 | 11 | 2,672,821 | T | C | 10.2 | 10.2 | 2.3x10 ⁻¹¹ | 1.07 (1.05-1.10) | N | N | N | 74,117 | 823,997 | 0.96 |
| <i>KCNQ1</i> | rs231361 | 11 | 2,691,500 | A | G | 25.6 | 25.6 | 5.0x10 ⁻²⁵ | 1.08 (1.07-1.10) | N | N | N | 74,117 | 823,997 | 0.96 |
| <i>KCNQ1</i> | rs2283220 | 11 | 2,755,548 | A | G | 69.0 | 31.0 | 1.4x10 ⁻⁹ | 1.05 (1.03-1.06) | N | N | N | 74,117 | 823,997 | 0.93 |
| <i>KCNQ1</i> | rs234853 | 11 | 2,850,828 | G | A | 24.8 | 24.8 | 6.8x10 ⁻¹⁶ | 1.08 (1.06-1.10) | N | N | N | 74,116 | 823,996 | 0.97 |
| <i>KCNQ1</i> | rs2237895 | 11 | 2,857,194 | C | A | 42.6 | 42.6 | 6.0x10 ⁻⁵² | 1.12 (1.11-1.14) | N | N | Y | 74,116 | 823,996 | 0.94 |
| <i>KCNQ1</i> | rs2237897 | 11 | 2,858,546 | C | T | 95.4 | 4.57 | 8.4x10 ⁻³² | 1.23 (1.19-1.27) | N | N | N | 74,117 | 823,997 | 0.93 |
| <i>KCNQ1</i> | rs445084 | 11 | 2,908,754 | G | A | 36.1 | 36.1 | 1.7x10 ⁻⁶ | 1.03 (1.02-1.05) | N | N | N | 74,115 | 823,996 | 0.96 |
| <i>PDE3B</i> | rs141521721 | 11 | 14,763,828 | A | C | 2.36 | 2.36 | 2.7x10 ⁻⁸ | 1.13 (1.08-1.17) | N | Y | Y | 74,117 | 823,997 | 0.96 |
| <i>KCNJ11</i> | rs5213 | 11 | 17,408,404 | C | T | 36.2 | 36.2 | 3.5x10 ⁻²⁷ | 1.07 (1.06-1.09) | N | N | Y | 74,115 | 823,995 | 0.99 |
| <i>KCNJ11</i> | rs67254669 | 11 | 17,470,143 | G | A | 0.110 | 0.110 | 1.1x10 ⁻⁸ | 1.89 (1.52-2.35) | N | N | N | 55,757 | 521,088 | 0.82 |
| <i>METTL15</i> | rs4923543 | 11 | 28,534,898 | A | G | 33.2 | 33.2 | 4.5x10 ⁻⁸ | 1.04 (1.03-1.06) | Y | Y | Y | 50,400 | 523,886 | 1.00 |
| <i>QSER1</i> | rs7943101 | 11 | 32,460,873 | T | C | 16.1 | 16.1 | 8.5x10 ⁻⁶ | 1.04 (1.02-1.06) | N | Y | N | 74,117 | 823,997 | 0.99 |
| <i>QSER1</i> | rs145678014 | 11 | 32,927,778 | G | T | 95.7 | 4.33 | 2.0x10 ⁻¹⁰ | 1.11 (1.07-1.14) | N | Y | Y | 74,117 | 823,997 | 0.94 |
| <i>QSER1</i> | rs528122639 | 11 | 33,091,735 | A | G | 0.0900 | 0.0900 | 1.6x10 ⁻⁷ | 2.09 (1.59-2.76) | N | Y | N | 62,871 | 788,941 | 0.69 |
| <i>PDHX</i> | rs286925 | 11 | 34,642,668 | A | G | 18.2 | 18.2 | 5.0x10 ⁻⁶ | 1.04 (1.02-1.06) | N | Y | N | 74,116 | 823,996 | 0.99 |
| <i>PDHX</i> | rs2767036 | 11 | 34,982,148 | C | A | 29.1 | 29.1 | 3.3x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,115 | 823,996 | 1.00 |
| <i>HSD17B12</i> | rs1061810 | 11 | 43,877,934 | A | C | 28.8 | 28.8 | 6.0x10 ⁻¹³ | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>CRY2</i> | rs7115753 | 11 | 45,912,013 | A | G | 44.9 | 44.9 | 3.8x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,115 | 823,995 | 0.98 |

| <i>CELF1</i> | rs7124681 | 11 | 47,529,947 | A | C | 41.0 | 41.0 | 5.1x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,038 | 819,490 | 1.00 |
|---------------------|-------------|----|-------------|---|---|--------|--------|-----------------------|------------------|---|---|---|--------|---------|------|
| <i>MAP3K11</i> | rs1783541 | 11 | 65,294,799 | T | C | 20.4 | 20.4 | 2.0x10 ⁻¹⁴ | 1.06 (1.05-1.08) | N | N | Y | 74,116 | 823,996 | 0.98 |
| <i>CCND1</i> | rs61881115 | 11 | 68,997,225 | G | A | 83.8 | 16.2 | 4.1x10 ⁻⁷ | 1.05 (1.03-1.06) | N | Y | N | 74,117 | 823,996 | 0.97 |
| <i>CCND1</i> | rs11820019 | 11 | 69,448,758 | T | C | 97.3 | 2.67 | 5.1x10 ⁻¹² | 1.16 (1.11-1.20) | N | Y | Y | 74,117 | 823,997 | 0.88 |
| <i>CENTD2/ARAP1</i> | rs77464186 | 11 | 72,460,398 | A | C | 83.6 | 16.4 | 4.7x10 ⁻³³ | 1.11 (1.09-1.13) | N | N | Y | 74,116 | 823,997 | 0.99 |
| <i>MTNR1B</i> | rs10830963 | 11 | 92,708,710 | G | C | 27.7 | 27.7 | 4.8x10 ⁻⁴³ | 1.10 (1.09-1.12) | N | N | Y | 74,117 | 823,996 | 0.97 |
| <i>MTNR1B</i> | rs57235767 | 11 | 93,013,531 | C | T | 70.6 | 29.4 | 5.9x10 ⁻¹⁰ | 1.04 (1.03-1.06) | N | N | N | 74,117 | 823,996 | 0.99 |
| <i>ETS1</i> | rs10893829 | 11 | 128,042,575 | T | C | 85.3 | 14.7 | 1.3x10 ⁻¹⁰ | 1.06 (1.04-1.08) | N | Y | N | 74,117 | 823,997 | 1.00 |
| <i>ETS1</i> | rs10750397 | 11 | 128,234,144 | A | G | 28.2 | 28.2 | 8.3x10 ⁻¹³ | 1.05 (1.04-1.07) | N | Y | N | 74,117 | 823,996 | 1.00 |
| <i>ETS1</i> | rs67232546 | 11 | 128,398,938 | T | C | 20.7 | 20.7 | 1.3x10 ⁻¹¹ | 1.06 (1.04-1.07) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>ETS1</i> | rs112595469 | 11 | 128,583,975 | T | C | 2.84 | 2.84 | 6.2x10 ⁻⁶ | 1.10 (1.05-1.14) | N | Y | N | 74,117 | 823,997 | 0.88 |
| <i>CCND2</i> | rs10848958 | 12 | 4,031,104 | C | T | 80.4 | 19.6 | 1.5x10 ⁻⁷ | 1.04 (1.03-1.06) | N | N | N | 74,117 | 823,997 | 0.94 |
| <i>CCND2</i> | rs11063028 | 12 | 4,300,172 | C | T | 18.0 | 18.0 | 8.5x10 ⁻¹¹ | 1.06 (1.04-1.07) | N | N | N | 74,116 | 823,997 | 1.00 |
| <i>CCND2</i> | rs4238013 | 12 | 4,376,089 | C | T | 20.9 | 20.9 | 3.2x10 ⁻¹¹ | 1.06 (1.04-1.07) | N | N | N | 74,115 | 823,995 | 0.94 |
| <i>CCND2</i> | rs3217792 | 12 | 4,384,696 | C | T | 91.3 | 8.69 | 2.6x10 ⁻²¹ | 1.12 (1.10-1.15) | N | N | N | 74,117 | 823,997 | 0.85 |
| <i>CCND2</i> | rs76895963 | 12 | 4,384,844 | T | G | 98.0 | 1.98 | 1.4x10 ⁻⁶⁹ | 1.62 (1.54-1.71) | N | N | Y | 74,117 | 823,997 | 0.72 |
| <i>CCND2</i> | rs3217860 | 12 | 4,399,050 | G | A | 25.8 | 25.8 | 3.9x10 ⁻⁹ | 1.05 (1.03-1.06) | N | N | N | 74,117 | 823,997 | 0.94 |
| <i>CDKN1B</i> | rs2066827 | 12 | 12,871,099 | G | T | 23.5 | 23.5 | 4.2x10 ⁻⁸ | 1.05 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 0.78 |
| <i>ITPR2</i> | rs718314 | 12 | 26,453,283 | G | A | 25.3 | 25.3 | 8.4x10 ⁻¹¹ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>KLHDC5</i> | rs10842994 | 12 | 27,965,150 | C | T | 80.5 | 19.5 | 4.1x10 ⁻²⁰ | 1.08 (1.06-1.09) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>HMGAA2</i> | rs2258238 | 12 | 66,221,060 | T | A | 10.4 | 10.4 | 4.5x10 ⁻²¹ | 1.10 (1.08-1.13) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>HMGAA2</i> | rs1042725 | 12 | 66,358,347 | T | C | 49.0 | 49.0 | 1.8x10 ⁻¹³ | 1.05 (1.03-1.06) | N | N | N | 74,116 | 823,996 | 1.00 |
| <i>TSPAN8/LGR5</i> | rs1796330 | 12 | 71,522,953 | G | C | 57.1 | 42.9 | 2.2x10 ⁻¹⁴ | 1.05 (1.04-1.06) | N | N | Y | 74,117 | 823,996 | 1.00 |
| <i>USP44</i> | rs2197973 | 12 | 95,928,560 | T | C | 53.8 | 46.3 | 3.6x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,115 | 823,996 | 1.00 |
| <i>RMST</i> | rs759111467 | 12 | 97,562,756 | A | G | 0.0300 | 0.0300 | 1.7x10 ⁻⁶ | 3.07 (1.94-4.85) | N | Y | N | 47,869 | 741,820 | 0.81 |
| <i>RMST</i> | rs557027608 | 12 | 97,779,248 | A | G | 0.0600 | 0.0600 | 2.7x10 ⁻⁷ | 2.34 (1.69-3.24) | N | Y | N | 42,508 | 733,542 | 0.88 |
| <i>RMST</i> | rs77864822 | 12 | 97,848,775 | A | G | 93.2 | 6.76 | 1.1x10 ⁻⁸ | 1.08 (1.05-1.11) | N | Y | Y | 74,117 | 823,997 | 0.97 |
| <i>WSCD2</i> | rs1426371 | 12 | 108,629,780 | G | A | 73.9 | 26.1 | 8.2x10 ⁻¹² | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.95 |
| <i>KSR2</i> | rs34965774 | 12 | 118,412,373 | A | G | 14.4 | 14.4 | 2.0x10 ⁻⁹ | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>KSR2</i> | rs12578639 | 12 | 118,489,636 | A | T | 82.8 | 17.2 | 2.2x10 ⁻⁶ | 1.04 (1.02-1.06) | N | Y | N | 74,117 | 823,997 | 0.97 |
| <i>HNF1A</i> | rs11065299 | 12 | 121,297,815 | A | G | 7.54 | 7.54 | 5.8x10 ⁻⁷ | 1.06 (1.04-1.09) | N | N | N | 74,117 | 823,997 | 0.96 |
| <i>HNF1A</i> | rs73226260 | 12 | 121,380,541 | G | A | 96.7 | 3.31 | 5.9x10 ⁻¹¹ | 1.13 (1.09-1.17) | N | N | N | 74,117 | 823,997 | 0.96 |
| <i>HNF1A</i> | rs1800574 | 12 | 121,416,864 | T | C | 2.96 | 2.96 | 1.7x10 ⁻¹² | 1.14 (1.10-1.19) | N | N | N | 74,117 | 823,997 | 0.94 |
| <i>HNF1A</i> | rs56348580 | 12 | 121,432,117 | G | C | 68.9 | 31.1 | 2.3x10 ⁻¹³ | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>HNF1A</i> | rs28638142 | 12 | 121,501,461 | A | C | 4.42 | 4.42 | 2.9x10 ⁻⁶ | 1.08 (1.04-1.11) | N | N | N | 74,117 | 823,997 | 0.95 |
| <i>HNF1A</i> | rs73224262 | 12 | 121,882,395 | T | C | 0.680 | 0.680 | 9.1x10 ⁻⁷ | 1.24 (1.14-1.34) | N | N | N | 74,117 | 823,997 | 0.75 |
| <i>MPHOSPH9</i> | rs4148856 | 12 | 123,450,765 | C | G | 78.1 | 21.9 | 1.7x10 ⁻¹⁰ | 1.05 (1.03-1.07) | N | N | Y | 74,115 | 823,995 | 0.99 |
| <i>ZNF664</i> | rs7978610 | 12 | 124,468,572 | G | C | 66.6 | 33.5 | 2.0x10 ⁻⁸ | 1.27 (1.17-1.38) | Y | Y | Y | 50,400 | 523,888 | 1.00 |
| <i>ZNF664</i> | rs825452 | 12 | 124,509,177 | A | G | 60.3 | 39.7 | 2.4x10 ⁻⁶ | 1.04 (1.02-1.05) | Y | Y | N | 50,400 | 523,887 | 0.99 |
| <i>FBRSL1</i> | rs12811407 | 12 | 133,069,698 | A | G | 33.1 | 33.1 | 1.7x10 ⁻¹² | 1.05 (1.04-1.07) | N | Y | Y | 74,117 | 823,996 | 0.92 |
| <i>RNF6</i> | rs34584161 | 13 | 26,776,999 | A | G | 76.0 | 24.0 | 2.2x10 ⁻¹⁰ | 1.05 (1.03-1.06) | N | N | Y | 74,117 | 823,997 | 0.97 |
| <i>HMGB1</i> | rs11842871 | 13 | 31,042,452 | G | T | 73.5 | 26.6 | 1.2x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.96 |
| <i>KL</i> | rs576674 | 13 | 33,554,302 | G | A | 16.9 | 16.9 | 8.3x10 ⁻¹⁰ | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>DLEU1</i> | rs963740 | 13 | 51,096,095 | A | T | 71.3 | 28.7 | 2.1x10 ⁻⁸ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,996 | 1.00 |
| <i>PCDH17</i> | rs9537803 | 13 | 58,366,634 | C | T | 27.7 | 27.7 | 4.6x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,996 | 1.00 |
| <i>PCDH17</i> | rs9569864 | 13 | 58,965,435 | C | T | 82.5 | 17.5 | 8.7x10 ⁻⁸ | 1.05 (1.03-1.06) | N | Y | N | 74,115 | 823,997 | 1.00 |
| <i>SRGAP2D</i> | rs9563615 | 13 | 59,077,406 | A | T | 71.0 | 29.0 | 6.4x10 ⁻¹¹ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>SRGAP2D</i> | rs76251711 | 13 | 59,184,234 | G | A | 1.26 | 1.26 | 2.3x10 ⁻⁶ | 1.16 (1.09-1.23) | N | Y | N | 73,705 | 823,607 | 0.87 |
| <i>SPRY2</i> | rs1359790 | 13 | 80,717,156 | G | A | 72.0 | 28.0 | 2.4x10 ⁻³¹ | 1.09 (1.07-1.10) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>IRS2</i> | rs7987740 | 13 | 109,947,213 | T | C | 60.9 | 39.1 | 4.0x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>IRS2</i> | rs4771648 | 13 | 110,431,626 | G | A | 66.9 | 33.2 | 8.9x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | N | 74,117 | 823,997 | 0.99 |
| <i>SLC7A7</i> | rs17122772 | 14 | 23,288,935 | G | C | 22.8 | 22.8 | 1.6x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.91 |
| <i>AKAP6</i> | rs17522122 | 14 | 33,302,882 | T | G | 47.4 | 47.4 | 3.2x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,116 | 823,995 | 0.98 |
| <i>CLEC14A</i> | rs8017808 | 14 | 38,848,419 | G | T | 74.3 | 25.7 | 2.1x10 ⁻⁸ | 1.04 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| <i>NRXN3</i> | rs17836088 | 14 | 79,932,041 | C | G | 21.7 | 21.7 | 6.7x10 ⁻¹⁴ | 1.06 (1.04-1.08) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>SMEK1</i> | rs8010382 | 14 | 91,963,722 | G | A | 42.1 | 42.1 | 6.5x10 ⁻⁹ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,996 | 0.94 |
| <i>MARK3</i> | rs62007683 | 14 | 103,894,071 | G | T | 65.3 | 34.7 | 3.1x10 ⁻⁸ | 1.04 (1.02-1.05) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>RASGRP1</i> | rs8032939 | 15 | 38,834,033 | C | T | 24.6 | 24.6 | 3.5x10 ⁻¹⁴ | 1.06 (1.04-1.07) | N | N | N | 74,116 | 823,997 | 0.99 |
| <i>RASGRP1</i> | rs34715063 | 15 | 38,873,115 | C | T | 12.4 | 12.4 | 2.3x10 ⁻¹⁹ | 1.10 (1.07-1.12) | N | N | Y | 74,117 | 823,997 | 0.89 |
| <i>LTK</i> | rs11070332 | 15 | 41,809,205 | A | G | 35.8 | 35.8 | 1.1x10 ⁻¹³ | 1.05 (1.04-1.06) | N | Y | Y | 74,117 | 823,995 | 0.98 |
| <i>LTK</i> | rs543786825 | 15 | 42,201,410 | T | C | 0.0400 | 0.0400 | 4.7x10 ⁻⁶ | 3.15 (1.93-5.15) | N | Y | N | 45,873 | 756,610 | 0.75 |
| <i>ONECUT1</i> | rs2456530 | 15 | 53,091,553 | T | C | 12.7 | 12.7 | 5.4x10 ⁻⁹ | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>WDR72</i> | rs528350911 | 15 | 53,747,228 | G | C | 0.680 | 0.680 | 2.1x10 ⁻⁸ | 1.27 (1.17-1.38) | N | Y | Y | 74,117 | 823,997 | 0.75 |
| <i>TCF12</i> | rs117483894 | 15 | 57,456,802 | G | A | 3.69 | 3.69 | 3.9x10 ⁻⁸ | 1.10 (1.06-1.13) | N | Y | Y | 74,115 | 823,996 | 0.99 |
| <i>C2CD4A/B</i> | rs8037894 | 15 | 62,394,264 | G | C | 56.6 | 43.4 | 2.6x10 ⁻¹³ | 1.05 (1.03-1.06) | N | N | Y | 74,115 | 823,996 | 1.00 |
| <i>USP3</i> | rs7178762 | 15 | 63,871,292 | C | T | 46.0 | 46.0 | 5.4x10 ⁻¹⁰ | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,997 | 1.00 |

| | | | | | | | | | | | | | | | |
|--------------------|-------------|----|------------|---|---|--------|--------|-----------------------|------------------|---|---|---|--------|---------|------|
| <i>MAP2K5</i> | rs4776970 | 15 | 68,080,886 | A | T | 64.1 | 35.9 | 5.0×10^{-9} | 1.04 (1.03-1.05) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>PTPN9</i> | rs13737 | 15 | 75,932,129 | G | T | 75.9 | 24.1 | 5.6×10^{-10} | 1.05 (1.03-1.06) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>HMG20A</i> | rs1005752 | 15 | 77,818,128 | A | C | 71.5 | 28.5 | 2.5×10^{-29} | 1.08 (1.07-1.10) | N | N | Y | 74,115 | 823,995 | 0.99 |
| <i>AP3S2</i> | rs4932265 | 15 | 90,423,293 | T | C | 26.7 | 26.7 | 4.2×10^{-20} | 1.07 (1.05-1.08) | N | N | Y | 74,115 | 823,995 | 0.99 |
| <i>PRC1</i> | rs12910825 | 15 | 91,511,260 | G | A | 36.1 | 36.1 | 1.6×10^{-15} | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,995 | 0.99 |
| <i>ITFG3</i> | rs6600191 | 16 | 295,795 | T | C | 82.5 | 17.5 | 9.3×10^{-13} | 1.06 (1.05-1.08) | N | N | Y | 74,117 | 823,997 | 1.00 |
| <i>CLUAP1</i> | rs3751837 | 16 | 3,583,173 | T | C | 22.0 | 22.0 | 1.4×10^{-8} | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,997 | 0.98 |
| <i>ATP2A1</i> | rs8046545 | 16 | 28,915,217 | G | A | 35.9 | 35.9 | 1.9×10^{-8} | 1.04 (1.02-1.05) | N | Y | Y | 74,115 | 823,995 | 0.98 |
| <i>FAM57B</i> | rs11642430 | 16 | 30,045,789 | G | C | 39.9 | 39.9 | 2.2×10^{-9} | 1.04 (1.03-1.05) | N | Y | Y | 74,116 | 823,996 | 0.99 |
| <i>FAM57B</i> | rs199795270 | 16 | 30,419,384 | C | G | 0.650 | 0.650 | 1.2×10^{-6} | 1.25 (1.14-1.36) | N | Y | N | 72,819 | 821,143 | 0.75 |
| <i>FTO</i> | rs4281707 | 16 | 53,501,946 | G | A | 54.4 | 45.6 | 3.2×10^{-10} | 1.04 (1.03-1.05) | N | N | N | 74,117 | 823,997 | 0.99 |
| <i>FTO</i> | rs78020297 | 16 | 53,758,720 | A | G | 5.17 | 5.17 | 6.5×10^{-9} | 1.09 (1.06-1.12) | N | N | N | 74,117 | 823,997 | 0.95 |
| <i>FTO</i> | rs1421085 | 16 | 53,800,954 | C | T | 41.5 | 41.5 | 3.1×10^{-84} | 1.13 (1.12-1.15) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>NFAT5</i> | rs862320 | 16 | 69,651,866 | C | T | 57.8 | 42.2 | 3.9×10^{-11} | 1.04 (1.03-1.06) | N | N | Y | 74,116 | 823,996 | 1.00 |
| <i>BCAR1</i> | rs72802342 | 16 | 75,234,872 | C | A | 92.3 | 7.69 | 4.0×10^{-32} | 1.17 (1.14-1.20) | N | N | Y | 74,117 | 823,997 | 0.95 |
| <i>BCAR1</i> | rs3115960 | 16 | 75,516,534 | G | C | 37.0 | 37.0 | 2.8×10^{-6} | 1.03 (1.02-1.05) | N | N | N | 74,115 | 823,995 | 0.96 |
| <i>CMIP</i> | rs2925979 | 16 | 81,534,790 | T | C | 30.0 | 30.0 | 1.4×10^{-14} | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>SPG7</i> | rs12920022 | 16 | 89,564,055 | A | T | 15.8 | 15.8 | 3.4×10^{-9} | 1.05 (1.04-1.07) | N | Y | Y | 74,117 | 823,997 | 0.94 |
| <i>ZEEF1</i> | rs1043246 | 17 | 3,828,086 | G | C | 15.7 | 15.7 | 7.9×10^{-7} | 1.05 (1.03-1.07) | N | N | N | 74,117 | 823,997 | 0.73 |
| <i>ZEEF1</i> | rs3826482 | 17 | 3,860,356 | A | T | 57.6 | 42.4 | 2.1×10^{-7} | 1.03 (1.02-1.05) | N | N | N | 74,116 | 823,996 | 0.94 |
| <i>ZEEF1</i> | rs1377807 | 17 | 4,045,440 | C | G | 31.2 | 31.2 | 4.2×10^{-13} | 1.05 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.99 |
| <i>ATP1B2</i> | rs1641523 | 17 | 7,549,681 | C | T | 42.8 | 42.8 | 1.2×10^{-10} | 1.05 (1.04-1.07) | Y | Y | Y | 50,400 | 523,886 | 0.97 |
| <i>ATP1B2</i> | rs62059712 | 17 | 7,740,170 | T | C | 91.8 | 8.16 | 4.9×10^{-6} | 1.07 (1.04-1.10) | Y | Y | N | 50,402 | 523,888 | 0.90 |
| <i>GLP2R</i> | rs7222481 | 17 | 9,785,187 | C | G | 32.4 | 32.4 | 1.4×10^{-8} | 1.04 (1.03-1.05) | N | N | Y | 74,117 | 823,996 | 1.00 |
| <i>RAI1</i> | rs4925109 | 17 | 17,661,802 | A | G | 31.6 | 31.6 | 2.8×10^{-12} | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>NF1</i> | rs71372253 | 17 | 29,413,019 | C | T | 6.42 | 6.42 | 4.4×10^{-8} | 1.08 (1.05-1.10) | N | Y | Y | 74,117 | 823,997 | 0.89 |
| <i>HNF1B</i> | rs10962 | 17 | 36,046,451 | C | G | 22.6 | 22.6 | 9.9×10^{-10} | 1.05 (1.03-1.07) | N | N | N | 74,117 | 823,997 | 0.89 |
| <i>HNF1B</i> | rs2189301 | 17 | 36,063,685 | G | A | 87.2 | 12.8 | 6.5×10^{-8} | 1.05 (1.03-1.08) | N | N | N | 74,117 | 823,997 | 0.98 |
| <i>HNF1B</i> | rs10908278 | 17 | 36,099,952 | T | A | 48.1 | 48.1 | 6.4×10^{-36} | 1.08 (1.07-1.10) | N | N | Y | 74,116 | 823,997 | 0.96 |
| <i>MLX</i> | rs34855406 | 17 | 40,731,411 | C | G | 27.7 | 27.7 | 2.3×10^{-12} | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>TLIL6</i> | rs35895680 | 17 | 47,060,322 | C | A | 67.8 | 32.2 | 2.5×10^{-15} | 1.06 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.96 |
| <i>KIF2B</i> | rs569511541 | 17 | 52,140,805 | G | A | 0.0200 | 0.0200 | 1.5×10^{-8} | 7.63 (3.78-15.4) | N | Y | Y | 37,528 | 715,995 | 0.69 |
| <i>ACE</i> | rs2727301 | 17 | 61,965,043 | T | C | 75.4 | 24.6 | 1.3×10^{-6} | 1.04 (1.02-1.05) | N | Y | N | 74,117 | 823,996 | 0.96 |
| <i>ACE</i> | rs60276348 | 17 | 62,203,304 | T | C | 14.0 | 14.0 | 2.6×10^{-8} | 1.05 (1.03-1.07) | N | Y | Y | 74,117 | 823,997 | 0.93 |
| <i>BPTF</i> | rs11657492 | 17 | 65,648,427 | G | T | 10.0 | 10.0 | 5.6×10^{-8} | 1.06 (1.04-1.08) | N | N | N | 74,117 | 823,997 | 0.96 |
| <i>BPTF</i> | rs558308082 | 17 | 65,820,153 | C | G | 0.0800 | 0.0800 | 6.6×10^{-6} | 2.04 (1.49-2.77) | N | N | N | 60,898 | 785,469 | 0.73 |
| <i>BPTF</i> | rs61676547 | 17 | 65,892,507 | C | G | 19.2 | 19.2 | 2.9×10^{-11} | 1.06 (1.04-1.07) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>LAMA1</i> | rs7240767 | 18 | 7,070,642 | C | T | 37.6 | 37.6 | 1.6×10^{-8} | 1.04 (1.02-1.05) | N | N | Y | 74,115 | 823,997 | 0.98 |
| <i>COMMID9</i> | rs62080313 | 18 | 36,278,709 | C | T | 12.3 | 12.3 | 1.0×10^{-8} | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,996 | 1.00 |
| <i>TCF4</i> | rs76197067 | 18 | 52,604,955 | G | A | 0.0500 | 0.0500 | 8.1×10^{-6} | 2.68 (1.74-4.12) | N | Y | N | 39,596 | 474,053 | 0.94 |
| <i>TCF4</i> | rs72926932 | 18 | 53,050,646 | C | A | 8.39 | 8.39 | 1.0×10^{-14} | 1.09 (1.07-1.12) | N | Y | Y | 74,117 | 823,997 | 1.00 |
| <i>TCF4</i> | rs28719468 | 18 | 53,452,144 | C | T | 15.9 | 15.9 | 1.9×10^{-6} | 1.04 (1.02-1.06) | N | Y | N | 74,117 | 823,997 | 1.00 |
| <i>WDR7</i> | rs17684074 | 18 | 54,675,384 | G | C | 74.0 | 26.0 | 2.9×10^{-8} | 1.04 (1.03-1.06) | N | Y | Y | 74,116 | 823,997 | 0.99 |
| <i>GRP</i> | rs9957145 | 18 | 56,876,228 | G | A | 82.9 | 17.1 | 8.1×10^{-9} | 1.05 (1.03-1.07) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>MC4R</i> | rs523288 | 18 | 57,848,369 | T | A | 23.8 | 23.8 | 7.6×10^{-13} | 1.05 (1.04-1.07) | N | N | Y | 74,116 | 823,997 | 1.00 |
| <i>MC4R</i> | rs74452128 | 18 | 58,056,566 | C | A | 97.6 | 2.37 | 1.0×10^{-9} | 1.15 (1.10-1.20) | N | N | N | 74,117 | 823,997 | 0.97 |
| <i>BCL2A</i> | rs10469140 | 18 | 60,668,270 | G | A | 48.5 | 48.5 | 6.6×10^{-6} | 1.03 (1.02-1.04) | N | N | N | 74,115 | 823,995 | 0.99 |
| <i>BCL2A</i> | rs12454712 | 18 | 60,845,884 | T | C | 61.4 | 38.6 | 4.6×10^{-13} | 1.05 (1.04-1.06) | N | N | Y | 74,116 | 823,996 | 0.87 |
| <i>UHFR1</i> | rs7249758 | 19 | 4,948,862 | A | G | 20.4 | 20.4 | 3.4×10^{-9} | 1.05 (1.03-1.07) | N | Y | Y | 74,117 | 823,996 | 0.98 |
| <i>PTPRS</i> | rs116953931 | 19 | 5,224,998 | A | G | 3.71 | 3.71 | 6.4×10^{-6} | 1.08 (1.04-1.12) | N | Y | N | 74,117 | 823,997 | 0.94 |
| <i>INSR</i> | rs75253922 | 19 | 7,240,848 | C | T | 19.1 | 19.1 | 2.7×10^{-8} | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.98 |
| <i>MAP2K7</i> | rs4804833 | 19 | 7,970,635 | A | G | 39.0 | 39.0 | 7.7×10^{-13} | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,997 | 0.96 |
| <i>FARSA</i> | rs755734872 | 19 | 12,938,471 | T | C | 0.0500 | 0.0500 | 8.3×10^{-6} | 2.37 (1.62-3.46) | N | Y | N | 37,418 | 480,876 | 0.81 |
| <i>FARSA</i> | rs3111316 | 19 | 13,038,415 | A | G | 58.9 | 41.2 | 6.3×10^{-13} | 1.05 (1.03-1.06) | N | Y | Y | 74,115 | 823,997 | 0.97 |
| <i>TM6SF2</i> | rs8107974 | 19 | 19,388,500 | T | A | 7.69 | 7.69 | 3.3×10^{-15} | 1.10 (1.07-1.12) | N | N | Y | 74,116 | 823,996 | 0.99 |
| <i>TM6SF2</i> | rs188247550 | 19 | 19,396,616 | T | C | 1.95 | 1.95 | 5.2×10^{-6} | 1.15 (1.08-1.22) | N | N | N | 73,705 | 823,607 | 0.66 |
| <i>PEPD</i> | rs10406327 | 19 | 33,890,838 | C | G | 52.3 | 47.7 | 3.8×10^{-8} | 1.04 (1.02-1.05) | N | N | Y | 74,117 | 823,996 | 0.99 |
| <i>TOMM40/APOE</i> | rs745903616 | 19 | 44,938,870 | A | G | 0.130 | 0.130 | 8.3×10^{-6} | 1.61 (1.30-1.98) | N | N | N | 61,325 | 782,180 | 0.82 |
| <i>TOMM40/APOE</i> | rs429358 | 19 | 45,411,941 | T | C | 84.6 | 15.4 | 2.6×10^{-18} | 1.08 (1.06-1.10) | N | N | Y | 74,117 | 823,997 | 0.92 |
| <i>GIPR</i> | rs10406431 | 19 | 46,157,019 | A | G | 56.3 | 43.8 | 9.6×10^{-14} | 1.05 (1.04-1.06) | N | N | Y | 74,117 | 823,996 | 0.95 |
| <i>GIPR</i> | rs2238689 | 19 | 46,178,661 | C | T | 41.8 | 41.8 | 5.4×10^{-9} | 1.04 (1.03-1.05) | N | N | N | 74,116 | 823,996 | 0.94 |
| <i>GIPR</i> | rs533172266 | 19 | 46,351,837 | T | C | 0.0600 | 0.0600 | 3.7×10^{-6} | 2.33 (1.63-3.33) | N | N | N | 47,435 | 495,952 | 0.70 |
| <i>ZC3H4</i> | rs3810291 | 19 | 47,569,003 | A | G | 67.3 | 32.7 | 8.9×10^{-12} | 1.05 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 0.94 |
| <i>NKX2.2</i> | rs13041756 | 20 | 21,466,795 | C | T | 10.7 | 10.7 | 1.4×10^{-8} | 1.06 (1.04-1.08) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| <i>RALY</i> | rs2268078 | 20 | 32,596,704 | A | G | 65.7 | 34.3 | 2.3×10^{-10} | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,995 | 0.98 |
| <i>HNF4A</i> | rs76811102 | 20 | 42,905,415 | T | C | 4.24 | 4.24 | 1.1×10^{-7} | 1.09 (1.06-1.13) | N | N | N | 74,117 | 823,997 | 0.83 |

| | | | | | | | | | | | | | | | |
|--------------------|-------------|----|------------|---|---|-------|------|-----------------------|------------------|---|---|---|--------|---------|------|
| <i>HNF4A</i> | rs4810426 | 20 | 43,001,721 | T | C | 10.6 | 10.6 | 3.1x10 ⁻¹⁷ | 1.09 (1.07-1.12) | N | N | N | 74,117 | 823,997 | 0.98 |
| <i>HNF4A</i> | rs191830490 | 20 | 43,023,355 | G | A | 99.4 | 0.59 | 2.2x10 ⁻⁶ | 1.24 (1.13-1.36) | N | N | N | 73,967 | 822,104 | 0.87 |
| <i>HNF4A</i> | rs1800961 | 20 | 43,042,364 | T | C | 3.53 | 3.53 | 2.3x10 ⁻²² | 1.18 (1.15-1.23) | N | N | Y | 74,117 | 823,997 | 0.98 |
| <i>HNF4A</i> | rs11696357 | 20 | 43,233,649 | A | G | 93.4 | 6.60 | 9.9x10 ⁻⁶ | 1.06 (1.04-1.09) | N | N | N | 74,117 | 823,997 | 0.75 |
| <i>EYA2</i> | rs560716466 | 20 | 45,317,678 | A | G | 0.310 | 0.31 | 9.8x10 ⁻⁶ | 1.36 (1.19-1.56) | N | Y | N | 69,739 | 808,452 | 0.71 |
| <i>EYA2</i> | rs6063048 | 20 | 45,598,564 | G | A | 72.5 | 27.5 | 2.2x10 ⁻¹¹ | 1.05 (1.03-1.06) | N | Y | Y | 74,117 | 823,996 | 0.99 |
| <i>CEPB</i> | rs11699802 | 20 | 48,832,135 | C | T | 53.6 | 46.4 | 1.8x10 ⁻¹¹ | 1.04 (1.03-1.06) | N | Y | Y | 74,116 | 823,996 | 0.98 |
| <i>TSHZ2</i> | rs34454109 | 20 | 51,223,594 | A | T | 77.1 | 22.9 | 7.1x10 ⁻⁹ | 1.04 (1.03-1.06) | N | Y | Y | 74,115 | 823,997 | 0.99 |
| <i>GNAS</i> | rs6070625 | 20 | 57,394,628 | G | C | 51.7 | 48.3 | 5.3x10 ⁻¹⁴ | 1.05 (1.04-1.06) | N | Y | Y | 74,117 | 823,995 | 1.00 |
| <i>GNAS</i> | rs862016 | 20 | 57,551,099 | G | A | 7.83 | 7.83 | 1.1x10 ⁻⁷ | 1.07 (1.04-1.09) | N | Y | N | 74,115 | 823,995 | 0.94 |
| <i>ZBTB46</i> | rs6011155 | 20 | 62,450,664 | T | C | 63.0 | 37.0 | 6.3x10 ⁻⁶ | 1.04 (1.02-1.05) | Y | Y | N | 50,402 | 523,887 | 0.96 |
| <i>TCEA2</i> | rs59944054 | 20 | 62,693,175 | A | G | 23.8 | 23.8 | 1.5x10 ⁻⁸ | 1.06 (1.04-1.08) | Y | Y | Y | 50,402 | 523,888 | 0.97 |
| <i>MTMR3/ASCC2</i> | rs6518681 | 22 | 30,609,554 | G | A | 91.4 | 8.64 | 1.1x10 ⁻¹² | 1.09 (1.06-1.11) | N | N | Y | 74,115 | 823,995 | 0.98 |
| <i>YWHAH</i> | rs117001013 | 22 | 32,348,841 | C | T | 91.2 | 8.83 | 1.7x10 ⁻⁸ | 1.07 (1.04-1.09) | N | Y | Y | 74,117 | 823,997 | 0.99 |
| <i>EP300</i> | rs5758223 | 22 | 41,489,920 | A | G | 71.7 | 28.3 | 3.8x10 ⁻⁸ | 1.04 (1.03-1.05) | N | Y | Y | 74,116 | 823,996 | 1.00 |
| <i>PNPLA3</i> | rs738408 | 22 | 44,324,730 | T | C | 22.6 | 22.6 | 1.4x10 ⁻¹⁰ | 1.05 (1.03-1.07) | N | N | Y | 74,116 | 823,996 | 0.98 |
| <i>PIM3</i> | rs1801645 | 22 | 50,356,850 | C | T | 27.5 | 27.5 | 1.5x10 ⁻⁸ | 1.04 (1.02-1.05) | N | N | Y | 74,116 | 823,996 | 0.85 |
| <i>PIM3</i> | rs112915006 | 22 | 50,604,696 | G | A | 5.12 | 5.12 | 9.8x10 ⁻⁷ | 1.08 (1.05-1.11) | N | N | N | 74,117 | 823,997 | 0.96 |

RAF: risk allele frequency; MAF: minor allele frequency; OR: Odds ratio; CI: confidence interval.

