

Supplementary Material

List of Abbreviations and Acronyms

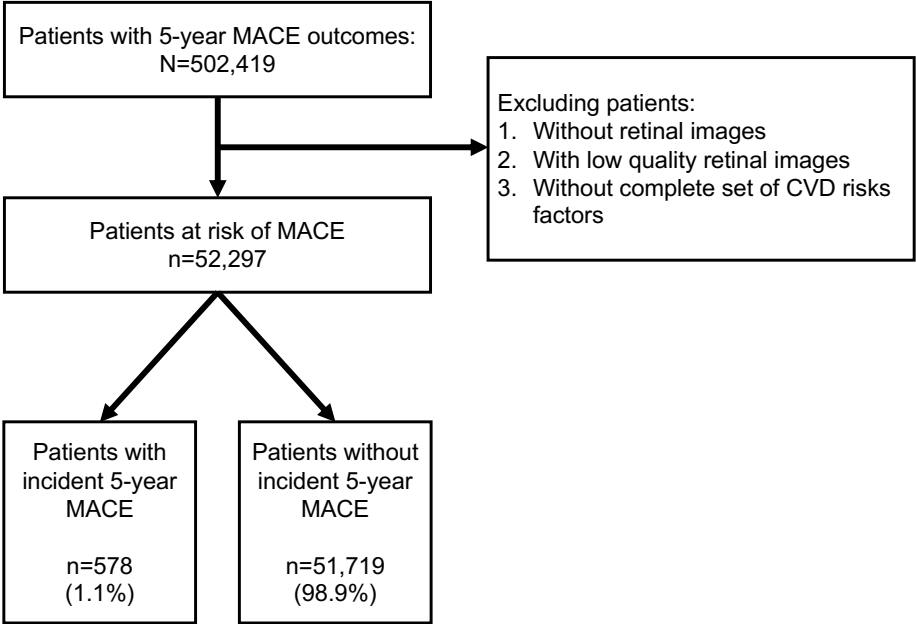
|          |                                      |
|----------|--------------------------------------|
| BMI      | Body Mass Index                      |
| CVD      | Cardiovascular Disease               |
| DBP      | Diastolic Blood Pressure             |
| SBP      | Systolic Blood Pressure              |
| LDL      | Low-Density Lipoprotein cholesterol  |
| HDL      | High-Density Lipoprotein cholesterol |
| Trig     | Triglycerides                        |
| HbA1c    | Hemoglobin A1c                       |
| MACE     | Major Adverse Cardiac Events         |
| SCORE    | Systematic COronary Risk Evaluation  |
| AUC      | Area Under Curve                     |
| PCE      | Pooled Cohort Equation               |
| FRA      | Framingham risk score                |
| DL       | Deep Learning                        |
| WHO      | World Health Organization            |
| IQR      | Interquartile Range                  |
| BCE Loss | Binary Cross Entropy Loss            |
| MSE      | Mean Squared Error                   |
| NRI      | Net Reclassification Index           |
| IDI      | Integrate Discrimination Index       |
| RGB      | Red Green Blue                       |

List of data augmentation in the training

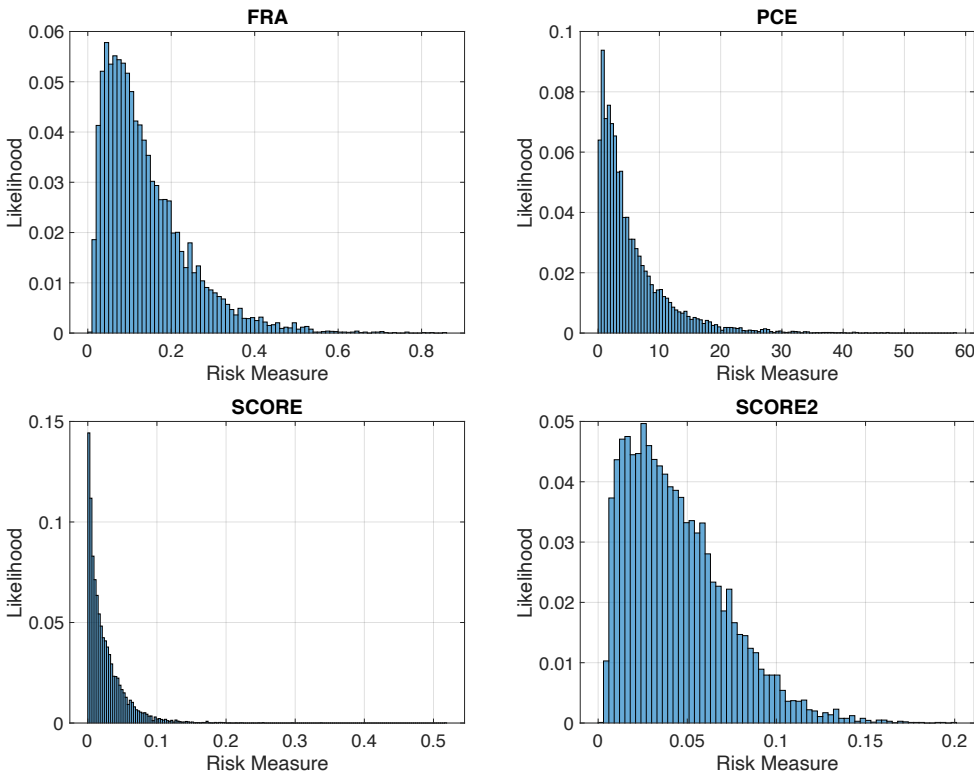
All data augmentation transformation are from Pytorch library.

