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Machine learning adds to clinical and CAC assessments in predicting 10-year CHD and CVD deaths

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Abstract

Objectives: We aimed to evaluate whether machine learning (ML) of non-contrast CT and clinical variables improve the prediction of atherosclerotic cardiovascular disease (ASCVD) and coronary heart disease (CHD) deaths as compared to coronary artery calcium (CAC) Agatston scoring and clinical data.

Background: The CAC score provides a measure of the global burden of coronary atherosclerosis, and its long-term prognostic utility has been consistently shown to have incremental value over clinical risk assessment. However, current approaches fail to integrate all available CT and clinical variables for comprehensive risk assessment.

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Methods: The study included data from 66,636 asymptomatic individuals (54±11 years, 67% Male) without established ASCVD undergoing CAC scanning and followed for CVD and CHD deaths at 10 years. Clinical risk assessment employed the ASCVD risk score. For ML we used an ensemble boosting approach to fit a predictive classifier for outcomes followed by automated feature selection using information gain ratio. The model building process used all available clinical and CT data, including the CAC score, the number, volume and density of CAC plaques, and extracoronary scores, comprising a total of 77 variables. We evaluated our overall proposed model (ML all) using a 10-fold cross-validation framework on the population data and area under the curve (AUC) as metrics. The prediction performance was also compared with two traditional scores (ASCVD risk and CAC score) and two additional models that were trained using all the clinical data (ML Clinical) and CT variables (ML CT).

Results: AUC by ML All (0.845) for predicting CVD death was superior compared to that obtained by clinical data alone (0.821), CAC score alone (0.781) and ML-CT alone (0.804) ($p<0.001$ for all). Similarly, for predicting CHD death, AUC by ML All (0.860) was superior to the other analyses (0.835 for clinical data, 0.816 for CAC, and 0.827 for ML-CT, $p<0.001$).

Conclusions: The comprehensive ML model was superior to clinical risk factors, CAC scores, and a ML model fitted using CT variables alone in prediction of both CVD and CHD deaths.

Condensed Abstract:

The study included data from 66,636 asymptomatic individuals without established atherosclerotic cardiovascular disease (ASCVD) undergoing coronary artery calcium (CAC) scanning and followed for cardiovascular disease (CVD) and coronary heart disease (CHD) deaths at 10 years. Comprehensive machine learning (ML) used 77 clinical and CT variables, including: the number, volume and density of CAC plaques, CAC and extracoronary scores, among others. Risk estimation by ML was superior to the current traditional risk equation and CAC score for prediction of CVD and CHD deaths and demonstrated high concordance between ML-predicted and actual observed risk for both CVD and CHD deaths.

Keywords

Machine learning; Coronary artery calcification; Coronary heart disease death; Cardiovascular disease death; Pooled cohort equation

Introduction

In clinical practice traditional risk factors obtained from population-based studies are used to predict cardiovascular disease (CVD) events. For example, the 2013 ACC/AHA atherosclerotic cardiovascular disease (ASCVD) risk estimator has improved risk assessment of cardiovascular diseases compared to previous risk algorithms¹. However, current clinical models still misclassify future risk assessment (1–4). Several studies, for instance, have shown that the ASCVD risk estimator and all other current risk scores overestimate actual observed risk (1–4).

Coronary artery calcification is a robust marker of coronary atherosclerosis. The coronary artery calcium (CAC) score measured by non-contrast cardiac-gated computed tomography

(CT) provides a measure of the global burden of coronary atherosclerosis, reflecting the effect all measured and unmeasured risk factors causing coronary atherosclerosis in an individual patient. Its long-term prognostic value has been shown consistently to provide independent predictive information to assess clinical risk of CVD and coronary heart disease (CHD) events^(5–9). In addition, other CT variables such as the total number of calcified coronary lesions, plaque density and thoracic aorta calcification have been demonstrated to add to CAC assessment in prediction of CVD events (10,11). However, current models fail to integrate all available CT and clinical variables for comprehensive risk assessment.

Machine learning (ML) is the scientific field that enables data-driven predictions by learning from data. ML builds models that can learn from training samples to subsequently perform prediction tasks in unseen samples. ML techniques have showed equal or better performance than humans in medical tasks such as diagnosis, decision-making and risk prediction in cardiology (12–16). This is the first study, to our knowledge, to assess the prognostic value of ML to estimate CVD and coronary heart disease (CHD) deaths among asymptomatic individuals integrating clinical and CAC data. We hypothesized that comprehensive ML of CAC and other variables from non-contrast cardiac CT can better predict CHD and CVD deaths than current state-of-the art methods for risk prediction. The aim of the current study was then to evaluate whether ML, considering all available clinical and cardiac CT imaging variables, predicts CVD and CHD deaths more accurately than existing assessments.

Methods

Study population

The CAC Consortium is a large multicenter observational cohort study of patients who have undergone CAC scanning for clinical purposes and is designed to determine the cause-specific death including CVD, CHD and non-CVD deaths.