Supplementary Table 3 | Summary of comparison of effect estimates from BMI-adjusted and BMI-unadjusted analyses models obtained from the 28 studies the contributed to both analyses.

| Nearest gene | Index variant | Chromosome | Position (Build 37 bp) | Risk allele | Other allele | RAF (%) | MAF (%) | BMI-unadjusted | | | | BMI-unadjusted | | | | Heterogeneity p-value |
|---------------------|---------------|------------|------------------------|-------------|--------------|---------|---------|----------------|----------|-----------------------|------------------|----------------|----------|-----------------------|------------------|-----------------------|
| | | | | | | | | Cases | Controls | p-value | OR (95% CI) | Cases | Controls | p-value | OR (95% CI) | |
| <i>MACF1</i> | rs3768321 | 1 | 40,035,928 | T | G | 20.0 | 20.0 | 50,791 | 526,120 | 3.3x10 ⁻²⁰ | 1.09 (1.07-1.11) | 50,401 | 523,887 | 9.6x10 ⁻¹⁶ | 1.08 (1.06-1.10) | 0.05 |
| <i>FAF1</i> | rs58432198 | 1 | 51,256,091 | C | T | 88.1 | 11.9 | 50,791 | 521,613 | 3.9x10 ⁻⁸ | 1.07 (1.04-1.09) | 50,322 | 519,380 | 1.4x10 ⁻⁹ | 1.08 (1.05-1.10) | 0.1 |
| <i>PATJ</i> | rs12140153 | 1 | 62,579,891 | G | T | 90.5 | 9.49 | 50,791 | 526,121 | 1.2x10 ⁻⁸ | 1.08 (1.05-1.10) | 50,402 | 523,888 | 0.005 | 1.04 (1.01-1.07) | 9.3x10 ⁻¹⁰ |
| <i>DENND2C</i> | rs184660829 | 1 | 115,144,899 | C | T | 0.0200 | 0.0200 | 26,326 | 457,160 | 8.6x10 ⁻⁸ | 7.88 (3.70-16.8) | 26,115 | 455,486 | 9.8x10 ⁻⁷ | 6.42 (3.05-13.5) | 0.2 |
| <i>PTGFRN</i> | rs1127215 | 1 | 117,532,790 | C | T | 58.4 | 41.6 | 50,791 | 526,119 | 5.1x10 ⁻¹⁰ | 1.05 (1.03-1.06) | 50,402 | 523,888 | 6.2x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.3 |
| <i>NOTCH2</i> | rs1493694 | 1 | 120,526,982 | T | C | 10.9 | 10.9 | 50,791 | 526,121 | 8.0x10 ⁻¹⁴ | 1.09 (1.07-1.12) | 50,402 | 523,888 | 1.7x10 ⁻¹⁵ | 1.10 (1.07-1.13) | 0.1 |
| <i>FAM63A</i> | rs10305745 | 1 | 150,786,038 | A | G | 1.45 | 1.45 | 50,069 | 523,661 | 1.1x10 ⁻⁶ | 1.32 (1.18-1.48) | 49,681 | 521,430 | 4.5x10 ⁻⁶ | 1.32 (1.17-1.48) | 0.9 |
| <i>FAM63A</i> | rs145904381 | 1 | 151,017,991 | T | C | 98.7 | 1.33 | 50,791 | 526,121 | 2.5x10 ⁻⁷ | 1.20 (1.12-1.29) | 50,402 | 523,888 | 3.4x10 ⁻⁷ | 1.20 (1.12-1.29) | 0.9 |
| <i>SEC16B</i> | rs539515 | 1 | 177,889,025 | C | A | 19.8 | 19.8 | 50,790 | 526,120 | 2.6x10 ⁻¹⁰ | 1.06 (1.04-1.08) | 50,401 | 523,887 | 0.07 | 1.02 (1.00-1.04) | 2.7x10 ⁻²² |
| <i>DSTYK</i> | rs12048743 | 1 | 205,114,873 | G | C | 44.2 | 44.2 | 50,791 | 526,120 | 1.6x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,887 | 3.3x10 ⁻⁷ | 1.04 (1.02-1.05) | 1 |
| <i>SRGAP2</i> | rs9430095 | 1 | 206,593,900 | C | G | 49.4 | 49.4 | 50,790 | 526,120 | 0.00007 | 1.03 (1.02-1.04) | 50,401 | 523,886 | 0.00006 | 1.03 (1.02-1.05) | 0.8 |
| <i>PROX1</i> | rs79687284 | 1 | 214,150,821 | C | G | 3.48 | 3.48 | 50,791 | 526,121 | 4.3x10 ⁻¹⁹ | 1.21 (1.16-1.26) | 50,402 | 523,888 | 6.3x10 ⁻²¹ | 1.23 (1.18-1.28) | 0.2 |
| <i>PROX1</i> | rs340874 | 1 | 214,159,256 | C | T | 55.6 | 44.5 | 50,791 | 526,121 | 3.8x10 ⁻²¹ | 1.07 (1.06-1.09) | 50,402 | 523,888 | 4.8x10 ⁻²¹ | 1.07 (1.06-1.09) | 0.6 |
| <i>PROX1</i> | rs114526150 | 1 | 214,175,531 | G | T | 2.25 | 2.25 | 50,790 | 526,121 | 0.01 | 1.06 (1.01-1.12) | 50,402 | 523,888 | 0.009 | 1.07 (1.02-1.12) | 0.8 |
| <i>LYPLA1</i> | rs553014999 | 1 | 219,584,164 | C | T | 0.130 | 0.130 | 26,999 | 463,806 | 0.0007 | 2.22 (1.40-3.51) | 26,788 | 462,132 | 0.002 | 2.07 (1.31-3.28) | 0.5 |
| <i>LYPLA1</i> | rs2820446 | 1 | 219,748,818 | C | G | 70.6 | 29.5 | 50,791 | 526,121 | 2.7x10 ⁻¹⁰ | 1.05 (1.04-1.07) | 50,402 | 523,888 | 5.8x10 ⁻¹⁴ | 1.06 (1.05-1.08) | 0.002 |
| <i>ABCBL10</i> | rs348330 | 1 | 229,672,955 | G | A | 36.1 | 36.1 | 50,789 | 526,120 | 8.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 50,400 | 523,887 | 3.7x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.6 |
| <i>GNG4</i> | rs2913167 | 1 | 235,690,800 | G | A | 63.2 | 36.8 | 50,791 | 526,119 | 3.4x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,401 | 523,887 | 1.5x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.2 |
| <i>TMEM18</i> | rs62107261 | 2 | 422,144 | T | C | 95.4 | 4.64 | 50,791 | 526,121 | 3.6x10 ⁻⁹ | 1.11 (1.08-1.16) | 50,402 | 523,888 | 0.1 | 1.03 (0.99-1.07) | 3.9x10 ⁻²¹ |
| <i>TMEM18</i> | rs35913461 | 2 | 653,575 | C | T | 82.9 | 17.1 | 50,791 | 526,120 | 3.7x10 ⁻⁹ | 1.06 (1.04-1.08) | 50,402 | 523,888 | 0.3 | 1.01 (0.99-1.03) | 1.7x10 ⁻²⁶ |
| <i>FAM49A</i> | rs11680058 | 2 | 16,574,669 | A | G | 86.3 | 13.7 | 50,789 | 526,119 | 6.7x10 ⁻⁶ | 1.06 (1.03-1.08) | 50,400 | 523,886 | 7.9x10 ⁻⁶ | 1.06 (1.03-1.08) | 0.9 |
| <i>DTNB</i> | rs17802463 | 2 | 25,643,221 | G | T | 73.1 | 26.9 | 50,791 | 526,121 | 5.7x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,401 | 523,888 | 9.9x10 ⁻⁷ | 1.04 (1.02-1.06) | 1 |
| <i>GCKR</i> | rs1260326 | 2 | 27,730,940 | C | T | 60.7 | 39.3 | 50,791 | 526,121 | 9.6x10 ⁻²⁵ | 1.08 (1.06-1.09) | 50,402 | 523,888 | 7.4x10 ⁻¹⁹ | 1.07 (1.05-1.09) | 0.01 |
| <i>THADA</i> | rs28525376 | 2 | 43,207,872 | G | T | 42.2 | 42.2 | 50,790 | 526,119 | 0.009 | 1.02 (1.00-1.03) | 50,401 | 523,886 | 0.0008 | 1.03 (1.01-1.04) | 0.07 |
| <i>THADA</i> | rs6708643 | 2 | 43,430,440 | A | G | 50.1 | 49.9 | 50,790 | 526,119 | 2.1x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,401 | 523,886 | 3.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.009 |
| <i>THADA</i> | rs80147536 | 2 | 43,698,028 | A | T | 90.4 | 9.57 | 50,790 | 526,121 | 3.8x10 ⁻²⁵ | 1.14 (1.11-1.17) | 50,400 | 523,888 | 2.3x10 ⁻²⁶ | 1.15 (1.12-1.17) | 0.3 |
| <i>BNIPL</i> | rs10193538 | 2 | 58,981,064 | T | G | 61.0 | 39.0 | 50,790 | 526,120 | 2.5x10 ⁻⁹ | 1.05 (1.03-1.06) | 50,400 | 523,887 | 0.0002 | 1.03 (1.01-1.04) | 1.8x10 ⁻⁶ |
| <i>BNIPL</i> | rs6545714 | 2 | 59,307,725 | G | A | 39.2 | 39.2 | 50,790 | 526,120 | 4.3x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,400 | 523,887 | 0.003 | 1.02 (1.01-1.04) | 0.00001 |
| <i>BCL11A</i> | rs243024 | 2 | 60,583,665 | A | G | 46.0 | 46.0 | 50,790 | 526,120 | 7.2x10 ⁻¹⁵ | 1.06 (1.04-1.07) | 50,400 | 523,888 | 4.5x10 ⁻¹⁵ | 1.06 (1.05-1.08) | 0.6 |
| <i>CEP68</i> | rs2249105 | 2 | 65,287,896 | A | G | 63.4 | 36.6 | 50,791 | 526,121 | 1.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 50,402 | 523,888 | 8.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.8 |
| <i>CEP68</i> | rs2052261 | 2 | 65,355,270 | G | A | 30.4 | 30.4 | 50,790 | 526,120 | 0.0001 | 1.03 (1.02-1.05) | 50,401 | 523,887 | 0.00005 | 1.03 (1.02-1.05) | 0.6 |
| <i>CEP68</i> | rs2028150 | 2 | 65,655,012 | C | G | 59.8 | 40.2 | 50,791 | 526,121 | 5.1x10 ⁻¹¹ | 1.05 (1.03-1.07) | 50,401 | 523,888 | 7.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.8 |
| <i>TMEM127</i> | rs79046683 | 2 | 96,913,918 | T | G | 0.480 | 0.480 | 35,029 | 479,149 | 2.5x10 ⁻⁷ | 2.12 (1.59-2.81) | 34,775 | 477,429 | 3.0x10 ⁻⁸ | 2.34 (1.73-3.16) | 0.1 |
| <i>DDX18</i> | rs562386202 | 2 | 118,071,061 | G | A | 0.0600 | 0.0600 | 26,326 | 457,160 | 3.4x10 ⁻⁸ | 3.20 (2.12-4.85) | 26,115 | 455,486 | 1.8x10 ⁻⁸ | 3.31 (2.18-5.01) | 0.7 |
| <i>GLI2</i> | rs11688931 | 2 | 121,318,166 | C | G | 84.9 | 15.1 | 50,791 | 526,121 | 2.2x10 ⁻⁹ | 1.06 (1.04-1.08) | 50,402 | 523,888 | 2.1x10 ⁻¹⁰ | 1.07 (1.05-1.09) | 0.2 |
| <i>GLI2</i> | rs11688682 | 2 | 121,347,612 | G | C | 72.8 | 27.2 | 50,791 | 526,120 | 5.4x10 ⁻¹² | 1.06 (1.04-1.08) | 50,401 | 523,887 | 1.6x10 ⁻¹³ | 1.07 (1.05-1.09) | 0.2 |
| <i>GLI2</i> | rs6647705 | 2 | 121,378,852 | T | C | 96.7 | 3.30 | 50,791 | 526,121 | 0.0004 | 1.08 (1.04-1.13) | 50,402 | 523,888 | 0.0005 | 1.08 (1.03-1.13) | 0.9 |
| <i>PABPC1P2</i> | rs35999103 | 2 | 147,861,633 | T | C | 15.5 | 15.5 | 50,790 | 526,121 | 6.1x10 ⁻⁸ | 1.06 (1.04-1.08) | 50,401 | 523,888 | 3.6x10 ⁻⁶ | 1.05 (1.03-1.07) | 0.2 |
| <i>CYTIP</i> | rs134246680 | 2 | 158,339,550 | A | G | 93.7 | 6.27 | 50,791 | 526,120 | 1.0x10 ⁻⁷ | 1.08 (1.05-1.11) | 50,402 | 523,887 | 1.4x10 ⁻⁹ | 1.10 (1.06-1.13) | 0.05 |
| <i>RBMS1</i> | rs3772071 | 2 | 161,135,544 | T | C | 71.4 | 28.7 | 50,789 | 526,119 | 1.0x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,400 | 523,887 | 0.00005 | 1.03 (1.02-1.05) | 0.01 |
| <i>GRB14/COBLL1</i> | rs10195252 | 2 | 165,513,091 | T | C | 58.6 | 41.4 | 50,791 | 526,121 | 7.4x10 ⁻¹⁸ | 1.07 (1.05-1.08) | 50,402 | 523,888 | 5.7x10 ⁻²⁶ | 1.08 (1.07-1.10) | 1.3x10 ⁻⁶ |
| <i>GRB14/COBLL1</i> | rs13024606 | 2 | 165,573,194 | T | C | 4.72 | 4.72 | 50,791 | 526,121 | 0.0002 | 1.07 (1.03-1.11) | 50,402 | 523,888 | 0.001 | 1.06 (1.02-1.10) | 0.3 |
| <i>CRYBA2</i> | rs113414093 | 2 | 219,859,171 | A | G | 5.14 | 5.14 | 50,791 | 526,121 | 1.6x10 ⁻⁶ | 1.10 (1.06-1.14) | 50,402 | 523,888 | 6.6x10 ⁻⁹ | 1.12 (1.08-1.17) | 0.01 |
| <i>IRS1</i> | rs2972144 | 2 | 227,101,411 | G | A | 63.9 | 36.2 | 50,789 | 526,120 | 3.7x10 ⁻³⁷ | 1.10 (1.09-1.12) | 50,400 | 523,887 | 2.9x10 ⁻⁴⁰ | 1.11 (1.09-1.13) | 0.05 |
| <i>PPARG</i> | rs11709077 | 3 | 12,336,507 | G | A | 87.7 | 12.4 | 50,791 | 526,121 | 9.0x10 ⁻¹⁸ | 1.10 (1.08-1.12) | 50,402 | 523,888 | 5.0x10 ⁻²⁵ | 1.13 (1.10-1.15) | 9.8x10 ⁻⁶ |
| <i>PPARG</i> | rs17819328 | 3 | 12,489,342 | G | T | 42.5 | 42.5 | 50,791 | 526,121 | 2.6x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,402 | 523,888 | 1.2x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.7 |
| <i>UBE2E2</i> | rs35352848 | 3 | 23,455,582 | T | C | 78.8 | 21.2 | 50,791 | 526,121 | 2.9x10 ⁻²⁰ | 1.09 (1.07-1.11) | 50,402 | 523,888 | 5.4x10 ⁻²² | 1.09 (1.07-1.11) | 0.2 |
| <i>UBE2E2</i> | rs17013314 | 3 | 23,510,044 | G | A | 3.13 | 3.13 | 50,790 | 526,120 | 1.0x10 ⁻⁸ | 1.13 (1.09-1.18) | 50,401 | 523,888 | 6.1x10 ⁻⁸ | 1.13 (1.08-1.18) | 0.7 |
| <i>KIF9</i> | rs11926707 | 3 | 46,925,539 | C | T | 62.6 | 37.4 | 50,789 | 526,119 | 4.4x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 1.7x10 ⁻⁹ | 1.05 (1.03-1.06) | 0.1 |
| <i>KIF9</i> | rs75423501 | 3 | 47,242,923 | G | A | 10.1 | 10.1 | 50,710 | 521,612 | 1.1x10 ⁻⁶ | 1.07 (1.04-1.09) | 50,321 | 519,379 | 0.0005 | 1.05 (1.02-1.08) | 0.005 |
| <i>RBM6</i> | rs4688760 | 3 | 49,980,596 | T | C | 68.4 | 31.6 | 50,789 | 526,120 | 3.7x10 ⁻⁹ | 1.05 (1.03-1.06) | 50,401 | 523,887 | 0.0001 | 1.03 (1.01-1.04) | 2.8x10 ⁻⁸ |
| <i>RFT1</i> | rs2581787 | 3 | 53,127,677 | T | | | | | | | | | | | | |

| | | | | | | | | | | | | | | | | |
|-----------------|-------------|---|-------------|---|---|-------|-------|--------|---------|-----------------------|------------------|--------|---------|-----------------------|------------------|-----------------------|
| <i>TMCC1</i> | rs559138871 | 3 | 129,470,067 | T | C | 0.200 | 0.200 | 44,190 | 499,486 | 4.6x10 ⁻⁷ | 1.61 (1.34-1.94) | 43,891 | 497,703 | 1.3x10 ⁻⁶ | 1.59 (1.32-1.92) | 0.8 |
| <i>TSC22D2</i> | rs62271373 | 3 | 150,066,540 | A | T | 5.53 | 5.53 | 50,791 | 526,121 | 1.4x10 ⁻⁷ | 1.09 (1.06-1.12) | 50,402 | 523,888 | 2.5x10 ⁻⁹ | 1.10 (1.07-1.14) | 0.07 |
| <i>MBNL1</i> | rs13065698 | 3 | 152,086,533 | A | G | 60.0 | 40.0 | 50,791 | 526,121 | 2.9x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,402 | 523,887 | 0.00005 | 1.03 (1.02-1.05) | 0.3 |
| <i>MBNL1</i> | rs74653713 | 3 | 152,417,881 | C | A | 95.7 | 4.29 | 50,790 | 526,121 | 1.5x10 ⁻⁹ | 1.12 (1.08-1.17) | 50,401 | 523,888 | 8.3x10 ⁻¹¹ | 1.14 (1.09-1.18) | 0.2 |
| <i>MBNL1</i> | rs35497231 | 3 | 152,433,628 | C | T | 31.7 | 31.7 | 50,789 | 526,119 | 4.2x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,401 | 523,886 | 2.8x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.7 |
| <i>EGFEM1P</i> | rs7629630 | 3 | 168,218,841 | A | T | 85.7 | 14.3 | 50,791 | 526,121 | 1.8x10 ⁻⁶ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 5.8x10 ⁻⁶ | 1.05 (1.03-1.07) | 0.8 |
| <i>SLC2A2</i> | rs9873618 | 3 | 170,733,076 | G | A | 71.0 | 29.0 | 50,791 | 526,120 | 1.2x10 ⁻¹⁵ | 1.07 (1.05-1.08) | 50,402 | 523,888 | 2.7x10 ⁻²¹ | 1.08 (1.06-1.10) | 0.0002 |
| <i>ABCC5</i> | rs2872246 | 3 | 183,738,460 | A | C | 45.4 | 45.4 | 50,791 | 526,120 | 3.0x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,401 | 523,887 | 0.0001 | 1.03 (1.01-1.04) | 0.01 |
| <i>IGF2BP2</i> | rs6780171 | 3 | 185,503,456 | A | T | 31.4 | 31.4 | 50,789 | 526,120 | 4.4x10 ⁻⁴⁵ | 1.12 (1.10-1.13) | 50,401 | 523,887 | 2.4x10 ⁻⁵¹ | 1.13 (1.11-1.15) | 0.003 |
| <i>IGF2BP2</i> | rs150111048 | 3 | 185,514,421 | G | A | 23.9 | 23.9 | 45,267 | 501,592 | 0.05 | 1.21 (1.00-1.47) | 45,638 | 500,506 | 0.1 | 1.17 (0.96-1.42) | 0.4 |
| <i>IGF2BP2</i> | rs11717959 | 3 | 185,541,213 | G | T | 62.1 | 37.9 | 50,791 | 526,121 | 3.3x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,401 | 523,888 | 1.4x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.1 |
| <i>IGF2BP2</i> | rs1516728 | 3 | 185,829,891 | A | T | 75.9 | 24.1 | 50,789 | 526,119 | 0.0004 | 1.03 (1.01-1.05) | 50,400 | 523,886 | 0.2 | 1.01 (0.99-1.03) | 1.2x10 ⁻⁶ |
| <i>ST6GAL1</i> | rs3887925 | 3 | 186,665,645 | T | C | 54.7 | 45.3 | 50,791 | 526,121 | 2.1x10 ⁻¹⁴ | 1.06 (1.04-1.07) | 50,402 | 523,888 | 3.7x10 ⁻¹⁶ | 1.06 (1.05-1.08) | 0.1 |
| <i>ST6GAL1</i> | rs7645517 | 3 | 186,675,277 | A | G | 5.76 | 5.76 | 50,791 | 526,120 | 0.0003 | 1.05 (1.02-1.08) | 50,402 | 523,887 | 0.006 | 1.05 (1.01-1.08) | 0.7 |
| <i>LPP</i> | rs4686471 | 3 | 187,740,899 | C | T | 61.0 | 39.0 | 50,789 | 526,119 | 1.1x10 ⁻¹⁹ | 1.07 (1.05-1.09) | 50,400 | 523,886 | 1.1x10 ⁻¹⁹ | 1.07 (1.06-1.09) | 0.6 |
| <i>PCGF3</i> | rs11827885 | 4 | 616,608 | C | T | 1.56 | 1.56 | 45,840 | 512,001 | 0.00009 | 1.17 (1.08-1.26) | 45,601 | 511,670 | 0.00002 | 1.19 (1.10-1.28) | 0.3 |
| <i>PCGF3</i> | rs1531583 | 4 | 744,972 | T | G | 4.58 | 4.58 | 50,791 | 526,121 | 1.1x10 ⁻¹⁰ | 1.13 (1.09-1.17) | 50,402 | 523,888 | 8.8x10 ⁻¹² | 1.14 (1.10-1.18) | 0.2 |
| <i>PCGF3</i> | rs35654957 | 4 | 1,010,077 | C | T | 36.7 | 36.7 | 50,790 | 526,120 | 0.0007 | 1.03 (1.01-1.04) | 50,401 | 523,887 | 0.0001 | 1.03 (1.01-1.05) | 0.2 |
| <i>MAEA</i> | rs56337234 | 4 | 1,784,403 | C | T | 50.3 | 49.7 | 50,790 | 526,119 | 1.2x10 ⁻¹⁵ | 1.06 (1.05-1.08) | 50,401 | 523,886 | 1.3x10 ⁻²¹ | 1.08 (1.06-1.09) | 0.00008 |
| <i>HTT</i> | rs362307 | 4 | 3,241,845 | T | C | 7.68 | 7.68 | 50,791 | 526,121 | 4.8x10 ⁻⁷ | 1.07 (1.04-1.10) | 50,402 | 523,888 | 0.0005 | 1.05 (1.02-1.08) | 0.001 |
| <i>WFS1</i> | rs1801212 | 4 | 6,302,519 | A | G | 70.9 | 29.1 | 50,791 | 526,121 | 1.3x10 ⁻³⁴ | 1.10 (1.09-1.12) | 50,402 | 523,886 | 1.3x10 ⁻³¹ | 1.10 (1.08-1.12) | 0.5 |
| <i>WFS1</i> | rs10937721 | 4 | 6,306,763 | C | G | 58.8 | 41.2 | 50,789 | 526,119 | 1.3x10 ⁻³³ | 1.09 (1.08-1.11) | 50,400 | 523,886 | 1.8x10 ⁻³¹ | 1.09 (1.08-1.11) | 0.6 |
| <i>LCORL</i> | rs12640250 | 4 | 17,792,869 | C | A | 71.5 | 28.5 | 50,789 | 526,119 | 0.00003 | 1.03 (1.02-1.05) | 50,400 | 523,886 | 0.0004 | 1.03 (1.01-1.05) | 0.2 |
| <i>GNPDA2</i> | rs10938398 | 4 | 45,186,139 | A | G | 42.9 | 42.9 | 50,790 | 526,120 | 1.4x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,400 | 523,887 | 0.1 | 1.01 (1.00-1.03) | 1.5x10 ⁻¹⁶ |
| <i>USP46</i> | rs2102278 | 4 | 52,818,664 | G | A | 31.9 | 31.9 | 50,790 | 526,119 | 3.4x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 0.0002 | 1.03 (1.01-1.05) | 0.06 |
| <i>USP46</i> | rs114447556 | 4 | 53,207,093 | T | C | 8.39 | 8.39 | 50,790 | 526,121 | 0.00002 | 1.06 (1.03-1.09) | 50,401 | 523,888 | 0.002 | 1.05 (1.02-1.08) | 0.02 |
| <i>SCD5</i> | rs12642790 | 4 | 83,578,271 | A | G | 33.8 | 33.8 | 50,790 | 526,120 | 2.5x10 ⁻⁹ | 1.05 (1.03-1.06) | 50,400 | 523,887 | 1.0x10 ⁻⁸ | 1.05 (1.03-1.06) | 0.9 |
| <i>FAM13A</i> | rs1903002 | 4 | 89,740,894 | G | C | 50.1 | 50.0 | 50,789 | 526,119 | 0.00003 | 1.03 (1.02-1.05) | 50,400 | 523,886 | 2.7x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.2 |
| <i>FAM13A</i> | rs576406049 | 4 | 89,857,291 | T | C | 0.130 | 0.130 | 39,467 | 496,986 | 2.2x10 ⁻⁶ | 1.82 (1.42-2.33) | 44,289 | 502,471 | 8.1x10 ⁻⁷ | 1.87 (1.46-2.39) | 0.6 |
| <i>SMARCAD1</i> | rs6821438 | 4 | 95,091,911 | A | G | 53.4 | 46.6 | 50,790 | 526,120 | 3.4x10 ⁻⁶ | 1.03 (1.02-1.05) | 50,401 | 523,887 | 0.0005 | 1.03 (1.01-1.04) | 0.02 |
| <i>SLC9B1</i> | rs1580278 | 4 | 104,140,848 | C | A | 47.3 | 47.3 | 50,789 | 526,119 | 1.9x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 6.1x10 ⁻⁷ | 1.04 (1.02-1.05) | 0.3 |
| <i>PABPC4L</i> | rs1296328 | 4 | 137,083,193 | A | C | 44.6 | 44.6 | 50,790 | 526,119 | 1.0x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 0.001 | 1.02 (1.01-1.04) | 0.0005 |
| <i>TMEM154</i> | rs7669833 | 4 | 153,513,369 | T | A | 70.5 | 29.6 | 50,790 | 526,121 | 5.8x10 ⁻¹¹ | 1.05 (1.04-1.07) | 50,401 | 523,888 | 4.4x10 ⁻¹³ | 1.06 (1.04-1.08) | 0.06 |
| <i>PDGFC</i> | rs28819812 | 4 | 157,652,753 | C | A | 67.7 | 32.3 | 50,789 | 526,121 | 3.6x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,401 | 523,888 | 7.7x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.4 |
| <i>ACSL1</i> | rs58730668 | 4 | 185,717,759 | T | C | 85.8 | 14.2 | 50,791 | 526,121 | 4.2x10 ⁻¹⁰ | 1.07 (1.05-1.09) | 50,402 | 523,888 | 2.5x10 ⁻¹¹ | 1.07 (1.05-1.10) | 0.2 |
| <i>ANKH</i> | rs3845281 | 5 | 14,610,134 | G | A | 90.4 | 9.61 | 50,791 | 526,121 | 0.00007 | 1.05 (1.03-1.08) | 50,402 | 523,888 | 0.0003 | 1.05 (1.02-1.07) | 0.5 |
| <i>ANKH</i> | rs78408340 | 5 | 14,751,305 | C | T | 99.4 | 0.620 | 47,969 | 516,095 | 6.7x10 ⁻¹⁴ | 1.47 (1.33-1.63) | 47,581 | 513,864 | 5.6x10 ⁻¹⁴ | 1.48 (1.34-1.64) | 0.9 |
| <i>ANKH</i> | rs17250977 | 5 | 14,753,745 | G | A | 3.76 | 3.76 | 50,791 | 526,121 | 1.2x10 ⁻¹² | 1.15 (1.11-1.20) | 50,402 | 523,888 | 9.5x10 ⁻¹⁴ | 1.16 (1.12-1.21) | 0.3 |
| <i>ANKH</i> | rs68881532 | 5 | 14,768,092 | C | G | 90.4 | 9.60 | 50,791 | 526,121 | 1.6x10 ⁻¹² | 1.09 (1.07-1.12) | 50,402 | 523,888 | 6.6x10 ⁻¹⁴ | 1.10 (1.07-1.13) | 0.2 |
| <i>ANKH</i> | rs76549217 | 5 | 14,768,766 | T | C | 2.95 | 2.95 | 50,791 | 526,121 | 1.8x10 ⁻⁶ | 1.12 (1.07-1.17) | 50,402 | 523,888 | 1.1x10 ⁻⁷ | 1.13 (1.08-1.19) | 0.2 |
| <i>MRPS30</i> | rs62368490 | 5 | 44,534,364 | T | C | 3.13 | 3.13 | 50,712 | 521,614 | 0.00003 | 1.10 (1.05-1.15) | 50,323 | 519,381 | 0.0001 | 1.09 (1.04-1.14) | 0.6 |
| <i>MRPS30</i> | rs6884702 | 5 | 44,682,589 | G | A | 39.3 | 39.3 | 50,711 | 521,614 | 6.1x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,323 | 519,380 | 5.7x10 ⁻⁷ | 1.04 (1.02-1.05) | 0.7 |
| <i>ITGA1</i> | rs17261179 | 5 | 51,791,225 | T | C | 51.7 | 48.3 | 50,790 | 526,120 | 1.6x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,401 | 523,887 | 0.00002 | 1.03 (1.02-1.05) | 0.3 |
| <i>ITGA1</i> | rs3811978 | 5 | 52,100,489 | G | A | 16.7 | 16.7 | 50,790 | 526,120 | 8.3x10 ⁻⁸ | 1.05 (1.03-1.07) | 50,401 | 523,887 | 9.1x10 ⁻⁹ | 1.06 (1.04-1.08) | 0.3 |
| <i>ITGA1</i> | rs62357230 | 5 | 52,315,682 | A | G | 3.39 | 3.39 | 50,791 | 526,121 | 0.00004 | 1.09 (1.04-1.13) | 50,401 | 523,888 | 0.00003 | 1.09 (1.05-1.14) | 0.6 |
| <i>ARL15</i> | rs62370480 | 5 | 52,774,510 | A | G | 22.0 | 22.0 | 50,790 | 526,121 | 4.8x10 ⁻⁷ | 1.05 (1.03-1.06) | 50,401 | 523,888 | 1.7x10 ⁻⁷ | 1.05 (1.03-1.07) | 0.5 |
| <i>ARL15</i> | rs702634 | 5 | 53,271,420 | A | G | 69.0 | 31.0 | 50,790 | 526,121 | 1.1x10 ⁻⁹ | 1.05 (1.03-1.07) | 50,401 | 523,887 | 9.7x10 ⁻¹⁴ | 1.06 (1.05-1.08) | 0.0007 |
| <i>ARL15</i> | rs279744 | 5 | 53,412,620 | C | A | 69.1 | 30.9 | 50,791 | 526,121 | 0.00008 | 1.03 (1.02-1.05) | 50,402 | 523,888 | 0.0001 | 1.03 (1.02-1.05) | 1 |
| <i>ANKRD55</i> | rs465002 | 5 | 55,808,475 | T | C | 74.2 | 25.8 | 50,789 | 526,119 | 6.0x10 ⁻²⁰ | 1.08 (1.06-1.10) | 50,401 | 523,886 | 7.6x10 ⁻²¹ | 1.08 (1.06-1.10) | 0.3 |
| <i>ANKRD55</i> | rs2431115 | 5 | 55,848,669 | A | G | 40.2 | 40.2 | 50,789 | 526,119 | 0.004 | 1.02 (1.01-1.04) | 50,400 | 523,886 | 0.0007 | 1.03 (1.01-1.04) | 0.2 |
| <i>ANKRD55</i> | rs9687832 | 5 | 55,861,595 | A | G | 19.8 | 19.8 | 50,791 | 526,121 | 2.2x10 ⁻¹² | 1.07 (1.05-1.09) | 50,401 | 523,888 | 8.0x10 ⁻¹³ | 1.07 (1.05-1.09) | 0.5 |
| <i>ANKRD55</i> | rs96844 | 5 | 56,196,604 | G | A | 26.2 | 26.2 | 50,790 | 526,119 | 2.8x10 ⁻⁸ | 1.05 (1.03-1.06) | 50,400 | 523,887 | 4.2x10 ⁻⁷ | 1.04 (1.03-1.06) | 0.5 |
| <i>PIK3R1</i> | rs4976033 | 5 | 67,714,246 | G | A | 41.1 | 41.1 | 50,790 | 526,120 | 2.7x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,401 | 523,887 | 1.0x10 ⁻⁹ | 1.05 (1.03-1.06) | 0.01 |
| <i>POCS</i> | rs2307111 | 5 | 75,003,678 | T | C | 60.5 | 39.5 | 50,791 | 526,121 | 6.5x10 ⁻¹³ | 1.05 (1.04-1.07) | 50,402 | 523,888 | 0.0005 | 1.03 (1.01-1.04) | 4.5x10 ⁻¹⁵ |