1. Resize and center crop
2. Color jitter set to (0.2,0.2,0.2)
3. Random Affine transformation set to (180,(0.1,0.1),(0.8,1.2),5,0)
4. Random horizontal flip
5. Random Vertical flip
6. Random Grayscale
7. Gaussian blur set to (5, sigma=(0.1, 2.0))

Supplement Figure 1. Diagram illustrates the flow of data.



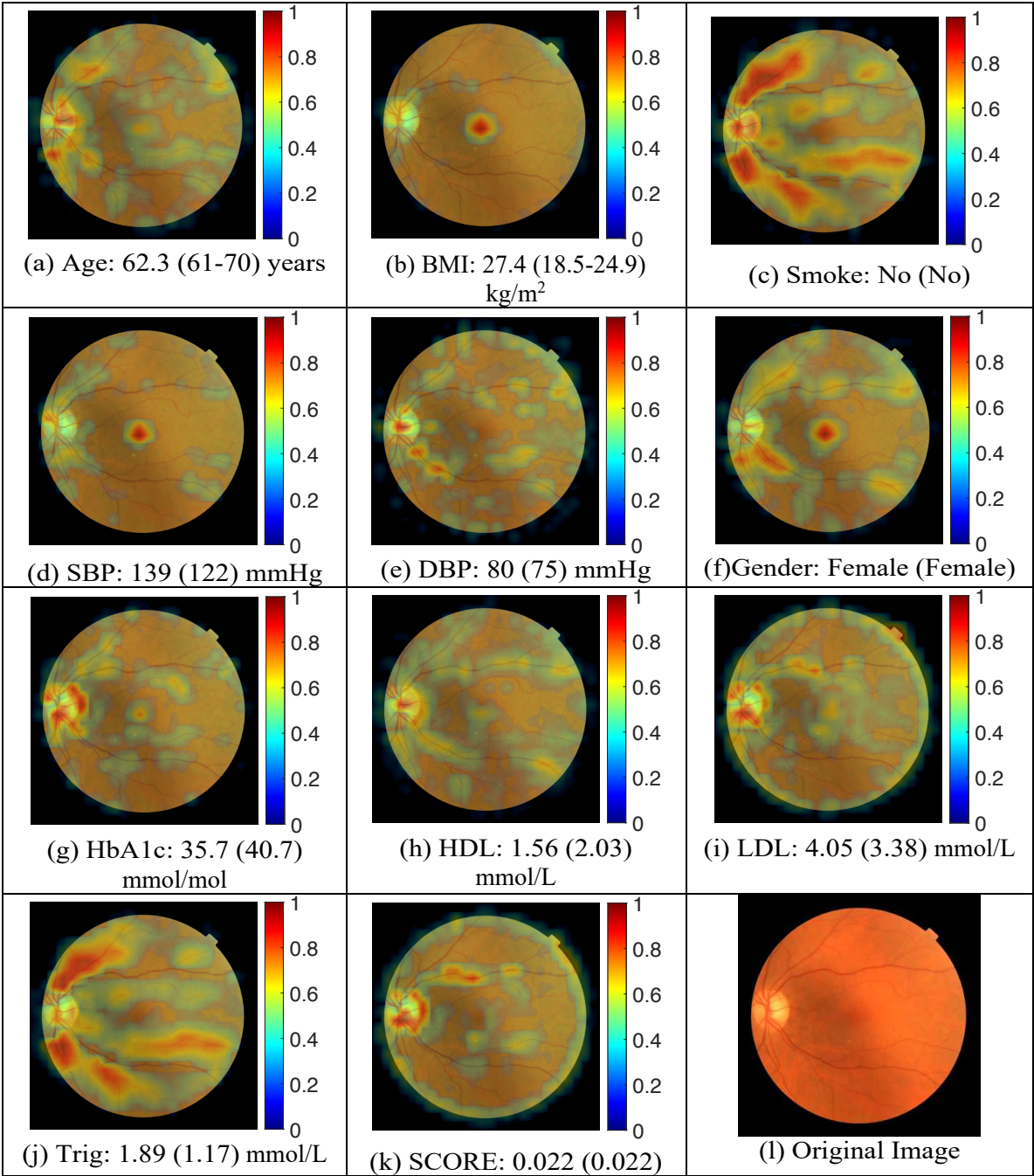
Supplement Figure 2. Distribution of PCE, FRS and SCORE, SCORE2