Details regarding the CAC consortium have been described previously (17). The CAC consortium includes 66,636 asymptomatic individuals (54 ± 11 years, 67% Male) without known CHD who non-contrast cardiac CT for detecting CAC at 4 high volume centers in the U.S (Harbor UCLA Medical Center, Torrance, California; Cedars Sinai Medical Center, Los Angeles, California; Columbus, Ohio; and Minneapolis Heart Institute, Minneapolis, Minnesota). All sites had at least 10 years' experience to exam CAC scanning, provided >5000 scans per site and can complete >90% of clinical demographics which were required for the study. Inclusion criteria were patients with ≥ 18 years old, asymptomatic, no history of CHD, and who underwent CAC scanning. Exclusion criteria were missing data of scan identifiers (n=2650), no-dedicated CAC score (n=4669), no-CAC scanning (n=4833), uncertain date of birth (n=150), uncertain data of scan (n=11), or insufficient data for follow-up (n=10,320).

Each institution obtained Institutional Review Board approval and all participants provided informed consent.

Clinical demographics

Clinical demographics and laboratory data were collected at the time of CAC scanning or at a clinical visit associated with the scan. Hypertension, diabetes and dyslipidemia were defined when individuals self-reported diagnosis made by their physicians, had testing at the time of the scan visit, or had been treated by medications for these diseases. Dyslipidemia was also defined when LDL-C >160 mg/dL, HDL-C <40 mg/dL in men and <50 mg/dL in women, or fasting triglycerides >150 mg/dL were present. Never, former, or current smoking was recorded for smoking status. Family history of CHD was defined as premature family history (<55 years in old in a male relative and <65 years old in a female relative) at the Columbus, Ohio site, or the presence of a first-degree relative with a history of premature CHD at other 3 sites. The ASCVD risk score was calculated by using the PCE (17). Total cholesterol and high-density lipoprotein (220 mg/dL and 40mg/dL for patients with untreated dyslipidemia; 190mg/dL and 60mg/dL for patients without dyslipidemia; 180mg/dL and 50mg/dL for dyslipidemic patients with treatment) were used to calculate ASCVD risk score. When historical risk-factors did not include measurements regarding blood pressure, 150mmHg and 90mmHg were used as systolic and diastolic blood pressure for hypertensive patients without treatment. 135 mmHg and 85mmHg for hypertensive patients with treatment, and 120mmH and 80mmHg for patients without hypertension were used for calculating ASCVD risk score.

Study Follow-up

Death was defined by patient identifiers including social security number, name, date of birth through the Social Security Death Index (SSDI) Death Master File and followed through June 1st 2014. Cause of death was determined by coded death certificates through the National Death Index service. The maximum follow-up time was truncated at 10 years to investigate actual 10 years CVD/CHD risk prediction by ML.

CT protocol and interpretation of CAC

CAC scans were performed in accordance with standard protocols (18). Because of the current study nature to investigate >10-year death for patients, electron beam tomography was obtained in approximately 93% of patients. In total, approximately 13%, 38%, 38% and 3.5% of patients were scanned with the Imatron C-100 scanner, the C-150, the C-300, and the e-Speed scanner (GE-Imatron), respectively. More recent data at two sites was collected using multidetector CT in 7% of patients on a 4-slice MDCT scanner (Somatom Volume Zoom, Siemens Medical Solutions) and the General Electric LightSpeed VCT 64-slice platform (GE Healthcare). Due to the long-term follow-up, most of the scans (>90%) were performed by electron beam tomography and the rest were scanned by multidetector CT. CAC and extracoronary calcification including thoracic aortic calcification (TAC), aortic valve calcification (AVC) and mitral valve calcification (MVC) were scored using Agatston method (19). Besides, CAC scores as well as volume score and mean CAC densities for left main and other main three vessels were also available. Additional information regarding CAC including total number of CAC plaques, CAC volume scores, CAC density, TAC scores, AVC scores and MVC scores were available in 68%, 51%, 30%, 51%, 15% and 15% of the cohort, respectively.

Machine learning

Figure 1 illustrates the steps followed to train and evaluate the proposed model (ML all) using a 10-fold cross validation framework in the study population. First, the overall population was randomly divided into 10 equally sized non-overlapping groups. One group containing 10% was retained as the test set and the other 90% were used as the training set; second, a feature selection was performed using the training set and information gain; third, a data-driven model was fitted using the training set and an ensemble boosting approach (LogitBoost); fourth, the prediction performance was evaluated using the test set. The cross-validation procedure then looped 10 times over the various groups, each time performing variable selection and model building, and using different training and test sets - meaning that none of the data points were used for model training and evaluation at the same time. We used this validation procedure seeking to maximize the use of training and validation data, avoid the testing of hypothesis suggested by arbitrary splitting of data, and reduce the variance in prediction error. Once finished, as fifthly step, the predictions of the corresponding 10 models were stacked to assess the overall prediction performance of CHD and CVD deaths.