| | | | | | | | | | | | | | | | | |
|-----------|-------------|---|-------------|---|---|--------|--------|--------|---------|-----------------------|------------------|--------|---------|-----------------------|------------------|-----------------------|
| VEGFA | rs11967262 | 6 | 43,760,327 | G | C | 48.6 | 48.6 | 50,790 | 526,119 | 3.5x10 ⁻⁹ | 1.04 (1.03-1.06) | 50,401 | 523,886 | 3.7x10 ⁻¹³ | 1.06 (1.04-1.07) | 0.0007 |
| VEGFA | rs6458354 | 6 | 43,814,190 | C | T | 28.9 | 28.9 | 50,790 | 526,119 | 3.4x10 ⁻⁹ | 1.05 (1.03-1.07) | 50,401 | 523,886 | 7.4x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.1 |
| TFAP2B | rs3798519 | 6 | 50,788,778 | C | A | 18.4 | 18.4 | 50,791 | 526,121 | 4.7x10 ⁻¹² | 1.07 (1.05-1.09) | 50,402 | 523,888 | 0.0002 | 1.04 (1.02-1.06) | 2.4x10 ⁻¹¹ |
| TFAP2B | rs2465043 | 6 | 51,180,765 | G | A | 64.4 | 35.6 | 50,790 | 526,121 | 0.00002 | 1.03 (1.02-1.05) | 50,401 | 523,888 | 0.004 | 1.02 (1.01-1.04) | 0.004 |
| SLC25A5P1 | rs555402748 | 6 | 67,387,490 | T | C | 0.0400 | 0.0400 | 36,772 | 475,836 | 1.6x10 ⁻⁷ | 3.67 (2.25-5.96) | 36,490 | 474,088 | 1.1x10 ⁻⁸ | 4.20 (2.57-6.88) | 0.2 |
| BEND3 | rs4946812 | 6 | 107,431,688 | G | A | 67.4 | 32.6 | 50,791 | 526,121 | 8.5x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,402 | 523,888 | 1.8x10 ⁻⁶ | 1.04 (1.02-1.06) | 1 |
| CENPW | rs11759026 | 6 | 126,792,095 | G | A | 23.2 | 23.2 | 50,791 | 526,121 | 6.1x10 ⁻¹⁴ | 1.07 (1.05-1.09) | 50,402 | 523,888 | 1.2x10 ⁻¹⁶ | 1.08 (1.06-1.10) | 0.03 |
| SOGA3 | rs2800733 | 6 | 127,416,930 | A | G | 71.7 | 28.4 | 50,791 | 526,121 | 4.3x10 ⁻¹² | 1.06 (1.04-1.07) | 50,402 | 523,888 | 2.4x10 ⁻¹⁴ | 1.07 (1.05-1.08) | 0.05 |
| SLC35D3 | rs9494624 | 6 | 137,300,960 | A | G | 29.0 | 29.0 | 50,790 | 526,119 | 2.0x10 ⁻⁶ | 1.04 (1.02-1.06) | 50,401 | 523,886 | 8.3x10 ⁻⁸ | 1.05 (1.03-1.06) | 0.1 |
| MIR3668 | rs2982521 | 6 | 139,835,329 | A | T | 38.0 | 38.0 | 50,790 | 526,120 | 8.1x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,887 | 8.8x10 ⁻⁹ | 1.05 (1.03-1.06) | 0.03 |
| MIR3668 | rs616279 | 6 | 140,249,466 | A | G | 73.8 | 26.2 | 50,710 | 521,612 | 0.00002 | 1.04 (1.02-1.05) | 50,321 | 519,379 | 4.7x10 ⁻⁶ | 1.04 (1.02-1.06) | 0.4 |
| SLC22A3 | rs474513 | 6 | 160,770,312 | A | G | 51.7 | 48.3 | 50,791 | 526,121 | 8.1x10 ⁻⁹ | 1.04 (1.03-1.06) | 50,402 | 523,888 | 1.3x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.009 |
| QKI | rs4709746 | 6 | 164,133,001 | C | T | 86.8 | 13.2 | 50,791 | 526,121 | 8.7x10 ⁻⁹ | 1.07 (1.04-1.09) | 50,402 | 523,888 | 3.5x10 ⁻¹¹ | 1.08 (1.05-1.10) | 0.03 |
| DGKB | rs17168486 | 7 | 14,898,282 | T | C | 18.1 | 18.1 | 50,791 | 526,120 | 3.1x10 ⁻¹⁴ | 1.08 (1.06-1.10) | 50,401 | 523,887 | 7.5x10 ⁻¹⁴ | 1.08 (1.06-1.10) | 0.9 |
| DGKB | rs10228066 | 7 | 15,063,569 | T | C | 53.7 | 46.3 | 50,790 | 526,119 | 2.9x10 ⁻²¹ | 1.07 (1.06-1.09) | 50,401 | 523,887 | 2.2x10 ⁻²¹ | 1.07 (1.06-1.09) | 0.5 |
| DGKB | rs2908334 | 7 | 15,206,239 | T | C | 63.1 | 36.9 | 50,791 | 526,120 | 0.05 | 1.01 (1.00-1.03) | 50,401 | 523,887 | 0.08 | 1.01 (1.00-1.03) | 0.8 |
| IGF2BP3 | rs78840640 | 7 | 23,434,606 | G | C | 2.0 | 2.20 | 50,791 | 526,121 | 3.7x10 ⁻⁶ | 1.12 (1.07-1.18) | 50,402 | 523,888 | 3.1x10 ⁻⁷ | 1.14 (1.09-1.20) | 0.2 |
| IGF2BP3 | rs4279506 | 7 | 23,512,896 | G | C | 61.0 | 39.0 | 50,790 | 526,121 | 1.5x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,401 | 523,888 | 2.8x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.9 |
| JAZF1 | rs1708302 | 7 | 28,198,677 | C | T | 51.2 | 48.8 | 50,790 | 526,120 | 6.4x10 ⁻³¹ | 1.09 (1.07-1.10) | 50,401 | 523,887 | 8.7x10 ⁻⁴¹ | 1.11 (1.09-1.12) | 2.0x10 ⁻⁶ |
| CRHR2 | rs917195 | 7 | 30,728,452 | C | T | 77.0 | 23.0 | 50,791 | 526,121 | 7.8x10 ⁻⁸ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 1.4x10 ⁻⁶ | 1.04 (1.03-1.06) | 0.3 |
| GCK | rs878521 | 7 | 44,255,643 | A | G | 24.5 | 24.5 | 50,791 | 526,121 | 2.8x10 ⁻¹¹ | 1.06 (1.04-1.08) | 50,402 | 523,888 | 8.7x10 ⁻¹³ | 1.06 (1.05-1.08) | 0.1 |
| GCK | rs116913033 | 7 | 44,365,549 | C | T | 83.0 | 17.0 | 50,791 | 526,121 | 0.00002 | 1.04 (1.02-1.06) | 50,402 | 523,888 | 0.0002 | 1.04 (1.02-1.06) | 0.3 |
| FBXL13 | rs56376556 | 7 | 102,038,318 | T | C | 5.33 | 5.33 | 50,791 | 526,121 | 4.3x10 ⁻⁷ | 1.10 (1.06-1.14) | 50,402 | 523,888 | 1.9x10 ⁻⁶ | 1.09 (1.05-1.13) | 0.7 |
| FBXL13 | rs11496066 | 7 | 102,486,254 | T | C | 81.8 | 18.2 | 50,791 | 526,121 | 7.3x10 ⁻⁸ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 1.6x10 ⁻⁷ | 1.05 (1.03-1.07) | 0.9 |
| RELN | rs624284205 | 7 | 102,987,583 | G | T | 8.19 | 8.19 | 50,791 | 526,120 | 8.8x10 ⁻⁷ | 1.07 (1.04-1.10) | 50,402 | 523,887 | 1.3x10 ⁻⁶ | 1.07 (1.04-1.10) | 0.9 |
| RELN | rs39328 | 7 | 103,444,978 | T | C | 43.3 | 43.3 | 50,791 | 526,121 | 4.5x10 ⁻⁶ | 1.03 (1.02-1.05) | 50,402 | 523,888 | 0.005 | 1.02 (1.01-1.04) | 0.0002 |
| CTTNBP2 | rs6976111 | 7 | 117,495,667 | A | C | 31.3 | 31.3 | 50,790 | 526,121 | 3.9x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,402 | 523,888 | 4.9x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.4 |
| KLF14 | rs268382 | 7 | 130,027,037 | C | A | 32.7 | 32.7 | 50,791 | 526,121 | 0.0001 | 1.03 (1.01-1.05) | 50,402 | 523,887 | 0.001 | 1.03 (1.01-1.04) | 0.4 |
| KLF14 | rs1562396 | 7 | 130,457,914 | G | A | 31.9 | 31.9 | 50,790 | 526,121 | 8.1x10 ⁻¹⁴ | 1.06 (1.04-1.08) | 50,401 | 523,888 | 1.1x10 ⁻¹⁶ | 1.07 (1.05-1.09) | 0.02 |
| AOC1 | rs62492368 | 7 | 150,537,635 | A | G | 30.8 | 30.8 | 50,791 | 526,121 | 2.4x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 3.8x10 ⁻¹⁰ | 1.05 (1.04-1.07) | 0.8 |
| MNX1 | rs6459733 | 7 | 156,930,550 | G | C | 67.3 | 32.7 | 50,791 | 526,121 | 7.1x10 ⁻¹⁷ | 1.07 (1.05-1.08) | 50,402 | 523,888 | 1.0x10 ⁻¹⁴ | 1.06 (1.05-1.08) | 0.4 |
| MSRA | rs17689007 | 8 | 9,974,824 | G | A | 53.3 | 46.7 | 50,711 | 521,614 | 2.5x10 ⁻¹¹ | 1.05 (1.04-1.07) | 50,323 | 519,380 | 3.2x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.00004 |
| XKR6 | rs57327348 | 8 | 10,808,687 | A | T | 78.2 | 21.8 | 50,712 | 521,614 | 1.1x10 ⁻⁷ | 1.05 (1.03-1.07) | 50,323 | 519,381 | 0.0003 | 1.03 (1.02-1.05) | 0.0006 |
| LPL | rs10096633 | 8 | 19,830,921 | C | T | 87.7 | 12.3 | 50,791 | 526,121 | 2.9x10 ⁻⁹ | 1.07 (1.05-1.09) | 50,402 | 523,888 | 2.1x10 ⁻¹⁰ | 1.08 (1.05-1.10) | 0.2 |
| PURG | rs10954772 | 8 | 30,863,938 | T | C | 31.4 | 31.4 | 50,789 | 526,119 | 6.0x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 0.0003 | 1.03 (1.01-1.05) | 0.0002 |
| ANK1 | rs13262861 | 8 | 41,508,577 | C | A | 82.9 | 17.1 | 50,791 | 526,121 | 3.2x10 ⁻²⁷ | 1.11 (1.09-1.13) | 50,402 | 523,888 | 7.4x10 ⁻²⁷ | 1.11 (1.09-1.14) | 0.8 |
| ANK1 | rs4736819 | 8 | 41,509,915 | T | C | 55.4 | 44.6 | 50,789 | 526,119 | 1.1x10 ⁻²⁴ | 1.08 (1.06-1.09) | 50,401 | 523,886 | 1.3x10 ⁻²² | 1.08 (1.06-1.09) | 0.7 |
| ANK1 | rs14876668 | 8 | 41,552,046 | C | T | 3.78 | 3.78 | 50,791 | 526,121 | 3.2x10 ⁻¹⁰ | 1.13 (1.09-1.18) | 50,402 | 523,888 | 1.4x10 ⁻¹⁰ | 1.14 (1.10-1.19) | 0.5 |
| TP53INP1 | rs11786992 | 8 | 95,685,147 | A | C | 64.4 | 35.6 | 50,791 | 526,121 | 3.8x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,402 | 523,888 | 1.1x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.001 |
| TP53INP1 | rs10097617 | 8 | 95,961,626 | T | C | 48.5 | 48.5 | 50,791 | 526,120 | 5.6x10 ⁻¹² | 1.05 (1.04-1.07) | 50,402 | 523,886 | 2.0x10 ⁻¹¹ | 1.05 (1.04-1.07) | 1 |
| TP53INP1 | rs187936726 | 8 | 96,092,422 | G | A | 2.39 | 2.39 | 50,791 | 526,121 | 1.3x10 ⁻⁶ | 1.14 (1.08-1.20) | 50,402 | 523,888 | 1.7x10 ⁻⁶ | 1.14 (1.08-1.21) | 0.8 |
| CPQ | rs149364428 | 8 | 97,737,741 | A | G | 1.04 | 1.04 | 50,641 | 524,228 | 1.2x10 ⁻¹¹ | 1.32 (1.22-1.43) | 50,252 | 521,995 | 3.8x10 ⁻¹⁰ | 1.30 (1.20-1.41) | 0.4 |
| TRHR | rs12680028 | 8 | 110,123,183 | C | G | 53.4 | 46.6 | 50,791 | 526,121 | 2.4x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,888 | 6.6x10 ⁻⁶ | 1.03 (1.02-1.05) | 0.2 |
| SLC30A8 | rs3802177 | 8 | 118,185,025 | G | A | 68.5 | 31.5 | 50,790 | 526,120 | 3.1x10 ⁻⁴⁵ | 1.12 (1.1-1.140) | 50,401 | 523,888 | 3.8x10 ⁻⁵³ | 1.13 (1.11-1.15) | 0.0004 |
| SLC30A8 | rs80244329 | 8 | 118,404,672 | G | A | 97.8 | 2.19 | 50,791 | 526,121 | 0.00005 | 1.12 (1.06-1.18) | 50,402 | 523,888 | 0.00002 | 1.13 (1.07-1.19) | 0.5 |
| CASC11 | rs17772814 | 8 | 128,711,742 | G | A | 91.5 | 8.49 | 50,791 | 526,121 | 1.0x10 ⁻⁶ | 1.07 (1.04-1.11) | 50,402 | 523,888 | 4.0x10 ⁻⁸ | 1.09 (1.05-1.12) | 0.1 |
| PVT1 | rs1561927 | 8 | 129,568,078 | C | T | 26.9 | 26.9 | 50,791 | 526,121 | 1.4x10 ⁻⁶ | 1.04 (1.02-1.06) | 50,402 | 523,888 | 2.9x10 ⁻⁶ | 1.04 (1.02-1.06) | 0.9 |
| BOP1 | rs4977213 | 8 | 145,507,304 | C | T | 37.5 | 37.5 | 50,789 | 526,119 | 6.6x10 ⁻¹² | 1.05 (1.04-1.07) | 50,400 | 523,886 | 1.2x10 ⁻¹⁵ | 1.06 (1.05-1.08) | 0.003 |
| BOP1 | rs12719778 | 8 | 145,879,883 | T | C | 53.8 | 46.2 | 50,791 | 526,120 | 2.4x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,886 | 8.3x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.5 |
| GLIS3 | rs510807 | 9 | 3,965,689 | A | C | 49.1 | 49.1 | 50,789 | 526,119 | 0.0005 | 1.03 (1.01-1.04) | 50,400 | 523,887 | 0.01 | 1.02 (1.00-1.03) | 0.04 |
| GLIS3 | rs79103584 | 9 | 4,243,045 | T | A | 98.6 | 1.38 | 50,791 | 526,120 | 0.0001 | 1.14 (1.06-1.21) | 50,402 | 523,887 | 0.00002 | 1.15 (1.08-1.23) | 0.3 |
| GLIS3 | rs10974438 | 9 | 4,291,928 | C | A | 35.7 | 35.7 | 50,790 | 526,121 | 1.7x10 ⁻¹³ | 1.06 (1.04-1.07) | 50,401 | 523,887 | 3.2x10 ⁻¹⁴ | 1.06 (1.04-1.08) | 0.4 |
| HAUS6 | rs7022807 | 9 | 19,067,833 | G | A | 40.1 | 40.1 | 50,790 | 526,121 | 4.9x10 ⁻⁹ | 1.04 (1.03-1.06) | 50,401 | 523,888 | 3.2x10 ⁻¹⁰ | 1.05 (1.03-1.06) | 0.2 |
| FOCAD | rs7867635 | 9 | 20,241,069 | C | T | 41.2 | 41.2 | 50,789 | 526,119 | 2.4x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 0.00006 | 1.03 (1.02-1.05) | 0.2 |
| FOCAD | rs7847880 | 9 | 20,662,703 | C | T | 84.3 | 15.7 | 50,789 | 526,119 | 0.0002 | 1.04 (1.02-1.06) | 50,400 | | | | |

| | | | | | | | | | | | | | | | | |
|----------------------|-----------------|----|-------------|---|---|--------|--------|--------|---------|------------------------|------------------|--------|---------|------------------------|------------------|-----------------------|
| <i>GPSM1</i> | rs11793035 | 9 | 139,507,212 | C | T | 33.1 | 33.1 | 50,789 | 526,119 | 7.2x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 2.8x10 ⁻⁷ | 1.04 (1.03-1.06) | 0.09 |
| <i>GPSM1</i> | 9:139737088:G:A | 9 | 139,737,088 | A | G | 0.0700 | 0.0700 | 31,447 | 464,429 | 7.1x10 ⁻⁶ | 2.73 (1.76-4.23) | 31,207 | 462,716 | 0.0003 | 2.33 (1.48-3.66) | 0.1 |
| <i>CDC123/CAMK1D</i> | rs11257655 | 10 | 12,307,894 | T | C | 21.8 | 21.8 | 50,791 | 526,121 | 3.8x10 ⁻²⁴ | 1.09 (1.07-1.11) | 50,402 | 523,888 | 7.7x10 ⁻²⁶ | 1.10 (1.08-1.12) | 0.2 |
| <i>NEUROG3</i> | rs177045 | 10 | 71,321,279 | G | A | 31.6 | 31.6 | 50,791 | 526,121 | 2.2x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 6.6x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 1 |
| <i>NEUROG3</i> | rs61850200 | 10 | 71,321,658 | C | G | 27.7 | 27.7 | 50,791 | 526,121 | 0.07 | 1.02 (1.00-1.03) | 50,401 | 523,888 | 0.09 | 1.01 (1.00-1.03) | 0.8 |
| <i>NEUROG3</i> | rs41277236 | 10 | 71,332,301 | T | C | 4.31 | 4.31 | 50,791 | 526,121 | 2.1x10 ⁻⁹ | 1.12 (1.08-1.17) | 50,402 | 523,888 | 1.7x10 ⁻⁹ | 1.13 (1.08-1.17) | 0.7 |
| <i>NEUROG3</i> | rs549498088 | 10 | 71,347,311 | T | C | 0.600 | 0.600 | 45,059 | 507,254 | 6.7x10 ⁻⁸ | 1.65 (1.37-1.98) | 44,507 | 503,870 | 2.4x10 ⁻⁷ | 1.65 (1.36-1.99) | 1 |
| <i>NEUROG3</i> | rs2642588 | 10 | 71,466,578 | G | T | 70.2 | 29.8 | 50,791 | 526,121 | 2.3x10 ⁻⁹ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 1.2x10 ⁻⁹ | 1.05 (1.03-1.07) | 0.6 |
| <i>ZMIZ1</i> | rs703972 | 10 | 80,952,826 | G | C | 53.3 | 46.7 | 50,791 | 526,120 | 1.9x10 ⁻²⁴ | 1.08 (1.06-1.09) | 50,402 | 523,887 | 3.4x10 ⁻²⁶ | 1.08 (1.07-1.10) | 0.1 |
| <i>ZMIZ1</i> | rs1317617 | 10 | 81,096,589 | G | A | 79.8 | 20.2 | 50,791 | 526,121 | 0.0002 | 1.03 (1.02-1.05) | 50,402 | 523,888 | 0.0002 | 1.04 (1.02-1.05) | 0.7 |
| <i>PTEN</i> | rs11202627 | 10 | 89,769,340 | T | C | 15.2 | 15.2 | 50,791 | 526,121 | 2.0x10 ⁻⁸ | 1.06 (1.04-1.08) | 50,402 | 523,888 | 4.7x10 ⁻⁸ | 1.06 (1.04-1.08) | 0.9 |
| <i>HHX/IDE</i> | rs7078559 | 10 | 93,924,663 | T | C | 57.8 | 42.2 | 50,790 | 526,121 | 1.8x10 ⁻¹⁰ | 1.05 (1.03-1.06) | 50,400 | 523,888 | 4.7x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.09 |
| <i>HHX/IDE</i> | rs10882101 | 10 | 94,462,427 | T | C | 58.7 | 41.3 | 50,790 | 526,121 | 5.6x10 ⁻⁵⁰ | 1.12 (1.10-1.13) | 50,401 | 523,888 | 1.6x10 ⁻⁵⁷ | 1.13 (1.11-1.15) | 0.0007 |
| <i>HHX/IDE</i> | rs1112718 | 10 | 94,479,107 | A | G | 59.8 | 40.2 | 50,791 | 526,120 | 2.7x10 ⁻⁴⁷ | 1.11 (1.10-1.13) | 50,402 | 523,886 | 5.5x10 ⁻⁵⁵ | 1.13 (1.11-1.14) | 0.0005 |
| <i>TCF7L2</i> | rs536643418 | 10 | 114,699,835 | G | C | 0.520 | 0.520 | 47,555 | 512,283 | 3.8x10 ⁻⁸ | 1.65 (1.38-1.98) | 47,249 | 510,491 | 1.7x10 ⁻⁷ | 1.66 (1.37-2.00) | 1 |
| <i>TCF7L2</i> | rs140242150 | 10 | 114,702,962 | A | G | 0.500 | 0.500 | 46,759 | 509,075 | 1.2x10 ⁻⁷ | 1.39 (1.23-1.58) | 47,044 | 513,490 | 1.8x10 ⁻⁸ | 1.43 (1.26-1.62) | 0.4 |
| <i>TCF7L2</i> | rs7918400 | 10 | 114,703,136 | T | C | 47.6 | 47.6 | 50,791 | 526,119 | 0.4 | 1.01 (0.99-1.02) | 50,402 | 523,886 | 0.04 | 1.02 (1.00-1.03) | 0.009 |
| <i>TCF7L2</i> | rs184509201 | 10 | 114,740,337 | C | G | 98.2 | 1.82 | 50,791 | 526,121 | 3.1x10 ⁻¹¹ | 1.20 (1.14-1.27) | 50,402 | 523,888 | 1.1x10 ⁻¹¹ | 1.22 (1.15-1.29) | 0.5 |
| <i>TCF7L2</i> | rs180988137 | 10 | 114,751,173 | G | A | 1.04 | 1.04 | 50,791 | 526,121 | 2.4x10 ⁻¹⁶ | 1.38 (1.28-1.49) | 50,402 | 523,888 | 1.4x10 ⁻¹⁹ | 1.44 (1.33-1.56) | 0.02 |
| <i>TCF7L2</i> | rs7903146 | 10 | 114,758,349 | C | T | 70.6 | 29.5 | 50,791 | 526,121 | 6.3x10 ⁻³⁶⁰ | 1.38 (1.36-1.41) | 50,402 | 523,888 | 6.2x10 ⁻⁴²² | 1.43 (1.41-1.46) | 1.5x10 ⁻²² |
| <i>TCF7L2</i> | rs78025551 | 10 | 114,757,956 | C | G | 85.1 | 14.9 | 50,791 | 526,121 | 9.9x10 ⁻⁵⁴ | 1.17 (1.15-1.20) | 50,402 | 523,888 | 1.2x10 ⁻⁵⁶ | 1.18 (1.16-1.21) | 0.1 |
| <i>TCF7L2</i> | rs34855922 | 10 | 114,871,594 | A | G | 71.6 | 28.4 | 50,791 | 526,121 | 0.0001 | 1.03 (1.01-1.04) | 50,401 | 523,888 | 0.0001 | 1.03 (1.01-1.05) | 0.9 |
| <i>WDR11</i> | rs72631105 | 10 | 122,915,345 | A | G | 19.0 | 19.0 | 50,791 | 526,121 | 3.1x10 ⁻⁸ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 3.7x10 ⁻⁹ | 1.06 (1.04-1.08) | 0.2 |
| <i>PLEKH1</i> | rs2280141 | 10 | 124,193,181 | T | G | 51.6 | 48.4 | 50,790 | 526,120 | 2.7x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,401 | 523,887 | 3.7x10 ⁻¹⁰ | 1.05 (1.03-1.06) | 0.06 |
| <i>INS/IGF2</i> | rs12802972 | 11 | 1,704,596 | A | G | 42.8 | 42.8 | 50,789 | 526,121 | 0.0002 | 1.03 (1.01-1.04) | 50,401 | 523,888 | 0.0001 | 1.03 (1.01-1.05) | 0.7 |
| <i>INS/IGF2</i> | rs11042596 | 11 | 2,118,860 | G | T | 66.5 | 33.5 | 50,789 | 526,119 | 0.00002 | 1.03 (1.02-1.05) | 50,400 | 523,886 | 0.00005 | 1.03 (1.02-1.05) | 0.7 |
| <i>INS/IGF2</i> | rs555759341 | 11 | 2,151,761 | C | G | 0.490 | 0.490 | 46,197 | 507,005 | 7.8x10 ⁻⁷ | 1.48 (1.27-1.73) | 45,890 | 505,220 | 8.4x10 ⁻⁷ | 1.49 (1.27-1.74) | 0.9 |
| <i>INS/IGF2</i> | rs571342427 | 11 | 2,182,519 | C | T | 0.150 | 0.150 | 41,154 | 495,389 | 1.7x10 ⁻⁸ | 1.86 (1.50-2.31) | 40,855 | 493,606 | 5.4x10 ⁻⁸ | 1.83 (1.47-2.27) | 0.7 |
| <i>INS/IGF2</i> | rs4929965 | 11 | 2,197,286 | A | G | 38.3 | 38.3 | 50,789 | 526,120 | 2.0x10 ⁻²³ | 1.08 (1.06-1.09) | 50,401 | 523,886 | 1.6x10 ⁻¹⁹ | 1.07 (1.06-1.09) | 0.1 |
| <i>KCNQ1</i> | rs4930091 | 11 | 2,372,356 | T | C | 75.9 | 24.1 | 50,789 | 526,119 | 0.0002 | 1.03 (1.02-1.05) | 50,400 | 523,886 | 0.0004 | 1.03 (1.01-1.05) | 0.8 |
| <i>KCNQ1</i> | rs2281364 | 11 | 2,579,163 | A | G | 94.7 | 5.32 | 50,791 | 526,121 | 0.00008 | 1.07 (1.03-1.10) | 50,402 | 523,888 | 0.0003 | 1.06 (1.03-1.10) | 0.6 |
| <i>KCNQ1</i> | rs80102379 | 11 | 2,634,177 | G | T | 98.2 | 1.78 | 50,791 | 526,121 | 0.0009 | 1.11 (1.04-1.17) | 50,402 | 523,888 | 0.0006 | 1.11 (1.05-1.18) | 0.7 |
| <i>KCNQ1</i> | rs231349 | 11 | 2,672,821 | T | C | 10.2 | 10.2 | 50,791 | 526,121 | 4.8x10 ⁻⁷ | 1.06 (1.04-1.09) | 50,402 | 523,888 | 4.0x10 ⁻⁸ | 1.07 (1.04-1.10) | 0.2 |
| <i>KCNQ1</i> | rs231361 | 11 | 2,691,500 | A | G | 25.6 | 25.6 | 50,791 | 526,121 | 2.4x10 ⁻¹⁵ | 1.07 (1.05-1.09) | 50,402 | 523,888 | 6.9x10 ⁻¹⁸ | 1.08 (1.06-1.10) | 0.05 |
| <i>KCNQ1</i> | rs2283220 | 11 | 2,755,548 | A | G | 69.0 | 31.0 | 50,791 | 526,121 | 1.4x10 ⁻⁹ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 4.9x10 ⁻¹¹ | 1.06 (1.04-1.07) | 0.1 |
| <i>KCNQ1</i> | rs234853 | 11 | 2,850,828 | G | A | 24.8 | 24.8 | 50,790 | 526,120 | 0.00001 | 1.04 (1.02-1.05) | 50,401 | 523,887 | 0.00002 | 1.04 (1.02-1.06) | 0.9 |
| <i>KCNQ1</i> | rs2237895 | 11 | 2,857,194 | C | A | 42.6 | 42.6 | 50,790 | 526,120 | 3.6x10 ⁻⁴² | 1.11 (1.09-1.12) | 50,401 | 523,887 | 9.9x10 ⁻⁴⁷ | 1.12 (1.10-1.13) | 0.01 |
| <i>KCNQ1</i> | rs2237897 | 11 | 2,858,546 | C | T | 95.4 | 4.57 | 50,791 | 526,121 | 1.3x10 ⁻²⁸ | 1.23 (1.19-1.27) | 50,402 | 523,888 | 1.7x10 ⁻³¹ | 1.25 (1.20-1.30) | 0.07 |
| <i>KCNQ1</i> | rs445084 | 11 | 2,908,754 | G | A | 36.1 | 36.1 | 50,789 | 526,120 | 0.00005 | 1.03 (1.02-1.05) | 50,401 | 523,887 | 2.4x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.1 |
| <i>PDE3B</i> | rs141521721 | 11 | 14,763,828 | A | C | 2.36 | 2.36 | 50,791 | 526,121 | 6.7x10 ⁻⁸ | 1.14 (1.09-1.20) | 50,402 | 523,888 | 3.4x10 ⁻⁷ | 1.14 (1.08-1.19) | 0.7 |
| <i>KCNJ11</i> | rs5213 | 11 | 17,408,404 | C | T | 36.2 | 36.2 | 50,789 | 526,119 | 6.8x10 ⁻¹⁸ | 1.07 (1.05-1.08) | 50,400 | 523,886 | 6.5x10 ⁻²⁵ | 1.08 (1.07-1.10) | 0.00002 |
| <i>KCNJ11</i> | rs67254669 | 11 | 17,470,143 | G | A | 0.110 | 0.110 | 43,879 | 501,587 | 3.2x10 ⁻⁸ | 1.93 (1.53-2.43) | 43,580 | 499,804 | 1.8x10 ⁻⁸ | 1.98 (1.56-2.50) | 0.7 |
| <i>METTL15</i> | rs4923543 | 11 | 28,534,898 | A | G | 33.2 | 33.2 | 50,790 | 526,120 | 4.8x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 4.5x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.02 |
| <i>QSER1</i> | rs7943101 | 11 | 32,460,873 | T | C | 16.1 | 16.1 | 50,791 | 526,121 | 0.00005 | 1.04 (1.02-1.06) | 50,402 | 523,888 | 0.00007 | 1.04 (1.02-1.06) | 1 |
| <i>QSER1</i> | rs145678014 | 11 | 32,927,778 | G | T | 95.7 | 4.33 | 50,791 | 526,121 | 1.6x10 ⁻⁸ | 1.11 (1.07-1.15) | 50,402 | 523,888 | 1.6x10 ⁻⁷ | 1.10 (1.06-1.15) | 0.5 |
| <i>QSER1</i> | rs528122639 | 11 | 33,091,735 | A | G | 0.0900 | 0.0900 | 37,282 | 489,749 | 1.3x10 ⁻⁶ | 2.11 (1.16-2.85) | 41,170 | 492,220 | 0.00001 | 1.97 (1.45-2.68) | 0.3 |
| <i>PDHX</i> | rs286925 | 11 | 34,642,668 | A | G | 18.2 | 18.2 | 50,790 | 526,120 | 0.00004 | 1.04 (1.02-1.06) | 50,401 | 523,887 | 0.0001 | 1.04 (1.02-1.06) | 0.7 |
| <i>PDHX</i> | rs2767036 | 11 | 34,982,148 | C | A | 29.1 | 29.1 | 50,789 | 526,120 | 8.8x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,400 | 523,887 | 3.9x10 ⁻⁷ | 1.04 (1.03-1.06) | 0.7 |
| <i>HSD17B12</i> | rs1061810 | 11 | 43,877,934 | A | C | 28.8 | 28.8 | 50,791 | 526,121 | 3.9x10 ⁻⁹ | 1.05 (1.03-1.06) | 50,402 | 523,888 | 0.00008 | 1.03 (1.02-1.05) | 0.00006 |
| <i>CRY2</i> | rs7115753 | 11 | 45,912,013 | A | G | 44.9 | 44.9 | 50,789 | 526,119 | 2.6x10 ⁻⁸ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 9.9x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.8 |
| <i>CELF1</i> | rs7124681 | 11 | 47,529,947 | A | C | 41.0 | 41.0 | 50,712 | 521,614 | 2.8x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,323 | 519,381 | 0.008 | 1.02 (1.01-1.04) | 1.1x10 ⁻⁷ |
| <i>MAP3K11</i> | rs1783541 | 11 | 65,294,799 | T | C | 20.4 | 20.4 | 50,790 | 526,120 | 5.2x10 ⁻¹² | 1.06 (1.05-1.08) | 50,401 | 523,887 | 4.4x10 ⁻⁹ | 1.06 (1.04-1.07) | 0.05 |
| <i>CCND1</i> | rs61881115 | 11 | 68,997,225 | G | A | 83.8 | 16.2 | 50,791 | 526,120 | 1.2x10 ⁻⁶ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 5.3x10 ⁻⁶ | 1.05 (1.03-1.07) | 0.6 |
| <i>CCND1</i> | rs11820019 | 11 | 69,448,758 | T | C | 97.3 | 2.67 | 50,791 | 526,121 | 9. | | | | | | |

| HMGA2 | rs1042725 | 12 | 66,358,347 | T | C | 49.0 | 49.0 | 50,790 | 526,120 | 8.0x10 ⁻¹⁵ | 1.06 (1.04-1.07) | 50,401 | 523,887 | 2.9x10 ⁻¹⁵ | 1.06 (1.05-1.08) | 0.5 |
|--------------------|-------------|----|-------------|---|---|--------|--------|--------|---------|-----------------------|------------------|--------|---------|-----------------------|------------------|-----------------------|
| <i>TSPAN8/LGR5</i> | rs1796330 | 12 | 71,522,953 | G | C | 57.1 | 42.9 | 50,791 | 526,120 | 2.5x10 ⁻¹¹ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 1.0x10 ⁻¹¹ | 1.05 (1.04-1.07) | 0.5 |
| <i>USP44</i> | rs2197973 | 12 | 95,928,560 | T | C | 53.8 | 46.3 | 50,789 | 526,120 | 9.5x10 ⁻⁹ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 6.6x10 ⁻⁶ | 1.03 (1.02-1.05) | 0.02 |
| <i>RMST</i> | rs759111467 | 12 | 97,562,756 | A | G | 0.0300 | 0.0300 | 29,460 | 449,523 | 0.02 | 2.08 (1.10-3.95) | 29,264 | 448,156 | 0.03 | 2.03 (1.05-3.90) | 0.9 |
| <i>RMST</i> | rs557027608 | 12 | 97,779,248 | A | G | 0.0600 | 0.0600 | 21,106 | 438,604 | 0.002 | 2.11 (1.31-3.42) | 20,927 | 436,945 | 0.0009 | 2.23 (1.39-3.59) | 0.6 |
| <i>RMST</i> | rs77864822 | 12 | 97,848,775 | A | G | 93.2 | 6.76 | 50,791 | 526,121 | 4.0x10 ⁻⁶ | 1.07 (1.04-1.10) | 50,402 | 523,888 | 2.5x10 ⁻⁷ | 1.08 (1.05-1.11) | 0.2 |
| <i>WSCD2</i> | rs1426371 | 12 | 108,629,780 | G | A | 73.9 | 26.1 | 50,791 | 526,121 | 3.3x10 ⁻¹¹ | 1.06 (1.04-1.07) | 50,402 | 523,888 | 3.5x10 ⁻⁹ | 1.05 (1.03-1.07) | 0.2 |
| <i>KSR2</i> | rs34965774 | 12 | 118,412,373 | A | G | 14.4 | 14.4 | 50,791 | 526,121 | 0.00002 | 1.05 (1.02-1.07) | 50,402 | 523,888 | 0.00002 | 1.05 (1.03-1.07) | 0.6 |
| <i>KSR2</i> | rs12578639 | 12 | 118,489,636 | A | T | 82.8 | 17.2 | 50,791 | 526,121 | 0.00001 | 1.04 (1.02-1.06) | 50,402 | 523,888 | 1.3x10 ⁻⁷ | 1.05 (1.03-1.07) | 0.02 |
| <i>HNF1A</i> | rs11065299 | 12 | 121,297,815 | G | G | 7.54 | 7.54 | 50,791 | 526,121 | 3.0x10 ⁻⁷ | 1.07 (1.05-1.10) | 50,402 | 523,888 | 0.00001 | 1.06 (1.03-1.09) | 0.1 |
| <i>HNF1A</i> | rs73226260 | 12 | 121,380,541 | G | A | 96.7 | 3.31 | 50,791 | 526,121 | 1.6x10 ⁻⁸ | 1.12 (1.08-1.17) | 50,402 | 523,888 | 4.3x10 ⁻⁹ | 1.13 (1.09-1.18) | 0.4 |
| <i>HNF1A</i> | rs1800574 | 12 | 121,416,864 | T | C | 2.96 | 2.96 | 50,791 | 526,121 | 3.5x10 ⁻¹¹ | 1.15 (1.10-1.20) | 50,402 | 523,888 | 5.7x10 ⁻¹² | 1.16 (1.11-1.21) | 0.3 |
| <i>HNF1A</i> | rs56348580 | 12 | 121,432,117 | G | C | 68.9 | 31.1 | 50,791 | 526,121 | 6.5x10 ⁻¹³ | 1.06 (1.04-1.07) | 50,402 | 523,888 | 2.0x10 ⁻¹⁰ | 1.05 (1.04-1.07) | 0.1 |
| <i>HNF1A</i> | rs28638142 | 12 | 121,501,461 | A | C | 4.42 | 4.42 | 50,791 | 526,121 | 2.1x10 ⁻⁶ | 1.09 (1.05-1.13) | 50,402 | 523,888 | 5.0x10 ⁻⁶ | 1.09 (1.05-1.13) | 0.9 |
| <i>HNF1A</i> | rs73224262 | 12 | 121,882,395 | T | C | 0.680 | 0.680 | 50,791 | 526,121 | 0.00002 | 1.23 (1.12-1.35) | 50,402 | 523,888 | 5.0x10 ⁻⁶ | 1.25 (1.14-1.38) | 0.4 |
| <i>MPHOSPH9</i> | rs4148856 | 12 | 123,450,765 | G | G | 78.1 | 21.9 | 50,789 | 526,119 | 1.4x10 ⁸ | 1.05 (1.03-1.07) | 50,400 | 523,886 | 2.4x10 ⁻⁹ | 1.06 (1.04-1.07) | 0.4 |
| <i>ZNF664</i> | rs7978610 | 12 | 124,468,572 | G | C | 66.6 | 33.5 | 50,789 | 526,121 | 9.5x10 ⁻⁶ | 1.03 (1.02-1.05) | 50,400 | 523,888 | 7.7x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 0.00004 |
| <i>ZNF664</i> | rs825452 | 12 | 124,509,177 | A | G | 60.3 | 39.7 | 50,790 | 526,120 | 0.00005 | 1.03 (1.02-1.05) | 50,400 | 523,887 | 8.9x10 ⁻⁹ | 1.04 (1.03-1.06) | 0.00006 |
| <i>FBRSL1</i> | rs12811407 | 12 | 133,069,698 | A | G | 33.1 | 33.1 | 50,791 | 526,120 | 2.4x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 5.4x10 ⁻⁸ | 1.04 (1.03-1.06) | 0.1 |
| <i>RNF6</i> | rs34584161 | 13 | 26,776,999 | A | G | 76.0 | 24.0 | 50,791 | 526,121 | 1.9x10 ⁻⁷ | 1.05 (1.03-1.06) | 50,402 | 523,888 | 5.2x10 ⁻⁹ | 1.05 (1.03-1.07) | 0.09 |
| <i>HMGB1</i> | rs11842871 | 13 | 31,042,452 | G | T | 73.5 | 26.6 | 50,791 | 526,121 | 7.9x10 ⁻⁸ | 1.05 (1.03-1.06) | 50,402 | 523,888 | 0.00001 | 1.04 (1.02-1.06) | 0.06 |
| <i>KL</i> | rs576674 | 13 | 33,554,302 | G | A | 16.9 | 16.9 | 50,791 | 526,121 | 2.6x10 ⁻⁶ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 1.3x10 ⁻⁷ | 1.05 (1.03-1.08) | 0.1 |
| <i>DLEU1</i> | rs963740 | 13 | 51,096,095 | A | T | 71.3 | 28.7 | 50,791 | 526,120 | 4.7x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,402 | 523,887 | 1.4x10 ⁻⁸ | 1.05 (1.03-1.06) | 0.09 |
| <i>PCDH17</i> | rs9537803 | 13 | 58,366,634 | C | T | 27.7 | 27.7 | 50,789 | 526,120 | 6.1x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,400 | 523,887 | 0.0003 | 1.03 (1.01-1.05) | 0.006 |
| <i>PCDH17</i> | rs9569864 | 13 | 58,965,435 | C | T | 82.5 | 17.5 | 50,789 | 526,121 | 9.7x10 ⁻¹⁰ | 1.06 (1.04-1.08) | 50,400 | 523,888 | 2.9x10 ⁻⁷ | 1.05 (1.03-1.07) | 0.07 |
| <i>SRGAP2D</i> | rs9563615 | 13 | 59,077,406 | A | T | 71.0 | 29.0 | 50,791 | 526,121 | 9.8x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 50,402 | 523,887 | 4.0x10 ⁻⁶ | 1.04 (1.02-1.06) | 0.002 |
| <i>SRGAP2D</i> | rs76251711 | 13 | 59,184,234 | G | A | 1.26 | 1.26 | 50,379 | 525,731 | 0.00006 | 1.16 (1.08-1.24) | 49,990 | 523,498 | 0.00009 | 1.15 (1.07-1.24) | 1 |
| <i>SPRY2</i> | rs1359790 | 13 | 80,717,156 | G | A | 72.0 | 28.0 | 50,790 | 526,121 | 3.8x10 ⁻²⁵ | 1.09 (1.07-1.10) | 50,402 | 523,888 | 6.5x10 ⁻²⁸ | 1.10 (1.08-1.11) | 0.06 |
| <i>IRS2</i> | rs7987740 | 13 | 109,947,213 | T | C | 60.9 | 39.1 | 50,790 | 526,120 | 3.0x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,887 | 3.4x10 ⁻⁶ | 1.04 (1.02-1.05) | 0.4 |
| <i>IRS2</i> | rs4771648 | 13 | 110,431,626 | G | A | 66.9 | 33.2 | 50,791 | 526,121 | 1.8x10 ⁻⁶ | 1.04 (1.02-1.05) | 50,402 | 523,888 | 2.2x10 ⁻⁶ | 1.04 (1.02-1.06) | 0.8 |
| <i>SLC7A7</i> | rs17122772 | 14 | 23,288,935 | G | C | 22.8 | 22.8 | 50,791 | 526,121 | 5.7x10 ⁻⁸ | 1.05 (1.03-1.07) | 50,402 | 523,888 | 8.0x10 ⁻⁹ | 1.05 (1.04-1.07) | 0.3 |
| <i>AKAP6</i> | rs17522122 | 14 | 33,302,882 | T | G | 47.4 | 47.4 | 50,790 | 526,119 | 9.4x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,400 | 523,886 | 0.007 | 1.02 (1.01-1.04) | 3.2x10 ⁻⁶ |
| <i>CLEC14A</i> | rs8017808 | 14 | 38,848,419 | G | T | 74.3 | 25.7 | 50,791 | 526,121 | 3.5x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,402 | 523,888 | 3.7x10 ⁻⁸ | 1.05 (1.03-1.07) | 0.2 |
| <i>NRXN3</i> | rs17836088 | 14 | 79,932,041 | C | G | 21.7 | 21.7 | 50,790 | 526,121 | 6.9x10 ⁻¹³ | 1.07 (1.05-1.08) | 50,402 | 523,888 | 0.00002 | 1.04 (1.02-1.06) | 1.3x10 ⁻⁹ |
| <i>SMEK1</i> | rs8010382 | 14 | 91,963,722 | G | A | 42.1 | 42.1 | 50,791 | 526,120 | 1.5x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,401 | 523,887 | 2.6x10 ⁻⁷ | 1.04 (1.03-1.06) | 0.9 |
| <i>MARK3</i> | rs62007683 | 14 | 103,894,071 | G | T | 65.3 | 34.7 | 50,790 | 526,120 | 3.1x10 ⁻⁷ | 1.04 (1.02-1.06) | 50,401 | 523,887 | 0.00009 | 1.03 (1.02-1.05) | 0.02 |
| <i>RASGRP1</i> | rs8032939 | 15 | 38,834,033 | C | T | 24.6 | 24.6 | 50,790 | 526,121 | 3.6x10 ⁻⁸ | 1.05 (1.03-1.06) | 50,402 | 523,888 | 9.2x10 ⁻⁷ | 1.04 (1.03-1.06) | 0.3 |
| <i>RASGRP1</i> | rs34715063 | 15 | 38,873,115 | C | T | 12.4 | 12.4 | 50,791 | 526,121 | 6.3x10 ⁻¹³ | 1.09 (1.06-1.11) | 50,402 | 523,888 | 3.7x10 ⁻¹³ | 1.09 (1.06-1.11) | 0.7 |
| <i>LTK</i> | rs11070332 | 15 | 41,809,205 | A | G | 35.8 | 35.8 | 50,791 | 526,119 | 9.4x10 ⁻¹² | 1.05 (1.04-1.07) | 50,402 | 523,887 | 9.6x10 ⁻¹⁰ | 1.05 (1.03-1.07) | 0.2 |
| <i>LTK</i> | rs543786825 | 15 | 42,201,410 | T | C | 0.0400 | 0.0400 | 27,464 | 464,313 | 1.9x10 ⁻⁶ | 3.52 (2.10-5.91) | 27,253 | 462,639 | 2.2x10 ⁻⁷ | 3.96 (2.35-6.67) | 0.3 |
| <i>ONECUT1</i> | rs2456530 | 15 | 53,091,553 | T | C | 12.7 | 12.7 | 50,791 | 526,121 | 1.3x10 ⁻⁹ | 1.07 (1.05-1.09) | 50,402 | 523,888 | 1.1x10 ⁻⁶ | 1.06 (1.03-1.08) | 0.02 |
| <i>WDR72</i> | rs528350911 | 15 | 53,747,228 | G | C | 0.680 | 0.680 | 50,791 | 526,121 | 0.00001 | 1.25 (1.13-1.38) | 50,402 | 523,888 | 8.1x10 ⁻⁸ | 1.32 (1.19-1.46) | 0.02 |
| <i>TCF12</i> | rs117483894 | 15 | 57,456,802 | G | A | 3.69 | 3.69 | 50,789 | 526,120 | 2.0x10 ⁻⁶ | 1.10 (1.06-1.14) | 50,400 | 523,887 | 0.00006 | 1.09 (1.04-1.13) | 0.2 |
| <i>C2CD4A/B</i> | rs8037894 | 15 | 62,394,264 | G | C | 56.6 | 43.4 | 50,789 | 526,120 | 1.7x10 ⁹ | 1.05 (1.03-1.05) | 50,400 | 523,887 | 4.4x10 ⁻¹⁰ | 1.05 (1.03-1.06) | 0.5 |
| <i>USP3</i> | rs7178762 | 15 | 63,871,292 | C | T | 46.0 | 46.0 | 50,791 | 526,121 | 6.9x10 ⁻⁹ | 1.04 (1.03-1.06) | 50,402 | 523,888 | 4.4x10 ⁻⁹ | 1.04 (1.03-1.06) | 0.6 |
| <i>MAP2K5</i> | rs4776970 | 15 | 68,080,886 | A | T | 64.1 | 35.9 | 50,791 | 526,121 | 5.4x10 ⁻⁷ | 1.04 (1.02-1.05) | 50,402 | 523,888 | 0.08 | 1.01 (1.00-1.03) | 2.0x10 ⁻¹² |
| <i>PTPN9</i> | rs13737 | 15 | 75,932,129 | G | T | 75.9 | 24.1 | 50,791 | 526,121 | 5.2x10 ⁻¹⁰ | 1.05 (1.04-1.07) | 50,402 | 523,888 | 1.1x10 ⁻⁸ | 1.05 (1.03-1.07) | 0.5 |
| <i>HMG20A</i> | rs1005752 | 15 | 77,818,128 | A | C | 71.5 | 28.5 | 50,789 | 526,119 | 1.4x10 ⁻²¹ | 1.08 (1.06-1.10) | 50,400 | 523,886 | 7.8x10 ⁻¹⁸ | 1.07 (1.06-1.09) | 0.1 |
| <i>AP3S2</i> | rs4932265 | 15 | 90,423,293 | T | C | 26.7 | 26.7 | 50,789 | 526,119 | 6.6x10 ⁻¹⁷ | 1.07 (1.05-1.09) | 50,400 | 523,886 | 7.7x10 ⁻²⁰ | 1.08 (1.06-1.10) | 0.02 |
| <i>PRC1</i> | rs12910825 | 15 | 91,511,260 | G | A | 36.1 | 36.1 | 50,790 | 526,119 | 8.1x10 ⁻¹⁶ | 1.06 (1.05-1.08) | 50,401 | 523,886 | 2.2x10 ⁻¹⁴ | 1.06 (1.04-1.08) | 0.5 |
| <i>ITFG3</i> | rs6600191 | 16 | 295,795 | T | C | 82.5 | 17.5 | 50,791 | 526,121 | 1.6x10 ⁻¹² | 1.07 (1.05-1.09) | 50,402 | 523,888 | 3.8x10 ⁻¹¹ | 1.07 (1.05-1.09) | 0.5 |
| <i>CLUAP1</i> | rs3751837 | 16 | 3,583,173 | T | C | 22.0 | 22.0 | 50,789 | 526,121 | 6.1x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,400 | 523,888 | 0.0009 | 1.03 (1.01-1.05) | 0.0005 |
| <i>ATP2A1</i> | rs8046545 | 16 | 28,915,217 | G | A | 35.9 | 35.9 | 50,789 | 526,119 | 1.5x10 ⁻⁷ | 1.04 (1.03-1.06) | 50,400 | 523,886 | 0.1 | 1.01 (1.00-1.03) | 9.1x10 ⁻¹⁶ |
| <i>FAM57B</i> | rs11642430 | 16 | 30,045,789 | G | C | 39.9 | 39 | | | | | | | | | |

RAF: risk allele frequency; MAF: minor allele frequency; OR: Odds ratio; CI: confidence interval.