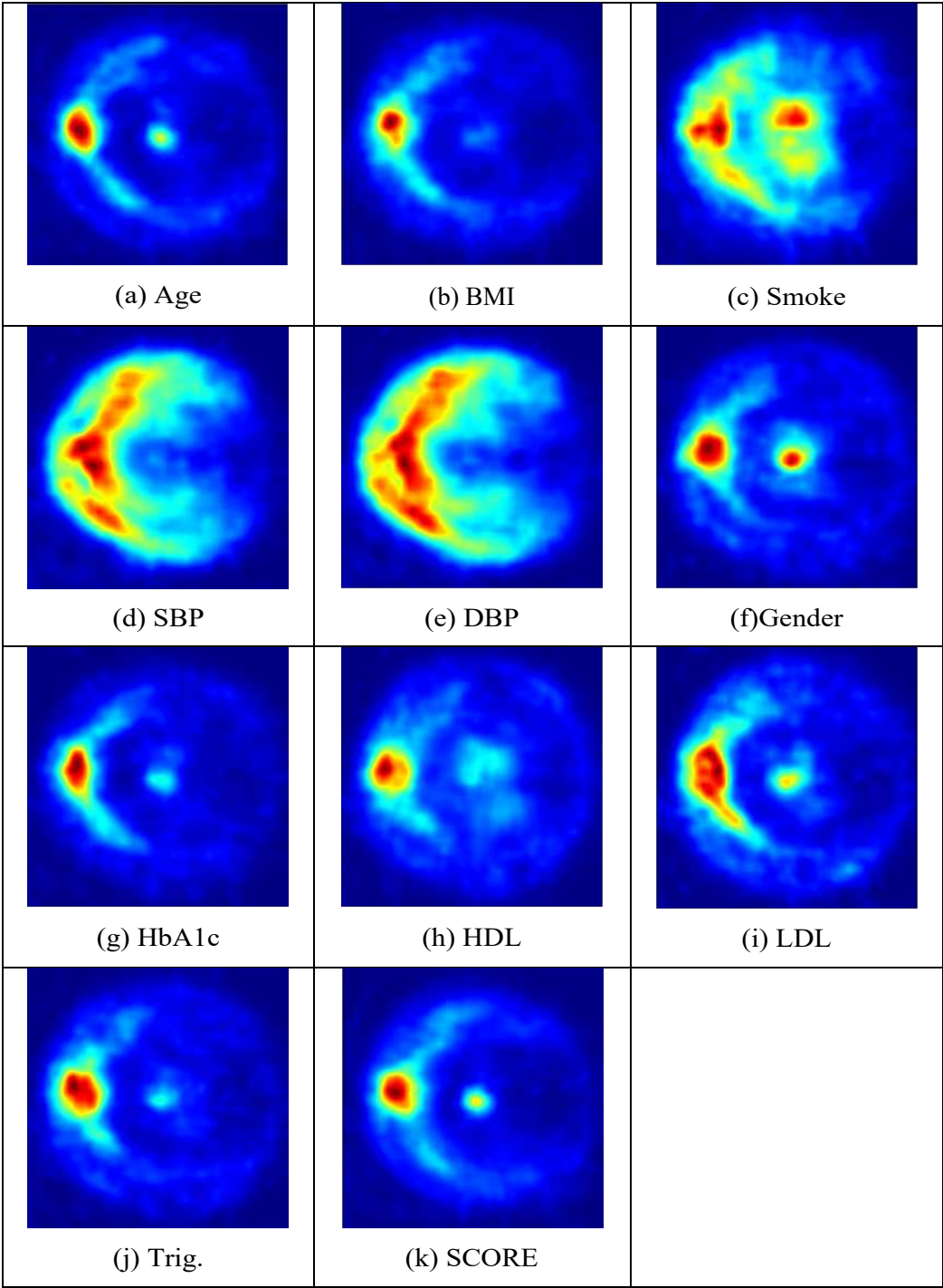
### Visualization of the model in the first stage on the retinal image

A modified version of IGOS<sup>18</sup> is implemented in this work (**Figures 3&4**), where the original IGOS algorithm only supports visualization for the classification task, but we are doing regression on CVD risk factors. The IGOS algorithm uses the regression results from our XMACE model and traces back to the image pixel that has a higher impact on the prediction (**Figure 3**). **Figure 4** shows a visualization of attention map aggregate from 3000 samples. From the visualization, we found the disk, fovea and the vessel took the most of attention in the risk prediction. Risk factors such as age, BMI, smoking, gender, HbA1c, HDL cholesterol, LDL cholesterol, and triglycerides have high level of attention at the optic disc and the fovea. Systolic and diastolic blood pressure were linked to the arcade retinal vessels. The statuses of the patient's health condition are visualized by a performance polygon diagram (**Figure 5 - 8**). In the figure, the severity of four health indicator: diabetes, hypertension, dyslipidemia, and obesity and the estimated smoke habit are included in the diagram.