Variable selection—A total of 77 variables, including 46 clinical variables (e.g. ASCVD risk score, age, sex, race, body mass index, hypertension, diabetes, hyperlipidemia, current smoking, family history of CHD, smoking years, and medication information) and 31 CT variables (derived from CAC scans) (Table 2), were available to train the model. We firstly used information gain to select the best attributes for the classifier using those variables that resulted in an information gain $> 1e-5$. Information gain is a measure of the amount of information gained from the data by attribute (20,21).

Model building—We used an ensemble boosting approach to fit a predictive classifier for cardiovascular outcomes (22). This boosting method called LogitBoost is tree-based learning technique that combines the predictions of many weak classifiers to produce a single powerful prediction: A weak learner is fit in each iteration seeking to reduce the misclassification error of previous iterations. For a given patient, the outcome of ML model - called the ML score - was then the probability risk of having CVD and CHD mortality. It is also worth adding that this technique is suitable to deal with missing data (21). This technique uses non-missing data to establish a ranking of surrogate variables: The first surrogate is the feature that best describe the training data while the second surrogate does the second-best description, and so on. It then imputes missing data, either in the training or test phase, using the ranking of surrogate variables in order, if the first surrogate variable is missing. ML and feature selection were implemented in the open-source Waikato Environment for Knowledge Analysis (WEKA) platform 3.8.0 (University of Waikato, Hamilton, New Zealand).

Prediction models—We trained two additional ML models to compare to our proposed model (ML All) following the steps previously described: 1) a first model trained with all the clinical variables (ML Clinical); and 2) a second model trained with all CT variables (ML CT). These two ML models were trained and evaluated using the same folds and cross-validation procedure followed for the ML All model to subsequently enable paired

comparisons. We also compared our ML ALL model with a logistic regression model (LR-3F) trained with age, ASCVD and CAC scores to predict both CHD and CVD deaths. This analysis was done to determine the benefit of combining all variables using ML technique such as LogitBoost.

Statistical analyses—We compared the prediction performance of our ML model with the ASCVD risk and CAC score: using AUC as metric to evaluate the overall performance of the ML models and traditional scores. Youden index was also provided to summarize performance predictions. Pairwise comparisons were performed between the ASCVD risk score, CAC score alone, and ML models using DeLong test (23). The ML models were assessed by sex and the Brier scores were computed between predicted and observed CVD and CHD deaths (24). Additionally, ML All model was compared with traditional scores in single random stratified partitioning of our population data. Statistical calculations were performed in R software version 3.4 using the pROC package for DeLong analysis (25).

Results

Table 1 summarizes clinical characteristics of patients in the current study. Mean age was 54 ± 11 years and 67% were males. Most of the study cohort were white. The mean 10-year ASCVD risk score was $7.4 \pm 8.9\%$. Hyperlipidemia and family history of CHD were the most common cardiovascular risk factors, following hypertension, current smoking and diabetes, respectively.

The endpoints of the current study are CHD death ($n=524$), and CVD death ($n=971$) including death from CHD ($n=524$, 54%), stroke ($n=160$, 17%), congestive heart failure ($n=51$, 5%), and other circulatory disease ($n=236$, 24%). The variable rankings (first 50 variables) for prediction of CVD and CHD deaths are listed in Figure 2 (Figure 2a: CVD death, and Figure 2b: CHD death). ASCVD risk score was the feature that obtained the highest gain for predicting both types of deaths, followed by age and CAC score. Similarly, these variables were the most used to train the weak classifiers in the ML All model for predicting both types of risk.

AUCs for predicting the 10-year CVD and CHD deaths are shown in Central Illustration (a and b respectively). For CVD death, AUC for ML Clinical was significantly higher than ML for ASCVD risk alone (AUC; 0.826 vs. 0.821, $p < 0.001$) and ML CT significantly improved the prediction compared to CAC alone (0.804 vs. 0.781, $p < 0.001$). In addition, AUC for ML All (0.845) was higher than that for ML clinical (0.826, $p < 0.001$) and ML CT (0.804, $p < 0.001$) (Central Illustration-a). With respect to CHD death, AUCs for ML Clinical and ASCVD risk alone were comparable (0.835 vs. 0.834, $p = 0.639$). ML CT significantly improved the prediction compared to CAC alone (0.827 vs. 0.816, $p = 0.025$). AUC for ML All (0.860) was higher than that for ML Clinical ($p < 0.001$) or ML CT ($p < 0.001$) (Central Illustration-b). The separate results in male and female cohorts are provided in Supplemental Figure 1. Tables 3 and 4 provide the Youden index for each model and traditional scores, where our ML All model obtained the highest Youden index for both CVH and CHD deaths.

A comparative plot of observed and ML predicted risks is shown in Figure 3. An excellent Brier score (<0.05) was shown for prediction of CVD and CHD deaths, resulting in precise ML-prediction of CVD (Figure 3a) and CHD death (Figure 3b) compared to observed deaths.

In the sub-analyses to investigate if ML ALL improved the prognostic predictions compared to conventional analysis, ML ALL improved the prognostic prediction of a logistic regression model trained with age, ASCVD and CAC scores (LR-3F) (Supplemental Figure 2).