Supplementary Table 4 | Summary of sex-differentiated analysis.

| Nearest gene | Index variant | Chromosome | Position (Build 37 bp) | Risk allele | Other allele | Male-specific meta-analysis | | | | Female-specific meta-analysis | | | | Sex-differentiated meta-analysis | | | |
|---------------------|---------------|------------|---------------------------|-------------|--------------|-----------------------------|----------|-----------------------|------------------|-------------------------------|----------|-----------------------|------------------|----------------------------------|----------|-----------------------|-----------------------|
| | | | | | | Cases | Controls | p-value | OR (95% CI) | Cases | Controls | p-value | OR (95% CI) | Cases | Controls | p-value | Heterogeneity p-value |
| <i>SEC16B</i> | rs539515 | 1 | 177,889,025 | C | A | 41,842 | 383,764 | 0.003 | 1.03 (1.01-1.05) | 30,049 | 434,330 | 1.7x10 ⁻¹¹ | 1.08 (1.06-1.11) | 74,116 | 823,996 | 1.9x10 ⁻¹² | 0.002 |
| <i>SRGAP2</i> | rs9430095 | 1 | 206,593,900 | C | G | 41,842 | 383,763 | 5.1x10 ⁻⁸ | 1.05 (1.03-1.07) | 30,048 | 434,330 | 0.06 | 1.02 (1.00-1.04) | 74,116 | 823,996 | 6.0x10 ⁻⁸ | 0.03 |
| <i>ABCB10</i> | rs348330 | 1 | 229,672,955 | G | A | 41,841 | 383,762 | 4.5x10 ⁻¹⁴ | 1.07 (1.05-1.09) | 30,048 | 434,330 | 0.002 | 1.03 (1.01-1.05) | 74,115 | 823,996 | 3.3x10 ⁻¹⁵ | 0.01 |
| <i>THADA</i> | rs80147536 | 2 | 43,698,028 | A | T | 41,842 | 383,762 | 9.6x10 ⁻²⁵ | 1.16 (1.13-1.20) | 30,048 | 434,329 | 2.1x10 ⁻⁷ | 1.09 (1.06-1.13) | 74,116 | 823,997 | 1.6x10 ⁻²⁹ | 0.004 |
| <i>GLI2</i> | rs11688931 | 2 | 121,318,166 | C | G | 41,843 | 383,764 | 0.0003 | 1.04 (1.02-1.07) | 30,049 | 434,331 | 6.0x10 ⁻⁹ | 1.08 (1.06-1.11) | 74,117 | 823,997 | 5.9x10 ⁻¹¹ | 0.05 |
| <i>GLI2</i> | rs11688682 | 2 | 121,347,612 | G | C | 41,843 | 383,763 | 0.00001 | 1.04 (1.02-1.07) | 30,049 | 434,330 | 5.2x10 ⁻¹¹ | 1.08 (1.05-1.1) | 74,117 | 823,996 | 3.5x10 ⁻¹⁴ | 0.03 |
| <i>CYTIP</i> | rs13426680 | 2 | 158,339,550 | A | G | 41,843 | 383,763 | 0.001 | 1.06 (1.02-1.10) | 30,049 | 434,330 | 1.5x10 ⁻⁹ | 1.13 (1.09-1.18) | 74,117 | 823,996 | 5.6x10 ⁻¹¹ | 0.01 |
| <i>GRB14/COBLL1</i> | rs10195252 | 2 | 165,513,091 | T | C | 41,843 | 383,764 | 3.8x10 ⁻⁸ | 1.05 (1.03-1.07) | 30,050 | 434,331 | 4.4x10 ⁻¹⁵ | 1.08 (1.06-1.10) | 74,117 | 823,997 | 1.1x10 ⁻²⁰ | 0.02 |
| <i>TSC22D2</i> | rs62271373 | 3 | 150,066,540 | A | T | 41,843 | 383,763 | 0.002 | 1.06 (1.02-1.10) | 30,050 | 434,331 | 4.5x10 ⁻⁸ | 1.13 (1.08-1.18) | 74,117 | 823,997 | 2.6x10 ⁻⁹ | 0.04 |
| <i>RREB1</i> | rs9379084 | 6 | 7,231,843 | G | A | 41,843 | 383,764 | 3.2x10 ⁻⁸ | 1.08 (1.05-1.11) | 30,050 | 434,331 | 6.2x10 ⁻¹⁴ | 1.13 (1.09-1.16) | 74,117 | 823,997 | 1.3x10 ⁻¹⁹ | 0.04 |
| <i>VEGFA</i> | rs6458354 | 6 | 43,814,190 | C | T | 41,842 | 383,763 | 2.5x10 ⁻¹¹ | 1.06 (1.04-1.08) | 30,048 | 434,329 | 0.002 | 1.03 (1.01-1.06) | 74,116 | 823,995 | 1.6x10 ⁻¹² | 0.05 |
| <i>SLC25A51P1</i> | rs555402748 | 6 | 67,387,490 | T | C | 20,276 | 211,157 | 0.02 | 2.22 (1.14-4.32) | 9,791 | 251,388 | 1.1x10 ⁻⁷ | 8.97 (4.00-20.2) | 45,657 | 492,696 | 4.8x10 ⁻⁸ | 0.009 |
| <i>KLF14</i> | rs2268382 | 7 | 130,027,037 | C | A | 41,843 | 383,764 | 0.1 | 1.01 (1.00-1.03) | 30,050 | 434,331 | 0.00002 | 1.04 (1.02-1.07) | 74,117 | 823,997 | 0.00004 | 0.03 |
| <i>KLF14</i> | rs1562396 | 7 | 130,457,914 | G | A | 41,842 | 383,763 | 0.0001 | 1.04 (1.02-1.06) | 30,049 | 434,331 | 8.9x10 ⁻¹⁶ | 1.09 (1.07-1.11) | 74,116 | 823,997 | 4.8x10 ⁻¹⁸ | 0.0005 |
| <i>ANK1</i> | rs13262861 | 8 | 41,508,577 | C | A | 41,843 | 383,764 | 2.3x10 ⁻²³ | 1.12 (1.10-1.15) | 30,050 | 434,330 | 4.8x10 ⁻⁶ | 1.06 (1.04-1.09) | 74,117 | 823,997 | 7.7x10 ⁻²⁷ | 0.001 |
| <i>ANK1</i> | rs4736819 | 8 | 41,509,915 | T | C | 41,841 | 383,763 | 1.4x10 ⁻²¹ | 1.09 (1.07-1.10) | 30,050 | 434,330 | 0.0006 | 1.03 (1.01-1.05) | 74,115 | 823,995 | 4.4x10 ⁻²³ | 0.0002 |
| <i>HAUS6</i> | rs7022807 | 9 | 19,067,833 | G | A | 41,842 | 383,763 | 0.001 | 1.03 (1.01-1.05) | 30,050 | 434,330 | 1.7x10 ⁻⁸ | 1.06 (1.04-1.08) | 74,116 | 823,997 | 5.1x10 ⁻¹⁰ | 0.03 |
| <i>MTND2P8</i> | rs11137820 | 9 | 81,359,113 | C | G | 41,843 | 383,763 | 2.6x10 ⁻⁸ | 1.05 (1.03-1.07) | 30,048 | 434,330 | 0.08 | 1.02 (1.00-1.04) | 74,116 | 823,996 | 3.9x10 ⁻⁸ | 0.02 |
| <i>TLE1</i> | rs2796441 | 9 | 84,308,948 | G | A | 41,843 | 383,764 | 1.1x10 ⁻¹⁸ | 1.08 (1.06-1.10) | 30,049 | 434,331 | 7.2x10 ⁻⁷ | 1.05 (1.03-1.07) | 74,116 | 823,997 | 5.5x10 ⁻²³ | 0.04 |
| <i>INS/IGF2</i> | rs11042596 | 11 | 2,118,860 | G | T | 41,841 | 383,762 | 0.02 | 1.02 (1.00-1.04) | 30,048 | 434,329 | 6.8x10 ⁻⁷ | 1.05 (1.03-1.08) | 74,115 | 823,995 | 2.6x10 ⁻⁷ | 0.03 |
| <i>KCNQ1</i> | rs2237895 | 11 | 2,857,194 | C | A | 41,843 | 383,763 | 3.4x10 ⁻³⁵ | 1.12 (1.10-1.13) | 30,049 | 434,330 | 9.9x10 ⁻¹³ | 1.08 (1.05-1.10) | 74,116 | 823,996 | 4.2x10 ⁻⁴⁵ | 0.007 |
| <i>MAP3K11</i> | rs1783541 | 11 | 65,294,799 | T | C | 41,843 | 383,764 | 0.0009 | 1.04 (1.01-1.06) | 30,050 | 434,330 | 5.7x10 ⁻¹⁴ | 1.10 (1.07-1.12) | 74,116 | 823,996 | 2.2x10 ⁻¹⁵ | 0.0006 |
| <i>CDKN1B</i> | rs2066827 | 12 | 12,871,099 | G | T | 41,842 | 383,763 | 2.3x10 ⁻⁸ | 1.06 (1.04-1.08) | 30,050 | 434,330 | 0.04 | 1.03 (1.00-1.05) | 74,116 | 823,996 | 2.2x10 ⁻⁸ | 0.04 |
| <i>IRS2</i> | rs7987740 | 13 | 109,947,213 | T | C | 41,843 | 383,764 | 0.004 | 1.03 (1.01-1.04) | 30,048 | 434,331 | 1.1x10 ⁻⁷ | 1.05 (1.03-1.08) | 74,116 | 823,996 | 1.0x10 ⁻⁸ | 0.04 |
| <i>RASGRP1</i> | rs8032939 | 15 | 38,834,033 | C | T | 41,843 | 383,764 | 0.01 | 1.02 (1.00-1.04) | 30,049 | 434,331 | 2.3x10 ⁻⁹ | 1.07 (1.05-1.09) | 74,116 | 823,997 | 8.3x10 ⁻¹⁰ | 0.004 |
| <i>TCF12</i> | rs117483894 | 15 | 57,456,802 | G | A | 41,843 | 383,763 | 0.04 | 1.05 (1.00-1.10) | 30,049 | 434,329 | 7.0x10 ⁻⁹ | 1.16 (1.10-1.22) | 74,115 | 823,996 | 6.0x10 ⁻⁹ | 0.003 |
| <i>CMIP</i> | rs2925979 | 16 | 81,534,790 | T | C | 41,843 | 383,764 | 0.004 | 1.03 (1.01-1.05) | 30,050 | 434,331 | 2.6x10 ⁻¹⁷ | 1.09 (1.07-1.12) | 74,117 | 823,997 | 4.0x10 ⁻¹⁸ | 8.3x10 ⁻⁶ |
| <i>PTPRS</i> | rs116953931 | 19 | 5,224,998 | A | G | 41,843 | 383,764 | 0.05 | 1.05 (1.00-1.10) | 30,050 | 434,331 | 0.00001 | 1.12 (1.07-1.18) | 74,117 | 823,997 | 7.8x10 ⁻⁶ | 0.04 |
| <i>FARSA</i> | rs3111316 | 19 | 13,038,415 | A | G | 41,841 | 383,763 | 0.002 | 1.03 (1.01-1.04) | 30,048 | 434,330 | 7.3x10 ⁻¹³ | 1.07 (1.05-1.10) | 74,115 | 823,997 | 5.6x10 ⁻¹⁴ | 0.0007 |
| <i>TM6SF2</i> | rs8107974 | 19 | 19,388,500 | T | A | 41,843 | 383,763 | 3.9x10 ⁻¹³ | 1.12 (1.09-1.16) | 30,050 | 434,331 | 0.002 | 1.06 (1.02-1.10) | 74,116 | 823,996 | 2.5x10 ⁻¹⁴ | 0.02 |
| <i>GIPR</i> | rs533172266 | 19 | 46,351,837 | T | C | 19,837 | 207,502 | 0.03 | 1.75 (1.05-2.91) | 16,833 | 269,826 | 1.7x10 ⁻⁶ | 3.81 (2.20-6.59) | 47,435 | 495,952 | 1.1x10 ⁻⁶ | 0.04 |

RAF: risk allele frequency; MAF: minor allele frequency; OR: Odds ratio; CI: confidence interval.

Supplementary Table 5 | Summary of 99% credible sets for 380 distinct T2D association signals.

| Nearest gene | Index variant | Chromosome | Position (Build 37 bp) | Risk allele | Other allele | RAF (%) | MAF (%) | P-value | 99% genetic credible set | | | | 99% functional credible set | | | |
|---------------------|---------------|------------|------------------------|-------------|--------------|---------|---------|-----------------------|--------------------------|---------------|---------------------|--------------------|-----------------------------|---------------|---------------------|--------------------|
| | | | | | | | | | SNPs | Interval (bp) | Interval start (bp) | Interval stop (bp) | SNPs | Interval (bp) | Interval start (bp) | Interval stop (bp) |
| <i>MACF1</i> | rs3768321 | 1 | 40,035,928 | T | G | 20.0 | 20.0 | 2.6x10 ⁻²⁶ | 72 | 441,918 | 39,618,556 | 40,060,473 | 57 | 39,736,334 | 40,038,571 | 302,238 |
| <i>FAF1</i> | rs58432198 | 1 | 51,256,091 | C | T | 88.1 | 11.9 | 2.1x10 ⁻¹⁰ | 233 | 588,366 | 50,920,161 | 51,508,526 | 221 | 50,920,161 | 51,493,428 | 573,268 |
| <i>PATJ</i> | rs12140153 | 1 | 62,579,891 | G | T | 90.5 | 9.49 | 1.3x10 ⁻⁸ | 1 | 1 | 62,579,891 | 62,579,891 | 1 | 62,579,891 | 62,579,891 | 1 |
| <i>DENND2C</i> | rs184660829 | 1 | 115,144,899 | C | T | 0.0200 | 0.0200 | 2.5x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>PTGFRN</i> | rs1127215 | 1 | 117,532,790 | C | T | 58.4 | 41.6 | 1.6x10 ⁻¹³ | 3 | 3,333 | 117,529,458 | 117,532,790 | 3 | 117,529,458 | 117,532,790 | 3,333 |
| <i>NOTCH2</i> | rs1493694 | 1 | 120,526,982 | T | C | 10.9 | 10.9 | 2.7x10 ⁻¹⁶ | 48 | 116,417 | 120,437,718 | 120,554,134 | 50 | 120,437,718 | 120,554,134 | 116,417 |
| <i>FAM63A</i> | rs10305745 | 1 | 150,786,038 | A | G | 1.45 | 1.45 | 4.4x10 ⁻⁶ | 723 | 763,915 | 150,518,927 | 151,282,841 | 374 | 150,521,099 | 151,282,841 | 761,743 |
| <i>FAM63A</i> | rs145904381 | 1 | 151,017,991 | T | C | 98.7 | 1.33 | 2.6x10 ⁻⁸ | 21 | 823,456 | 150,651,482 | 151,474,937 | 36 | 150,521,099 | 151,474,937 | 953,839 |
| <i>SEC16B</i> | rs539515 | 1 | 177,889,025 | C | A | 19.8 | 19.8 | 1.6x10 ⁻¹⁰ | 44 | 116,173 | 177,797,347 | 177,913,519 | 41 | 177,793,822 | 177,913,519 | 119,698 |
| <i>DSTYK</i> | rs12048743 | 1 | 205,114,873 | G | C | 44.2 | 44.2 | 3.5x10 ⁻⁹ | 55 | 222,669 | 205,039,295 | 205,261,963 | 50 | 205,039,295 | 205,261,963 | 222,669 |
| <i>SRGAP2</i> | rs9430095 | 1 | 206,593,900 | C | G | 49.4 | 49.4 | 1.9x10 ⁻⁸ | 24 | 130,362 | 206,512,359 | 206,642,720 | 26 | 206,512,359 | 206,642,720 | 130,362 |
| <i>PROX1</i> | rs79687284 | 1 | 214,150,821 | C | G | 3.48 | 3.48 | 2.6x10 ⁻¹⁶ | 2 | 377 | 214,150,445 | 214,150,821 | 2 | 214,150,445 | 214,150,821 | 377 |
| <i>PROX1</i> | rs340874 | 1 | 214,159,256 | C | T | 55.6 | 44.5 | 1.6x10 ⁻²² | 1 | 1 | 214,159,256 | 214,159,256 | 1 | 214,159,256 | 214,159,256 | 1 |
| <i>PROX1</i> | rs114526150 | 1 | 214,175,531 | G | T | 2.25 | 2.25 | 4.2x10 ⁻⁷ | 69 | 293,959 | 213,946,973 | 214,240,931 | 67 | 214,147,076 | 214,240,931 | 93,856 |
| <i>LYPLAL1</i> | rs553014999 | 1 | 219,584,164 | C | T | 0.130 | 0.130 | 7.7x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>LYPLAL1</i> | rs2820446 | 1 | 219,748,818 | C | G | 70.6 | 29.5 | 3.3x10 ⁻¹⁶ | 14 | 36,482 | 219,726,100 | 219,762,581 | 13 | 219,730,006 | 219,762,581 | 32,576 |
| <i>ABCB10</i> | rs348330 | 1 | 229,672,955 | G | A | 36.1 | 36.1 | 2.7x10 ⁻¹⁴ | 1 | 1 | 229,672,955 | 229,672,955 | 1 | 229,672,955 | 229,672,955 | 1 |
| <i>GNG4</i> | rs291367 | 1 | 235,690,800 | G | A | 63.2 | 36.8 | 4.7x10 ⁻¹⁰ | 65 | 171,172 | 235,534,750 | 235,705,921 | 45 | 235,534,750 | 235,705,921 | 171,172 |
| <i>TMEM18</i> | rs62107261 | 2 | 422,144 | T | C | 95.4 | 4.64 | 3.8x10 ⁻¹² | 3 | 13,432 | 408,713 | 422,144 | 4 | 408,713 | 466,003 | 57,291 |
| <i>TMEM18</i> | rs35913461 | 2 | 653,575 | C | T | 82.9 | 17.1 | 1.6x10 ⁻¹¹ | 175 | 32,690 | 621,185 | 653,874 | 170 | 601,905 | 653,874 | 51,970 |
| <i>FAM49A</i> | rs11680058 | 2 | 16,574,669 | A | G | 86.3 | 13.7 | 1.4x10 ⁻⁸ | 1 | 1 | 16,574,669 | 16,574,669 | 1 | 16,574,669 | 16,574,669 | 1 |
| <i>DTNB</i> | rs17802463 | 2 | 25,643,221 | G | T | 73.1 | 26.9 | 2.9x10 ⁻⁸ | 44 | 293,360 | 25,513,652 | 25,807,011 | 42 | 25,513,652 | 25,807,011 | 293,360 |
| <i>GCKR</i> | rs1260326 | 2 | 27,730,940 | C | T | 60.7 | 39.3 | 6.5x10 ⁻²⁵ | 3 | 11,664 | 27,730,940 | 27,742,603 | 4 | 27,598,097 | 27,742,603 | 144,507 |
| <i>THADA</i> | rs28525376 | 2 | 43,207,872 | G | T | 42.2 | 42.2 | 2.7x10 ⁻⁶ | 768 | 508,789 | 43,198,473 | 43,707,261 | 708 | 43,198,473 | 43,705,862 | 507,390 |
| <i>THADA</i> | rs6708643 | 2 | 43,430,440 | A | G | 50.1 | 49.9 | 3.9x10 ⁻⁸ | 20 | 25,273 | 43,421,344 | 43,446,616 | 19 | 43,421,344 | 43,446,616 | 25,273 |
| <i>THADA</i> | rs80147536 | 2 | 43,698,028 | A | T | 90.4 | 9.57 | 2.7x10 ⁻²⁹ | 155 | 236,769 | 43,528,057 | 43,764,825 | 144 | 43,528,057 | 43,764,825 | 236,769 |
| <i>BNIPL</i> | rs10193538 | 2 | 58,981,064 | T | G | 61.0 | 39.0 | 8.9x10 ⁻⁹ | 76 | 225,883 | 58,868,909 | 59,094,791 | 74 | 58,868,909 | 59,094,190 | 225,282 |
| <i>BNIPL</i> | rs6545714 | 2 | 59,307,725 | G | A | 39.2 | 39.2 | 8.9x10 ⁻⁹ | 31 | 372,883 | 59,291,172 | 59,664,054 | 30 | 59,291,172 | 59,664,054 | 372,883 |
| <i>BCL11A</i> | rs243024 | 2 | 60,583,665 | A | G | 46.0 | 46.0 | 2.5x10 ⁻²⁰ | 8 | 7,084 | 60,579,624 | 60,586,707 | 8 | 60,579,624 | 60,586,707 | 7,084 |
| <i>CEP68</i> | rs2249105 | 2 | 65,287,896 | A | G | 63.4 | 36.6 | 2.2x10 ⁻¹⁴ | 3 | 8,385 | 65,287,896 | 65,296,280 | 3 | 65,287,896 | 65,296,280 | 3,835 |
| <i>CEP68</i> | rs2052261 | 2 | 65,355,270 | G | A | 30.4 | 30.4 | 2.5x10 ⁻⁶ | 1,551 | 929,304 | 64,855,857 | 65,785,160 | 263 | 64,860,118 | 65,756,072 | 895,955 |
| <i>CEP68</i> | rs2028150 | 2 | 65,655,012 | C | G | 59.8 | 40.2 | 2.3x10 ⁻¹² | 24 | 42,056 | 65,642,097 | 65,684,152 | 22 | 65,642,097 | 65,684,152 | 42,056 |
| <i>TMEM127</i> | rs79046683 | 2 | 96,913,918 | T | G | 0.480 | 0.480 | 3.0x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>DDX18</i> | rs562386202 | 2 | 118,071,061 | G | A | 0.0600 | 0.0600 | 4.2x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>GLI2</i> | rs11688931 | 2 | 121,318,166 | C | G | 84.9 | 15.1 | 8.2x10 ⁻⁶ | 2,400 | 970,450 | 120,847,642 | 121,818,091 | 1,903 | 120,847,642 | 121,818,091 | 970,450 |
| <i>GLI2</i> | rs11688682 | 2 | 121,347,612 | G | C | 72.8 | 27.2 | 4.2x10 ⁻⁹ | 4 | 2,981 | 121,347,612 | 121,350,592 | 3 | 121,347,612 | 121,349,906 | 2,295 |
| <i>GLI2</i> | rs6647705 | 2 | 121,378,852 | T | C | 96.7 | 3.30 | 2.6x10 ⁻⁶ | 351 | 947,985 | 120,882,782 | 121,830,766 | 238 | 120,980,712 | 121,832,984 | 852,273 |
| <i>PABPC1P2</i> | rs35999103 | 2 | 147,861,633 | T | C | 15.5 | 15.5 | 9.7x10 ⁻⁹ | 53 | 146,388 | 147,807,932 | 147,954,319 | 54 | 147,807,932 | 147,954,319 | 146,388 |
| <i>CYTIP</i> | rs13426680 | 2 | 158,339,550 | A | G | 93.7 | 6.27 | 6.7x10 ⁻¹⁰ | 26 | 100,208 | 158,309,480 | 158,409,687 | 27 | 158,309,480 | 158,409,687 | 100,208 |
| <i>RBMS1</i> | rs3772071 | 2 | 161,135,544 | T | C | 71.4 | 28.7 | 1.2x10 ⁻¹¹ | 30 | 246,462 | 161,087,411 | 161,333,872 | 24 | 161,087,411 | 161,265,099 | 177,689 |
| <i>GRB14/COBLL1</i> | rs10195252 | 2 | 165,513,091 | T | C | 58.6 | 41.4 | 6.0x10 ⁻²⁵ | 5 | 31,273 | 165,508,389 | 165,539,661 | 5 | 165,508,389 | 165,539,661 | 31,273 |
| <i>GRB14/COBLL1</i> | rs13024606 | 2 | 165,573,194 | T | C | 4.72 | 4.72 | 1.7x10 ⁻⁸ | 41 | 818,947 | 165,144,110 | 165,963,056 | 35 | 165,470,735 | 165,702,813 | 232,079 |
| <i>CRYBA2</i> | rs113414093 | 2 | 219,859,171 | A | G | 5.14 | 5.14 | 6.6x10 ⁻⁹ | 793 | 995,525 | 219,361,518 | 220,357,042 | 170 | 219,390,845 | 220,349,385 | 958,541 |
| <i>IRS1</i> | rs2972144 | 2 | 227,101,411 | G | A | 63.9 | 36.2 | 2.1x10 ⁻⁴⁶ | 30 | 34,288 | 227,083,411 | 227,117,698 | 32 | 227,083,411 | 227,122,216 | 38,806 |
| <i>PPARG</i> | rs11709077 | 3 | 12,336,507 | G | A | 87.7 | 12.4 | 1.8x10 ⁻³⁶ | 11 | 67,131 | 12,329,783 | 12,396,913 | 8 | 12,329,783 | 12,393,125 | 63,343 |
| <i>PPARG</i> | rs17819328 | 3 | 12,489,342 | G | T | 42.5 | 42.5 | 4.8x10 ⁻¹⁶ | 8 | 10,383 | 12,488,882 | 12,499,264 | 8 | 12,488,882 | 12,499,264 | 10,383 |
| <i>UBE2E2</i> | rs35352848 | 3 | 23,455,582 | T | C | 78.8 | 21.2 | 1.3x10 ⁻¹⁷ | 5 | 2,291 | 23,454,790 | 23,457,080 | 5 | 23,454,790 | 23,457,080 | 2,291 |
| <i>UBE2E2</i> | rs17013314 | 3 | 23,510,044 | G | A | 3.13 | 3.13 | 8.4x10 ⁻⁹ | 209 | 385,895 | 23,252,089 | 23,637,983 | 208 | 23,252,089 | 23,637,983 | 385,895 |
| <i>KIF9</i> | rs11926707 | 3 | 46,925,539 | C | T | 62.6 | 37.4 | 2.1x10 ⁻⁸ | 838 | 999,083 | 46,426,391 | 47,425,473 | 248 | 46,447,326 | 47,425,473 | 978,148 |
| <i>KIF9</i> | rs75423501 | 3 | 47,242,923 | G | A | 10.1 | 10.1 | 7.5x10 ⁻⁶ | 1,083 | 681,388 | 46,743,934 | 47,425,321 | 954 | 46,743,934 | 47,425,473 | 681,540 |
| <i>RBM6</i> | rs4688760 | 3 | 49,980,596 | T | C | 68.4 | 31.6 | 3.5x10 ⁻¹⁰ | 155 | 438,284 | 49,736,565 | 50,174,848 | 164 | 49,735,746 | 50,174,848 | 439,103 |
| <i>RFT1</i> | rs2581787 | 3 | 53,127,677 | T | G | 56.3 | 43.7 | 2.4x10 ⁻⁸ | 43 | 151,248 | 52,989,383 | 53,140,630 | 35 | 53,103,796 | 53,139,977 | 36,182 |
| <i>CACNA2D3</i> | rs76263492 | 3 | 54,828,827 | T | G | 4.52 | 4.52 | 6.3x10 ⁻⁹ | 4 | 52,976 | 54,823,598 | 54,876,573 | 5 | 54,823,598 | 54,876,573 | 52,976 |
| <i>PSMD6</i> | rs3774723 | 3 | 63,962,339 | G | A | 84.4 | 15.6 | 1.6x10 ⁻¹³ | 20 | 151,397 | 63,853,423 | 64,004,819 | 18 | 63,853,423 | 64,004,819 | 151,397 |
| <i>PSMD6</i> | rs74368513 | 3 | 64,460,694 | G | A | 99.6 | 0.44 | 7.0x10 ⁻⁶ | 1,227 | 501,222 | 63,961,083 | 64,462,304 | 1,293 | 63,961,083 | 64,461,524 | 500,442 |

| | | | | | | | | | | | | | | | | |
|---------|-------------|---|-------------|---|---|-------|-------|-----------------------|-------|---------|-------------|-------------|-------|-------------|-------------|---------|
| TSC2D2 | rs62271373 | 3 | 150,066,540 | A | T | 5.53 | 5.53 | 1.0x10 ⁻⁹ | 5 | 35,329 | 150,047,893 | 150,083,221 | 9 | 150,018,637 | 150,090,266 | 71,630 |
| MBNL1 | rs13065698 | 3 | 152,086,533 | A | G | 60.0 | 40.0 | 8.1x10 ⁻¹³ | 27 | 172,252 | 151,993,809 | 152,166,060 | 22 | 151,993,809 | 152,166,060 | 172,292 |
| MBNL1 | rs74653713 | 3 | 152,417,881 | C | A | 95.7 | 4.29 | 1.2x10 ⁻⁸ | 100 | 165,759 | 152,369,607 | 152,535,365 | 98 | 152,369,607 | 152,535,365 | 165,759 |
| MBNL1 | rs35497231 | 3 | 152,433,628 | C | T | 31.7 | 31.7 | 7.6x10 ⁻⁸ | 149 | 186,566 | 152,320,458 | 152,507,023 | 141 | 152,320,458 | 152,507,023 | 186,566 |
| EGFEM1P | rs7629630 | 3 | 168,218,841 | A | T | 85.7 | 14.3 | 2.5x10 ⁻⁸ | 29 | 56,542 | 168,212,861 | 168,269,402 | 22 | 168,212,861 | 168,268,634 | 55,774 |
| SLC2A2 | rs9873618 | 3 | 170,733,076 | G | A | 71.0 | 29.0 | 4.8x10 ⁻²¹ | 14 | 105,168 | 170,627,909 | 170,733,076 | 11 | 170,665,296 | 170,733,076 | 67,781 |
| ABCC5 | rs2872246 | 3 | 183,738,460 | A | C | 45.4 | 45.4 | 1.5x10 ⁻⁸ | 100 | 347,159 | 183,411,802 | 183,758,960 | 86 | 183,563,571 | 183,946,205 | 382,635 |
| IGF2BP2 | rs6780171 | 3 | 185,503,456 | A | T | 31.4 | 31.4 | 9.0x10 ⁻⁵⁶ | 41 | 39,163 | 185,495,320 | 185,534,482 | 40 | 185,495,320 | 185,534,482 | 39,163 |
| IGF2BP2 | rs150111048 | 3 | 185,514,421 | G | A | 23.9 | 23.9 | 2.7x10 ⁻⁷ | 243 | 963,206 | 185,028,894 | 185,992,099 | 2 | 185,514,421 | 185,653,097 | 138,677 |
| IGF2BP2 | rs11717959 | 3 | 185,541,213 | G | T | 62.1 | 37.9 | 3.0x10 ⁻⁶ | 1,528 | 960,667 | 185,042,049 | 186,002,715 | 317 | 185,042,049 | 185,996,480 | 954,432 |
| IGF2BP2 | rs1516728 | 3 | 185,829,891 | A | T | 75.9 | 24.1 | 6.3x10 ⁻⁶ | 979 | 672,325 | 185,330,391 | 186,002,715 | 162 | 185,336,716 | 185,996,480 | 659,765 |
| ST6GAL1 | rs3887925 | 3 | 186,665,645 | T | C | 54.7 | 45.3 | 3.1x10 ⁻²² | 2 | 8,494 | 186,665,645 | 186,674,138 | 1 | 186,665,645 | 186,665,645 | 1 |
| ST6GAL1 | rs7645517 | 3 | 186,675,277 | A | G | 5.76 | 5.76 | 2.5x10 ⁻⁸ | 9 | 8,743 | 186,667,713 | 186,676,455 | 9 | 186,657,420 | 186,675,277 | 17,858 |
| LPP | rs4686471 | 3 | 187,740,899 | C | T | 61.0 | 39.0 | 1.7x10 ⁻²⁰ | 3 | 1,320 | 187,740,523 | 187,741,842 | 3 | 187,740,523 | 187,741,842 | 1,320 |
| PCGF3 | rs111827885 | 4 | 616,608 | C | T | 1.56 | 1.56 | 8.4x10 ⁻⁶ | 3,104 | 870,284 | 245,447 | 1,115,730 | 2,420 | 245,447 | 1,115,730 | 870,284 |
| PCGF3 | rs15131583 | 4 | 744,972 | T | G | 4.58 | 4.58 | 3.5x10 ⁻¹⁴ | 26 | 42,130 | 702,843 | 744,972 | 23 | 702,843 | 744,972 | 42,130 |
| PCGF3 | rs35654957 | 4 | 1,010,077 | C | T | 36.7 | 36.7 | 4.2x10 ⁻⁷ | 426 | 731,751 | 512,932 | 1,244,682 | 145 | 593,216 | 1,244,682 | 651,467 |
| MAEA | rs56337234 | 4 | 1,784,403 | C | T | 50.3 | 49.7 | 8.6x10 ⁻¹⁸ | 5 | 2,920 | 1,781,686 | 1,784,605 | 5 | 1,781,686 | 1,784,605 | 2,920 |
| HTT | rs362307 | 4 | 3,241,845 | T | C | 7.68 | 7.68 | 1.1x10 ⁻⁹ | 13 | 70,762 | 3,241,845 | 3,312,606 | 2 | 3,241,845 | 3,243,804 | 1,960 |
| WFS1 | rs1801212 | 4 | 6,302,519 | A | G | 70.9 | 29.1 | 4.5x10 ⁻⁶ | 3,997 | 995,219 | 5,806,789 | 6,802,007 | 2,013 | 5,809,127 | 6,800,680 | 991,554 |
| WFS1 | rs10937721 | 4 | 6,306,763 | C | G | 58.8 | 41.2 | 1.5x10 ⁻⁸ | 108 | 41,845 | 6,280,449 | 6,322,293 | 108 | 6,271,043 | 6,322,293 | 51,251 |
| LCRL | rs12640250 | 4 | 17,792,869 | C | A | 71.5 | 28.5 | 3.7x10 ⁻⁸ | 74 | 422,736 | 17,792,869 | 18,215,604 | 38 | 17,792,869 | 18,046,499 | 253,631 |
| GNPDA2 | rs10938398 | 4 | 45,186,139 | A | G | 42.9 | 42.9 | 3.6x10 ⁻¹² | 5 | 10,449 | 45,175,691 | 45,186,139 | 5 | 45,175,691 | 45,186,139 | 10,449 |
| USP46 | rs2102278 | 4 | 52,818,664 | G | A | 31.9 | 31.9 | 3.7x10 ⁻⁸ | 254 | 621,369 | 52,661,630 | 53,282,998 | 177 | 52,661,630 | 52,942,986 | 281,357 |
| USP46 | rs114447556 | 4 | 53,207,093 | T | C | 8.39 | 8.39 | 8.4x10 ⁻⁶ | 1,695 | 610,184 | 52,708,220 | 53,318,403 | 1,405 | 52,708,220 | 53,318,403 | 610,184 |
| SCD5 | rs12642790 | 4 | 83,578,271 | A | G | 33.8 | 33.8 | 4.4x10 ⁻¹⁰ | 32 | 27,770 | 83,561,503 | 83,589,272 | 7 | 83,578,271 | 83,587,562 | 9,292 |
| FAM13A | rs1903002 | 4 | 89,740,894 | G | C | 50.1 | 50.0 | 2.7x10 ⁻⁸ | 52 | 64,455 | 89,704,367 | 89,768,821 | 47 | 89,706,643 | 89,768,821 | 62,179 |
| FAM13A | rs576406049 | 4 | 89,857,291 | T | C | 0.130 | 0.130 | 1.6x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | |
| SMARCA1 | rs6821438 | 4 | 95,091,911 | A | G | 53.4 | 46.6 | 4.0x10 ⁻¹¹ | 138 | 312,989 | 95,000,394 | 95,313,382 | 152 | 95,012,459 | 95,318,346 | 305,888 |
| SLC9B1 | rs1580278 | 4 | 104,140,848 | C | A | 47.3 | 47.3 | 2.2x10 ⁻¹⁰ | 373 | 549,326 | 103,675,108 | 104,224,433 | 350 | 103,675,108 | 104,140,848 | 465,741 |
| PABPC4L | rs1296328 | 4 | 137,083,193 | A | C | 44.6 | 44.6 | 3.5x10 ⁻⁸ | 57 | 132,082 | 136,966,779 | 137,098,860 | 63 | 136,966,779 | 137,098,860 | 132,082 |
| TMEM154 | rs7669833 | 4 | 153,513,369 | T | A | 70.5 | 29.6 | 1.2x10 ⁻¹⁴ | 36 | 24,961 | 153,495,515 | 153,520,475 | 28 | 153,495,515 | 153,520,475 | 24,961 |
| PDGFC | rs28819812 | 4 | 157,652,753 | C | A | 67.7 | 32.3 | 2.2x10 ⁻⁸ | 120 | 131,233 | 157,614,725 | 157,745,957 | 122 | 157,614,725 | 157,757,755 | 143,031 |
| ACSL1 | rs58730668 | 4 | 185,717,759 | T | C | 85.8 | 14.2 | 1.3x10 ⁻¹³ | 46 | 17,143 | 185,713,608 | 185,730,750 | 44 | 185,713,608 | 185,730,750 | 17,143 |
| ANKH | rs3845281 | 5 | 14,610,134 | G | A | 90.4 | 9.61 | 2.3x10 ⁻¹¹ | 41 | 137,017 | 14,591,039 | 14,728,055 | 40 | 14,591,039 | 14,728,055 | 137,017 |
| ANKH | rs78408340 | 5 | 14,751,305 | C | T | 99.4 | 0.62 | 7.8x10 ⁻¹³ | 2 | 4,615 | 14,751,305 | 14,755,919 | 2 | 14,751,305 | 14,755,919 | 4,615 |
| ANKH | rs17250977 | 5 | 14,753,745 | G | A | 3.76 | 3.76 | 2.0x10 ⁻¹¹ | 2 | 14,781 | 14,738,965 | 14,753,745 | 2 | 14,738,965 | 14,753,745 | 14,781 |
| ANKH | rs6885132 | 5 | 14,768,092 | C | G | 90.4 | 9.60 | 1.7x10 ⁻⁸ | 74 | 285,404 | 14,568,231 | 14,853,634 | 41 | 14,763,925 | 14,835,599 | 71,675 |
| ANKH | rs76549217 | 5 | 14,768,766 | T | C | 2.95 | 2.95 | 3.0x10 ⁻¹⁰ | 1 | 1 | 14,768,766 | 14,768,766 | 1 | 14,768,766 | 14,768,766 | 1 |
| MRPS30 | rs62368490 | 5 | 44,534,364 | T | C | 3.13 | 3.13 | 3.4x10 ⁻⁶ | 763 | 588,240 | 44,446,074 | 45,034,313 | 237 | 44,446,133 | 45,030,781 | 584,649 |
| MRPS30 | rs6884702 | 5 | 44,682,589 | G | A | 39.3 | 39.3 | 1.5x10 ⁻¹⁰ | 247 | 302,421 | 44,642,670 | 44,945,090 | 227 | 44,642,670 | 44,945,090 | 302,421 |
| ITGA1 | rs17261179 | 5 | 51,791,225 | T | C | 51.7 | 48.3 | 1.3x10 ⁻⁸ | 25 | 74,215 | 51,720,684 | 51,794,898 | 26 | 51,720,684 | 51,794,898 | 74,215 |
| ITGA1 | rs3811978 | 5 | 52,100,489 | G | A | 16.7 | 16.7 | 7.7x10 ⁻¹¹ | 26 | 46,295 | 52,072,194 | 52,118,488 | 24 | 52,072,194 | 52,118,488 | 46,295 |
| ITGA1 | rs62357230 | 5 | 52,315,682 | A | G | 3.39 | 3.39 | 5.9x10 ⁻⁶ | 1,480 | 998,483 | 51,816,708 | 52,815,190 | 1,086 | 51,816,708 | 52,811,204 | 994,497 |
| ARL15 | rs62370480 | 5 | 52,774,510 | A | G | 22.0 | 22.0 | 2.0x10 ⁻⁶ | 1,652 | 999,128 | 52,275,165 | 53,274,292 | 150 | 52,297,932 | 53,268,072 | 970,141 |
| ARL15 | rs702634 | 5 | 53,271,420 | A | G | 69.0 | 31.0 | 7.7x10 ⁻¹⁴ | 3 | 4,882 | 53,271,420 | 53,276,301 | 3 | 53,271,420 | 53,276,301 | 4,882 |
| ARL15 | rs279744 | 5 | 53,412,620 | C | A | 69.1 | 30.9 | 3.1x10 ⁻⁸ | 97 | 192,436 | 53,290,571 | 53,483,006 | 97 | 53,290,571 | 53,483,006 | 192,436 |
| ANKRD55 | rs465002 | 5 | 55,808,475 | T | C | 74.2 | 25.8 | 6.1x10 ⁻³⁸ | 13 | 11,122 | 55,799,184 | 55,810,305 | 12 | 55,799,184 | 55,810,305 | 11,122 |
| ANKRD55 | rs2431115 | 5 | 55,848,669 | A | G | 40.2 | 40.2 | 3.9x10 ⁻¹⁰ | 16 | 19,035 | 55,834,402 | 55,853,436 | 16 | 55,834,402 | 55,853,436 | 19,035 |
| ANKRD55 | rs9687832 | 5 | 55,861,595 | A | G | 19.8 | 19.8 | 1.7x10 ⁻²⁰ | 8 | 5,520 | 55,856,375 | 55,861,894 | 8 | 55,856,375 | 55,861,894 | 5,520 |
| ANKRD55 | rs96844 | 5 | 56,196,604 | G | A | 26.2 | 26.2 | 5.4x10 ⁻⁸ | 62 | 286,150 | 55,989,823 | 56,275,972 | 17 | 56,106,474 | 56,261,335 | 154,862 |
| PIK3R1 | rs4976033 | 5 | 67,714,246 | G | A | 41.1 | 41.1 | 1.0x10 ⁻⁹ | 115 | 884,675 | 67,328,752 | 68,213,426 | 44 | 67,511,340 | 67,824,690 | 313,351 |
| POCS | rs2307111 | 5 | 75,003,678 | T | C | 60.5 | 39.5 | 2.1x10 ⁻¹⁶ | 12 | 81,234 | 74,934,009 | 75,015,242 | 9 | 74,965,122 | 75,010,002 | 44,881 |
| ZBED3 | rs4457053 | 5 | 76,424,949 | G | A | 30.4 | 30.4 | 8.4x10 ⁻¹⁸ | 5 | 10,056 | 76,424,949 | 76,435,004 | 5 | 76,424,949 | 76,435,004 | 10,056 |
| DMGDH | rs1316776 | 5 | 78,430,607 | C | A | 64.8 | 35.2 | 2.6x10 ⁻¹² | 65 | 153,804 | 78,416,416 | 78,570,219 | 65 | 78,416,416 | 78,602,863 | 186,448 |
| RASA1 | rs7719891 | 5 | 86,577,352 | G | A | 25.9 | 25.9 | 2.4x10 ⁻⁸ | 41 | 563,995 | 86,500,273 | 87,064,267 | 29 | 86,500,273 | 87,059,812 | 559,540 |
| SLCO6A1 | rs138337556 | 5 | 101,232,944 | G | A | 0.360 | 0.360 | 4.7x10 ⁻⁹ | 3 | 52,249 | 101,662,386 | 101,714,634 | 3 | 101,662,386 | 101,714,634 | 52,249 |
| PAM | rs78408340 | 5 | 102,338,739 | G | C | 0.830 | 0.830 | 2.1x10 ⁻²⁴ | 1 | 1 | 102,338,739 | 102,338,739 | 1 | 102,338,739 | 102,338,739 | 1 |
| PAM | | | | | | | | | | | | | | | | |