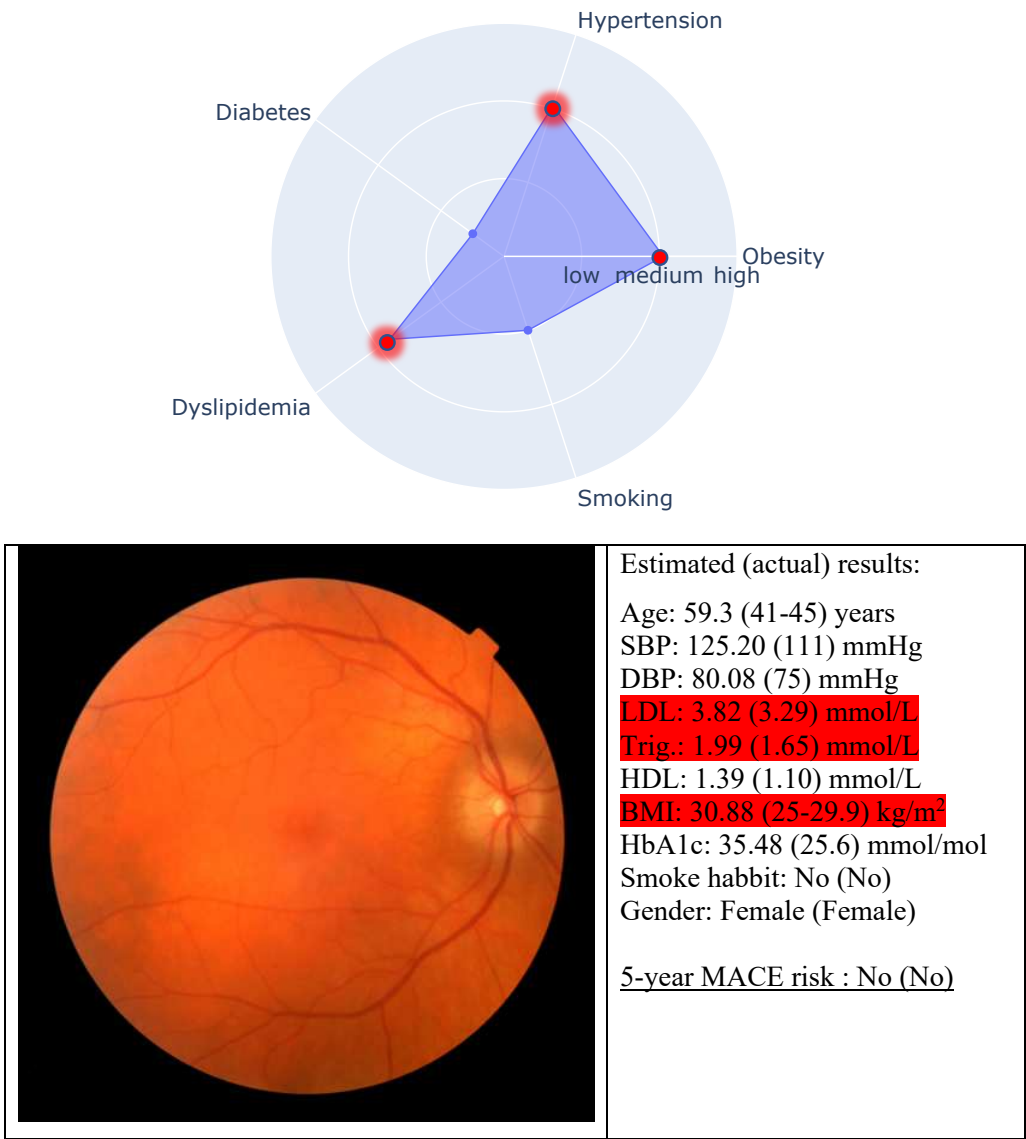
Supplement Figure 3. IGOS Visualization of attention map on features and the estimation value (actual value) of a single patient.