When the ML All model was compared with traditional scores in single random stratified partitioning of the population data into training (80%) and test (20%) sets, we obtained the same tendency of variable rankings and performance predictions for both CVD and CHD death (Supplemental Figure 3) and the AUC for ML was higher than that for ASCVD risk and CAC score (Supplemental Figure 4).

Discussion

In a large multicenter cohort of 66,636 asymptomatic individuals undergoing clinical CAC scans, we have showed that ML models that integrates all clinical and non-contrast CT imaging variables can obtain superior prognostic performance than traditional scores to predict both CVD and CHD deaths. In this study, we developed a ML method to show the added prognostic value of integrating clinical and non-contrast CT imaging variables, providing a ranking of the most significant variables to predict CVD and CHD deaths. Our comprehensive ML model obtained high concordance between the ML risk score and actual observed risk for both CVD and CHD deaths, suggesting prospective clinical implementations of ML models to assess the risk for both types of deaths. We have demonstrated that age, ASCVD risk and CAC score showed highest gains to predict events and that additional variables provide significant, incremental value in the overall prediction performance.

The standard approach to analyze the predictive value of CAC scanning along with clinical information has been performed on a limited number of variables. The ASCVD risk score, for example, is a parsimonious score based on a handful of these available clinical variables. Also, prognosis with CAC scanning is typically assessed alone without consideration of variables such as CAC score in individual vessels, number of plaques, or extracoronary calcification. While a comprehensive method for integrating all available variables from the CAC scan has been advocated (26), such a method has not been developed. Further, a comprehensive integration of all available clinical information with a more complete analysis of variables from non-contrast CT has also not been proposed. In the current study, we developed a ML method that integrating clinical and non-contrast cardiac CT imaging data outperformed the traditional scores and single-variable-type ML models on the prognostic prediction for CVD and CHD death: Similarly by gender, we showed that our ML method outperformed the prognostic prediction in females and males for CVD death. In the current study the prognostic improvement of the ML method was significant for CHD death in males but not in females. The reason for this finding is not clear. A prior study from

the CAC Consortium revealed that after adjustment for ASCVD risk, CAC scores added significantly prognostic accuracy in both males and females for prediction of CVD death (reports for CHD death were not made) (27).

Conventional risk assessment involves classification of patients into few predefined risk categories. Without precise quantitative estimates, however, crude risk categorization with arbitrary thresholds may misclassify patients compared to continuous risk predictions (28). In contrast, ML can give a precise risk calibration for specific patient. Recent studies have motivated the development of comprehensive risk assessment by ML with imaging based on coronary CT angiography or myocardial single photon emission CT (29,30). In this study, we used a similar method to maximize non-contrast CT information for cardiovascular risk prediction in a large asymptomatic population. Our findings suggest that future risk prediction models based on all available information can achieve a more accurate and precise model to identify risk that could be implemented clinically to improve the clinical use of CAC scanning in risk assessment and guiding management decisions (14,15,30). The ML approach is likely to become a routine tool for risk assessment using CAC scanning with the evolution of the electronic medical record and its integration with imaging data in future clinical practice.

Limitations

Our study has some limitations. Although the available CT data contained several variables in addition to the CAC score, multiple CT variables were not available in all patients, introducing uncertainties that may affect the prediction performance of the ML model. Additional variables of prognostic importance such as epicardial adipose tissue were not contained in the database. Our outcome variables were CVD and CHD deaths, since the CAC Consortium database did not contain regarding nonfatal cardiac events. CHD and CVD deaths overlap, with the CHD deaths being included in the latter. By including the latter, we are able to show that CAC is as predictive for both CVD and CHD events. There were a relatively small number of CHD deaths in females. This may possibly explain that the improvement did not reach statistical significance in AUC of ML All over ML Clinical for CHD death in females. We used CVD and CHD deaths for the primary outcomes in the current study since the CAC Consortium was designed to determine cause-specific death including CVD, CHD and non-CVD deaths among asymptomatic patients undergoing CAC.

Conclusion

The current study demonstrated that a ML approaches that integrate clinical and non-contrast CT variables can provide better risk assessment of CVD and CHD death than the combination of the traditional ASCVD and CAC scores. As clinical data from digital medical records become available for seamless integration with imaging data, the ML approach is likely to become a routine tool for clinical risk assessment using CAC scanning in the future.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

CAC	Coronary artery calcium score
ML	Machine learning
ASCVD	Atherosclerotic cardiovascular disease
CVD	Cardiovascular disease
CHD	Coronary heart disease
PCE	Pooled cohort equation
LR	Logistic regression

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Clinical Perspectives:**COMPETENCY IN MEDICAL KNOWLEDGE:**

The integration of clinical and non-contrast CT imaging variables into ML methods enables to develop risk models that have superior prognostic prediction of CVD and CHD death than the traditional clinical risk scores such as ASCVD and CAC scores.

