Supplementary Table S1. Comparison of Baseline Characteristics between the Selected Individuals for this Study and the Full Type 2 Diabetes Population in GoDARTS.

|  | Current Study <br> $(\mathrm{n}=5,152)$ | GoDARTS Type 2 Diabetes <br> Cases (n=8,698) |
| :--- | :---: | :---: |
| Age (years) | 65 | 67 |
| Male (\%) | 56.1 | 56.3 |
| Body Mass Index (kg/m2) | 32 | 31 |
| Smoking history (\%) | 51 | 63 |
| Systolic blood pressure (mmHg) | 77 | 141 |
| Diastolic blood pressure (mmHg) | 59.6 | 60.7 |
| Glycated Haemoglobin (mmol/mol) | 7.6 | 7.7 |
| Glycated Haemoglobin (\%) | 4.4 | 4.3 |
| Total Cholesterol (mmol/l) | 1.3 | 1.3 |
| HDL-Cholesterol (mmol/l) |  |  |

GoDARTS Type 2 diabetes baseline data taken from Hebert et al., 2018 (reference 20 in the manuscript)

Supplementary Table S2. Association between Retinal Risk Score and Major Adverse Cardiovascular Events Across Genetic and Clinical Risk Tertiles

| Adjusted Hazard Ratio (95\% CI) |  | $\mathbf{p}$ value |
| :--- | :---: | :---: |
|  |  | Interaction p value |
| Polygenic Risk Score Tertile |  |  |
| $\mathbf{1}$ | $1.17(1.03-1.32)$ | $\mathbf{0 . 0 1 4}$ |
| $\mathbf{2}$ | $1.08(0.97-1.22)$ | 0.17 |
| $\mathbf{3}$ | $1.08(0.97-1.19)$ | 0.15 |
| Pooled Cohort Equations Risk Score Tertile |  |  |
| $\mathbf{1}$ | $1.17(1.01-1.35)$ | $\mathbf{0 . 0 3 1}$ |
| $\mathbf{2}$ | $1.18(1.06-1.31)$ | $\mathbf{0 . 0 0 2}$ |
| $\mathbf{3}$ | $1.03(0.93-1.14)$ | 0.62 |
| Multivariable model included: age, sex, glycated haemoglobin, duration of diabetes, retinal risk score |  |  |
| PCE and CHD PRS. All hazard ratios per standard deviation increase. |  |  |

Supplementary Table S3. Area Under the Curve, Integrated Discrimination Improvement and Continuous Net Reclassification for Retinal Risk Score and CHD PRS versus the Pooled Cohort Equations ASCVD Risk Score for Prediction of Major Adverse Cardiovascular Events at 10 Years.

|  | Model 1 | Model 2 |  | Model 3 |  | Model 4 |  | Model 5 |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | PCE only | Retinal Risk | p | CHD PRS + | p | CHD PRS + | p | CHD PRS + | $p$ value |
|  | (baseline) | Score + Age + | value | Age + Sex | value | Retinal Risk | value | Retinal Risk |  |
|  |  | Sex |  |  |  | Score + Age + |  | Score + ASCVD |  |
|  |  |  |  |  |  | Sex |  | Risk Score + Age |  |
|  |  |  |  |  |  |  |  | + Sex |  |
| AUC | 0.658 | 0.663 | 0.33 | 0.683 | <0.001 | 0.686 | <0.001 | 0.690 | <0.001 |
|  | (0.640-0.677) | (0.645-0.681) |  | (0.665-0.701) |  | (0.667-0.704) |  | (0.672-0.708) |  |
| IDI (vs. | - | 0.001 | 0.79 | 0.018 | <0.001 | 0.019 | <0.001 | 0.023 | <0.001 |
| PCE only) |  | (-0.003-0.004) |  | (0.012-0.023) |  | (0.013-0.025) |  | (0.018-0.028) |  |
| Continuous | - | 0.080 | 0.024 | 0.216 | <0.001 | 0.240 | <0.001 | 0.319 | <0.001 |
| NRI (vs. |  | (0.010-0.150) |  | (0.147-0.285) |  | (0.171-0.309) |  | (0.251-0.388) |  |
| PCE only) |  |  |  |  |  |  |  |  |  |

$\overline{\text { AUC - Area Under the Curve; PCE - Pooled Cohort Equations Atherosclerotic Cardiovascular Disease Risk Score; PRS - Coronary Heart }}$ Disease Genetic Risk Score; IDI - Integrated Discrimination Improvement; NRI - Net Reclassification Index.