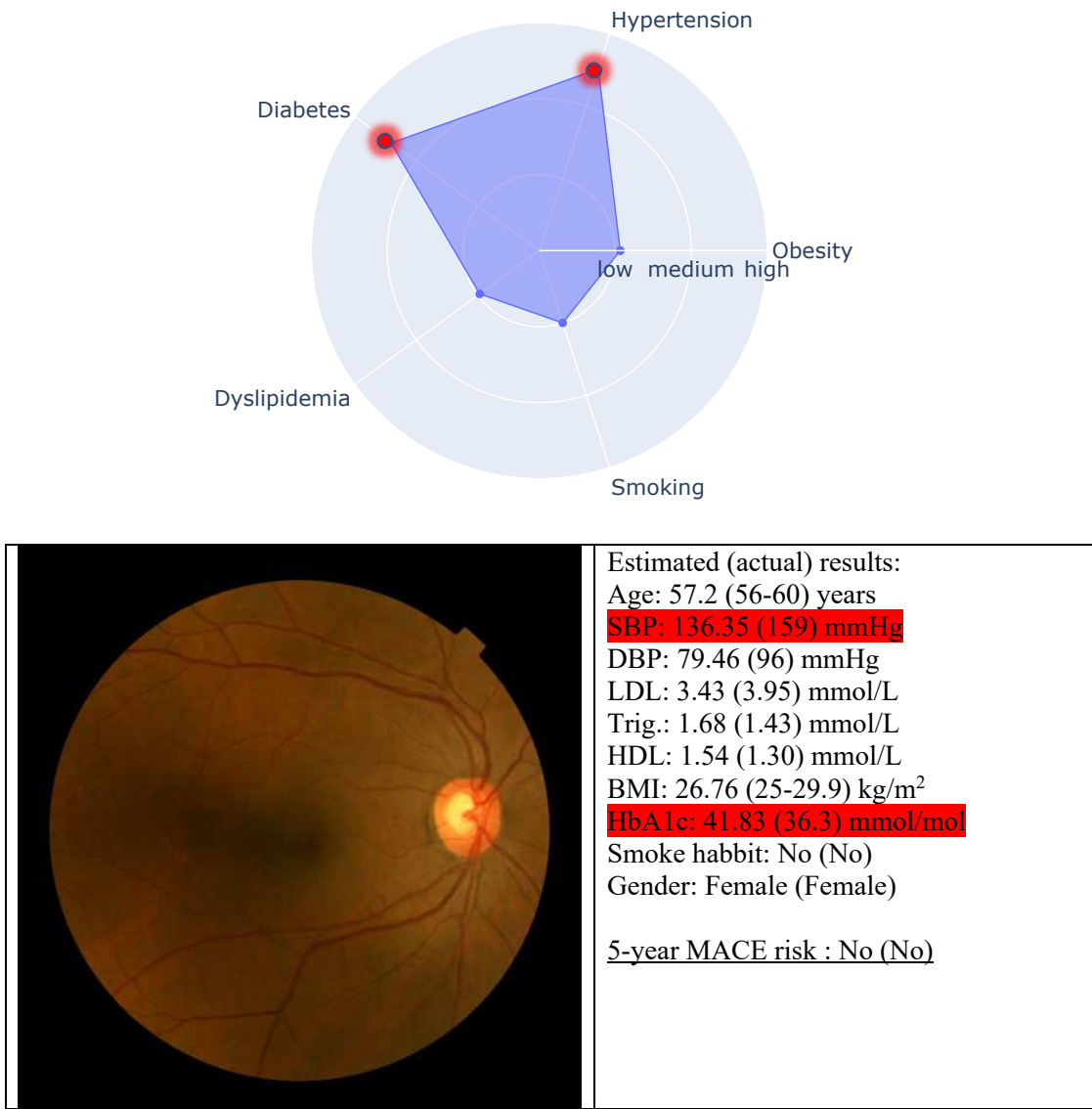
Supplement Figure 4. Aggregated attention map.



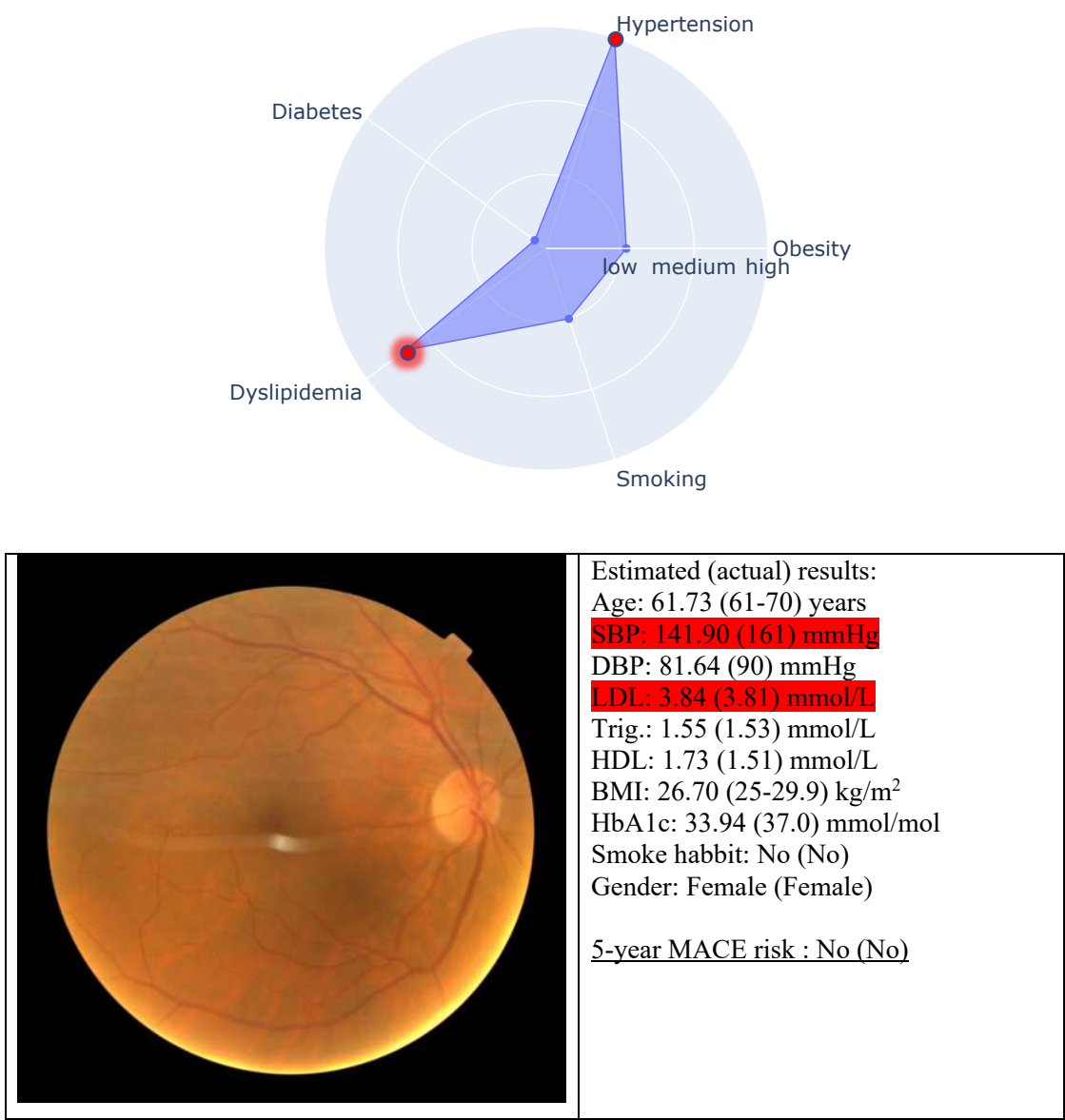
Supplement Figure 5. Cardiovascular disease (CVD) risk profiling based on the retinal image: a patient case with high risk of CVD due to obesity and dyslipidemia.