| | | | | | | | | | | | | | | | | |
|------------|-------------|---|-------------|---|---|--------|--------|------------------------|-------|---------|-------------|-------------|-------|-------------|-------------|---------|
| VEGFA | rs6458354 | 6 | 43,814,190 | C | T | 28.9 | 28.9 | 2.1x10 ⁻¹² | 21 | 25,134 | 43,804,808 | 43,829,941 | 20 | 43,805,502 | 43,829,941 | 24,440 |
| TFAP2B | rs3798519 | 6 | 50,788,778 | C | A | 18.4 | 18.4 | 2.6x10 ⁻¹² | 37 | 195,798 | 50,784,880 | 50,940,677 | 27 | 50,784,880 | 50,937,412 | 152,533 |
| TFAP2B | rs2465043 | 6 | 51,180,765 | G | A | 64.4 | 35.6 | 2.9x10 ⁻⁶ | 1,171 | 607,518 | 50,681,022 | 51,288,539 | 974 | 50,681,022 | 51,288,539 | 607,518 |
| SLC25A51P1 | rs555402748 | 6 | 67,387,490 | T | C | 0.0400 | 0.0400 | 4.6x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| BEND3 | rs4946812 | 6 | 107,431,688 | G | A | 67.4 | 32.6 | 8.2x10 ⁻⁹ | 17 | 23,028 | 107,422,239 | 107,445,266 | 17 | 107,422,239 | 107,445,266 | 23,028 |
| CENPW | rs11759026 | 6 | 126,792,095 | G | A | 23.2 | 23.2 | 2.4x10 ⁻¹⁸ | 2 | 172,416 | 126,792,095 | 126,964,510 | 2 | 126,792,095 | 126,964,510 | 172,416 |
| SOGA3 | rs2800733 | 6 | 127,416,930 | A | G | 71.7 | 28.4 | 6.0x10 ⁻¹¹ | 31 | 29,188 | 127,399,045 | 127,428,232 | 31 | 127,399,045 | 127,437,399 | 38,355 |
| SLC35D3 | rs9494624 | 6 | 137,300,960 | A | G | 29.0 | 29.0 | 6.1x10 ⁻⁹ | 34 | 17,765 | 137,287,702 | 137,305,466 | 33 | 137,287,702 | 137,302,880 | 15,179 |
| MIR3668 | rs2982521 | 6 | 139,835,329 | A | T | 38.0 | 38.0 | 1.3x10 ⁻⁹ | 14 | 7,569 | 139,829,560 | 139,837,128 | 14 | 139,829,560 | 139,837,128 | 7,569 |
| MIR3668 | rs616279 | 6 | 140,249,466 | A | G | 73.8 | 26.2 | 6.7x10 ⁻⁷ | 106 | 585,706 | 139,749,571 | 140,335,276 | 109 | 139,750,827 | 140,335,276 | 584,450 |
| SLC22A3 | rs474513 | 6 | 160,770,312 | A | G | 51.7 | 48.3 | 8.1x10 ⁻¹⁰ | 28 | 83,862 | 160,767,905 | 160,851,766 | 24 | 160,769,423 | 160,776,325 | 6,903 |
| QKI | rs4709746 | 6 | 164,133,001 | C | T | 86.8 | 13.2 | 5.8x10 ⁻⁹ | 7 | 29,759 | 164,103,243 | 164,133,001 | 7 | 164,103,243 | 164,133,001 | 29,759 |
| DGKB | rs17168486 | 7 | 14,898,282 | T | C | 18.1 | 18.1 | 2.3x10 ⁻¹⁷ | 6 | 62,871 | 14,859,137 | 14,922,007 | 5 | 14,859,137 | 14,922,007 | 62,871 |
| DGKB | rs10228066 | 7 | 15,063,569 | T | C | 53.7 | 46.3 | 1.1x10 ⁻²⁸ | 17 | 2,774 | 15,062,694 | 15,065,467 | 14 | 15,062,694 | 15,065,074 | 2,381 |
| DGKB | rs2908334 | 7 | 15,206,239 | T | C | 63.1 | 36.9 | 5.9x10 ⁻⁶ | 2,548 | 855,887 | 14,706,444 | 15,562,330 | 1,829 | 14,706,444 | 15,562,302 | 855,859 |
| IGF2BP3 | rs78840640 | 7 | 23,434,606 | G | C | 2.20 | 2.20 | 2.8x10 ⁻⁶ | 1,958 | 921,440 | 23,013,118 | 23,934,557 | 184 | 23,021,797 | 23,925,750 | 903,954 |
| IGF2BP3 | rs4279506 | 7 | 23,512,896 | G | C | 61.0 | 39.0 | 4.8x10 ⁻⁸ | 388 | 989,817 | 23,013,118 | 24,002,934 | 146 | 23,052,426 | 24,002,934 | 950,509 |
| JAZF1 | rs1708302 | 7 | 28,198,677 | C | T | 51.2 | 48.8 | 1.1x10 ⁻⁴⁸ | 6 | 7,818 | 28,192,280 | 28,200,097 | 6 | 28,192,280 | 28,200,097 | 7,818 |
| CRHR2 | rs917195 | 7 | 30,728,452 | C | T | 77.0 | 23.0 | 4.2x10 ⁻¹¹ | 4 | 638 | 30,727,920 | 30,728,557 | 2 | 30,728,452 | 30,728,557 | 106 |
| GCK | rs878521 | 7 | 44,255,643 | A | G | 24.5 | 24.5 | 1.9x10 ⁻¹³ | 2 | 23,866 | 44,231,778 | 44,255,643 | 1 | 44,255,643 | 44,255,643 | 1 |
| GCK | rs116913033 | 7 | 44,365,549 | C | T | 83.0 | 17.0 | 7.1x10 ⁻⁶ | 1,573 | 889,226 | 43,866,240 | 44,755,465 | 439 | 43,878,911 | 44,747,499 | 868,589 |
| FBXL13 | rs56376556 | 7 | 102,038,318 | T | C | 5.33 | 5.33 | 1.7x10 ⁻⁶ | 217 | 534,233 | 101,998,515 | 102,532,747 | 95 | 101,998,515 | 102,400,109 | 401,595 |
| FBXL13 | rs11496066 | 7 | 102,486,254 | T | C | 81.8 | 18.2 | 1.1x10 ⁻⁸ | 176 | 981,655 | 101,998,515 | 102,980,169 | 121 | 101,998,515 | 102,902,565 | 904,051 |
| RELN | rs62482405 | 7 | 102,987,583 | G | T | 8.19 | 8.19 | 6.9x10 ⁻⁶ | 1,320 | 996,423 | 102,489,118 | 103,485,540 | 346 | 102,553,366 | 103,473,515 | 920,150 |
| RELN | rs39328 | 7 | 103,444,978 | T | C | 43.3 | 43.3 | 3.7x10 ⁻⁸ | 66 | 990,544 | 102,946,088 | 103,936,631 | 35 | 103,006,613 | 103,531,911 | 525,299 |
| CTTNBP2 | rs6976111 | 7 | 117,495,667 | A | C | 31.3 | 31.3 | 1.2x10 ⁻⁸ | 36 | 508,226 | 117,086,613 | 117,594,838 | 15 | 117,495,667 | 117,590,626 | 94,960 |
| KLF14 | rs2268382 | 7 | 130,027,037 | C | A | 32.7 | 32.7 | 7.4x10 ⁻⁶ | 844 | 567,254 | 129,958,098 | 130,525,351 | 233 | 129,964,539 | 130,506,439 | 541,901 |
| KLF14 | rs1562396 | 7 | 130,457,914 | G | A | 31.9 | 31.9 | 9.9x10 ⁻¹⁸ | 26 | 40,188 | 130,427,388 | 130,467,575 | 26 | 130,430,969 | 130,467,575 | 36,607 |
| AOC1 | rs62492368 | 7 | 150,537,635 | A | G | 30.8 | 30.8 | 1.1x10 ⁻¹⁰ | 42 | 37,872 | 150,504,840 | 150,542,711 | 51 | 150,504,840 | 150,542,711 | 37,872 |
| MNX1 | rs6459733 | 7 | 156,930,550 | G | C | 67.3 | 32.7 | 2.4x10 ⁻¹⁷ | 78 | 102,375 | 156,930,550 | 157,032,924 | 76 | 156,930,550 | 157,032,924 | 102,375 |
| MSRA | rs17689007 | 8 | 9,974,824 | G | A | 53.3 | 46.7 | 2.5x10 ⁻⁹ | 20 | 22,672 | 9,973,718 | 9,996,389 | 18 | 9,973,718 | 9,996,389 | 22,672 |
| XKR6 | rs57327348 | 8 | 10,808,687 | A | T | 78.2 | 21.8 | 4.5x10 ⁻⁸ | 160 | 618,694 | 10,630,568 | 11,249,261 | 145 | 10,630,568 | 11,161,310 | 530,743 |
| LPL | rs10096633 | 8 | 19,830,921 | C | T | 87.7 | 12.3 | 1.1x10 ⁻¹² | 51 | 115,012 | 19,819,328 | 19,934,339 | 46 | 19,819,328 | 19,934,339 | 115,012 |
| PURG | rs10954772 | 8 | 30,863,938 | T | C | 31.4 | 31.4 | 1.8x10 ⁻⁹ | 20 | 28,404 | 30,835,535 | 30,863,938 | 21 | 30,828,375 | 30,863,938 | 35,564 |
| ANK1 | rs13262861 | 8 | 41,508,577 | C | A | 82.9 | 17.1 | 4.0x10 ⁻¹² | 9 | 16,323 | 41,508,577 | 41,524,899 | 2 | 41,508,577 | 41,522,991 | 14,415 |
| ANK1 | rs4736819 | 8 | 41,509,915 | T | C | 55.4 | 44.6 | 5.4x10 ⁻⁷ | 1,474 | 996,457 | 41,009,939 | 42,006,395 | 99 | 41,073,424 | 41,976,334 | 902,911 |
| ANK1 | rs14876658 | 8 | 41,552,046 | C | T | 3.78 | 3.78 | 5.7x10 ⁻⁷ | 384 | 942,125 | 41,062,358 | 42,004,482 | 147 | 41,073,424 | 42,003,850 | 930,427 |
| TP53INP1 | rs11786992 | 8 | 95,685,147 | A | C | 64.4 | 35.6 | 5.3x10 ⁻⁶ | 1,909 | 723,324 | 95,461,791 | 96,185,114 | 1,415 | 95,461,969 | 96,184,234 | 722,266 |
| TP53INP1 | rs10097617 | 8 | 95,961,626 | T | C | 48.5 | 48.5 | 3.3x10 ⁻¹¹ | 14 | 56,937 | 95,911,477 | 95,968,413 | 10 | 95,926,890 | 95,968,413 | 41,524 |
| TP53INP1 | rs187936726 | 8 | 96,092,422 | G | A | 2.39 | 2.39 | 5.4x10 ⁻⁶ | 2,329 | 868,216 | 95,593,358 | 96,461,573 | 1,627 | 95,595,702 | 96,461,573 | 865,872 |
| CPQ | rs149364428 | 8 | 97,737,741 | A | G | 1.04 | 1.04 | 1.8x10 ⁻¹² | 2 | 13,312 | 97,724,430 | 97,737,741 | 2 | 97,724,430 | 97,737,741 | 13,312 |
| TRHR | rs12680028 | 8 | 110,123,183 | C | G | 53.4 | 46.6 | 2.5x10 ⁻⁸ | 117 | 111,250 | 110,029,315 | 110,140,564 | 119 | 110,029,315 | 110,140,564 | 111,250 |
| SLC30A8 | rs3802177 | 8 | 118,185,025 | G | A | 68.5 | 31.5 | 1.1x10 ⁻⁵⁵ | 5 | 19,238 | 118,184,783 | 118,204,020 | 2 | 118,184,783 | 118,185,025 | 243 |
| SLC30A8 | rs80244329 | 8 | 118,404,672 | G | A | 97.8 | 2.19 | 6.9x10 ⁻⁶ | 1,770 | 780,008 | 117,905,012 | 118,685,019 | 1,674 | 117,905,012 | 118,684,994 | 779,983 |
| CASC11 | rs17772814 | 8 | 128,711,742 | G | A | 91.5 | 8.49 | 5.4x10 ⁻¹⁰ | 1 | 1 | 128,711,742 | 128,711,742 | 1 | 128,711,742 | 128,711,742 | 1 |
| PVT1 | rs1561927 | 8 | 129,568,078 | C | T | 26.9 | 26.9 | 1.5x10 ⁻⁹ | 55 | 32,546 | 129,538,595 | 129,571,140 | 52 | 129,538,595 | 129,571,140 | 32,546 |
| BOP1 | rs4977213 | 8 | 145,507,304 | C | T | 37.5 | 37.5 | 9.1x10 ⁻¹⁴ | 13 | 38,733 | 145,507,304 | 145,546,036 | 13 | 145,507,304 | 145,546,036 | 38,733 |
| BOP1 | rs12719778 | 8 | 145,879,883 | T | C | 53.8 | 46.2 | 5.0x10 ⁻⁹ | 76 | 301,061 | 145,706,276 | 146,007,336 | 74 | 145,639,726 | 146,007,336 | 367,611 |
| GLIS3 | rs510807 | 9 | 3,965,689 | A | C | 49.1 | 49.1 | 1.4x10 ⁻⁶ | 2,455 | 673,711 | 3,791,962 | 4,465,672 | 2,087 | 3,792,088 | 4,464,795 | 672,708 |
| GLIS3 | rs79103584 | 9 | 4,243,045 | T | A | 98.6 | 1.38 | 4.4x10 ⁻⁶ | 1,893 | 950,944 | 3,792,088 | 4,743,031 | 1,257 | 3,792,767 | 4,743,022 | 950,256 |
| GLIS3 | rs10974438 | 9 | 4,291,928 | C | A | 35.7 | 35.7 | 1.5x10 ⁻¹⁴ | 2 | 6,662 | 4,291,928 | 4,298,589 | 2 | 4,291,928 | 4,298,589 | 6,662 |
| HAUS6 | rs7022807 | 9 | 19,067,833 | G | A | 40.1 | 40.1 | 2.7x10 ⁻¹⁰ | 47 | 68,665 | 19,030,303 | 19,098,967 | 40 | 19,034,737 | 19,098,967 | 64,231 |
| FOCAD | rs7867635 | 9 | 20,241,069 | C | T | 41.2 | 41.2 | 4.0x10 ⁻⁸ | 544 | 994,178 | 19,744,948 | 20,739,125 | 340 | 19,769,828 | 20,739,125 | 969,298 |
| FOCAD | rs7847880 | 9 | 20,662,703 | C | T | 84.3 | 15.7 | 2.1x10 ⁻⁶ | 898 | 577,250 | 20,162,828 | 20,740,077 | 441 | 20,162,828 | 20,740,077 | 577,250 |
| CDKN2A/B | rs1412830 | 9 | 22,043,612 | C | T | 62.8 | 37.2 | 9.1x10 ⁻⁸ | 66 | 155,129 | 21,966,221 | 22,121,349 | 54 | 21,966,221 | 22,121,349 | 155,129 |
| CDKN2A/B | rs76011118 | 9 | 22,133,773 | A | G | 3.41 | 3.41 | 1.4x10 ⁻⁷ | 357 | 983,310 | 21,644,592 | 22,627,901 | 55 | 21,794,359 | 22,588,426 | 794,068 |
| CDKN2A/B | rs10811660 | 9 | 22,134,068 | G | A | 82.8 | 17.2 | 1.4x10 ⁻¹¹⁵ | 5 | 1,556 | 22,132,698 | 22,134,253 | 3 | 22,134,068 | 22,134,253 | 186 |
| CDKN2A/B | rs10757283 | 9 | 22,134,172 | T | C | 43.0 | 43.0 | 1.7x10 ⁻⁴¹ | 5 | 1,007 | 22,133,645 | | | | | |

| GPSM1 | 9:139737088:G:A | 9 | 139,737,088 | A | G | 0.0700 | 0.0700 | 7.9x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
|---------------|-----------------|----|-------------|---|---|--------|--------|------------------------|-------|---------|-------------|-------------|-------|-------------|-------------|---------|------------|------------|-----|
| CDC123/CAMK1D | rs11257655 | 10 | 12,307,894 | T | C | 21.8 | 21.8 | 1.5x10 ⁻³² | 1 | 1 | 12,307,894 | 12,307,894 | 1 | 12,307,894 | 12,307,894 | 1 | 12,307,894 | 12,307,894 | 1 |
| NEUROG3 | rs177045 | 10 | 71,321,279 | G | A | 31.6 | 31.6 | 6.6x10 ⁻¹⁸ | 2 | 337 | 71,320,943 | 71,321,279 | 3 | 71,320,943 | 71,332,204 | 11,262 | | | |
| NEUROG3 | rs61850200 | 10 | 71,321,658 | C | G | 27.7 | 27.7 | 7.3x10 ⁻⁶ | 2,560 | 854,835 | 70,966,757 | 71,821,591 | 1,198 | 70,966,757 | 71,820,982 | 854,226 | | | |
| NEUROG3 | rs41277236 | 10 | 71,332,301 | T | C | 4.31 | 4.31 | 1.5x10 ⁻⁶ | 2,519 | 865,527 | 70,966,757 | 71,832,283 | 3 | 71,332,301 | 71,446,372 | 114,072 | | | |
| NEUROG3 | rs549498088 | 10 | 71,347,311 | T | C | 0.600 | 0.600 | 4.7x10 ⁻⁷ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| NEUROG3 | rs2642588 | 10 | 71,466,578 | G | T | 70.2 | 29.8 | 2.2x10 ⁻¹⁴ | 18 | 26,683 | 71,464,178 | 71,490,860 | 19 | 71,464,178 | 71,490,860 | 26,683 | | | |
| ZMZ1 | rs703972 | 10 | 80,952,826 | G | C | 53.3 | 46.7 | 1.7x10 ⁻²⁹ | 10 | 10,832 | 80,942,620 | 80,953,451 | 10 | 80,942,620 | 80,953,451 | 10,832 | | | |
| ZMZ1 | rs1317617 | 10 | 81,096,589 | G | A | 79.8 | 20.2 | 1.8x10 ⁻⁶ | 1,583 | 855,529 | 80,596,754 | 81,452,282 | 861 | 80,598,052 | 81,452,282 | 854,231 | | | |
| PTEN | rs11202627 | 10 | 89,769,340 | T | C | 15.2 | 15.2 | 4.7x10 ⁻⁸ | 122 | 817,608 | 89,269,373 | 90,086,980 | 67 | 89,344,293 | 89,769,340 | 425,048 | | | |
| HHEX/IDE | rs7078559 | 10 | 93,924,663 | T | C | 57.8 | 42.2 | 4.1x10 ⁻⁷ | 175 | 678,129 | 93,677,338 | 94,355,951 | 122 | 93,667,030 | 94,134,467 | 467,438 | | | |
| HHEX/IDE | rs10882101 | 10 | 94,462,427 | T | C | 58.7 | 41.3 | 1.4x10 ⁻⁸ | 10 | 21,703 | 94,444,793 | 94,466,495 | 10 | 94,430,497 | 94,466,427 | 35,931 | | | |
| HHEX/IDE | rs1112718 | 10 | 94,479,107 | A | G | 59.8 | 40.2 | 5.0x10 ⁻⁷ | 1,828 | 998,391 | 93,980,048 | 94,978,438 | 754 | 93,980,048 | 94,972,243 | 992,196 | | | |
| TCF7L2 | rs536643418 | 10 | 114,699,835 | G | C | 0.520 | 0.520 | 2.6x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| TCF7L2 | rs140242150 | 10 | 114,702,962 | A | G | 0.500 | 0.500 | 2.2x10 ⁻⁸ | 4 | 81,965 | 114,702,962 | 114,784,926 | 5 | 114,702,962 | 114,899,115 | 196,154 | | | |
| TCF7L2 | rs7918400 | 10 | 114,703,136 | T | C | 47.6 | 47.6 | 2.0x10 ⁻¹⁵ | 4 | 9,547 | 114,702,608 | 114,712,154 | 5 | 114,702,608 | 114,712,154 | 9,547 | | | |
| TCF7L2 | rs184509201 | 10 | 114,740,337 | C | G | 98.2 | 1.82 | 1.2x10 ⁻¹³ | 9 | 6,502 | 114,736,670 | 114,743,171 | 8 | 114,736,670 | 114,743,171 | 6,502 | | | |
| TCF7L2 | rs180988137 | 10 | 114,751,173 | G | A | 1.04 | 1.04 | 6.1x10 ⁻⁶ | 1,979 | 984,191 | 114,266,338 | 115,250,528 | 1,608 | 114,266,992 | 115,250,528 | 983,537 | | | |
| TCF7L2 | rs7903146 | 10 | 114,758,349 | C | T | 70.6 | 29.5 | 5.8x10 ⁻⁴⁴⁷ | 3 | 4,279 | 114,754,071 | 114,758,349 | 2 | 114,754,071 | 114,758,349 | 4,279 | | | |
| TCF7L2 | rs78025551 | 10 | 114,757,956 | C | G | 85.1 | 14.9 | 1.6x10 ⁻⁷ | 339 | 985,504 | 114,271,035 | 115,256,538 | 133 | 114,301,588 | 115,240,149 | 938,562 | | | |
| TCF7L2 | rs34855922 | 10 | 114,871,594 | A | G | 71.6 | 28.4 | 5.5x10 ⁻¹² | 2 | 2,813 | 114,871,594 | 114,874,406 | 2 | 114,871,594 | 114,874,406 | 2,813 | | | |
| WDR11 | rs72631105 | 10 | 122,915,345 | A | G | 19.0 | 19.0 | 3.7x10 ⁻⁹ | 468 | 994,351 | 122,417,658 | 123,412,008 | 669 | 122,422,709 | 123,412,008 | 989,300 | | | |
| PLEKHA1 | rs2280141 | 10 | 124,193,181 | T | G | 51.6 | 48.4 | 1.4x10 ⁻¹³ | 25 | 59,193 | 124,139,393 | 124,198,585 | 26 | 124,134,803 | 124,198,585 | 63,783 | | | |
| INS/IGF2 | rs12802972 | 11 | 1,704,596 | A | G | 42.8 | 42.8 | 1.5x10 ⁻⁶ | 591 | 989,157 | 1,215,046 | 2,204,202 | 574 | 1,219,797 | 2,204,202 | 984,406 | | | |
| INS/IGF2 | rs11042596 | 11 | 2,118,860 | G | T | 66.5 | 33.5 | 2.0x10 ⁻⁸ | 144 | 893,553 | 1,665,146 | 2,558,698 | 118 | 1,858,632 | 2,418,136 | 559,505 | | | |
| INS/IGF2 | rs555759341 | 11 | 2,151,761 | C | G | 0.490 | 0.490 | 3.6x10 ⁻⁸ | 43 | 694,695 | 1,864,004 | 2,558,698 | 186 | 1,653,505 | 2,558,698 | 905,194 | | | |
| INS/IGF2 | rs571342427 | 11 | 2,182,519 | C | T | 0.150 | 0.150 | 1.0x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| KCNQ1 | rs4929965 | 11 | 2,197,286 | A | G | 38.3 | 38.3 | 4.0x10 ⁻²⁶ | 13 | 3,690 | 2,193,597 | 2,197,286 | 12 | 2,193,840 | 2,197,286 | 3,447 | | | |
| KCNQ1 | rs4930091 | 11 | 2,372,356 | C | T | 75.9 | 24.1 | 3.7x10 ⁻⁶ | 2,086 | 998,827 | 1,873,014 | 2,871,840 | 930 | 1,883,395 | 2,865,413 | 982,019 | | | |
| KCNQ1 | rs2283164 | 11 | 2,579,163 | A | G | 94.7 | 5.32 | 1.2x10 ⁻⁷ | 743 | 997,379 | 2,080,612 | 3,077,990 | 784 | 2,080,612 | 3,078,885 | 998,274 | | | |
| KCNQ1 | rs80102379 | 11 | 2,634,177 | G | T | 98.2 | 1.78 | 9.3x10 ⁻⁸ | 108 | 970,840 | 2,140,569 | 3,111,408 | 13 | 2,283,950 | 2,950,558 | 666,609 | | | |
| KCNQ1 | rs231349 | 11 | 2,672,821 | T | C | 10.2 | 10.2 | 2.3x10 ⁻¹¹ | 33 | 37,439 | 2,643,377 | 2,680,815 | 32 | 2,643,878 | 2,680,815 | 36,938 | | | |
| KCNQ1 | rs231361 | 11 | 2,691,500 | A | G | 25.6 | 25.6 | 5.0x10 ⁻²⁵ | 1 | 1 | 2,691,500 | 2,691,500 | 1 | 2,691,500 | 2,691,500 | 1 | | | |
| KCNQ1 | rs2283220 | 11 | 2,755,548 | A | G | 69.0 | 31.0 | 1.4x10 ⁻⁹ | 4 | 54 | 2,755,548 | 2,755,601 | 6 | 2,752,593 | 2,799,679 | 47,087 | | | |
| KCNQ1 | rs234853 | 11 | 2,850,828 | G | A | 24.8 | 24.8 | 6.8x10 ⁻¹⁶ | 8 | 14,095 | 2,843,803 | 2,857,897 | 9 | 2,840,424 | 2,857,897 | 17,474 | | | |
| KCNQ1 | rs2237895 | 11 | 2,857,194 | C | A | 42.6 | 42.6 | 6.0x10 ⁻⁵² | 1 | 1 | 2,857,194 | 2,857,194 | 1 | 2,857,194 | 2,857,194 | 1 | | | |
| KCNQ1 | rs2237897 | 11 | 2,858,546 | C | T | 95.4 | 4.57 | 8.4x10 ⁻³² | 3 | 197 | 2,858,440 | 2,858,636 | 3 | 2,858,440 | 2,858,636 | 197 | | | |
| KCNQ1 | rs445084 | 11 | 2,908,754 | G | A | 36.1 | 36.1 | 1.7x10 ⁻⁶ | 2,491 | 999,795 | 2,408,916 | 3,408,710 | 594 | 2,408,916 | 3,400,820 | 991,905 | | | |
| PDE3B | rs141521721 | 11 | 14,763,828 | A | C | 2.36 | 2.36 | 2.7x10 ⁻⁸ | 79 | 804,833 | 14,299,149 | 15,103,981 | 88 | 14,299,149 | 15,103,981 | 804,833 | | | |
| KCNJ11 | rs5213 | 11 | 17,408,404 | C | T | 36.2 | 36.2 | 3.5x10 ⁻²⁷ | 6 | 10,074 | 17,408,404 | 17,418,477 | 3 | 17,408,404 | 17,409,572 | 1,169 | | | |
| KCNJ11 | rs67254669 | 11 | 17,470,143 | G | A | 0.110 | 0.110 | 1.1x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| METTL15 | rs4923543 | 11 | 28,534,898 | A | G | 33.2 | 33.2 | 4.5x10 ⁻⁸ | 1,875 | 998,356 | 28,036,270 | 29,034,625 | 1,665 | 28,036,270 | 29,034,454 | 998,185 | | | |
| QSER1 | rs7943101 | 11 | 32,460,873 | T | C | 16.1 | 16.1 | 8.5x10 ⁻⁶ | 1,150 | 533,000 | 32,427,823 | 32,960,822 | 649 | 32,427,823 | 32,960,822 | 533,000 | | | |
| QSER1 | rs145678014 | 11 | 32,927,778 | G | T | 95.7 | 4.33 | 2.0x10 ⁻¹⁰ | 58 | 360,895 | 32,595,598 | 32,956,492 | 22 | 32,595,598 | 32,956,492 | 360,895 | | | |
| QSER1 | rs521822639 | 11 | 33,091,735 | A | G | 0.0900 | 0.0900 | 1.6x10 ⁻⁷ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| PDHX | rs286925 | 11 | 34,642,668 | A | G | 18.2 | 18.2 | 5.0x10 ⁻¹⁶ | 1,597 | 658,488 | 34,484,090 | 35,142,577 | 311 | 34,482,908 | 35,104,315 | 621,408 | | | |
| PDHX | rs2767036 | 11 | 34,982,148 | C | A | 29.1 | 29.1 | 3.3x10 ⁻⁸ | 273 | 272,253 | 34,759,989 | 35,032,241 | 254 | 34,759,989 | 35,032,241 | 272,253 | | | |
| HSD17B12 | rs1061610 | 11 | 43,877,934 | A | C | 28.8 | 28.8 | 6.0x10 ⁻¹³ | 67 | 125,964 | 43,752,522 | 43,878,485 | 66 | 43,752,522 | 43,878,485 | 125,964 | | | |
| CRY2 | rs7115753 | 11 | 45,912,013 | A | G | 44.9 | 44.9 | 3.8x10 ⁻⁹ | 29 | 74,557 | 45,837,457 | 45,912,013 | 22 | 45,839,889 | 45,912,013 | 72,125 | | | |
| CELF1 | rs7124681 | 11 | 47,529,947 | A | C | 41.0 | 41.0 | 5.1x10 ⁻⁹ | 223 | 603,428 | 47,385,350 | 47,988,777 | 185 | 47,437,033 | 48,001,082 | 564,050 | | | |
| MAP3K11 | rs1783541 | 11 | 65,294,799 | T | C | 20.4 | 20.4 | 2.0x10 ⁻¹⁴ | 12 | 79,725 | 65,257,527 | 65,337,251 | 11 | 65,263,895 | 65,337,251 | 73,357 | | | |
| CCND1 | rs61881115 | 11 | 68,997,225 | G | A | 83.8 | 16.2 | 4.1x10 ⁻⁷ | 97 | 521,852 | 68,951,753 | 69,473,604 | 121 | 68,951,753 | 69,489,910 | 538,158 | | | |
| CCND1 | rs11820019 | 11 | 69,448,758 | T | C | 97.3 | 2.67 | 5.1x10 ⁻¹² | 12 | 14,922 | 69,448,758 | 69,463,679 | 11 | 69,448,758 | 69,463,679 | 14,922 | | | |
| CENTD2/ARAP1 | rs77464186 | 11 | 72,460,398 | A | C | 83.6 | 16.4 | 4.7x10 ⁻³³ | 9 | 33,857 | 72,429,579 | 72,463,435 | 9 | 72,428,172 | 72,463,435 | 35,264 | | | |
| MTNR1B | rs10830963 | 11 | 92,708,710 | G | C | 27.7 | 27.7 | 4.8x10 ⁻⁴³ | 1 | 1 | 92,708,710 | 92,708,710 | 1 | 92,708,710 | 92,708,710 | 1 | | | |
| MTNR1B | rs57235767 | 11 | 93,013,531 | C | T | 70.6 | 29.4 | 5.9x10 ⁻¹⁰ | 63 | | | | | | | | | | |