TRANSLATIONAL OUTLOOK:

Breakthroughs in electronic medical records and integration systems with imaging workstations can facilitate the implementation of ML models as a routine tool for clinical risk assessment using clinical and CAC scanning in the future.

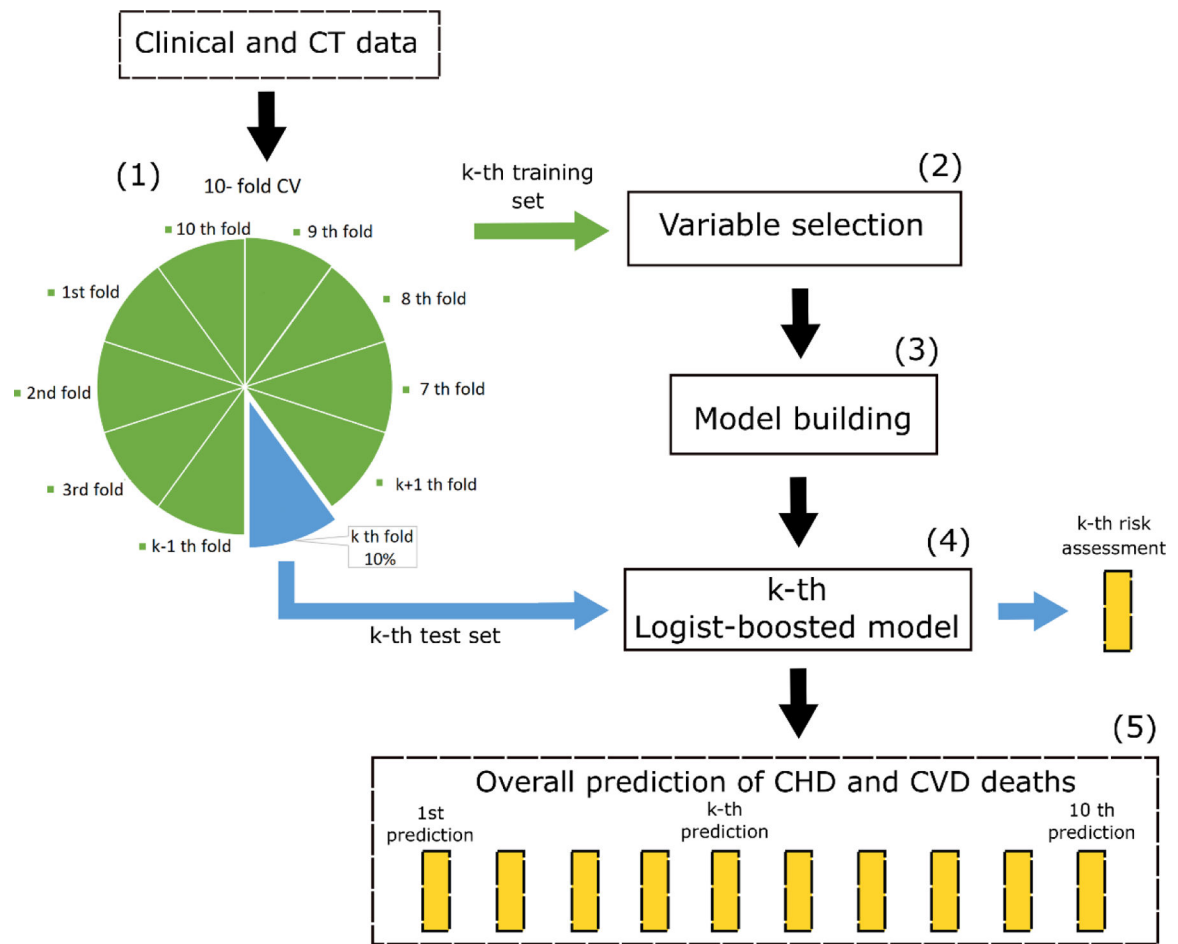


Figure 1. Workflow of the method used to train the proposed model.

- 1) Random split of population in 10 folds, 2) Selection of variables using information gain,
- 3) k-th model building using ensemble boosting, 4) Evaluation of prediction performance of k-th model, 5) evaluation of overall prediction of CHD and CVD events.