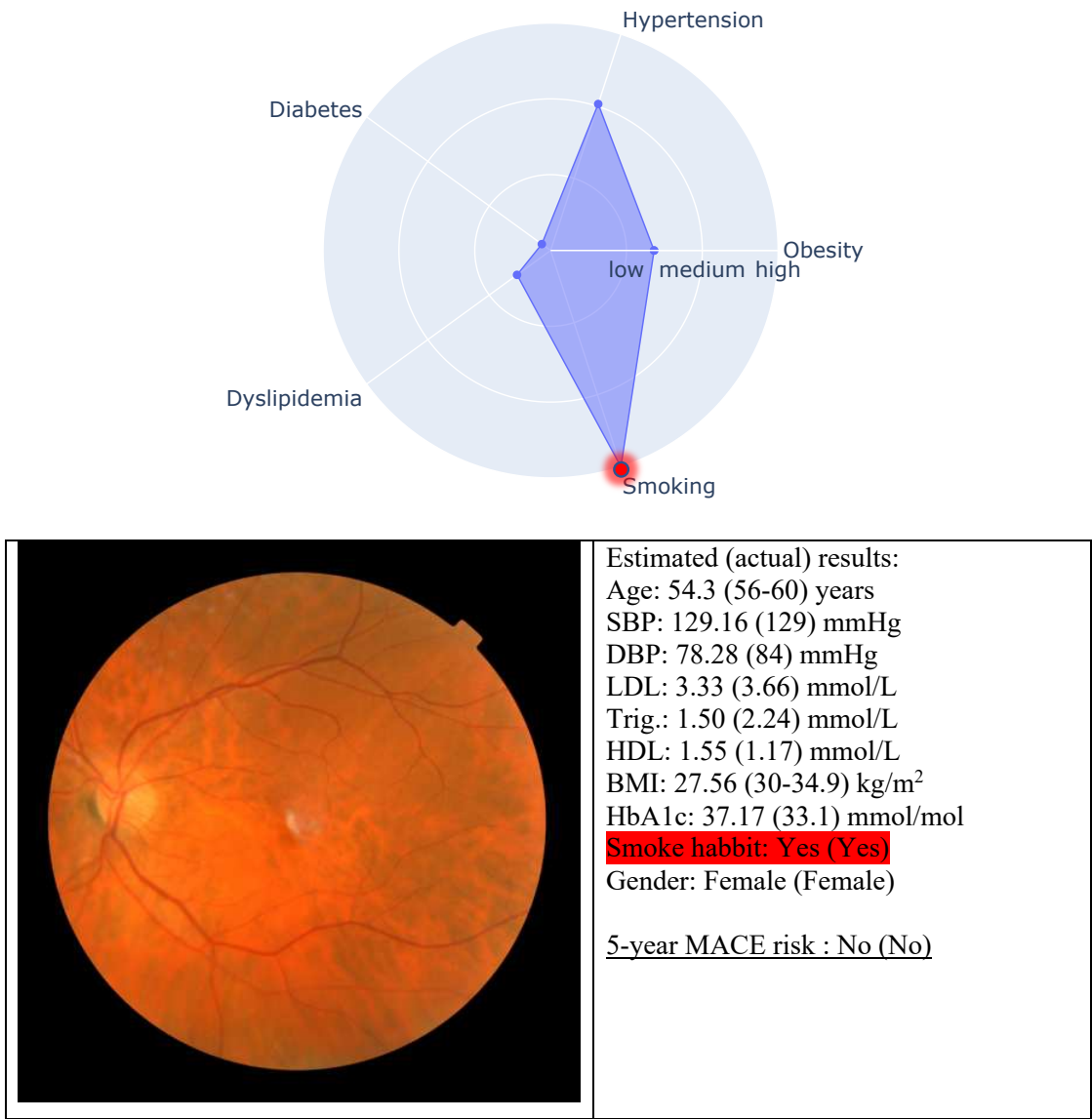
Supplement Figure 6. Cardiovascular disease (CVD) risk profiling based on the retinal image: a patient case with high risk of CVD due to diabetes and hypertension.