| | | | | | | | | | | | | | | | | |
|--------------------|-------------|----|-------------|---|---|--------|--------|-----------------------|-------|---------|-------------|-------------|-------|-------------|-------------|---------|
| <i>TSPAN8/LGR5</i> | rs1796330 | 12 | 71,522,953 | G | C | 57.1 | 42.9 | 2.2x10 ⁻¹⁴ | 119 | 131,397 | 71,408,690 | 71,540,086 | 116 | 71,409,261 | 71,538,310 | 129,050 |
| <i>USP44</i> | rs2197973 | 12 | 95,928,560 | T | C | 53.8 | 46.3 | 3.6x10 ⁻⁸ | 83 | 404,347 | 95,527,476 | 95,931,822 | 54 | 95,527,476 | 95,928,560 | 401,085 |
| <i>RMST</i> | rs759111467 | 12 | 97,562,756 | A | G | 0.0300 | 0.0300 | 1.7x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>RMST</i> | rs557027608 | 12 | 97,779,248 | A | G | 0.0600 | 0.0600 | 2.7x10 ⁻⁷ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>RMST</i> | rs77864822 | 12 | 97,848,775 | A | G | 93.2 | 6.76 | 1.1x10 ⁻⁸ | 18 | 24,363 | 97,843,343 | 97,867,705 | 20 | 97,843,343 | 97,868,906 | 25,564 |
| <i>WSCD2</i> | rs1426371 | 12 | 108,629,780 | G | A | 73.9 | 26.1 | 8.2x10 ⁻¹² | 3 | 20,147 | 108,609,634 | 108,629,780 | 3 | 108,609,634 | 108,629,780 | 20,147 |
| <i>KSR2</i> | rs34965774 | 12 | 118,412,373 | A | G | 14.4 | 14.4 | 2.0x10 ⁻⁹ | 10 | 20,919 | 118,394,008 | 118,414,926 | 10 | 118,394,008 | 118,414,926 | 20,919 |
| <i>KSR2</i> | rs12578639 | 12 | 118,489,636 | A | T | 82.8 | 17.2 | 2.2x10 ⁻⁶ | 993 | 919,999 | 117,991,656 | 118,911,654 | 78 | 118,204,979 | 118,843,443 | 638,465 |
| <i>HNF1A</i> | rs11065299 | 12 | 121,297,815 | A | G | 7.54 | 7.54 | 5.8x10 ⁻⁷ | 895 | 864,004 | 120,932,742 | 121,796,745 | 631 | 120,933,092 | 121,790,178 | 85,078 |
| <i>HNF1A</i> | rs73226260 | 12 | 121,380,541 | G | A | 96.7 | 3.31 | 5.9x10 ⁻¹¹ | 22 | 42,553 | 121,381,097 | 121,423,649 | 20 | 121,381,097 | 121,422,243 | 41,147 |
| <i>HNF1A</i> | rs1800574 | 12 | 121,416,864 | T | C | 2.96 | 2.96 | 1.7x10 ⁻¹² | 2 | 443 | 121,416,864 | 121,417,306 | 1 | 121,416,864 | 121,416,864 | 1 |
| <i>HNF1A</i> | rs56348580 | 12 | 121,432,117 | G | C | 68.9 | 31.1 | 2.3x10 ⁻¹³ | 5 | 27,423 | 121,429,194 | 121,456,616 | 3 | 121,432,117 | 121,455,589 | 23,473 |
| <i>HNF1A</i> | rs28638142 | 12 | 121,501,461 | A | C | 4.42 | 4.42 | 2.9x10 ⁻⁶ | 1,954 | 927,549 | 121,004,376 | 121,931,924 | 1,216 | 121,004,376 | 121,928,221 | 923,846 |
| <i>HNF1A</i> | rs73224262 | 12 | 121,882,395 | T | C | 0.680 | 0.680 | 9.1x10 ⁻⁷ | 116 | 539,850 | 121,382,697 | 121,922,546 | 36 | 121,386,617 | 121,922,546 | 535,930 |
| <i>MPHOSPH9</i> | rs4148856 | 12 | 123,450,765 | C | G | 78.1 | 21.9 | 1.7x10 ⁻¹⁰ | 208 | 612,745 | 123,296,204 | 123,908,948 | 180 | 123,296,204 | 123,897,177 | 600,974 |
| <i>ZNF664</i> | rs7978610 | 12 | 124,468,572 | G | C | 66.6 | 33.5 | 2.0x10 ⁻⁸ | 99 | 147,139 | 124,348,065 | 124,495,203 | 94 | 124,404,718 | 124,495,203 | 90,486 |
| <i>ZNF664</i> | rs825452 | 12 | 124,509,177 | A | G | 60.3 | 39.7 | 2.4x10 ⁻⁶ | 2,288 | 958,044 | 124,009,190 | 124,967,233 | 1,824 | 124,009,190 | 124,968,406 | 959,217 |
| <i>FBRSL1</i> | rs12811407 | 12 | 133,069,698 | A | G | 33.1 | 33.1 | 1.7x10 ⁻¹² | 11 | 21,899 | 133,066,392 | 133,088,290 | 8 | 133,066,392 | 133,088,290 | 21,899 |
| <i>RNF6</i> | rs34584161 | 13 | 26,776,999 | A | G | 76.0 | 24.0 | 2.2x10 ⁻¹⁰ | 16 | 22,638 | 26,776,255 | 26,798,892 | 11 | 26,776,999 | 26,797,320 | 20,322 |
| <i>HMGB1</i> | rs11842871 | 13 | 31,042,452 | G | T | 73.5 | 26.6 | 1.2x10 ⁻⁸ | 20 | 27,299 | 31,015,154 | 31,042,452 | 14 | 31,019,580 | 31,042,452 | 22,873 |
| <i>KL</i> | rs576674 | 13 | 33,554,302 | G | A | 16.9 | 16.9 | 8.3x10 ⁻¹⁰ | 9 | 8,204 | 33,554,302 | 33,562,505 | 7 | 33,554,302 | 33,562,505 | 8,204 |
| <i>DLEU1</i> | rs963740 | 13 | 51,096,095 | A | T | 71.3 | 28.7 | 2.1x10 ⁻⁸ | 73 | 514,058 | 50,738,924 | 51,252,981 | 5 | 51,087,007 | 51,096,095 | 9,089 |
| <i>PCDH17</i> | rs9537803 | 13 | 58,366,634 | C | T | 27.7 | 27.7 | 4.6x10 ⁻⁸ | 1,566 | 891,958 | 57,974,269 | 58,866,226 | 318 | 57,986,844 | 58,576,243 | 589,400 |
| <i>PCDH17</i> | rs9569864 | 13 | 58,965,435 | C | T | 82.5 | 17.5 | 8.7x10 ⁻⁸ | 270 | 922,369 | 58,472,262 | 59,394,630 | 157 | 58,472,262 | 59,184,234 | 711,973 |
| <i>SRGAP2D</i> | rs9563615 | 13 | 59,077,406 | A | T | 71.0 | 29.0 | 6.4x10 ⁻¹¹ | 3,909 | 998,840 | 58,578,463 | 59,577,302 | 2,051 | 58,866,706 | 59,576,788 | 710,083 |
| <i>SRGAP2D</i> | rs76251711 | 13 | 59,184,234 | G | A | 1.26 | 1.26 | 2.3x10 ⁻⁶ | 74 | 707,871 | 58,686,760 | 59,394,630 | 44 | 58,686,760 | 59,184,234 | 497,475 |
| <i>SPRY2</i> | rs1359790 | 13 | 80,717,156 | G | A | 72.0 | 28.0 | 2.4x10 ⁻³¹ | 10 | 11,842 | 80,705,315 | 80,717,156 | 9 | 80,705,315 | 80,717,156 | 11,842 |
| <i>IRS2</i> | rs7987740 | 13 | 109,947,213 | T | C | 60.9 | 39.1 | 4.0x10 ⁻⁸ | 71 | 182,606 | 109,833,363 | 110,015,968 | 26 | 109,850,598 | 110,004,571 | 153,974 |
| <i>IRS2</i> | rs4771648 | 13 | 110,431,626 | G | A | 66.9 | 33.2 | 8.9x10 ⁻⁸ | 28 | 294,886 | 110,151,979 | 110,446,864 | 8 | 110,420,832 | 110,446,864 | 26,033 |
| <i>SLC7A7</i> | rs17122772 | 14 | 23,288,935 | G | C | 22.8 | 22.8 | 1.6x10 ⁻⁸ | 5 | 8,393 | 23,280,797 | 23,289,189 | 4 | 23,288,935 | 23,289,189 | 255 |
| <i>AKAP6</i> | rs17522122 | 14 | 33,302,882 | T | G | 47.4 | 47.4 | 3.2x10 ⁻⁹ | 33 | 16,615 | 33,292,743 | 33,309,357 | 33 | 33,292,743 | 33,309,357 | 16,615 |
| <i>CLEC14A</i> | rs8017808 | 14 | 38,848,419 | G | T | 74.3 | 25.7 | 2.1x10 ⁻⁸ | 69 | 92,264 | 38,756,561 | 38,848,824 | 68 | 38,756,561 | 38,848,824 | 92,264 |
| <i>NRXN3</i> | rs17836088 | 14 | 79,932,041 | C | G | 21.7 | 21.7 | 6.7x10 ⁻¹⁴ | 16 | 54,707 | 79,890,456 | 79,945,162 | 16 | 79,890,456 | 79,945,162 | 54,707 |
| <i>SMEK1</i> | rs8010382 | 14 | 91,963,722 | G | A | 42.1 | 42.1 | 6.5x10 ⁻⁹ | 130 | 188,041 | 91,871,672 | 92,059,712 | 105 | 91,860,800 | 92,041,956 | 181,157 |
| <i>MARK3</i> | rs62007683 | 14 | 103,894,071 | G | T | 65.3 | 34.7 | 3.1x10 ⁻⁸ | 91 | 153,436 | 103,852,725 | 104,006,160 | 88 | 103,851,272 | 104,006,160 | 154,889 |
| <i>RASGRP1</i> | rs8032939 | 15 | 38,834,033 | C | T | 24.6 | 24.6 | 3.5x10 ⁻¹⁴ | 12 | 19,624 | 38,828,140 | 38,847,763 | 11 | 38,828,140 | 38,846,738 | 18,599 |
| <i>RASGRP1</i> | rs34715063 | 15 | 38,873,115 | C | T | 12.4 | 12.4 | 2.3x10 ⁻¹⁹ | 2 | 20,730 | 38,852,386 | 38,873,115 | 2 | 38,852,386 | 38,873,115 | 20,730 |
| <i>LTK</i> | rs11070332 | 15 | 41,809,205 | A | G | 35.8 | 35.8 | 1.1x10 ⁻¹³ | 9 | 31,697 | 41,801,512 | 41,833,208 | 14 | 41,801,512 | 41,836,234 | 34,723 |
| <i>LTK</i> | rs543786825 | 15 | 42,201,410 | T | C | 0.0400 | 0.0400 | 4.7x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | |
| <i>ONECUT1</i> | rs2456530 | 15 | 53,091,553 | T | C | 12.7 | 12.7 | 5.4x10 ⁻⁹ | 52 | 95,541 | 53,070,141 | 53,165,681 | 51 | 53,058,612 | 53,165,681 | 107,070 |
| <i>WDR72</i> | rs528350911 | 15 | 53,747,228 | G | C | 0.680 | 0.680 | 2.1x10 ⁻⁸ | 1 | 1 | 53,747,228 | 53,747,228 | 1 | 53,747,228 | 53,747,228 | 1 |
| <i>TCF12</i> | rs117483894 | 15 | 57,456,802 | G | A | 3.69 | 3.69 | 3.9x10 ⁻⁸ | 188 | 642,956 | 56,968,627 | 57,611,582 | 179 | 56,957,179 | 57,611,582 | 654,404 |
| <i>C2CD4A/B</i> | rs8037894 | 15 | 62,394,264 | G | C | 56.6 | 43.4 | 2.6x10 ⁻¹³ | 15 | 15,939 | 62,383,155 | 62,399,093 | 14 | 62,383,155 | 62,399,093 | 15,939 |
| <i>USP3</i> | rs7178762 | 15 | 63,871,292 | C | T | 46.0 | 46.0 | 5.4x10 ⁻¹⁰ | 109 | 331,410 | 63,800,065 | 64,131,474 | 108 | 63,800,065 | 64,131,474 | 331,410 |
| <i>MAP2K5</i> | rs4776970 | 15 | 68,080,886 | A | T | 64.1 | 35.9 | 5.0x10 ⁻⁹ | 49 | 167,502 | 67,937,755 | 68,105,256 | 39 | 67,937,755 | 68,120,644 | 182,890 |
| <i>PTPN9</i> | rs13737 | 15 | 75,932,129 | G | T | 75.9 | 24.1 | 5.6x10 ⁻¹⁰ | 76 | 251,778 | 75,681,006 | 75,932,777 | 50 | 75,693,635 | 75,932,777 | 239,143 |
| <i>HMG20A</i> | rs1005752 | 15 | 77,818,128 | A | C | 71.5 | 28.5 | 2.5x10 ⁻²⁹ | 24 | 121,044 | 77,711,719 | 77,832,762 | 20 | 77,711,719 | 77,828,668 | 116,950 |
| <i>AP3S2</i> | rs4932265 | 15 | 90,423,293 | T | C | 26.7 | 26.7 | 4.2x10 ⁻²⁰ | 46 | 81,457 | 90,368,067 | 90,449,523 | 48 | 90,368,067 | 90,449,523 | 81,457 |
| <i>PRC1</i> | rs12910825 | 15 | 91,511,260 | G | A | 36.1 | 36.1 | 1.6x10 ⁻¹⁵ | 25 | 30,176 | 91,506,452 | 91,536,627 | 37 | 91,503,752 | 91,551,876 | 48,125 |
| <i>ITFG3</i> | rs6600191 | 16 | 295,795 | T | C | 82.5 | 17.5 | 9.3x10 ⁻¹³ | 29 | 20,809 | 294,749 | 315,557 | 28 | 294,749 | 315,557 | 20,809 |
| <i>CLUAP1</i> | rs3751837 | 16 | 3,583,173 | T | C | 22.0 | 22.0 | 1.4x10 ⁻⁸ | 92 | 984,483 | 3,098,691 | 4,083,173 | 17 | 3,117,993 | 4,071,354 | 953,362 |
| <i>ATP2A1</i> | rs8046545 | 16 | 28,915,217 | G | A | 35.9 | 35.9 | 1.9x10 ⁻⁸ | 94 | 127,954 | 28,824,685 | 28,952,638 | 97 | 28,833,097 | 28,995,757 | 162,661 |
| <i>FAM57B</i> | rs11642430 | 16 | 30,045,789 | G | C | 39.9 | 39.9 | 2.2x10 ⁻⁹ | 58 | 159,038 | 29,958,216 | 30,117,253 | 61 | 29,958,216 | 30,117,253 | 159,038 |
| <i>FAM57B</i> | rs199795270 | 16 | 30,419,384 | C | G | 0.650 | 0.650 | 1.2x10 ⁻⁶ | 117 | 608,973 | 29,935,456 | 30,544,428 | 1 | 30,419,384 | 30,419,384 | 1 |
| <i>FTO</i> | rs4281707 | 16 | 53,501,946 | G | A | 54.4 | 45.6 | 3.2x10 ⁻¹⁰ | 73 | 115,821 | 53,429,173 | 53,544,993 | 69 | 53,429,173 | 53,544,993 | 115,821 |
| <i>FTO</i> | rs78020297 | 16 | 53,758,720 | A | G | 5.17 | 5.17 | 6.5x10 ⁻⁹ | 7 | 13,694 | 53,756,885 | 53,770,578 | 7 | 53,756,885 | 53,770,578 | 13,694 |
| <i>FTO</i> | rs1421085 | 16 | 53,800,954 | C | T | 41.5 | 41.5 | 3.1x10 ⁻⁸⁴ | 6 | 8,170 | 53,800,954 | 53,809,123 | 6 | 53,800,954 | 53,809,123 | 8,170 |

| | | | | | | | | | | | | | | | | |
|--------------------|-------------|----|------------|---|---|--------|--------|-----------------------|-------|---------|------------|------------|-------|------------|------------|---------|
| <i>HNF1B</i> | rs2189301 | 17 | 36,063,685 | G | A | 87.2 | 12.8 | 6.5x10 ⁻⁸ | 7 | 16,340 | 36,048,558 | 36,064,897 | 2 | 36,063,685 | 36,064,897 | 1,213 |
| <i>HNF1B</i> | rs10908278 | 17 | 36,099,952 | T | A | 48.1 | 48.1 | 6.4x10 ⁻³⁶ | 4 | 3,726 | 36,099,840 | 36,103,565 | 4 | 36,099,840 | 36,103,565 | 3,726 |
| <i>MLX</i> | rs34855406 | 17 | 40,731,411 | C | G | 27.7 | 27.7 | 2.3x10 ⁻¹² | 42 | 258,595 | 40,566,240 | 40,824,834 | 36 | 40,610,565 | 40,962,509 | 351,945 |
| <i>TTL6</i> | rs35895680 | 17 | 47,060,322 | C | A | 67.8 | 32.2 | 2.5x10 ⁻¹⁵ | 118 | 177,089 | 46,917,953 | 47,095,041 | 117 | 46,894,377 | 47,095,041 | 200,665 |
| <i>KIF2B</i> | rs569511541 | 17 | 52,140,805 | G | A | 0.0200 | 0.0200 | 1.5x10 ⁻⁸ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>ACE</i> | rs2727301 | 17 | 61,965,043 | T | C | 75.4 | 24.6 | 1.3x10 ⁻⁶ | 578 | 759,783 | 61,703,320 | 62,463,102 | 315 | 61,703,320 | 62,461,656 | 758,337 |
| <i>ACE</i> | rs60276348 | 17 | 62,203,304 | T | C | 14.0 | 14.0 | 2.6x10 ⁻⁸ | 27 | 95,013 | 62,114,233 | 62,209,245 | 27 | 62,117,733 | 62,209,245 | 91,513 |
| <i>BPTF</i> | rs11657492 | 17 | 65,648,427 | G | T | 10.0 | 10.0 | 5.6x10 ⁻⁸ | 22 | 25,359 | 65,634,625 | 65,659,983 | 22 | 65,634,625 | 65,659,983 | 25,359 |
| <i>BPTF</i> | rs58308082 | 17 | 65,820,153 | C | G | 0.0800 | 0.0800 | 6.6x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>BPTF</i> | rs61676547 | 17 | 65,892,507 | C | G | 19.2 | 19.2 | 2.9x10 ⁻¹¹ | 143 | 169,153 | 65,825,248 | 65,994,400 | 142 | 65,825,248 | 66,098,154 | 272,907 |
| <i>LAMA1</i> | rs7240767 | 18 | 7,070,642 | C | T | 37.6 | 37.6 | 1.6x10 ⁻⁸ | 24 | 19,825 | 7,063,196 | 7,083,020 | 29 | 7,062,395 | 7,083,020 | 20,626 |
| <i>COMM D9</i> | rs62080313 | 18 | 36,278,709 | C | T | 12.3 | 12.3 | 1.0x10 ⁻⁸ | 56 | 200,885 | 36,158,210 | 36,359,094 | 60 | 36,158,210 | 36,359,094 | 200,885 |
| <i>TCF4</i> | rs76197067 | 18 | 52,604,955 | G | A | 0.0500 | 0.0500 | 8.1x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>TCF4</i> | rs72926932 | 18 | 53,050,646 | C | A | 8.39 | 8.39 | 1.0x10 ⁻¹⁴ | 6 | 37,339 | 53,050,646 | 53,087,984 | 6 | 53,050,646 | 53,087,984 | 37,339 |
| <i>TCF4</i> | rs28719468 | 18 | 53,452,144 | C | T | 15.9 | 15.9 | 1.9x10 ⁻⁶ | 314 | 594,636 | 52,953,630 | 53,548,265 | 202 | 52,979,074 | 53,498,376 | 519,303 |
| <i>WDR7</i> | rs17684074 | 18 | 54,675,384 | G | C | 74.0 | 26.0 | 2.9x10 ⁻⁸ | 34 | 351,130 | 54,361,464 | 54,712,593 | 70 | 54,281,790 | 55,105,267 | 823,478 |
| <i>GRP</i> | rs9957145 | 18 | 56,876,228 | G | A | 82.9 | 17.1 | 8.1x10 ⁻⁹ | 17 | 5,510 | 56,876,228 | 56,881,737 | 18 | 56,876,228 | 56,883,319 | 7,092 |
| <i>MC4R</i> | rs523288 | 18 | 57,848,369 | T | A | 23.8 | 23.8 | 7.6x10 ⁻¹³ | 74 | 181,548 | 57,732,418 | 57,913,965 | 74 | 57,732,418 | 57,913,965 | 181,548 |
| <i>MC4R</i> | rs74452128 | 18 | 58,056,566 | C | A | 97.6 | 2.37 | 1.0x10 ⁻⁹ | 44 | 63,213 | 58,024,670 | 58,087,882 | 42 | 58,024,670 | 58,087,882 | 63,213 |
| <i>BCL2A</i> | rs10469140 | 18 | 60,668,270 | G | A | 48.5 | 48.5 | 6.6x10 ⁻⁶ | 2,545 | 999,354 | 60,345,952 | 61,345,305 | 1,780 | 60,345,952 | 61,167,061 | 821,110 |
| <i>BCL2A</i> | rs12454712 | 18 | 60,845,884 | T | C | 61.4 | 38.6 | 4.6x10 ⁻¹³ | 1 | 1 | 60,845,884 | 60,845,884 | 1 | 60,845,884 | 60,845,884 | 1 |
| <i>UHRF1</i> | rs7249758 | 19 | 4,948,862 | A | G | 20.4 | 20.4 | 3.4x10 ⁻⁹ | 28 | 154,224 | 4,920,414 | 5,074,637 | 33 | 4,920,414 | 5,086,808 | 166,395 |
| <i>PTPRS</i> | rs116953931 | 19 | 5,224,998 | A | G | 3.71 | 3.71 | 6.4x10 ⁻⁶ | 2,100 | 722,786 | 4,725,161 | 5,447,946 | 1,482 | 4,725,272 | 5,447,946 | 722,675 |
| <i>INSR</i> | rs75253922 | 19 | 7,240,848 | C | T | 19.1 | 19.1 | 2.7x10 ⁻⁸ | 34 | 32,352 | 7,216,908 | 7,249,259 | 35 | 7,216,908 | 7,249,259 | 32,352 |
| <i>MAP2K7</i> | rs4804833 | 19 | 7,970,635 | A | G | 39.0 | 39.0 | 7.7x10 ⁻¹³ | 5 | 10,263 | 7,968,168 | 7,978,430 | 4 | 7,968,168 | 7,976,529 | 8,362 |
| <i>FARSA</i> | rs755734872 | 19 | 12,938,471 | T | C | 0.0500 | 0.0500 | 8.3x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>FARSA</i> | rs3111316 | 19 | 13,038,415 | A | G | 58.9 | 41.2 | 6.3x10 ⁻¹³ | 39 | 95,738 | 12,999,458 | 13,095,195 | 34 | 13,000,247 | 13,095,195 | 94,949 |
| <i>TM6SF2</i> | rs8107974 | 19 | 19,388,500 | T | A | 7.69 | 7.69 | 3.3x10 ⁻¹⁵ | 6 | 80,993 | 19,379,549 | 19,460,541 | 6 | 19,379,549 | 19,460,541 | 80,993 |
| <i>TM6SF2</i> | rs188247550 | 19 | 19,396,616 | T | C | 1.95 | 1.95 | 5.2x10 ⁻⁶ | 1,571 | 989,557 | 18,889,077 | 19,887,633 | 646 | 18,897,440 | 19,888,375 | 990,936 |
| <i>PEPD</i> | rs10406327 | 19 | 33,890,838 | C | G | 52.3 | 47.7 | 3.8x10 ⁻⁸ | 117 | 238,226 | 33,786,208 | 34,024,433 | 93 | 33,792,631 | 34,015,935 | 223,305 |
| <i>TOMM40/APOE</i> | rs745903616 | 19 | 44,938,870 | A | G | 0.130 | 0.130 | 8.3x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>TOMM40/APOE</i> | rs429358 | 19 | 45,411,941 | T | C | 84.6 | 15.4 | 2.6x10 ⁻¹⁸ | 4 | 30,693 | 45,392,254 | 45,422,946 | 2 | 45,392,254 | 45,411,941 | 19,688 |
| <i>GIPR</i> | rs10406431 | 19 | 46,157,019 | A | G | 56.3 | 43.8 | 9.6x10 ⁻¹⁴ | 8 | 9,711 | 46,147,527 | 46,157,237 | 7 | 46,147,527 | 46,157,237 | 9,711 |
| <i>GIPR</i> | rs2238689 | 19 | 46,178,661 | C | T | 41.8 | 41.8 | 5.4x10 ⁻⁹ | 28 | 20,369 | 46,158,293 | 46,178,661 | 30 | 46,149,618 | 46,178,661 | 29,044 |
| <i>GIPR</i> | rs533172266 | 19 | 46,351,837 | T | C | 0.0600 | 0.0600 | 3.7x10 ⁻⁶ | N/A | N/A | N/A | N/A | N/A | N/A | N/A | N/A |
| <i>ZC3H4</i> | rs3810291 | 19 | 47,569,003 | A | G | 67.3 | 32.7 | 8.9x10 ⁻¹² | 31 | 56,674 | 47,564,643 | 47,621,316 | 29 | 47,564,968 | 47,621,316 | 56,349 |
| <i>NKX2.2</i> | rs13041756 | 20 | 21,466,795 | C | T | 10.7 | 10.7 | 1.4x10 ⁻⁸ | 19 | 73,584 | 21,451,139 | 21,524,722 | 22 | 21,451,139 | 21,524,722 | 73,584 |
| <i>RALY</i> | rs2268078 | 20 | 32,596,704 | A | G | 65.7 | 34.3 | 2.3x10 ⁻¹⁰ | 116 | 348,108 | 32,381,337 | 32,729,444 | 105 | 32,514,092 | 32,727,430 | 213,339 |
| <i>HNF4A</i> | rs76811102 | 20 | 42,905,415 | T | C | 4.24 | 4.24 | 1.1x10 ⁻⁷ | 135 | 839,354 | 42,552,496 | 43,391,849 | 322 | 42,552,496 | 43,391,849 | 839,354 |
| <i>HNF4A</i> | rs4810426 | 20 | 43,001,721 | T | C | 10.6 | 10.6 | 3.1x10 ⁻¹⁷ | 5 | 8,311 | 42,994,812 | 43,003,122 | 5 | 42,994,812 | 43,003,122 | 8,311 |
| <i>HNF4A</i> | rs191830490 | 20 | 43,023,355 | G | A | 99.4 | 0.590 | 2.2x10 ⁻⁶ | 1,779 | 980,737 | 42,542,425 | 43,523,161 | 31 | 42,552,496 | 43,419,867 | 867,372 |
| <i>HNF4A</i> | rs1800961 | 20 | 43,042,364 | T | C | 3.53 | 3.53 | 2.3x10 ⁻²² | 1 | 1 | 43,042,364 | 43,042,364 | 1 | 43,042,364 | 43,042,364 | 1 |
| <i>HNF4A</i> | rs11696357 | 20 | 43,233,649 | A | G | 93.4 | 6.60 | 9.9x10 ⁻⁶ | 1,995 | 807,927 | 42,733,946 | 43,541,872 | 1,572 | 42,733,946 | 43,541,872 | 807,927 |
| <i>EYA2</i> | rs560716466 | 20 | 45,317,678 | A | G | 0.310 | 0.310 | 9.8x10 ⁻⁶ | 2,135 | 718,554 | 45,099,069 | 45,817,622 | 558 | 45,101,445 | 45,816,733 | 715,289 |
| <i>EYA2</i> | rs6063048 | 20 | 45,598,564 | G | A | 72.5 | 27.5 | 2.2x10 ⁻¹¹ | 16 | 25,077 | 45,581,777 | 45,606,853 | 16 | 45,557,469 | 45,606,853 | 49,385 |
| <i>CEBPB</i> | rs11699802 | 20 | 48,832,135 | C | T | 53.6 | 46.4 | 1.8x10 ⁻¹¹ | 11 | 9,093 | 48,826,880 | 48,835,972 | 12 | 48,808,011 | 48,835,972 | 27,962 |
| <i>TSHZ2</i> | rs34454109 | 20 | 51,223,594 | A | T | 77.1 | 22.9 | 7.1x10 ⁻⁹ | 267 | 436,047 | 50,828,285 | 51,264,331 | 249 | 50,828,285 | 51,620,301 | 792,017 |
| <i>GNAS</i> | rs6070625 | 20 | 57,394,628 | G | C | 51.7 | 48.3 | 5.3x10 ⁻¹⁴ | 19 | 31,433 | 57,386,639 | 57,418,071 | 19 | 57,386,639 | 57,418,071 | 31,433 |
| <i>GNAS</i> | rs862016 | 20 | 57,551,099 | G | A | 7.83 | 7.83 | 1.1x10 ⁻⁷ | 20 | 29,678 | 57,550,273 | 57,579,950 | 23 | 57,236,717 | 57,579,950 | 343,234 |
| <i>ZBTB46</i> | rs6011155 | 20 | 62,450,664 | T | C | 63.0 | 37.0 | 6.3x10 ⁻⁶ | 1,618 | 724,304 | 62,193,402 | 62,917,705 | 972 | 62,194,128 | 62,917,047 | 722,920 |
| <i>TCEA2</i> | rs59944054 | 20 | 62,693,175 | A | G | 23.8 | 23.8 | 1.5x10 ⁻⁸ | 58 | 494,479 | 62,267,221 | 62,761,699 | 45 | 62,296,306 | 62,737,568 | 441,263 |
| <i>MTMR3/ASCC2</i> | rs6518681 | 22 | 30,609,554 | G | A | 91.4 | 8.64 | 1.1x10 ⁻¹² | 143 | 499,653 | 30,138,232 | 30,637,884 | 141 | 30,135,928 | 30,637,884 | 501,957 |
| <i>YWHAH</i> | rs117001013 | 22 | 32,348,841 | C | T | 91.2 | 8.83 | 1.7x10 ⁻⁸ | 315 | 215,643 | 32,344,403 | 32,560,045 | 299 | 32,344,403 | 32,555,730 | 211,328 |
| <i>EP300</i> | rs5758223 | 22 | 41,489,920 | A | G | 71.7 | 28.3 | 3.8x10 ⁻⁸ | 138 | 669,493 | 41,066,598 | 41,736,090 | 126 | 41,262,852 | 41,864,190 | 601,339 |
| <i>PNPLA3</i> | rs738408 | 22 | 44,324,730 | T | C | 22.6 | 22.6 | 1.4x10 ⁻¹⁰ | 52 | 70,725 | 44,324,727 | 44,395,451 | 29 | 44,324,727 | 44,395,451 | 70,725 |
| <i>PIM3</i> | rs1801645 | 22 | 50,356,850 | C | T | 27.5 | 27.5 | 1.5x10 ⁻⁸ | 50 | 109,631 | 50,351,977 | 50,461,607 | 48 | 50,351,977 | 50,460,687 | 108,711 |
| <i>PIM3</i> | rs112915006 | 22 | 50,604,696 | G | A | 5.12 | 5.12 | 9.8x10 ⁻⁷ | 1,140 | 748,200 | 50,107,943 | 50,856,142 | 862 | 50,107,943 | 50,843,852 | 735,910 |

RAF: risk allele frequency; MAF: minor allele frequency.

Supplementary Table 6 | Summary of comparison of HRC-based and 1000G-based 99% credible sets.

| Nearest gene | Index variant | 1000-G based 99% genetic credible set | | | | HRC-based 99% functional credible set | | | | Difference in interval (HRC-based - 1000G-based) | Difference in SNPs (HRC-based - 1000G-based) |
|----------------------|---------------|---------------------------------------|---------------|---------------------|--------------------|---------------------------------------|---------------|---------------------|--------------------|--|--|
| | | SNPs | Interval (bp) | Interval start (bp) | Interval stop (bp) | SNPs | Interval (bp) | Interval start (bp) | Interval stop (bp) | | |
| <i>MACF1</i> | rs3768321 | 251 | 536,277 | 39,551,488 | 40,087,764 | 72 | 441,918 | 39,618,556 | 40,060,473 | -94,359 | -179 |
| <i>FAF1</i> | rs58432198 | 1,903 | 970,055 | 50,809,304 | 51,779,358 | 233 | 588,366 | 50,920,161 | 51,508,526 | -381,689 | -1,670 |
| <i>NOTCH2</i> | rs1493694 | 73 | 134,811 | 120,436,751 | 120,571,561 | 48 | 116,417 | 120,437,718 | 120,554,134 | -18,394 | -25 |
| <i>PROX1</i> | rs340874 | 21 | 18,287 | 214,145,389 | 214,163,675 | 1 | 1 | 214,159,256 | 214,159,256 | -18,286 | -20 |
| <i>THADA</i> | rs80147536 | 162 | 295,211 | 43,469,615 | 43,764,825 | 155 | 236,769 | 43,528,057 | 43,764,825 | -58,442 | -7 |
| <i>BCL11A</i> | rs243024 | 26 | 60,316 | 60,533,913 | 60,594,228 | 8 | 7,084 | 60,579,624 | 60,586,707 | -53,232 | -18 |
| <i>RBMS1</i> | rs3772071 | 88 | 256,652 | 161,087,411 | 161,344,062 | 30 | 246,462 | 161,087,411 | 161,333,872 | -10,190 | -58 |
| <i>IRS1</i> | rs2972144 | 85 | 134,270 | 227,047,414 | 227,181,683 | 30 | 34,288 | 227,083,411 | 227,117,698 | -99,982 | -55 |
| <i>PPARG</i> | rs11709077 | 42 | 131,494 | 12,265,462 | 12,396,955 | 11 | 67,131 | 12,329,783 | 12,396,913 | -64,363 | -31 |
| <i>UBE2E2</i> | rs35352848 | 65 | 380,819 | 23,187,407 | 23,568,225 | 5 | 2,291 | 23,454,790 | 23,457,080 | -378,528 | -60 |
| <i>ADAMTS9</i> | rs9860730 | 14 | 30,677 | 64,699,445 | 64,730,121 | 23 | 29,697 | 64,700,425 | 64,730,121 | -980 | 9 |
| <i>ADCY5</i> | rs11708067 | 11 | 39,118 | 123,054,770 | 123,093,887 | 4 | 4,649 | 123,065,778 | 123,070,426 | -34,469 | -7 |
| <i>IGF2BP2</i> | rs6780171 | 30 | 32,656 | 185,497,635 | 185,530,290 | 41 | 39,163 | 185,495,320 | 185,534,482 | 6,507 | 11 |
| <i>ST6GAL1</i> | rs3887925 | 247 | 75,872 | 186,606,741 | 186,682,612 | 2 | 8,494 | 186,665,645 | 186,674,138 | -67,378 | -245 |
| <i>LPP</i> | rs4686471 | 324 | 79,292 | 187,677,977 | 187,757,268 | 3 | 1,320 | 187,740,523 | 187,741,842 | -77,972 | -321 |
| <i>MAEA</i> | rs56337234 | 2,719 | 1,754,840 | 510,985 | 2,265,824 | 5 | 2,920 | 1,781,686 | 1,784,605 | -1,751,920 | -2,714 |
| <i>WFS1</i> | rs10937721 | 87 | 57,401 | 6,263,996 | 6,321,396 | 108 | 41,845 | 6,280,449 | 6,322,293 | -15,556 | 21 |
| <i>TMEM154</i> | rs7669833 | 523 | 279,179 | 153,241,587 | 153,520,765 | 36 | 24,961 | 153,495,515 | 153,520,475 | -254,218 | -487 |
| <i>ACSL1</i> | rs58730668 | 67 | 48,312 | 185,684,247 | 185,732,558 | 46 | 17,143 | 185,713,608 | 185,730,750 | -31,169 | -21 |
| <i>ARL15</i> | rs702634 | 233 | 151,621 | 53,243,352 | 53,394,972 | 3 | 4,882 | 53,271,420 | 53,276,301 | -146,739 | -230 |
| <i>ANKRD55</i> | rs465002 | 20 | 21,450 | 55,794,632 | 55,816,081 | 13 | 11,122 | 55,799,184 | 55,810,305 | -10,328 | -7 |
| <i>ANKRD55</i> | rs9687832 | 10 | 5,520 | 55,856,375 | 55,861,894 | 8 | 5,520 | 55,856,375 | 55,861,894 | 0 | -2 |
| <i>ZBED3</i> | rs4457053 | 39 | 49,120 | 76,424,949 | 76,474,068 | 5 | 10,056 | 76,424,949 | 76,435,004 | -39,064 | -34 |
| <i>PAM</i> | rs115505614 | 2,842 | 1,202,395 | 101,624,383 | 102,826,777 | 8 | 285,050 | 102,301,358 | 102,586,407 | -917,345 | -2,834 |
| <i>RREB1</i> | rs9379084 | 315 | 160,806 | 7,166,829 | 7,327,634 | 1 | 1 | 7,231,843 | 7,231,843 | -160,805 | -314 |
| <i>CDKAL1</i> | rs7756992 | 6 | 30,073 | 20,673,880 | 20,703,952 | 6 | 13,117 | 20,673,880 | 20,686,996 | -16,956 | 0 |
| <i>CENPW</i> | rs11759026 | 125 | 412,763 | 126,657,472 | 127,070,234 | 2 | 172,416 | 126,792,095 | 126,964,510 | -240,347 | -123 |
| <i>SLC35D3</i> | rs9496424 | 40 | 18,391 | 137,287,076 | 137,305,466 | 34 | 17,765 | 137,287,702 | 137,305,466 | -626 | -6 |
| <i>DGKB</i> | rs17168486 | 7 | 61,806 | 14,862,102 | 14,923,907 | 6 | 62,871 | 14,859,137 | 14,922,007 | 1,065 | -1 |
| <i>DGKB</i> | rs10228066 | 59 | 18,333 | 15,047,280 | 15,065,612 | 17 | 2,774 | 15,062,694 | 15,065,467 | -15,559 | -42 |
| <i>JAZF1</i> | rs1708302 | 11 | 27,366 | 28,172,732 | 28,200,097 | 6 | 7,818 | 28,192,280 | 28,200,097 | -19,548 | -5 |
| <i>GCK</i> | rs878521 | 103 | 44,108 | 44,222,077 | 44,266,184 | 2 | 23,866 | 44,231,778 | 44,255,643 | -20,242 | -101 |
| <i>KLF14</i> | rs1562396 | 71 | 45,257 | 130,422,934 | 130,468,190 | 26 | 40,188 | 130,427,388 | 130,467,575 | -5,069 | -45 |
| <i>AOC1</i> | rs62492368 | 220 | 168,226 | 156,920,930 | 157,089,155 | 42 | 37,872 | 150,540,840 | 150,542,711 | -130,354 | -178 |
| <i>ANK1</i> | rs13262861 | 44 | 111,162 | 41,426,157 | 41,537,318 | 9 | 16,323 | 41,508,577 | 41,524,899 | -94,839 | -35 |
| <i>TP53INP1</i> | rs10097617 | 988 | 384,925 | 95,734,719 | 96,119,643 | 14 | 56,937 | 95,911,477 | 95,968,413 | -327,988 | -974 |
| <i>SLC30A8</i> | rs3802177 | 6 | 33,133 | 118,184,783 | 118,217,915 | 5 | 19,238 | 118,184,783 | 118,204,020 | -13,895 | -1 |
| <i>GLIS3</i> | rs10974438 | 359 | 68,430 | 4,243,162 | 4,311,591 | 2 | 6,662 | 4,291,928 | 4,298,589 | -61,768 | -357 |
| <i>CDKN2A/B</i> | rs1412830 | 105 | 128,333 | 21,997,015 | 22,125,347 | 66 | 155,129 | 21,966,221 | 22,121,349 | 26,796 | -39 |
| <i>CDKN2A/B</i> | rs10811660 | 6 | 1,556 | 22,132,698 | 22,134,253 | 5 | 1,556 | 22,132,698 | 22,134,253 | 0 | -1 |
| <i>CDKN2A/B</i> | rs10757283 | 5 | 1,007 | 22,133,645 | 22,134,651 | 5 | 1,007 | 22,133,645 | 22,134,651 | 0 | 0 |
| <i>TLE4</i> | rs17791513 | 192 | 130,813 | 81,881,363 | 82,012,175 | 22 | 60,142 | 81,891,987 | 81,952,128 | -70,671 | -170 |
| <i>TLE1</i> | rs2796441 | 8 | 18,348 | 84,293,453 | 84,311,800 | 2 | 2,853 | 84,308,948 | 84,311,800 | -15,495 | -6 |
| <i>ABO</i> | rs505922 | 40 | 17,936 | 136,137,065 | 136,155,000 | 17 | 15,736 | 136,139,265 | 136,155,000 | -2,200 | -23 |
| <i>CDC123/CAMK1D</i> | rs11257655 | 15 | 27,296 | 12,302,357 | 12,329,652 | 1 | 1 | 12,307,894 | 12,307,894 | -27,295 | -14 |
| <i>ZMIZ1</i> | rs703972 | 15 | 13,373 | 80,941,417 | 80,954,789 | 10 | 10,832 | 80,942,620 | 80,953,451 | -2,541 | -5 |
| <i>HHEX/IDE</i> | rs10882101 | 24 | 56,253 | 94,429,511 | 94,485,763 | 10 | 21,703 | 94,444,793 | 94,466,495 | -34,550 | -14 |
| <i>TCF7L2</i> | rs7903146 | 3 | 4,279 | 114,754,071 | 114,758,349 | 3 | 4,279 | 114,517,569 | 114,995,846 | 0 | 0 |
| <i>PLEKHA1</i> | rs2280141 | 25 | 59,193 | 124,139,393 | 124,198,585 | 25 | 59,193 | 21,935,597 | 21,972,286 | 0 | 0 |
| <i>KCNQ1</i> | rs231361 | 3 | 197 | 2,858,440 | 2,858,636 | 1 | 1 | 2,840,424 | 2,857,897 | -196 | -2 |
| <i>KCNJ11</i> | rs5213 | 22 | 53,506 | 17,368,381 | 17,421,886 | 6 | 10,074 | 17,408,404 | 17,418,477 | -43,432 | -16 |
| <i>HSD17B12</i> | rs1061810 | 74 | 126,013 | 43,752,522 | 43,878,534 | 67 | 125,964 | 43,752,522 | 43,878,485 | -49 | -7 |
| <i>MAP3K11</i> | rs1783541 | 83 | 314,376 | 65,217,626 | 65,532,001 | 12 | 79,725 | 65,257,527 | 65,337,251 | -234,651 | -71 |
| <i>CENTD2/ARAP1</i> | rs77464186 | 26 | 431,950 | 72,419,514 | 72,851,463 | 9 | 33,857 | 72,429,579 | 72,463,435 | -398,093 | -17 |
| <i>MNTR1B</i> | rs10830963 | 56 | 57,592 | 92,667,730 | 92,725,321 | 1 | 1 | 92,708,710 | 92,708,710 | -57,591 | -55 |
| <i>CCND2</i> | rs11063028 | 20 | 28,171 | 4,288,001 | 4,316,171 | 13 | 12,809 | 4,288,001 | 4,300,809 | -15,362 | -7 |
| <i>CCND2</i> | rs4238013 | 3 | 13,181 | 4,362,909 | 4,376,089 | 3 | 13,181 | 4,362,909 | 4,376,089 | 0 | 0 |
| <i>CCND2</i> | rs3217792 | 102 | 200,521 | 4,158,184 | 4,358,704 | 1 | 1 | 4,384,696 | 4,384,696 | -200,520 | -101 |
| <i>KLHDC5</i> | rs10842994 | 61 | 46,006 | 27,919,145 | 27,965,150 | 26 | 27,680 | 27,937,471 | 27,965,150 | -18,326 | -35 |
| <i>HMGAA2</i> | rs2258238 | 107 | 217,850 | 66,165,471 | 66,383,320 | 47 | 82,788 | 66,178,137 | 66,260,924 | -135,062 | -60 |
| <i>TSPAN8/LGR5</i> | rs1796330 | 536 | 252,482 | 71,413,705 | 71,666,186 | 119 | 131,397 | 71,408,690 | 71,540,086 | -121,085 | -417 |
| <i>HNF1A</i> | rs56348580 | 16 | 61,764 | 121,410,168 | 121,471,931 | 5 | 27,423 | 121,429,194 | 121,456,616 | -34,341 | -11 |
| <i>MPHOSPH9</i> | rs4148865 | 550 | 504,880 | 123,408,818 | 123,913,697 | 208 | 612,745 | 123,296,204 | 123,908,948 | 107,865 | -342 |
| <i>SPRY2</i> | rs1359790 | 9 | 13,340 | 80,705,315 | 80,718,654 | 10 | 11,842 | 80,705,315 | 80,717,156 | -1,498 | 1 |