Variable ranking for prediction of CVD death

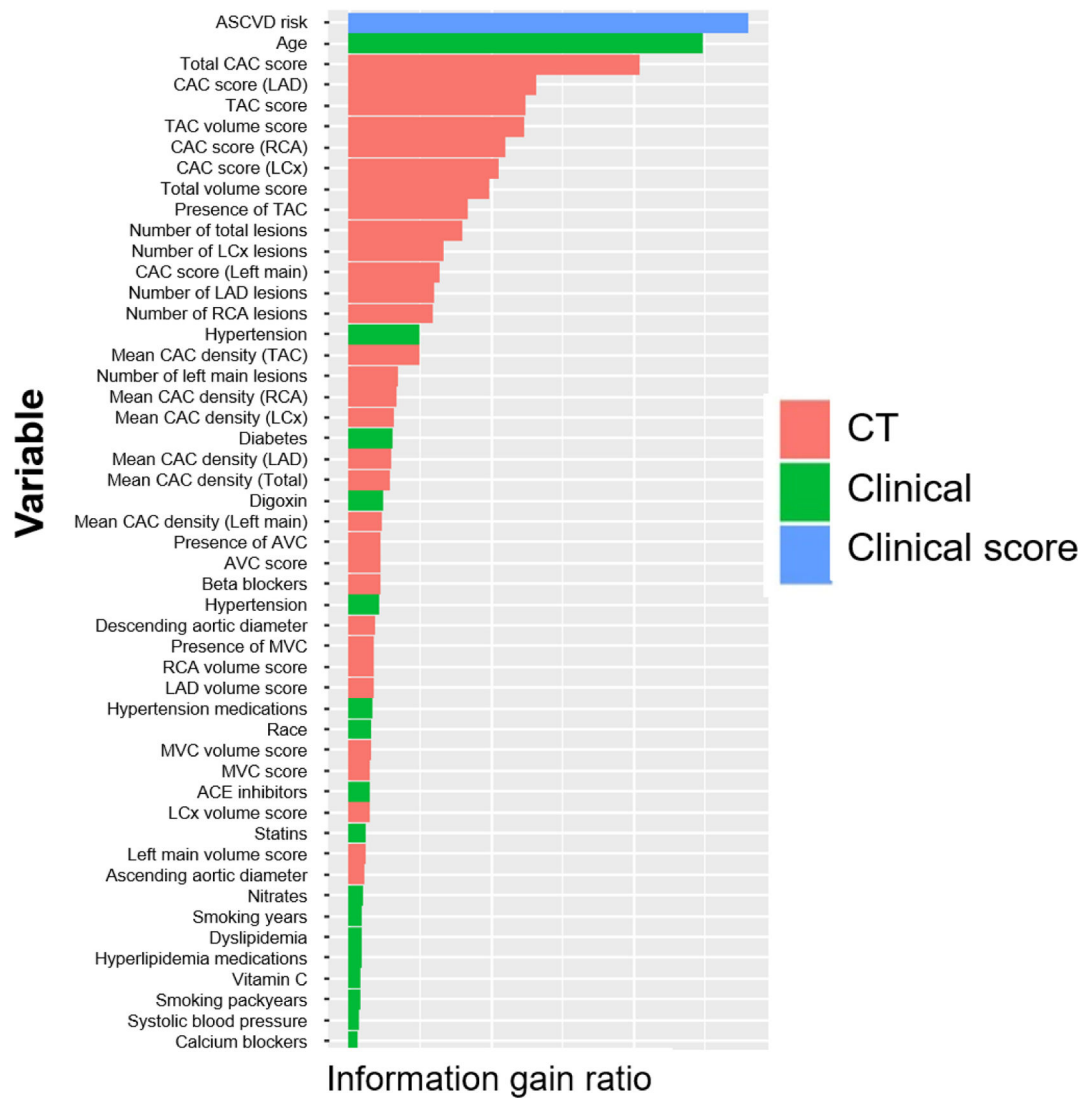


Figure 2a. Importance ranking of variables for prediction of CVD death.