Supplement Figure 7. Cardiovascular disease (CVD) risk profiling based on the retinal image: a case with high risk of CVD due to hypertension and dyslipidemia.



Supplement Figure 8. Cardiovascular disease (CVD) risk profiling based on the retinal image: a case with high risk of CVD due to hypertension and dyslipidemia.



Supplementary Table 1: Classification of risk levels

| Health conditions | Obesity | Diabetes | Dyslipidemia      | Hypertension | Smoking |
|-------------------|---------|----------|-------------------|--------------|---------|
| Risk indicator    | BMI     | HbA1c    | Total Cholesterol | SBP          | Smoking |
| Low risks         | 22      | 38       | 200               | 120          | False   |
| Medium risks      | 27      | 43       | 220               | 125          | -       |
| High risks        | 35      | 48       | 240               | 130          | True    |

Supplementary Code 1: Code for PCE calculation

```
def PCE(age,gender,cholesterol,HDL,SBP,smoker,diabetes):
    Sex=gender+1
    SystolicBloodPressure=SBP
    Age=age
    Race=0
    TotalCholesterol=cholesterol
    HDLCholesterol=HDL
    Smoker=smoker
    Diabetes=diabetes
    LogitFemale = -12.823110 + (0.106501 * Age) + (0.432440 * Race) + (0.000056 *
    SystolicBloodPressure**2) + (0.017666 * SystolicBloodPressure) + (0.731678 *
    OnHypertensionMed) + (0.943970 * Diabetes) + (1.009790 * Smoker) + (0.151318 *
    (TotalCholesterol / HDLCholesterol)) + (-0.008580 * Age * Race) + (-0.003647 *
    SystolicBloodPressure * OnHypertensionMed) + (0.006208 * SystolicBloodPressure * Race)
    + (0.152968 * Race * OnHypertensionMed) + (-0.000153 * Age * SystolicBloodPressure) +
    (0.115232 * Race * Diabetes) + (-0.092231 * Race * Smoker) + (0.070498 * Race *
    (TotalCholesterol / HDLCholesterol)) + (-0.000173 * Race * SystolicBloodPressure *
    OnHypertensionMed) + (-0.000094 * Age * SystolicBloodPressure * Race)
    riskFemale = 100 / (1 + np.exp(-LogitFemale))
    LogitMale = -11.679980 + (0.064200 * Age) + (0.482835 * Race) + (-0.000061 *
    SystolicBloodPressure**2) + (0.038950 * SystolicBloodPressure) + (2.055533 *
    OnHypertensionMed) + (0.842209 * Diabetes) + (0.895589 * Smoker) + (0.193307 *
    TotalCholesterol / HDLCholesterol) + (-0.014207 * SystolicBloodPressure *
    OnHypertensionMed) + (0.011609 * SystolicBloodPressure * Race) + (-0.119460 *
    OnHypertensionMed * Race) + (0.000025 * Age * SystolicBloodPressure) + (-0.077214 *
    Race * Diabetes) + (-0.226771 * Race * Smoker) + (-0.117749 * Race * TotalCholesterol
    / HDLCholesterol) + (0.004190 * Race * OnHypertensionMed * SystolicBloodPressure) + (-
    0.000199 * Race * Age * SystolicBloodPressure)
    riskMale = 100 / (1 + np.exp(-LogitMale))
    risk = (np.abs(Sex == 1) * riskFemale) + (np.abs(Sex == 2) * riskMale)
    return risk
```

**Supplementary Code 2: Code for FRS**

```
class FraminghamScoreT():
    def __init__(self):
        self.score='Framingham Score'

    def __woman_CHD(self,age,totalcholesterol,HDL,SBP,smoker,diabetes):
        bx=2.32888*math.log(age)+1.20904*math.log(totalcholesterol) - \
        0.70833*math.log(HDL)+2.76157*math.log(SBP)+0.52873*smoker+0.69154*diabetes
        s=1-0.95012**math.exp(bx-26.1931)
        return s

    def __man_CHD(self,age,totalcholesterol,HDL,SBP,smoker,diabetes):
        bx=3.06117*math.log(age)+1.12370*math.log(totalcholesterol) - \
        0.93263*math.log(HDL)+1.93303*math.log(SBP)+0.65451*smoker+0.57367*diabetes
        s=1-0.88936**math.exp(bx-23.9802)
        return s

    def riskModel(self,age,gender,totalcholesterol,HDL,SBP,smoker,diabetes):
        risk=0
        if gender==1:
            risk=self.__man_CHD(age,totalcholesterol,HDL,SBP,smoker,diabetes)
        else:
            risk=self.__woman_CHD(age,totalcholesterol,HDL,SBP,smoker,diabetes)
        return risk
```