| | | | | | | | | | | | |
|--------------------|-------------|-------|---------|------------|------------|-----|---------|------------|------------|----------|--------|
| <i>NRXN3</i> | rs17836088 | 537 | 387,539 | 79,689,186 | 80,076,724 | 16 | 54,707 | 79,890,456 | 79,945,162 | -332,832 | -521 |
| <i>C2CD4A/B</i> | rs8037894 | 46 | 263,439 | 62,117,975 | 62,381,413 | 15 | 15,939 | 62,383,155 | 62,399,093 | -247,500 | -31 |
| <i>HMG20A</i> | rs1005752 | 49 | 346,309 | 77,486,008 | 77,832,316 | 24 | 121,044 | 77,711,719 | 77,832,762 | -225,265 | -25 |
| <i>PRC1</i> | rs12910825 | 75 | 124,709 | 91,444,573 | 91,569,281 | 25 | 30,176 | 91,506,452 | 91,536,627 | -94,533 | -50 |
| <i>FTO</i> | rs1421085 | 64 | 43,402 | 53,799,507 | 53,842,908 | 6 | 8,170 | 53,800,954 | 53,809,123 | -35,232 | -58 |
| <i>BCAR1</i> | rs72802342 | 13 | 51,613 | 75,234,872 | 75,286,484 | 9 | 17,456 | 75,234,872 | 75,252,327 | -34,157 | -4 |
| <i>CMIP</i> | rs2925979 | 3 | 10,517 | 81,524,274 | 81,534,790 | 2 | 1,002 | 81,533,789 | 81,534,790 | -9,515 | -1 |
| <i>ZZEF1</i> | rs1377807 | 307 | 406,766 | 3,880,546 | 4,287,311 | 66 | 281,716 | 3,884,026 | 4,165,741 | -125,050 | -241 |
| <i>GLP2R</i> | rs7222481 | 87 | 41,165 | 9,765,514 | 9,806,678 | 79 | 40,195 | 9,766,484 | 9,806,678 | -970 | -8 |
| <i>HNF1B</i> | rs10908278 | 13 | 6,347 | 36,097,775 | 36,104,121 | 4 | 3,726 | 36,099,840 | 36,103,565 | -2,621 | -9 |
| <i>TTL6</i> | rs35895680 | 172 | 164,324 | 46,883,545 | 47,047,868 | 118 | 177,089 | 46,917,953 | 47,095,041 | 12,765 | -54 |
| <i>LAMA1</i> | rs7240767 | 28 | 20,514 | 7,062,395 | 7,082,908 | 24 | 19,825 | 7,063,196 | 7,083,020 | -689 | -4 |
| <i>MC4R</i> | rs523288 | 168 | 183,018 | 57,730,948 | 57,913,965 | 74 | 181,548 | 57,732,418 | 57,913,965 | -1,470 | -94 |
| <i>TM6SF2</i> | rs8107974 | 1,358 | 540,002 | 19,260,466 | 19,800,467 | 6 | 80,993 | 19,379,549 | 19,460,541 | -459,009 | -1,352 |
| <i>TOMM40/APOE</i> | rs745903616 | 4 | 11,006 | 45,411,941 | 45,422,946 | 4 | 30,693 | 45,392,254 | 45,422,946 | 19,687 | 0 |
| <i>GIPR</i> | rs10406431 | 30 | 30,276 | 46,148,386 | 46,178,661 | 8 | 9,711 | 46,147,527 | 46,157,237 | -20,565 | -22 |
| <i>HNF4A</i> | rs4810426 | 136 | 169,470 | 42,847,479 | 43,016,948 | 5 | 8,311 | 42,994,812 | 43,003,122 | -161,159 | -131 |
| <i>HNF4A</i> | rs1800961 | 1 | 1 | 43,042,364 | 43,042,364 | 1 | 1 | 43,042,364 | 43,042,364 | 0 | 0 |
| <i>MTMR3/ASCC2</i> | rs6518681 | 153 | 473,627 | 30,135,928 | 30,609,554 | 143 | 499,653 | 30,138,232 | 30,637,884 | 26,026 | -10 |

Supplementary Table 7 | Coding variants with high PPAs.

| Nearest gene | Index variant | Chromosome | Position (Build 37 bp) | Risk allele | Other allele | RAF (%) | MAF (%) | Posterior probability | | Missense variant with posterior probability >50% | |
|--------------------|---------------|------------|---------------------------|-------------|--------------|---------|---------|--------------------------|-----------------------------|--|--------------------|
| | | | | | | | | 99% genetic credible set | 99% functional credible set | Gene | Variant annotation |
| <i>PATJ</i> | rs12140153 | 1 | 62579891 | G | T | 90.5 | 9.49 | 0.996 | 1.000 | <i>PATJ</i> | p.Gly157Val |
| <i>GCKR</i> | rs1260326 | 2 | 27730940 | C | T | 60.7 | 39.3 | 0.897 | 0.899 | <i>GCKR</i> | p.Pro446Leu |
| <i>SCD5</i> | rs12642790 | 4 | 83578271 | A | G | 33.8 | 33.8 | 0.147 | 0.673 | <i>SCD5</i> | p.Glu197Gln |
| <i>ANKH</i> | rs78408340 | 5 | 14751305 | C | T | 99.4 | 0.620 | 0.770 | 0.963 | <i>ANKH</i> | p.Arg187Gln |
| <i>ANKH</i> | rs78408340 | 5 | 14751305 | C | T | 99.4 | 0.620 | 1.000 | 1.000 | <i>PAM</i> | p.Ser539Trp |
| <i>MRPS30</i> | rs62368490 | 5 | 44534364 | T | C | 3.13 | 3.13 | 0.010 | 0.522 | <i>MRPS30</i> | p.Glu128Gln |
| <i>POC5</i> | rs2307111 | 5 | 75003678 | T | C | 60.5 | 39.5 | 0.628 | 0.873 | <i>POC5</i> | p.His36Arg |
| <i>RREB1</i> | rs9379084 | 6 | 7231843 | G | A | 88.7 | 11.3 | 0.993 | 0.998 | <i>RREB1</i> | p.Asp1171Asn |
| <i>SLC30A8</i> | rs3802177 | 8 | 118185025 | G | A | 68.5 | 31.5 | 0.351 | 0.961 | <i>SLC30A8</i> | p.Arg276Trp |
| <i>NEUROG3</i> | rs41277236 | 10 | 71332301 | T | C | 4.31 | 4.31 | 0.737 | 0.987 | <i>NEUROG3</i> | p.Gly167Arg |
| <i>QSER1</i> | rs145678014 | 11 | 32927778 | G | T | 95.7 | 4.33 | 0.211 | 0.846 | <i>QSER1</i> | p.Arg1101Cys |
| <i>CDKN1B</i> | rs2066827 | 12 | 12871099 | G | T | 23.5 | 23.5 | 0.984 | 0.999 | <i>CDKN1B</i> | p.Val109Gly |
| <i>WSCD2</i> | rs1426371 | 12 | 108629780 | G | A | 73.9 | 26.1 | 0.440 | 0.753 | <i>WSCD2</i> | p.Thr113Ile |
| <i>HNF1A</i> | rs1800574 | 12 | 121416864 | T | C | 2.96 | 2.96 | 0.985 | 0.996 | <i>HNF1A</i> | p.Ala146Val |
| <i>HNF1A</i> | rs56348580 | 12 | 121432117 | G | C | 68.9 | 31.1 | 0.474 | 0.906 | <i>HNF1A</i> | p.Gly226Ala |
| <i>IRS2</i> | rs4771648 | 13 | 110431626 | G | A | 66.9 | 33.2 | 0.173 | 0.645 | <i>IRS2</i> | p.Gly1057Asp |
| <i>FAM57B</i> | rs199795270 | 16 | 30419384 | C | G | 0.650 | 0.650 | 0.661 | 0.993 | <i>ZNF771</i> | p.Glu4Gln |
| <i>TOMM40/APOE</i> | rs429358 | 19 | 45411941 | T | C | 84.6 | 15.4 | 0.896 | 0.973 | <i>APOE</i> | p.Cys130Arg |
| <i>HNF4A</i> | rs1800961 | 20 | 43042364 | T | C | 3.53 | 3.53 | 1.000 | 1.000 | <i>HNF4A</i> | p.Thr139Ile |

RAF: risk allele frequency; MAF: minor allele frequency.

Supplementary Table 8 | Non-coding credible set variants with >80% posterior probability of association.

| Locus | Index variant | Variant with highest PPA in 99% credible set | Chr | Pos | MAF (%) | PPA | | Chromatin states | | | | |
|----------------------|---------------|--|-----|-------------|---------|---------|------------|--------------------------------|-----------------|-----------------|-----------------|-----------------|
| | | | | | | Genetic | Functional | Islets | Islets | Adipose | Skeletal Muscle | Liver |
| <i>PROX1</i> | rs340874 | rs340874 | 1 | 214,159,256 | 44.5 | 1.000 | 1.000 | Active promoter | Active promoter | | | Active promoter |
| <i>ABCB10</i> | rs348330 | rs348330 | 1 | 229,672,955 | 36.1 | 1.000 | 1.000 | Strong enhancer | Active enhancer | | | |
| <i>ANKH</i> | rs76549217 | rs76549217 | 5 | 14,768,766 | 3.0 | 1.000 | 1.000 | Strong enhancer | Active enhancer | | Active enhancer | |
| <i>CDC123/CAMK1D</i> | rs11257655 | rs11257655 | 10 | 12,307,894 | 21.8 | 1.000 | 1.000 | Strong enhancer | Active enhancer | Weak enhancer | | Active enhancer |
| <i>KCNQ1</i> | rs231361 | rs231361 | 11 | 2,691,500 | 25.6 | 1.000 | 1.000 | Strong enhancer (No ATAC/Meth) | | | | |
| <i>MTRN1B</i> | rs10830963 | rs10830963 | 11 | 92,708,710 | 27.7 | 1.000 | 1.000 | Strong enhancer (No ATAC/Meth) | | | | |
| <i>CCND2</i> | rs76895963 | rs76895963 | 12 | 4,384,844 | 2.0 | 1.000 | 1.000 | Active promoter | Active promoter | | | |
| <i>BCL2A</i> | rs12454712 | rs12454712 | 18 | 60,845,884 | 38.6 | 1.000 | 1.000 | Weak enhancer (Methylation) | | | | |
| <i>CASC11</i> | rs17772814 | rs17772814 | 8 | 128,711,742 | 8.5 | 0.999 | 1.000 | Weak enhancer (Methylation) | | | | |
| <i>KCNQ1</i> | rs2237895 | rs2237895 | 11 | 2,857,194 | 42.6 | 0.999 | 0.998 | Strong enhancer (No ATAC/Meth) | | | | |
| <i>CCND2</i> | rs3217792 | rs3217792 | 12 | 4,384,696 | 8.7 | 0.999 | 0.998 | Active promoter | Active promoter | Active promoter | | |
| <i>WDR72</i> | rs528350911 | rs528350911 | 15 | 53,747,228 | 0.7 | 0.995 | 0.994 | | | | | Active enhancer |
| <i>FAM49A</i> | rs11680058 | rs11680058 | 2 | 16,574,669 | 13.7 | 0.994 | 1.000 | Strong enhancer | Active enhancer | | | |
| <i>LRFN2</i> | rs34298980 | rs34298980 | 6 | 40,409,243 | 49.7 | 0.991 | 0.999 | Strong enhancer | | | | |
| <i>TCF7L2</i> | rs140242150 | rs140242150 | 10 | 114702962 | 0.5 | 0.987 | 0.985 | | | | | |
| <i>CCND2</i> | rs3217860 | rs3217860 | 12 | 4399050 | 25.8 | 0.987 | 0.990 | Gene enhancer | | Genic enhancer | | |
| <i>GLI2</i> | rs11688682 | rs11688682 | 2 | 121347612 | 27.2 | 0.986 | 0.987 | | | Active enhancer | | |
| <i>ST6GAL1</i> | rs3887925 | rs3887925 | 3 | 186665645 | 45.3 | 0.985 | 0.993 | Weak enhancer (No ATAC/Meth.) | Active enhancer | Active enhancer | Weak enhancer | |
| <i>CAMK2B</i> | rs878521 | rs878521 | 7 | 44255643 | 24.5 | 0.984 | 0.997 | Strong enhancer | | | Active enhancer | |
| <i>HNF1B</i> | rs10962 | rs10962 | 17 | 36046451 | 22.6 | 0.984 | 0.981 | | | | | |
| <i>ANK1</i> | rs13262861 | rs13262861 | 8 | 41508577 | 17.1 | 0.973 | 0.988 | Active promoter | Active enhancer | | | |
| <i>GLIS3</i> | rs10974438 | rs10974438 | 9 | 4291928 | 35.7 | 0.973 | 0.885 | Strong enhancer (No ATAC/Meth) | Active enhancer | | | |
| <i>PTGFRN</i> | rs1127215 | rs1127215 | 1 | 117532790 | 41.6 | 0.972 | 0.874 | | | Active enhancer | Weak enhancer | |
| <i>HNF4A</i> | rs76811102 | rs76811102 | 20 | 42905415 | 4.2 | 0.965 | 0.943 | | | | | |
| <i>CRHR2</i> | rs917195 | rs917195 | 7 | 30728452 | 23.0 | 0.962 | 0.987 | Weak enhancer | | | | |
| <i>CMIP</i> | rs2925979 | rs2925979 | 16 | 81534790 | 30.0 | 0.960 | 0.931 | Weak enhancer (No ATAC/Meth.) | Weak enhancer | Active enhancer | Active enhancer | |
| <i>CDKN2B</i> | rs76011118 | rs76011118 | 9 | 22133773 | 3.4 | 0.949 | 0.952 | Weak enhancer (No ATAC/Meth.) | | | | |
| <i>TSC22D2</i> | rs62271373 | rs62271373 | 3 | 150066540 | 5.5 | 0.948 | 0.846 | | | Active enhancer | Weak enhancer | |
| <i>HTT</i> | rs362307 | rs362307 | 4 | 3241845 | 7.7 | 0.946 | 0.985 | | Active enhancer | Genic enhancer | Genic enhancer | |
| <i>INS_IGF2</i> | rs555759341 | rs555759341 | 11 | 2151761 | 0.5 | 0.943 | 0.722 | | | | | |
| <i>IGF2BP2</i> | rs150111048 | rs150111048 | 3 | 185514421 | 23.9 | 0.941 | 0.938 | Weak enhancer | Active enhancer | | | |
| <i>CDKAL1</i> | rs7756992 | rs7756992 | 6 | 20679709 | 27.4 | 0.929 | 0.167 | | | | | |
| <i>UBAP2</i> | rs12001437 | rs12001437 | 9 | 34074476 | 37.2 | 0.917 | 0.985 | Strong enhancer | Active enhancer | Weak enhancer | Weak enhancer | Weak enhancer |
| <i>CLUAP1</i> | rs3751837 | rs3751837 | 16 | 3583173 | 22.0 | 0.903 | 0.940 | | | | | |
| <i>KCNQ1</i> | rs2283164 | rs2283164 | 11 | 2579163 | 5.3 | 0.890 | 0.545 | | | | | |
| <i>CPO</i> | rs149364428 | rs149364428 | 8 | 97737741 | 1.0 | 0.867 | 0.673 | | | | | |
| <i>CACNA2D3</i> | rs76263492 | rs76263492 | 3 | 54828827 | 4.5 | 0.861 | 0.852 | | | | | |
| <i>NEUROG3</i> | rs177045 | rs177045 | 10 | 71321279 | 31.6 | 0.855 | 0.845 | | | | | |
| <i>FAM63A</i> | rs145904381 | rs145904381 | 1 | 151017991 | 1.3 | 0.836 | 0.597 | | | | | |
| <i>ANKH</i> | rs17250977 | rs17250977 | 5 | 14753745 | 3.8 | 0.835 | 0.836 | Gene enhancer | | | Active enhancer | |
| <i>TLE1</i> | rs2796441 | rs2796441 | 9 | 84308948 | 40.8 | 0.824 | 0.946 | Weak enhancer (Methylation) | | Weak enhancer | | |
| <i>KCNQ1</i> | rs2237897 | rs2237897 | 11 | 2858546 | 4.6 | 0.813 | 0.812 | | Active enhancer | Weak enhancer | Weak enhancer | |
| <i>WDR11</i> | rs72631105 | rs72631105 | 10 | 122915345 | 19.4 | 0.806 | 0.879 | Weak enhancer (No ATAC/Meth.) | Weak enhancer | | Weak enhancer | |
| <i>TCF7L2</i> | rs7903146 | rs7903146 | 10 | 114758349 | 29.4 | 0.592 | 0.971 | Strong enhancer | Active enhancer | Active enhancer | | |
| <i>HNF4A</i> | rs191830490 | rs191830490 | 20 | 43023355 | 0.6 | 0.519 | 0.956 | Strong enhancer | Active enhancer | Weak enhancer | | Active promoter |
| <i>VEGFA</i> | rs11967262 | rs11967262 | 6 | 43760327 | 48.6 | 0.634 | 0.950 | Strong enhancer | | Active enhancer | | |
| <i>ANKH</i> | rs6885132 | rs6885132 | 5 | 14768092 | 9.6 | 0.682 | 0.946 | Strong enhancer | Active enhancer | | Active enhancer | |

| | | | | | | | | | | | | |
|---------------|-------------|-------------|----|-----------|------|-------|-------|-----------------|-----------------|-----------------|-----------------|-----------------|
| <i>RAI1</i> | rs4925109 | rs4925109 | 17 | 17661802 | 31.6 | 0.562 | 0.908 | Strong enhancer | Active enhancer | | Weak enhancer | |
| <i>RNF6</i> | rs34584161 | rs34584161 | 13 | 26776999 | 24.0 | 0.672 | 0.898 | Strong enhancer | Active enhancer | | Weak enhancer | |
| <i>DGKB</i> | rs17168486 | rs17168486 | 7 | 14898282 | 18.1 | 0.577 | 0.898 | Weak promoter | | | | |
| <i>ASCL2</i> | rs80102379 | rs80102379 | 11 | 2634177 | 1.8 | 0.676 | 0.892 | Weak enhancer | | | | |
| <i>EYA2</i> | rs560716466 | rs560716466 | 20 | 45317678 | 0.3 | 0.396 | 0.890 | Active promoter |
| <i>DLEU1</i> | rs963740 | rs963740 | 13 | 51096095 | 28.7 | 0.503 | 0.879 | Strong enhancer | Active enhancer | Weak enhancer | Active enhancer | Active enhancer |
| <i>ADCY5</i> | rs11708067 | rs11708067 | 3 | 123065778 | 22.8 | 0.789 | 0.871 | Weak enhancer | Active enhancer | | Weak enhancer | |
| <i>CDKN2B</i> | rs10757283 | rs10757283 | 9 | 22134172 | 43.0 | 0.498 | 0.844 | Weak enhancer | | | | |
| <i>TBCE</i> | rs291367 | rs291367 | 1 | 235690800 | 36.8 | 0.240 | 0.842 | Weak enhancer | | | | Active enhancer |
| <i>ZBED3</i> | rs4457053 | rs7732130 | 5 | 76435004 | 30.3 | 0.276 | 0.834 | Strong enhancer | Active enhancer | Active enhancer | | |
| <i>MACF1</i> | rs3768321 | rs3768321 | 1 | 40035928 | 20.0 | 0.657 | 0.810 | | | | | Genic enhancer |

PPA: posterior probability of association; MAF: minor allele frequency.

¹Thurner, M. et al. Integration of human pancreatic islet genomic data refines regulatory mechanisms at Type 2 Diabetes susceptibility loci. bioRxiv (2017).

²Varshney, A. et al. Genetic regulatory signatures underlying islet gene expression and type 2 diabetes. Proc Natl Acad Sci U S A 114, 2301-2306 (2017).

Supplementary Table 9 | TCF7L2 credible set variants.

| Signal | Index variant | SNPs in 99% credible set (with PPA≥1%) | | Variant | Chr | Pos | Posterior probability of association | | Chromatin state | | | | | Transcription factor binding site ³ | RAF (%) | MAF (%) | |
|-----------|---------------|--|------------|-------------|-----|-----------|--------------------------------------|------------|-----------------------|-----------------------|----------------------|------------------------------|--------------------|--|---------|---------|--|
| | | Genetic | Functional | | | | Genetic | Functional | Islets_1 ¹ | Islets_2 ² | Adipose ² | Skeletal Muscle ² | Liver ² | | | | |
| | | | | | | | | | | | | | | | | | |
| Primary | rs7903146 | 3 (3) | 2 (2) | rs7903146 | 10 | 114758349 | 0.592 | 0.9705 | 5 | 10 | 10 | 13 | 13 | FOXA2, MAFB, NKX6.1, NKX2.2 | 29.4 | 29.4 | |
| | | | | rs35198068 | 10 | 114754784 | 0.045 | 0 | 12 | 13 | 9 | 13 | 14 | | 29.53 | 29.53 | |
| | | | | rs34872471 | 10 | 114754071 | 0.363 | 0.0262 | 12 | 13 | 10 | 13 | 14 | | 29.51 | 29.51 | |
| Secondary | rs536643418 | MAF<0.25%; not fine-mapped | | rs536643418 | 10 | 114699835 | - | - | 12 | 16 | 16 | 16 | 13 | | 0.52 | 0.52 | |
| | | rs140242150 | 4 (1) | rs140242150 | 10 | 114702962 | 0.987 | 0.9854 | 12 | 16 | 16 | 15 | 13 | | 0.5 | 0.5 | |
| | | | | rs7918400 | 10 | 114703136 | 0.712 | 0.5219 | 12 | 16 | 16 | 15 | 13 | | 52.39 | 47.61 | |
| | | rs7918400 | 4 (3) | rs72826045 | 10 | 114702608 | 0.263 | 0.1933 | 12 | 16 | 16 | 15 | 13 | | 52.37 | 47.63 | |
| | | | | rs7094871 | 10 | 114712154 | 0.01 | 0.158 | 1 | 6 | 6 | 6 | 6 | | 54.04 | 45.96 | |
| | | | | rs10885396 | 10 | 114711755 | 0.006 | 0.0984 | 1 | 6 | 6 | 6 | 6 | FOXA2 | 54.03 | 45.97 | |
| | | | | rs7094463 | 10 | 114711983 | 0 | 0.023 | 1 | 6 | 6 | 6 | 6 | | 53.19 | 46.81 | |
| | | | | rs144155527 | 10 | 114737633 | 0.171 | 0.6778 | 5 | 10 | 13 | 13 | 14 | FOXA2 | 98.17 | 1.83 | |
| | | | | rs116761056 | 10 | 114737324 | 0.044 | 0.1753 | 5 | 10 | 13 | 13 | 14 | NKX2.2, NKX6.1, PDX1 | 97.99 | 2.01 | |
| | | rs184509201 | 9 (8) | rs114322470 | 10 | 114736670 | 0.146 | 0.0462 | 2 | 10 | 13 | 13 | 14 | | 98.16 | 1.84 | |
| | | | | rs184509201 | 10 | 114740337 | 0.186 | 0.0292 | 12 | 13 | 10 | 13 | 14 | | 98.18 | 1.82 | |
| | | | | rs192408510 | 10 | 114739988 | 0.163 | 0.0257 | 12 | 13 | 10 | 13 | 14 | | 98.17 | 1.83 | |
| | | | | rs145034729 | 10 | 114741673 | 0.094 | 0.0148 | 12 | 13 | 10 | 13 | 14 | | 98.17 | 1.83 | |
| | | | | rs149954646 | 10 | 114741507 | 0.089 | 0.014 | 12 | 13 | 10 | 13 | 14 | | 98.17 | 1.83 | |
| | | | | rs147932983 | 10 | 114743171 | 0.08 | 0.0127 | 12 | 13 | 13 | 13 | 14 | | 98.15 | 1.85 | |
| | | | | rs182610315 | 10 | 114743091 | 0.027 | 0 | 12 | 13 | 13 | 13 | 14 | | 98.17 | 1.83 | |
| | | rs180988137 | 1979 (9) | rs180988137 | 10 | 114751173 | 0.289 | 0.3612 | 2 | 13 | 14 | 13 | 13 | | 1.04 | 1.04 | |
| | | | | rs146398962 | 10 | 114772182 | 0.133 | 0.1601 | 2 | 13 | 8 | 13 | 13 | | 0.56 | 0.56 | |
| | | | | rs180726800 | 10 | 114784926 | 0.064 | 0.0415 | 12 | 13 | 10 | 13 | 10 | | 1.95 | 1.95 | |
| | | | | rs151137175 | 10 | 114741473 | 0.056 | 0.0361 | 12 | 13 | 10 | 13 | 14 | | 0.42 | 0.42 | |
| | | | | rs145003494 | 10 | 114834411 | 0.02 | 0.0262 | 4 | 10 | 8 | 13 | 8 | | 1.03 | 1.03 | |
| | | | | rs4917646 | 10 | 114902811 | 0.003 | 0.0239 | 3 | 10 | 14 | 14 | 14 | | 21.13 | 21.13 | |
| | | | | rs186384225 | 10 | 114808119 | 0.002 | 0.019 | 5 | 10 | 8 | 10 | 10 | FOXA2 | 0.37 | 0.37 | |
| | | | | rs3814575 | 10 | 114898739 | 0.011 | 0.0177 | 2 | 13 | 11 | 13 | 13 | | 29.46 | 29.46 | |
| | | | | rs7080044 | 10 | 114899115 | 0.011 | 0.0172 | 2 | 13 | 8 | 13 | 13 | | 29.4 | 29.4 | |
| | | | | rs72828151 | 10 | 114968347 | 0.021 | 0.012 | 12 | 16 | 16 | 16 | 16 | | 0.53 | 0.53 | |
| Secondary | rs78025551 | 339 (6) | 133 (7) | rs10787478 | 10 | 114899446 | 0.013 | 0.01 | 12 | 13 | 11 | 13 | 13 | | 29.42 | 29.42 | |
| | | | | rs78025551 | 10 | 114757956 | 0.399 | 0.4119 | 4 | 10 | 10 | 13 | 13 | | 85.1 | 14.9 | |
| | | | | rs57225583 | 10 | 114771843 | 0.132 | 0.137 | 2 | 13 | 11 | 13 | 13 | | 85.23 | 14.77 | |
| | | | | rs11196183 | 10 | 114750264 | 0.106 | 0.1092 | 2 | 13 | 13 | 13 | 13 | | 85.08 | 14.92 | |
| | | | | rs7896811 | 10 | 114766717 | 0.101 | 0.1051 | 2 | 13 | 7 | 11 | 11 | | 85.32 | 14.68 | |
| | | | | rs11196182 | 10 | 114750157 | 0.093 | 0.0962 | 2 | 13 | 13 | 13 | 13 | | 85.03 | 14.97 | |
| | | | | rs114863326 | 10 | 114783775 | 0.069 | 0.0376 | 12 | 13 | 13 | 13 | 10 | | 85.28 | 14.72 | |
| Secondary | rs34855922 | 2 (2) | 2 (2) | rs11196191 | 10 | 114780633 | 0.001 | 0.0169 | 5 | 8 | 8 | 8 | 8 | NKX2.2 | 46.97 | 46.97 | |
| | | | | rs34855922 | 10 | 114871594 | 0.783 | 0.7793 | 2 | 13 | 10 | 13 | 8 | | 71.62 | 28.38 | |
| | | | | rs11196234 | 10 | 114874406 | 0.21 | 0.2123 | 2 | 13 | 10 | 13 | 8 | | 76.48 | 23.52 | |

PPA: posterior probability of association; MAF: minor allele frequency; 1: Active promoter; 2: Weak enhancer; 3: Weak enhan+ATAC+Meth; 4: Strong enhancer; 5: Strong enhan+ATAC+Meth; 6: 1_Active_TSS; 7: 8_Genic_enhancer; 8: 9_Active_enhancer_1; 9: 10_Active_enhancer_1; 10: 10_Active_enhancer_2; 11: 11_Weak_enhancer; 12: Heterochromatin; 13: 6_Weak_transcription; 14: 5_Strong_transcription; 15: 17_Weak_repressed_polycomb; 16: 18_Quiescent/low_signal.

¹Thurner, M. et al. Integration of human pancreatic islet genomic data refines regulatory mechanisms at Type 2 Diabetes susceptibility loci. *bioRxiv* (2017).

²Varshney, A. et al. Genetic regulatory signatures underlying islet gene expression and type 2 diabetes. *Proc Natl Acad Sci U S A* 114, 2301-2306 (2017).

³Pasquali, L. et al. Pancreatic islet enhancer clusters enriched in type 2 diabetes risk-associated variants. *Nat Genet* 46, 136-143 (2014).

Supplementary Table 10 | Summary results of LD Score regression analyses between T2D and various diseases, metabolic, and anthropometric traits.