Variable ranking for prediction of CHD death

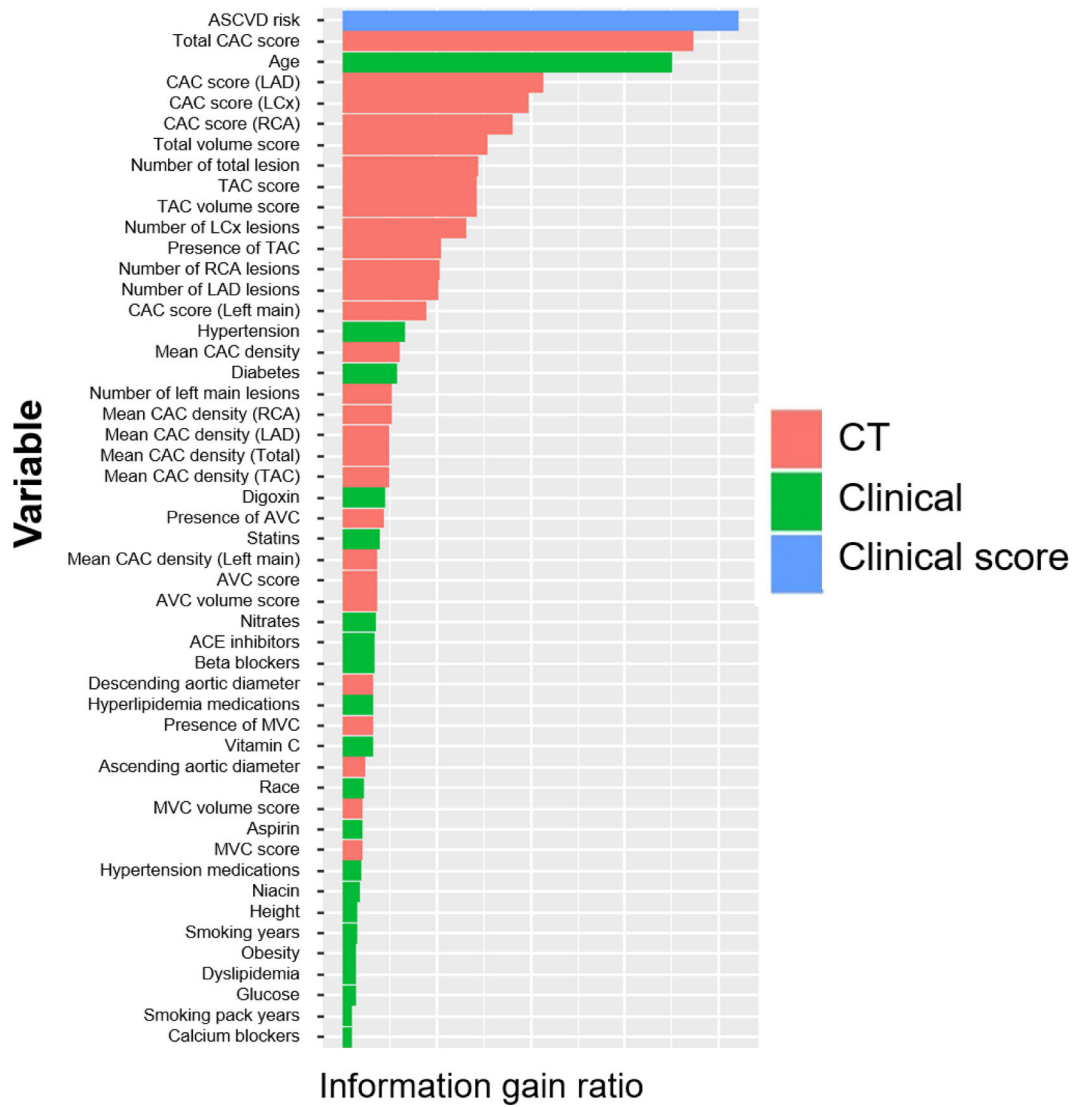


Figure 2b.
Importance ranking of variables for prediction of CHD death.