**Supplementary Code 3: Code for SCORE:**

```
class SCORE():
    def __init__(self):
        self.region='low risk'

    def __wRisk_CHD(self,cholesterol, SBP, smoker):
        beta_chol=0.24
        beta_sbp=0.018
        beta_smoker=0.71

        w=beta_chol*(cholesterol-6)+beta_sbp*(SBP-120)+beta_smoker*(smoker)
        return w

    def __wRisk_Non_CHD(self,cholesterol, SBP, smoker):
        beta_chol=0.02
        beta_sbp=0.022
        beta_smoker=0.63

        w=beta_chol*(cholesterol-6)+beta_sbp*(SBP-120)+beta_smoker*(smoker)
        return w
```

```
def __age_Model_note(self, age,alpha,p):
    return math.exp(-(math.exp(alpha))*((age-20)**p))

def __age_Men_CHD(self,age):
    alpha =-22.1
    p=4.71
    return self.__age_Model_note(age,alpha,p)

def __age_Women_CHD(self,age):
    alpha =-29.8
    p=6.36
    return self.__age_Model_note(age,alpha,p)

def __age_Men_Non_CHD(self,age):
    alpha =-26.7
    p=5.64
    return self.__age_Model_note(age,alpha,p)

def __age_Women_Non_CHD(self,age):
    alpha =-31.0
    p=6.62
    return self.__age_Model_note(age,alpha,p)

def __age_w_model_men_CHD(self,age,cholesterol,SBP,smoker):
    w=self.__wRisk_CHD(cholesterol, SBP, smoker)
    s0=self.__age_Men_CHD(age)
    s=s0**(math.exp(w))
    return s

def __age_w_model_men_Non_CHD(self,age,cholesterol,SBP,smoker):
    w=self.__wRisk_Non_CHD(cholesterol, SBP, smoker)
    s0=self.__age_Men_Non_CHD(age)
    s=s0**(math.exp(w))
    return s

def __age_w_model_women_CHD(self,age,cholesterol,SBP,smoker):
    w=self.__wRisk_CHD(cholesterol, SBP, smoker)
    s0=self.__age_Women_CHD(age)
    s=s0**(math.exp(w))
    return s

def __age_w_model_women_Non_CHD(self,age,cholesterol,SBP,smoker):
    w=self.__wRisk_Non_CHD(cholesterol, SBP, smoker)
    s0=self.__age_Women_Non_CHD(age)
    s=s0**(math.exp(w))
    return s

def riskModel(self,age,gender,cholesterol,SBP,smoker,years):
```

```

if gender==1:
    s=self.__age_w_model_men_CHD(age,cholesterol,SBP,smoker)
    sy=self.__age_w_model_men_CHD(age+years,cholesterol,SBP,smoker)
    srisk_CHD=1-sy/s
    s=self.__age_w_model_men_Non_CHD(age,cholesterol,SBP,smoker)
    sy=self.__age_w_model_men_Non_CHD(age+years,cholesterol,SBP,smoker)
    srisk_Non_CHD=1-sy/s
else:
    s=self.__age_w_model_women_CHD(age,cholesterol,SBP,smoker)
    sy=self.__age_w_model_women_CHD(age+years,cholesterol,SBP,smoker)
    srisk_CHD=1-sy/s

    s=self.__age_w_model_women_Non_CHD(age,cholesterol,SBP,smoker)
    sy=self.__age_w_model_women_Non_CHD(age+years,cholesterol,SBP,smoker)
    srisk_Non_CHD=1-sy/s

return srisk_CHD+srisk_Non_CHD

```

#### Supplementary Code 4: Code for SCORE2

```

class SCORE2():
    def __init__(self):
        self.region='low risk'

    def riskModel(self,age,gender,cholesterol,HDL,SBP,smoker,diabetes):
        self.cage=(age-60)/5
        self.csbp=(SBP-120)/20
        self.ctchol=(cholesterol-6)/1
        self.chdl=(HDL-1.3)/0.5
        self.smokewithAge=self.cage*smoker
        self.SBPwithAge=self.cage*self.csbp
        self.TotalCholesterolWithAge=self.cage*self.ctchol
        self.HDLwithAge=self.cage*self.chdl
        self.diabetewithAge=self.cage*diabetes

        if gender==1:

linearPred=0.3742*self.cage+0.6012*smoker+0.2777*self.csbp+0.6457*diabetes+0.1458*self
.ctchol \
        -0.2698*self.chdl-0.0755*self.smokewithAge-0.0255*self.SBPwithAge \
        -0.0281*self.TotalCholesterolWithAge+0.0426*self.HDLwithAge -
0.0983*self.diabetewithAge

        riskUncalibrated=1-math.pow(0.9605,linearPred)
        riskcalibrated2Low=1-math.exp(-math.exp(-0.5699+0.7476*math.log(-
math.log(1-riskUncalibrated))))
        else:

```

```
linearPred=0.4648*self.cage+0.7744*smoker+0.3131*self.csbp+0.8096*diabetes+0.1002*self
.ctchol \
    -0.2606*self.chdl-0.1088*self.smokewithAge-0.0277*self.SBPwithAge \
    -0.0226*self.TotalCholesterolWithAge+0.0613*self.HDLwithAge-
0.1272*self.diabetewithAge

    riskUncalibrated=1-math.pow(0.9776,linearPred)
    print(riskUncalibrated)
    riskcalibrated2Low=1-math.exp(-math.exp(-0.7380+0.7019*math.log(-
math.log(1-riskUncalibrated))))

    return riskcalibrated2Low
```