Bold indicates $\mathrm{p}<0.05$. All p values vs. Model 1.

Supplementary Table S4. List of Abbreviations.
AI - artificial intelligence
ASCVD - atherosclerotic cardiovascular disease
AUC - area under the curve
CHD - coronary heart disease
CRAE - central retinal arteriolar equivalent
CRVE - central retinal venular equivalent
CVD - cardiovascular disease
DRS - diabetes retinal screening
EHR - electronic health record
$\mathrm{FD}(\mathrm{a} / \mathrm{v})$ - fractal dimension (arterial/venous)
GoDARTS - Genetics of Diabetes Audit and Research Tayside Study
HbA1c - glycated haemoglobin
HR - hazard ratio
ICD - International Classification of Diseases
IDI - integrated discrimination index
MACE - major adverse cardiovascular events
MI - myocardial infarction
NRI - net reclassification index
PCE - pooled cohort equations
PRS - polygenic risk score
ROC - receiver operator characteristic
SNP - single nucleotide polymorphism
Tort (a/v) - tortuosity (arterial/venous)
VAMPIRE - Vascular Assessment and Measurement Platform for Images of the Retina

Supplementary Figure 1. Study Flowchart.

Cohort derivation.


Supplementary Figure 2. Correlations between Retinal Vascular Parameters and Continuous Clinical and Genomic Risk Factors.

Correlation matrix showing correlation coefficent (Pearson r) between Retinal Vascular Parameters and Continuous Clinical and Genomic Risk Factors.

Correlation coefficient

|  | Fractal <br> Dimension <br> (arterial) | Fractal <br> Dimension <br> (venous) | Tortuosity <br> (arterial) | Tortuosity <br> (venous) | CRAE | CRVE | Retinal <br> Risk <br> Score |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Age | -0.061 | -0.058 | -0.056 | 0.014 | -0.032 | -0.062 | 0.062 |
| SBP | -0.029 | 0.021 | -0.002 | 0.062 | -0.064 | -0.020 | 0.084 |
| DBP | 0.015 | 0.057 | 0.001 | -0.043 | -0.029 | 0.068 | -0.020 |
| HbA1c | -0.020 | 0.017 | 0.011 | 0.004 | 0.003 | 0.071 | 0.005 |
| TC | 0.026 | 0.042 | 0.004 | -0.008 | 0.012 | 0.055 | -0.025 |
| HDL | 0.008 | 0.032 | -0.009 | -0.006 | 0.027 | 0.003 | -0.020 |
| BMI | -0.025 | 0.012 | 0.049 | 0.029 | -0.002 | 0.037 | 0.031 |
| Diabetes <br> Duration | -0.037 | -0.009 | 0.013 | 0.062 | -0.044 | -0.052 | 0.074 |
| PCE | -0.067 | -0.049 | -0.038 | 0.034 | -0.068 | -0.055 | 0.094 |
| CHD PRS | -0.024 | -0.031 | 0.017 | -0.023 | 0.044 | 0.038 | -0.021 |

SBP - systolic blood pressure; DBP - diastolic blood pressure; HbA1c - glycated haemoglobin; TC total cholesterol; HDL - high-density lipoprotein cholesterol; BMI - body mass index; PCE - pooled cohort equations atherosclerotic cardiovascular risk score; CHD PRS - coronary heart disease polygenic risk score; CRAE - central retinal arterial equivalent; $C R V E$ - central retinal venous equivalent

Supplementary Figure 3. Association between Retinal Risk Score and Major Adverse Cardiovascular

Events.

Kaplan-Meier curve showing the association between tertiles of retinal risk score and time to first MACE incidence.