| Phenotype | Category | T2D-unadjusted for BMI | | | | T2D-adjusted for BMI | | | | Reference (PMID) |
|--|-----------------|-------------------------------|--------|--------|------------------------|-------------------------------|--------|---------|-----------------------|------------------|
| | | Genetic Correlation (r_g) | SE | z | P-value | Genetic Correlation (r_g) | SE | z | P-value | |
| Mothers age at death | aging | -0.3356 | 0.0598 | -5.61 | 2.0x10 ⁻⁸ | -0.2022 | 0.0628 | -3.2225 | 0.001 | 27015805 |
| Parents age at death | aging | -0.4033 | 0.0718 | -5.62 | 1.9x10 ⁻⁸ | -0.3183 | 0.0738 | -4.3096 | 0.00002 | 27015805 |
| Fathers age at death | aging | -0.4473 | 0.0611 | -7.32 | 2.5x10 ⁻¹³ | -0.2823 | 0.058 | -4.8647 | 1.1x10 ⁻⁶ | 27015805 |
| Waist circumference | anthropometric | 0.5917 | 0.0237 | 24.96 | 1.8x10 ⁻¹³⁷ | 0.2417 | 0.0296 | 8.1671 | 3.2x10 ⁻¹⁶ | 25673412 |
| Waist-to-hip ratio | anthropometric | 0.5649 | 0.0304 | 18.56 | 6.3x10 ⁻⁷⁷ | 0.3761 | 0.0328 | 11.4794 | 1.7x10 ⁻³⁰ | 25673412 |
| Overweight | anthropometric | 0.5608 | 0.0308 | 18.20 | 5.6x10 ⁻⁷⁴ | 0.1734 | 0.034 | 5.1067 | 3.3x10 ⁻⁷ | 23563607 |
| Obesity class 1 | anthropometric | 0.539 | 0.0305 | 17.69 | 5.0x10 ⁻⁷⁰ | 0.1437 | 0.0324 | 4.437 | 9.1x10 ⁻⁶ | 23563607 |
| Body mass index | anthropometric | 0.5098 | 0.0293 | 17.38 | 1.1x10 ⁻⁶⁷ | 0.0981 | 0.0313 | 3.1391 | 0.002 | 20935630 |
| Obesity class 2 | anthropometric | 0.5433 | 0.0376 | 14.45 | 2.7x10 ⁻⁴⁷ | 0.1385 | 0.0419 | 3.3097 | 0.0009 | 23563607 |
| Hip circumference | anthropometric | 0.4209 | 0.0292 | 14.43 | 3.4x10 ⁻⁴⁷ | 0.0554 | 0.0335 | 1.65 | 0.1 | 25673412 |
| Extreme bmi | anthropometric | 0.5291 | 0.045 | 11.76 | 6.2x10 ⁻³² | 0.1212 | 0.0492 | 2.4655 | 0.01 | 23563607 |
| Body fat | anthropometric | 0.4815 | 0.0451 | 10.69 | 1.2x10 ⁻²⁶ | 0.1295 | 0.0467 | 2.7768 | 0.006 | 26833246 |
| Obesity class 3 | anthropometric | 0.5385 | 0.0516 | 10.44 | 1.6x10 ⁻²⁵ | 0.1298 | 0.0531 | 2.4459 | 0.01 | 23563607 |
| Childhood obesity | anthropometric | 0.3307 | 0.0396 | 8.35 | 6.7x10 ⁻¹⁷ | 0.0655 | 0.0421 | 1.5543 | 0.1 | 22484627 |
| Extreme waist-to-hip ratio | anthropometric | 0.3998 | 0.0601 | 6.65 | 2.9x10 ⁻¹¹ | 0.4548 | 0.0669 | 6.7934 | 1.1x10 ⁻¹¹ | 23563607 |
| Height; Females at age 10 and males at age 12 | anthropometric | 0.1021 | 0.0371 | 2.75 | 0.006 | 0.0252 | 0.042 | 0.5997 | 0.5 | 23449627 |
| Sitting height ratio | anthropometric | 0.0511 | 0.0433 | 1.18 | 0.2 | 0.0471 | 0.0464 | 1.0163 | 0.3 | 25865494 |
| Infant head circumference | anthropometric | -0.0414 | 0.0546 | -0.76 | 0.4 | -0.1059 | 0.0658 | -1.609 | 0.1 | 22504419 |
| Extreme height | anthropometric | -0.049 | 0.0339 | -1.44 | 0.1 | -0.0318 | 0.039 | -0.8166 | 0.4 | 23563607 |
| Height_2010 | anthropometric | -0.0455 | 0.0228 | -2.00 | 0.05 | -0.0261 | 0.0258 | -1.0133 | 0.3 | 20881960 |
| Child birth length | anthropometric | -0.1064 | 0.0465 | -2.29 | 0.02 | -0.1003 | 0.052 | -1.9296 | 0.05 | 25281659 |
| Child birth weight | anthropometric | -0.1625 | 0.0506 | -3.21 | 0.001 | -0.2477 | 0.0596 | -4.1571 | 0.00003 | 23202124 |
| Difference in height between adolescence and adulthood; age 14 | anthropometric | -0.2176 | 0.0651 | -3.34 | 0.0008 | -0.082 | 0.0659 | -1.2443 | 0.2 | 23449627 |
| Difference in height between childhood and adulthood; age 8 | anthropometric | -0.1824 | 0.0485 | -3.76 | 0.0002 | -0.0792 | 0.0523 | -1.5139 | 0.1 | 23449627 |
| Birth weight | anthropometric | -0.21 | 0.0334 | -6.28 | 3.4x10 ⁻¹⁰ | -0.2934 | 0.0371 | -7.8986 | 2.8x10 ⁻¹⁵ | 27680694 |
| Asthma | autoimmune | 0.1706 | 0.056 | 3.05 | 0.002 | 0.0922 | 0.0647 | 1.4259 | 0.2 | 17611496 |
| Primary biliary cirrhosis | autoimmune | 0.0804 | 0.0479 | 1.68 | 0.09 | 0.0327 | 0.054 | 0.6064 | 0.5 | 26394269 |
| Systemic lupus erythematosus | autoimmune | 0.0704 | 0.0481 | 1.46 | 0.1 | 0.0619 | 0.0514 | 1.2047 | 0.2 | 26502338 |
| Rheumatoid Arthritis | autoimmune | 0.0023 | 0.0269 | 0.08 | 0.9 | -0.0413 | 0.0328 | -1.2588 | 0.2 | 24390342 |
| Crohns disease | autoimmune | -0.0105 | 0.0307 | -0.34 | 0.7 | -0.0652 | 0.0357 | -1.8249 | 0.07 | 26192919 |
| Eczema | autoimmune | -0.0258 | 0.0547 | -0.47 | 0.6 | -0.0591 | 0.0611 | -0.9668 | 0.3 | 26482879 |
| Inflammatory Bowel Disease (Euro) | autoimmune | -0.0365 | 0.0279 | -1.31 | 0.2 | -0.0606 | 0.034 | -1.7807 | 0.08 | 26192919 |
| Celiac disease | autoimmune | -0.0808 | 0.0474 | -1.70 | 0.09 | -0.0602 | 0.0588 | -1.0253 | 0.3 | 20190752 |
| Ulcerative colitis | autoimmune | -0.0602 | 0.0317 | -1.90 | 0.06 | -0.0476 | 0.0393 | -1.2109 | 0.2 | 26192919 |
| Lumbar spine bone mineral density | bone | 0.1088 | 0.0298 | 3.65 | 0.0003 | 0.1154 | 0.0346 | 3.337 | 0.0008 | 22504420 |
| Femoral neck bone mineral density | bone | 0.0893 | 0.0268 | 3.33 | 0.0009 | 0.0908 | 0.0317 | 2.8691 | 0.004 | 22504420 |
| Femoral Neck bone mineral density | bone | 0.0651 | 0.0341 | 1.91 | 0.06 | 0.0771 | 0.0396 | 1.9476 | 0.05 | 26367794 |
| Lumbar Spine bone mineral density | bone | 0.052 | 0.0347 | 1.50 | 0.1 | 0.0748 | 0.0411 | 1.8215 | 0.07 | 26367794 |
| Mean Hippocampus | brain_volume | 0.0361 | 0.0632 | 0.57 | 0.6 | 0.0723 | 0.0715 | 1.0102 | 0.3 | 25607358 |
| Mean Caudate | brain_volume | 0.013 | 0.0444 | 0.29 | 0.8 | 0.031 | 0.0532 | 0.5826 | 0.6 | 25607358 |
| Mean Pallidum | brain_volume | 0.0062 | 0.0592 | 0.11 | 0.9 | 0.0347 | 0.0665 | 0.5222 | 0.6 | 25607358 |
| Mean Thalamus | brain_volume | -0.0164 | 0.0624 | -0.26 | 0.8 | -0.0107 | 0.0698 | -0.1538 | 0.9 | 25607358 |
| Mean Putamen | brain_volume | -0.0212 | 0.047 | -0.45 | 0.7 | 0.0131 | 0.0505 | 0.2592 | 0.8 | 25607358 |
| ICV | brain_volume | -0.1789 | 0.057 | -3.14 | 0.002 | -0.1901 | 0.0643 | -2.9566 | 0.003 | 25607358 |
| Lung cancer | cancer | 0.1204 | 0.0422 | 2.85 | 0.004 | 0.0557 | 0.0449 | 1.2416 | 0.2 | 27488534 |
| Squamous cell lung cancer | cancer | 0.1572 | 0.0684 | 2.30 | 0.02 | 0.05 | 0.0723 | 0.6916 | 0.5 | 27488534 |
| Lung cancer (all) | cancer | 0.0982 | 0.0489 | 2.01 | 0.04 | 0.0461 | 0.0534 | 0.8639 | 0.4 | 24880342 |
| Coronary artery disease | cardiometabolic | 0.3971 | 0.0278 | 14.29 | 2.5x10 ⁻⁴⁶ | 0.3092 | 0.0318 | 9.7186 | 2.5x10 ⁻²² | 26343387 |
| Adiponectin | cardiometabolic | -0.1809 | 0.052 | -3.48 | 0.0005 | -0.2783 | 0.0606 | -4.5935 | 4.4x10 ⁻⁶ | 22479202 |
| Intelligence | cognitive | -0.1149 | 0.0245 | -4.68 | 2.8x10 ⁻⁶ | -0.0318 | 0.0273 | -1.1615 | 0.2 | 28530673 |
| Childhood IQ | education | -0.1922 | 0.0559 | -3.44 | 0.0006 | -0.1041 | 0.0654 | -1.592 | 0.1 | 2358156 |
| College completion | education | -0.2793 | 0.0305 | -9.17 | 4.6x10 ⁻²⁰ | -0.1422 | 0.0332 | -4.2839 | 0.00002 | 23722424 |
| Years of schooling 2016 | education | -0.2699 | 0.0199 | -13.56 | 7.4x10 ⁻⁴² | -0.1402 | 0.0224 | -6.2626 | 3.8x10 ⁻¹⁰ | 27225129 |
| Fasting insulin main effect | glycemic | 0.6217 | 0.0606 | 10.27 | 9.9x10 ⁻²⁵ | 0.4429 | 0.0647 | 6.8465 | 7.6x10 ⁻¹² | 22581228 |
| HbA1C | glycemic | 0.5697 | 0.0624 | 9.12 | 7.2x10 ⁻²⁰ | 0.5588 | 0.0648 | 8.6264 | 6.3x10 ⁻¹⁸ | 20858683 |
| Fasting glucose main effect | glycemic | 0.5799 | 0.0672 | 8.63 | 6.2x10 ⁻¹⁸ | 0.5583 | 0.068 | 8.211 | 2.2x10 ⁻¹⁶ | 22581228 |
| HOMA-IR | glycemic | 0.6632 | 0.0853 | 7.77 | 7.7x10 ⁻¹⁵ | 0.4777 | 0.0825 | 5.7922 | 6.9x10 ⁻⁹ | 20081858 |
| HOMA-B | glycemic | 0.2029 | 0.0602 | 3.37 | 0.0008 | 0.05 | 0.0632 | 0.7909 | 0.4 | 20081858 |
| Heart rate | haemotological | 0.0695 | 0.0312 | 2.22 | 0.03 | 0.093 | 0.0379 | 2.4543 | 0.01 | 23583979 |
| Mean platelet volume | haemotological | 0.0819 | 0.0388 | 2.11 | 0.03 | 0.0546 | 0.0446 | 1.226 | 0.2 | 22139419 |
| Platelet count | haemotological | 0.0386 | 0.0335 | 1.15 | 0.2 | 0.0563 | 0.0364 | 1.5448 | 0.1 | 22139419 |
| Leptin_not_adjBMI | hormone | 0.5036 | 0.0623 | 8.09 | 6.0x10 ⁻¹⁶ | 0.2126 | 0.0603 | 3.5269 | 0.0004 | 26833098 |
| Leptin adjBMI | hormone | 0.1797 | 0.0579 | 3.11 | 0.002 | 0.2029 | 0.0641 | 3.164 | 0.002 | 26833098 |
| Urinary albumin-to-creatinine ratio | kidney | 0.2556 | 0.059 | 4.33 | 0.00001 | 0.2561 | 0.0619 | 4.1348 | 0.00004 | 26631737 |
| Chronic Kidney Disease | kidney | 0.1128 | 0.0543 | 2.08 | 0.04 | 0.0115 | 0.06 | 0.1914 | 0.8 | 26831199 |
| Urinary albumin-to-creatinine ratio (non-diabetes) | kidney | 0.1133 | 0.058 | 1.95 | 0.05 | 0.1103 | 0.0619 | 1.7835 | 0.07 | 26631737 |
| Serum creatinine | kidney | 0.036 | 0.0273 | 1.32 | 0.2 | 0.071 | 0.0307 | 2.312 | 0.02 | 26831199 |
| Serum creatinine (non-diabetes) | kidney | 0.0264 | 0.0294 | 0.90 | 0.4 | 0.0644 | 0.033 | 1.9505 | 0.05 | 26831199 |
| Serum cystatin c | kidney | -0.1611 | 0.0607 | -2.66 | 0.008 | -0.0086 | 0.0426 | -0.2027 | 0.8 | 26831199 |
| Triglycerides | lipids | 0.4065 | 0.0537 | 7.56 | 3.9x10 ⁻¹⁴ | 0.3693 | 0.0569 | 6.4899 | 8.6x10 ⁻¹¹ | 20686565 |

| | | | | | | | | | | |
|--|---------------|---------|--------|--------|-----------------------|---------|--------|---------|-----------------------|----------|
| LDL cholesterol | lipids | 0.1253 | 0.049 | 2.56 | 0.01 | 0.1169 | 0.0491 | 2.3799 | 0.02 | 20686565 |
| Total Cholesterol | lipids | 0.0682 | 0.0385 | 1.77 | 0.08 | 0.0851 | 0.0428 | 1.9904 | 0.05 | 20686565 |
| HDL cholesterol | lipids | -0.475 | 0.0454 | -10.47 | 1.2×10^{-25} | -0.3683 | 0.043 | -8.5709 | 1.0×10^{-17} | 20686565 |
| Forced expiratory volume in 1 second (FEV1)/Forced Vital capacity(FVC) | lung_function | 0.0616 | 0.029 | 2.12 | 0.03 | 0.0203 | 0.0333 | 0.6087 | 0.5 | 28166213 |
| Forced expiratory volume in 1 second (FEV1) | lung_function | -0.1638 | 0.028 | -5.85 | 5.0×10^{-9} | -0.1516 | 0.0306 | -4.9565 | 7.2×10^{-7} | 28166213 |
| Forced Vital capacity(FVC) | lung_function | -0.2325 | 0.0277 | -8.39 | 4.7×10^{-17} | -0.2002 | 0.0303 | -6.6157 | 3.7×10^{-11} | 28166213 |
| Cholesterol esters in large VLDL | metabolites | 0.3324 | 0.0597 | 5.57 | 2.5×10^{-8} | 0.2151 | 0.0652 | 3.2998 | 0.001 | 27005778 |
| Total lipids in large VLDL | metabolites | 0.3492 | 0.0632 | 5.53 | 3.3×10^{-8} | 0.2207 | 0.0702 | 3.1456 | 0.002 | 27005778 |
| Glycoprotein acetyl; mainly a1-acid glycoprotein | metabolites | 0.4508 | 0.0835 | 5.40 | 6.6×10^{-8} | 0.2878 | 0.0763 | 3.7738 | 0.0002 | 27005778 |
| Total lipids in chylomicrons and largest VLDL particles | metabolites | 0.3598 | 0.0682 | 5.27 | 1.3×10^{-7} | 0.2238 | 0.0722 | 3.1021 | 0.002 | 27005778 |
| Total lipids in very large VLDL | metabolites | 0.3116 | 0.0605 | 5.15 | 2.6×10^{-7} | 0.1713 | 0.0665 | 2.5766 | 0.01 | 27005778 |
| Concentration of medium VLDL particles | metabolites | 0.3434 | 0.0675 | 5.08 | 3.7×10^{-7} | 0.2171 | 0.0703 | 3.0881 | 0.002 | 27005778 |
| Concentration of chylomicrons and largest VLDL particles | metabolites | 0.3627 | 0.0723 | 5.01 | 5.3×10^{-7} | 0.2426 | 0.0761 | 3.189 | 0.001 | 27005778 |
| Triglycerides in large VLDL | metabolites | 0.3695 | 0.0742 | 4.98 | 6.3×10^{-7} | 0.2328 | 0.079 | 2.9456 | 0.003 | 27005778 |
| Concentration of very large VLDL particles | metabolites | 0.3371 | 0.0678 | 4.97 | 6.6×10^{-7} | 0.1955 | 0.0703 | 2.7799 | 0.005 | 27005778 |
| Total lipids in medium VLDL | metabolites | 0.3542 | 0.0721 | 4.91 | 8.9×10^{-7} | 0.2253 | 0.0734 | 3.071 | 0.002 | 27005778 |
| Mean diameter for VLDL particles | metabolites | 0.3487 | 0.0722 | 4.83 | 1.4×10^{-6} | 0.2575 | 0.0772 | 3.3344 | 0.0009 | 27005778 |
| Triglycerides in very large VLDL | metabolites | 0.325 | 0.0676 | 4.81 | 1.5×10^{-6} | 0.185 | 0.0721 | 2.5663 | 0.01 | 27005778 |
| Concentration of large VLDL particles | metabolites | 0.3568 | 0.0742 | 4.81 | 1.5×10^{-6} | 0.2251 | 0.0762 | 2.9552 | 0.003 | 27005778 |
| Phospholipids in very large VLDL | metabolites | 0.3464 | 0.0742 | 4.67 | 3.0×10^{-6} | 0.191 | 0.0751 | 2.5425 | 0.01 | 27005778 |
| Isoleucine | metabolites | 0.511 | 0.1094 | 4.67 | 3.0×10^{-6} | 0.4027 | 0.1071 | 3.7607 | 0.0002 | 27005778 |
| Valine | metabolites | 0.4357 | 0.0936 | 4.66 | 3.2×10^{-6} | 0.3517 | 0.0938 | 3.7484 | 0.0002 | 27005778 |
| Concentration of small VLDL particles | metabolites | 0.3156 | 0.0681 | 4.63 | 3.6×10^{-6} | 0.1801 | 0.0674 | 2.6739 | 0.008 | 27005778 |
| Phospholipids in large VLDL | metabolites | 0.3459 | 0.0755 | 4.58 | 4.6×10^{-6} | 0.2099 | 0.0783 | 2.6804 | 0.007 | 27005778 |
| Phospholipids in chylomicrons and largest VLDL particles | metabolites | 0.3465 | 0.0761 | 4.56 | 5.2×10^{-6} | 0.1957 | 0.078 | 2.5091 | 0.01 | 27005778 |
| Free cholesterol in large VLDL | metabolites | 0.3082 | 0.0679 | 4.54 | 5.7×10^{-6} | 0.1717 | 0.0719 | 2.3889 | 0.02 | 27005778 |
| Triglycerides in medium VLDL | metabolites | 0.4306 | 0.095 | 4.53 | 5.8×10^{-6} | 0.2902 | 0.0932 | 3.1154 | 0.002 | 27005778 |
| Total cholesterol in large VLDL | metabolites | 0.306 | 0.0684 | 4.48 | 7.6×10^{-6} | 0.1703 | 0.0719 | 2.3679 | 0.02 | 27005778 |
| Triglycerides in chylomicrons and largest VLDL particles | metabolites | 0.3948 | 0.0908 | 4.35 | 0.00001 | 0.2379 | 0.0906 | 2.6257 | 0.009 | 27005778 |
| Triglycerides in small VLDL | metabolites | 0.3608 | 0.083 | 4.35 | 0.00001 | 0.2089 | 0.0789 | 2.6469 | 0.008 | 27005778 |
| Total lipids in small VLDL | metabolites | 0.305 | 0.0702 | 4.35 | 0.00001 | 0.1668 | 0.0681 | 2.4488 | 0.01 | 27005778 |
| Phospholipids in medium VLDL | metabolites | 0.363 | 0.0848 | 4.28 | 0.00002 | 0.2194 | 0.0815 | 2.6909 | 0.007 | 27005778 |
| Serum total triglycerides | metabolites | 0.3499 | 0.0842 | 4.16 | 0.00003 | 0.203 | 0.0803 | 2.528 | 0.01 | 27005778 |
| Triglycerides in very small VLDL | metabolites | 0.2897 | 0.0704 | 4.12 | 0.00004 | 0.1495 | 0.0685 | 2.1839 | 0.03 | 27005778 |
| Cholesterol esters in medium VLDL | metabolites | 0.2746 | 0.0668 | 4.11 | 0.00004 | 0.1563 | 0.067 | 2.3314 | 0.02 | 27005778 |
| Free cholesterol in medium VLDL | metabolites | 0.3337 | 0.0816 | 4.09 | 0.00004 | 0.1933 | 0.0799 | 2.4187 | 0.02 | 27005778 |
| Total cholesterol in medium VLDL | metabolites | 0.2793 | 0.0735 | 3.80 | 0.0001 | 0.1473 | 0.0722 | 2.0391 | 0.04 | 27005778 |
| Free cholesterol in small VLDL | metabolites | 0.2781 | 0.0757 | 3.67 | 0.0002 | 0.1313 | 0.073 | 1.7999 | 0.07 | 27005778 |
| Phospholipids in small VLDL | metabolites | 0.2693 | 0.0739 | 3.64 | 0.0003 | 0.1226 | 0.0704 | 1.7417 | 0.08 | 27005778 |
| Triglycerides in IDL | metabolites | 0.2403 | 0.0789 | 3.05 | 0.002 | 0.0916 | 0.0726 | 1.2627 | 0.2 | 27005778 |
| Concentration of very small VLDL particles | metabolites | 0.2181 | 0.0728 | 3.00 | 0.003 | 0.0757 | 0.0692 | 1.0933 | 0.3 | 27005778 |
| Total cholesterol in small VLDL | metabolites | 0.2376 | 0.0829 | 2.87 | 0.004 | 0.0899 | 0.0768 | 1.1705 | 0.2 | 27005778 |
| Total lipids in very small VLDL | metabolites | 0.2068 | 0.0764 | 2.71 | 0.007 | 0.0644 | 0.07 | 0.919 | 0.4 | 27005778 |
| Apolipoprotein B | metabolites | 0.3045 | 0.12 | 2.54 | 0.01 | 0.1482 | 0.0964 | 1.5373 | 0.1 | 27005778 |
| Alanine | metabolites | 0.1355 | 0.0658 | 2.06 | 0.04 | 0.1271 | 0.0789 | 1.6116 | 0.1 | 27005778 |
| Total lipids in small LDL | metabolites | 0.2276 | 0.1114 | 2.04 | 0.04 | 0.0651 | 0.0885 | 0.7357 | 0.5 | 27005778 |
| Concentration of small LDL particles | metabolites | 0.2051 | 0.1009 | 2.03 | 0.04 | 0.0525 | 0.0843 | 0.6234 | 0.5 | 27005778 |
| Total cholesterol in small LDL | metabolites | 0.2297 | 0.1271 | 1.81 | 0.07 | 0.0631 | 0.0966 | 0.6536 | 0.5 | 27005778 |
| Concentration of medium LDL particles | metabolites | 0.2051 | 0.114 | 1.80 | 0.07 | 0.046 | 0.0901 | 0.5101 | 0.6 | 27005778 |
| Phospholipids in medium LDL | metabolites | 0.1801 | 0.1017 | 1.77 | 0.08 | 0.0337 | 0.0855 | 0.3947 | 0.7 | 27005778 |
| Total lipids in medium LDL | metabolites | 0.1958 | 0.1122 | 1.74 | 0.08 | 0.0412 | 0.0886 | 0.4645 | 0.6 | 27005778 |
| Phospholipids in very small VLDL | metabolites | 0.1419 | 0.0856 | 1.66 | 0.1 | 0.0009 | 0.0743 | 0.0121 | 1.0 | 27005778 |
| Omega-3 fatty acids | metabolites | 0.1078 | 0.0659 | 1.64 | 0.1 | 0.1148 | 0.0751 | 1.5296 | 0.1 | 27005778 |
| Concentration of IDL particles | metabolites | 0.1686 | 0.1032 | 1.63 | 0.1 | 0.0064 | 0.0833 | 0.0766 | 0.9 | 27005778 |
| Cholesterol esters in medium LDL | metabolites | 0.1708 | 0.1047 | 1.63 | 0.1 | 0.0283 | 0.0866 | 0.3269 | 0.7 | 27005778 |
| Concentration of large LDL particles | metabolites | 0.1728 | 0.1114 | 1.55 | 0.1 | 0.0135 | 0.0877 | 0.1543 | 0.9 | 27005778 |
| Total cholesterol in LDL | metabolites | 0.1633 | 0.1102 | 1.48 | 0.1 | 0.0117 | 0.0869 | 0.1348 | 0.9 | 27005778 |
| Total lipids in large LDL | metabolites | 0.1603 | 0.1093 | 1.47 | 0.1 | 0.0075 | 0.0873 | 0.0859 | 0.9 | 27005778 |
| Total cholesterol in medium LDL | metabolites | 0.145 | 0.1 | 1.45 | 0.1 | -0.0003 | 0.0835 | -0.0032 | 1 | 27005778 |
| Total lipids in IDL | metabolites | 0.1422 | 0.1007 | 1.41 | 0.2 | -0.0124 | 0.0827 | -0.15 | 0.9 | 27005778 |
| Cholesterol esters in large LDL | metabolites | 0.1449 | 0.1036 | 1.40 | 0.2 | 0.0046 | 0.0863 | 0.053 | 1 | 27005778 |
| Phospholipids in IDL | metabolites | 0.1426 | 0.1088 | 1.31 | 0.2 | -0.0271 | 0.0855 | -0.3169 | 0.8 | 27005778 |
| Phospholipids in large LDL | metabolites | 0.133 | 0.1024 | 1.30 | 0.2 | -0.0233 | 0.083 | -0.2807 | 0.8 | 27005778 |
| Total cholesterol in IDL | metabolites | 0.0942 | 0.0938 | 1.00 | 0.3 | -0.0366 | 0.0824 | -0.4436 | 0.7 | 27005778 |
| Total cholesterol in large LDL | metabolites | 0.0908 | 0.0911 | 1.00 | 0.3 | -0.0489 | 0.0806 | -0.6067 | 0.5 | 27005778 |
| Free cholesterol in large LDL | metabolites | 0.0618 | 0.0863 | 0.72 | 0.5 | -0.0698 | 0.0798 | -0.8754 | 0.4 | 27005778 |
| 22:6 docosahexaenoic acid | metabolites | 0.0392 | 0.0615 | 0.64 | 0.5 | 0.0921 | 0.0732 | 1.2569 | 0.2 | 27005778 |
| Free cholesterol in IDL | metabolites | 0.0485 | 0.0811 | 0.60 | 0.5 | -0.082 | 0.0755 | -1.0862 | 0.3 | 27005778 |
| Description of average fatty acid chain length | metabolites | -0.0035 | 0.0651 | -0.05 | 1.0 | 0.062 | 0.0772 | 0.8033 | 0.4 | 27005778 |
| Creatinine | metabolites | -0.0272 | 0.0509 | -0.53 | 0.6 | -0.0509 | 0.0592 | -0.8605 | 0.4 | 27005778 |
| 18:2 linoleic acid (LA) | metabolites | -0.0467 | 0.0645 | -0.72 | 0.5 | -0.1437 | 0.0735 | -1.9543 | 0.05 | 27005778 |
| Citrate | metabolites | -0.0703 | 0.0576 | -1.22 | 0.2 | 0.0134 | 0.0711 | 0.1883 | 0.9 | 27005778 |
| Average number of double bonds in a fatty acid chain | metabolites | -0.0955 | 0.058 | -1.65 | 0.1 | -0.0076 | 0.059 | -0.1294 | 0.9 | 27005778 |
| Glutamine | metabolites | -0.2294 | 0.079 | -2.90 | 0.004 | -0.1762 | 0.0833 | -2.1138 | 0.03 | 27005778 |
| Apolipoprotein A-I | metabolites | -0.3422 | 0.094 | -3.64 | 0.0003 | -0.3002 | 0.1004 | -2.9906 | 0.003 | 27005778 |
| Phospholipids in medium HDL | metabolites | -0.313 | 0.0795 | -3.93 | 0.00008 | -0.2886 | 0.0917 | -3.1472 | 0.002 | 27005778 |
| Free cholesterol in medium HDL | metabolites | -0.4022 | 0.0867 | -4.64 | 3.5×10^{-6} | -0.3364 | 0.0935 | -3.5968 | 0.0003 | 27005778 |
| Phospholipids in very large HDL | metabolites | -0.4173 | 0.0836 | -4.99 | 6.0×10^{-7} | -0.2815 | 0.0758 | -3.7153 | 0.0002 | 27005778 |
| Total cholesterol in HDL | metabolites | -0.4825 | 0.0793 | -6.09 | 1.2×10^{-9} | -0.352 | 0.078 | -4.5136 | 6.4×10^{-6} | 27005778 |
| Total cholesterol in large HDL | metabolites | -0.5129 | 0.0793 | -6.46 | 1.0×10^{-10} | -0.3525 | 0.0738 | -4.7774 | 1.8×10^{-6} | 27005778 |

| | | | | | | | | | | |
|--------------------------------------|------------------|---------|--------|--------|-----------------------|---------|--------|---------|----------------------|----------|
| Mean diameter for HDL particles | metabolites | -0.4423 | 0.0673 | -6.57 | 5.1x10 ⁻¹¹ | -0.3209 | 0.0661 | -4.8538 | 1.2x10 ⁻⁶ | 27005778 |
| Free cholesterol in large HDL | metabolites | -0.5175 | 0.0778 | -6.65 | 2.9x10 ⁻¹¹ | -0.3631 | 0.0733 | -4.9548 | 7.2x10 ⁻⁷ | 27005778 |
| Cholesterol esters in large HDL | metabolites | -0.4714 | 0.0696 | -6.77 | 1.3x10 ⁻¹¹ | -0.3209 | 0.0674 | -4.7596 | 1.9x10 ⁻⁶ | 27005778 |
| Total lipids in large HDL | metabolites | -0.4651 | 0.0636 | -7.32 | 2.5x10 ⁻¹³ | -0.3276 | 0.0649 | -5.047 | 4.5x10 ⁻⁷ | 27005778 |
| Phospholipids in large HDL | metabolites | -0.4672 | 0.0625 | -7.48 | 7.6x10 ⁻¹⁴ | -0.3395 | 0.0652 | -5.2054 | 1.9x10 ⁻⁷ | 27005778 |
| Concentration of large HDL particles | metabolites | -0.461 | 0.0613 | -7.52 | 5.3x10 ⁻¹⁴ | -0.328 | 0.0644 | -5.0957 | 3.5x10 ⁻⁷ | 27005778 |
| Ferritin | metal | 0.2084 | 0.0683 | 3.05 | 0.002 | 0.1711 | 0.0743 | 2.304 | 0.02 | 25352340 |
| Parkinsons disease | neurological | 0.0586 | 0.0369 | 1.59 | 0.1 | 0.0148 | 0.0424 | 0.349 | 0.7 | 19915575 |
| Alzheimers disease | neurological | 0.0698 | 0.0656 | 1.06 | 0.3 | 0.0631 | 0.0679 | 0.9284 | 0.4 | 24162737 |
| Urate | other | 0.2838 | 0.0658 | 4.31 | 0.00002 | 0.1956 | 0.0526 | 3.7197 | 0.0002 | 23263486 |
| Neuroticism | personality | 0.0599 | 0.041 | 1.46 | 0.1 | 0.0555 | 0.0393 | 1.4124 | 0.2 | 27089181 |
| Neo-openness to experience | personality | -0.0181 | 0.0616 | -0.29 | 0.8 | 0.0232 | 0.0712 | 0.3252 | 0.7 | 21173776 |
| Depressive symptoms | psychiatric | 0.1982 | 0.0379 | 5.24 | 1.6x10 ⁻⁷ | 0.1167 | 0.0382 | 3.0561 | 0.002 | 27089181 |
| Major depressive disorder | psychiatric | 0.0528 | 0.0449 | 1.18 | 0.2 | 0.0664 | 0.0487 | 1.3615 | 0.2 | 22472876 |
| Subjective well being | psychiatric | -0.0377 | 0.0347 | -1.08 | 0.3 | -0.0287 | 0.0358 | -0.8023 | 0.4 | 27089181 |
| PGC cross-disorder analysis | psychiatric | -0.0323 | 0.0289 | -1.12 | 0.3 | -0.0049 | 0.0334 | -0.146 | 0.9 | 23453885 |
| Bipolar disorder | psychiatric | -0.0674 | 0.0328 | -2.06 | 0.04 | -0.0711 | 0.0369 | -1.9249 | 0.05 | 21926972 |
| Autism spectrum disorder | psychiatric | -0.0947 | 0.0432 | -2.19 | 0.03 | -0.0697 | 0.0481 | -1.4507 | 0.1 | 0 |
| Schizophrenia | psychiatric | -0.0596 | 0.0213 | -2.79 | 0.005 | -0.0426 | 0.0258 | -1.653 | 0.1 | 25056061 |
| Anorexia Nervosa | psychiatric | -0.1109 | 0.0281 | -3.95 | 0.00008 | -0.0572 | 0.0305 | -1.8733 | 0.06 | 24514567 |
| Number of children ever born | reproductive | 0.1187 | 0.03 | 3.96 | 0.00008 | 0.0429 | 0.035 | 1.2255 | 0.2 | 27798627 |
| Age at Menopause | reproductive | -0.0957 | 0.031 | -3.08 | 0.002 | -0.1021 | 0.0355 | -2.8773 | 0.004 | 26414677 |
| Age at Menarche | reproductive | -0.2274 | 0.023 | -9.90 | 4.1x10 ⁻²³ | -0.0987 | 0.0258 | -3.8247 | 0.0001 | 25231870 |
| Age of first birth | reproductive | -0.314 | 0.0259 | -12.12 | 8.1x10 ⁻³⁴ | -0.1597 | 0.0312 | -5.1217 | 3.0x10 ⁻⁷ | 27798627 |
| Insomnia | sleeping | 0.2512 | 0.0417 | 6.03 | 1.6x10 ⁻⁹ | 0.1602 | 0.0437 | 3.6705 | 0.0002 | 28604731 |
| Excessive daytime sleepiness | sleeping | 0.1513 | 0.0332 | 4.55 | 5.3x10 ⁻⁶ | 0.1136 | 0.0383 | 2.9646 | 0.003 | 27992416 |
| Chronotype | sleeping | 0.0696 | 0.0258 | 2.70 | 0.007 | 0.0541 | 0.0301 | 1.7961 | 0.07 | 27494321 |
| Sleep duration | sleeping | 0.0028 | 0.0402 | 0.07 | 0.9 | 0.0306 | 0.0405 | 0.7557 | 0.4 | 27494321 |
| Cigarettes smoked per day | smokingBehaviour | 0.3494 | 0.0676 | 5.17 | 2.4x10 ⁻⁷ | 0.2138 | 0.0627 | 3.4096 | 0.0007 | 20418890 |
| Ever vs never smoked | smokingBehaviour | 0.1626 | 0.0326 | 4.99 | 6.0x10 ⁻⁷ | 0.0641 | 0.0347 | 1.8478 | 0.06 | 20418890 |
| Age of smoking initiation | smokingBehaviour | -0.2022 | 0.0768 | -2.63 | 0.008 | -0.1532 | 0.083 | -1.8449 | 0.07 | 20418890 |
| Former vs Current smoker | smokingBehaviour | -0.3052 | 0.0525 | -5.81 | 6.1x10 ⁻⁹ | -0.1933 | 0.052 | -3.72 | 0.0002 | 20418890 |

SE: standard error, BMI: body mass index, HDL: high density lipoprotein, LDL: low density lipoprotein, ADHD: attention deficit hyperactive disorder, HOMA-IR: homeostasis model assessment of insulin resistance, HOMA-B: homeostasis model assessment of beta cell function.

iii. Supplementary Note

| COHORT | ACKNOWLEDGEMENTS |
|-------------------|---|
| DANISH-UCPH | The Novo Nordisk Foundation Center for Basic Metabolic Research is an independent Research Center at the University of Copenhagen partially funded by an unrestricted donation from the Novo Nordisk Foundation (www.metabol.ku.dk). Vejle Diabetes Biobank was supported by The Danish Research Fund and the National Danish Research Fund. |
| EGCUT | This study was funded by EU H2020 grants ePerMed and WIDENLIFE, Estonian Research Council Grant IUT20-60, IUT24-6, and European Union through the European Regional Development Fund Project No. 2014-2020.4.01.15-0012 GENTRANSMED. |
| FHS | Genotyping, quality control and calling of the Illumina HumanExome BeadChip in the Framingham Heart Study was supported by funding from the National Heart, Lung and Blood Institute Division of Intramural Research (Daniel Levy and Christopher J. O'Donnell, Principle Investigators). Also supported by National Institute for Diabetes and Digestive and Kidney Diseases (NIDDK) R01 DK078616 & U01 DK078616, NIDDK K24 DK080140 and American Diabetes Association Mentor-Based Postdoctoral Fellowship Award #7-09-MN-32, all to Dr. Meigs, and NIDDK Research Career Award K23 DK65978, a Massachusetts General Hospital Physician Scientist Development Award and a Doris Duke Charitable Foundation Clinical Scientist Development Award to Dr. Florez. |
| FUSION | The FUSION study was supported by DK093757, DK072193, DK062370, and ZIA-HG000024. |
| GCKD | The GCKD study was funded by the German Ministry of Research and Education (Bundesministerium für Bildung und Forschung, BMBF) and by the Foundation KfH Stiftung Präventivmedizin. Unregistered grants to support the study were provided by Bayer, Fresenius Medical Care and Amgen. Genotyping was supported by Bayer Pharma AG. |
| GENOA | Support for GENOA was provided by the National Heart, Lung and Blood Institute (HL054457, HL054464, HL054481, and HL087660) of the National Institutes of Health. |
| GERA | Data came from a grant, the Resource for Genetic Epidemiology Research in Adult Health and Aging (RC2 AG033067; Schaefer and Risch, PIs) awarded to the Kaiser Permanente Research Program on Genes, Environment, and Health (RPGEH) and the UCSF Institute for Human Genetics. The RPGEH was supported by grants from the Robert Wood Johnson Foundation, the Wayne and Gladys Valley Foundation, the Ellison Medical Foundation, Kaiser Permanente Northern California, and the Kaiser Permanente National and Northern California Community Benefit Programs. |
| GoDARTS | GoDARTS study was funded by The Wellcome Trust Study Cohort Wellcome Trust Functional Genomics Grant (2004-2008) (Grant No: 072960/2/03/2) and The Wellcome Trust Scottish Health Informatics Programme (SHIP) (2009-2012). (Grant No: 086113/Z/08/Z). |
| GoMAP and TEENAGE | GOMAP: This work was funded by the Wellcome Trust (098051). We thank all participants for their important contribution. We are grateful to Georgia Markou, Laiko General Hospital Diabetes Centre, Maria Emetsidou and Panagiota Fotinopoulou, Hippokratio General Hospital Diabetes Centre, Athina Karabela, Dafni Psychiatric Hospital, Eirini Glezou and Marios Mangioros, Dromokaiteio Psychiatric Hospital, Angela Rentari, Harokopio University of Athens, and Danielle Walker, Wellcome Trust Sanger Institute. TEENAGE: This research has been co-financed by the European Union (European Social Fund—ESF) and Greek national funds through the Operational Program “Education and Lifelong Learning” of the National Strategic Reference Framework (NSRF)—Research Funding Program: Heracleitus II. Investing in knowledge society through the European Social Fund. This work was funded by the Wellcome Trust (098051). We thank all study participants and their families as well as all volunteers for their contribution in this study. We thank the Sample Management and Genotyping Facilities staff at the Wellcome Trust Sanger Institute for sample preparation, quality control and genotyping. |

| | |
|----------------------------------|---|
| InterAct | The InterAct project (LSHM-CT-2006-037197) is a European-Community funded project under Framework Programme 6. We thank all EPIC participants and staff for their contribution to the study. We thank Nicola Kerrison (MRC Epidemiology Unit, Cambridge) for managing the data for the InterAct Project and staff from the Laboratory Team, Field Epidemiology Team, and Data Functional Group of the MRC Epidemiology Unit in Cambridge, UK, for carrying out sample preparation, DNA provision and quality control, genotyping, and data-handling work. |
| KORA | The KORA research platform (KORA, Cooperative Research in the Region of Augsburg) was initiated and financed by the Helmholtz Zentrum München – German Research Center for Environmental Health, which is funded by the German Federal Ministry of Education and Research and by the State of Bavaria. Furthermore, KORA research was supported within the Munich Center of Health Sciences (MC Health), Ludwig-Maximilians-Universität, as part of LMUinnovativ and by the German Center for Diabetes Research (DZD). |
| MESA | This research was supported by the Multi-Ethnic Study of Atherosclerosis (MESA and the MESA SHARE) projects and are conducted and supported by the National Heart, Lung, and Blood Institute (NHLBI) in collaboration with MESA investigators. Support for MESA is provided by contracts HHSN268201500003I, N01-HC-95159, N01-HC-95160, N01-HC-95161, N01-HC-95162, N01-HC-95163, N01-HC-95164, N01-HC-95165, N01-HC-95166, N01-HC-95167, N01-HC-95168, N01-HC-95169, UL1-TR-000040, UL1-TR-001079, UL1-TR-001420, UL1-TR-001881, and DK063491. Provision of exome chip genotyping was provided in part by support of NHLBI contract N02-HL-64278 and UCLA CTSI UL1-TR001881, and the S.Calif DRC DK063491. |
| METSIM | The METSIM study was supported by the Academy of Finland (contract 124243), the Finnish Heart Foundation, the Finnish Diabetes Foundation, Tekes (contract 1510/31/06), and the Commission of the European Community (HEALTH-F2-2007 201681), and the US National Institutes of Health grants DK093757, DK072193, DK062370, and ZIA- HG000024. |
| MGI | Our work was supported by NIH research grants HL117626 and HG007022. The Michigan Genomics Initiative was supported by internal research funds from the University of Michigan School of Public Health, the University of Michigan Medical School, and the University of Michigan President's Office. We are especially grateful to the generosity of all research participants. |
| Mount Sinai BioMe Biobank | The Mount Sinai BioMe Biobank is supported by The Andrea and Charles Bronfman Philantropies. |
| PIVUS and ULSAM | The PIVUS/ULSAM studies were supported by Wellcome Trust Grants WT098017, WT064890, WT090532, Uppsala University, Uppsala University Hospital, the Swedish Research Council and the Swedish Heart-Lung Foundation. |
| PROSPER | The PROSPER study was supported by an investigator initiated grant obtained from Bristol-Myers Squibb. Prof. Dr. J. W. Jukema is an Established Clinical Investigator of the Netherlands Heart Foundation (grant 2001 D 032). Support for genotyping was provided by the seventh framework program of the European commission (grant 223004) and by the Netherlands Genomics Initiative (Netherlands Consortium for Healthy Aging grant 050-060-810). |
| The Rotterdam Study | The authors are grateful to the Rotterdam Study participants, the staff involved with the Rotterdam Study, and the participating general practitioners and pharmacists. The Rotterdam Study is funded by Erasmus Medical Center and Erasmus University, Rotterdam, Netherlands Organization for the Health Research and Development (ZonMw), the Research Institute for Diseases in the Elderly (RIDE), the Ministry of Education, Culture and Science, the Ministry for Health, Welfare and Sports, the European Commission (DG XII), and the Municipality of Rotterdam. |
| WTCCC | Analysis and genotyping of these cohorts was supported by Wellcome Trust funding 090367, 098381, 090532, 083948, 085475, 101630 and 203141; MRC (G0601261), EU (Framework 7) HEALTH-F4-2007-201413, and NIDDK DK098032 and U01-DK105535. |

| | |
|---|---|
| UK Biobank | This study has been conducted using the UK Biobank Resource. |
| | |
| | |
| INDIVIDUAL | ACKNOWLEDGEMENTS |
| Andrew T Hattersley | AH is supported by a Wellcome Trust Senior Investigator award (Grant number 098395/Z/12/Z). |
| Anna L Gloyn | ALG is a Wellcome Trust Senior Fellow in Basic Biomedical Science. This work was funded by the Wellcome Trust (095101/Z/10/Z and 200837/Z/16/Z), National Institute for Health Research (NIHR) Oxford Biomedical Research Centre (BRC) and European Union Horizon 2020 Programme (T2D Systems). |
| Anna Köttgen and Matthias Wuttke | The work of AK and MW was supported by the German Research Foundation (KO 3598/3-1 to AK and CRC 1140). |
| Eleftheria Zeggini | Wellcome Trust (098051) |
| Erik Ingelsson | NIH/NIDDK 1R01DK106236-01A1 |
| James B Meigs | R01DK078616, U01DK078616, K24DK080140 |
| Jonathan Marchini | JM acknowledges support from the ERC (grant 617306). |
| Jose C Florez | JCF is a Massachusetts General Hospital Research Scholar. Parts of this work are supported by NIDDK U01 DK105554 and NIDDK K24 DK110550. |
| Josee Dupuis | R01DK078616, U01 DK078616 |
| Mark I McCarthy | MIM is a Wellcome Trust Senior Investigator (WT098381); and a National Institute of Health Research Senior Investigator. |
| Matthias Thurner | MT was supported by a Wellcome Trust Doctoral Studentship (099673/Z/12/Z). |
| Ruth J F Loos | RJFL is supported by the NIH (R01DK110113, U01HG007417, R01DK101855, R01DK107786). |