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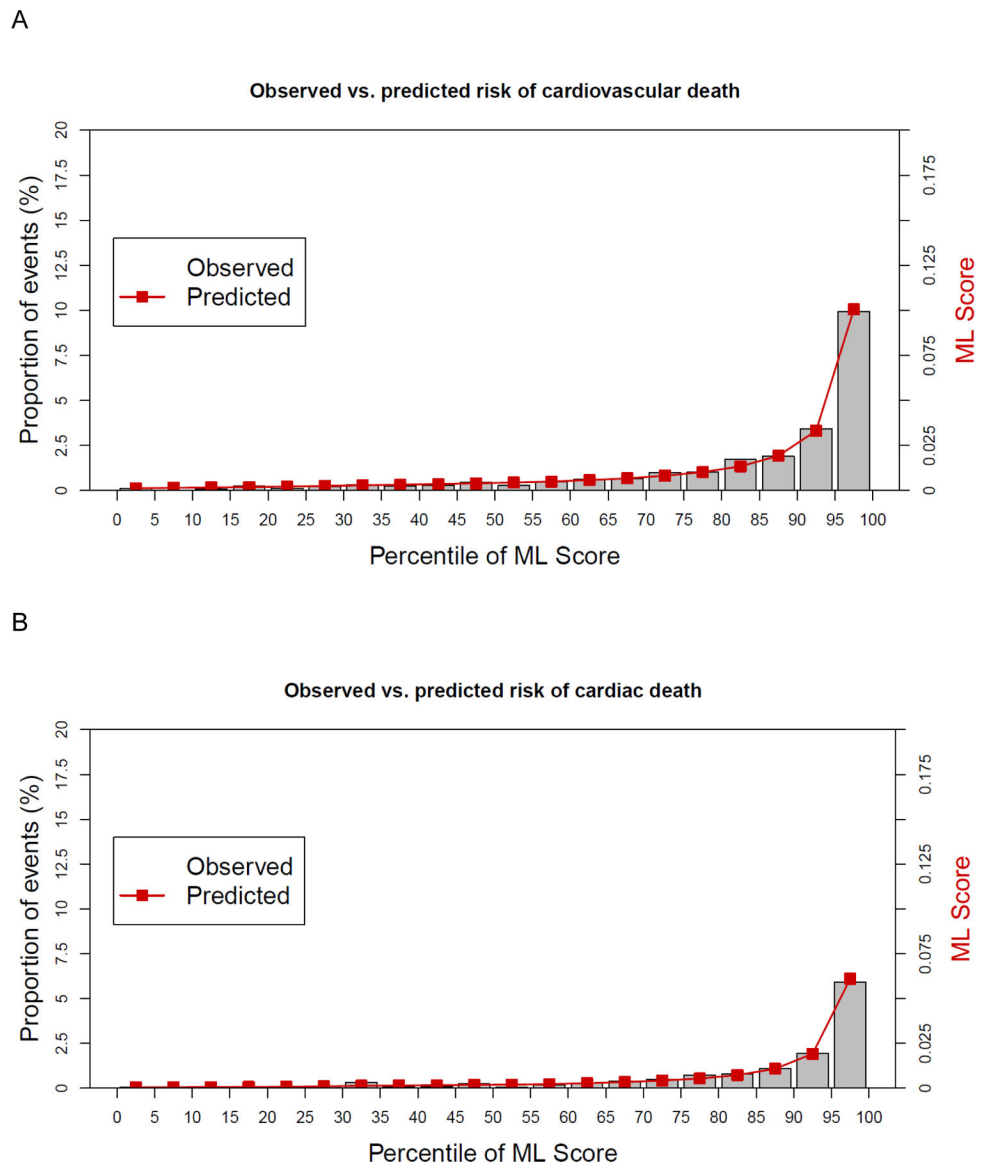
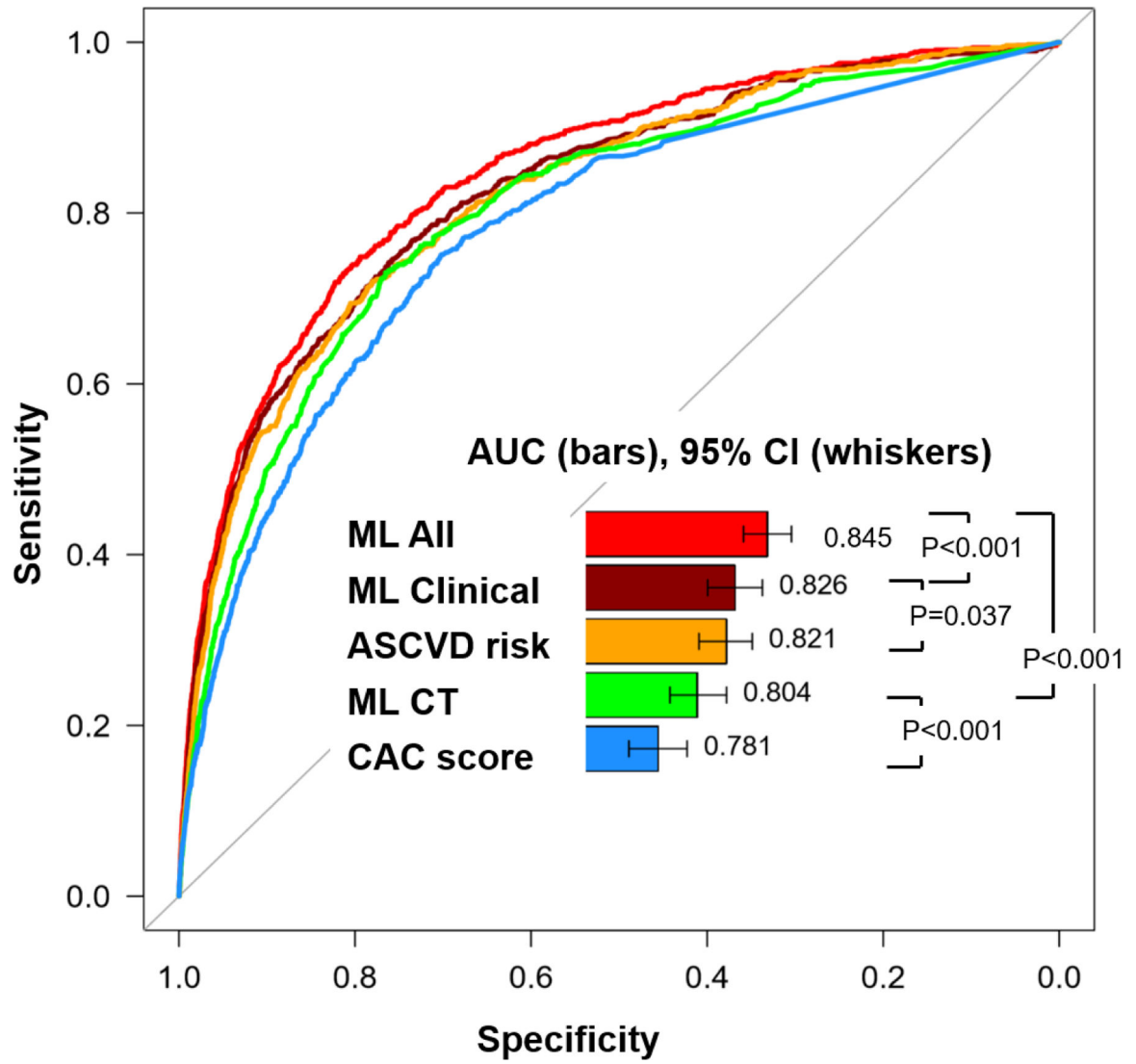


Figure 3. A comparative plot of observed and ML predicted risks for cardiovascular (A) and cardiac (B) death.

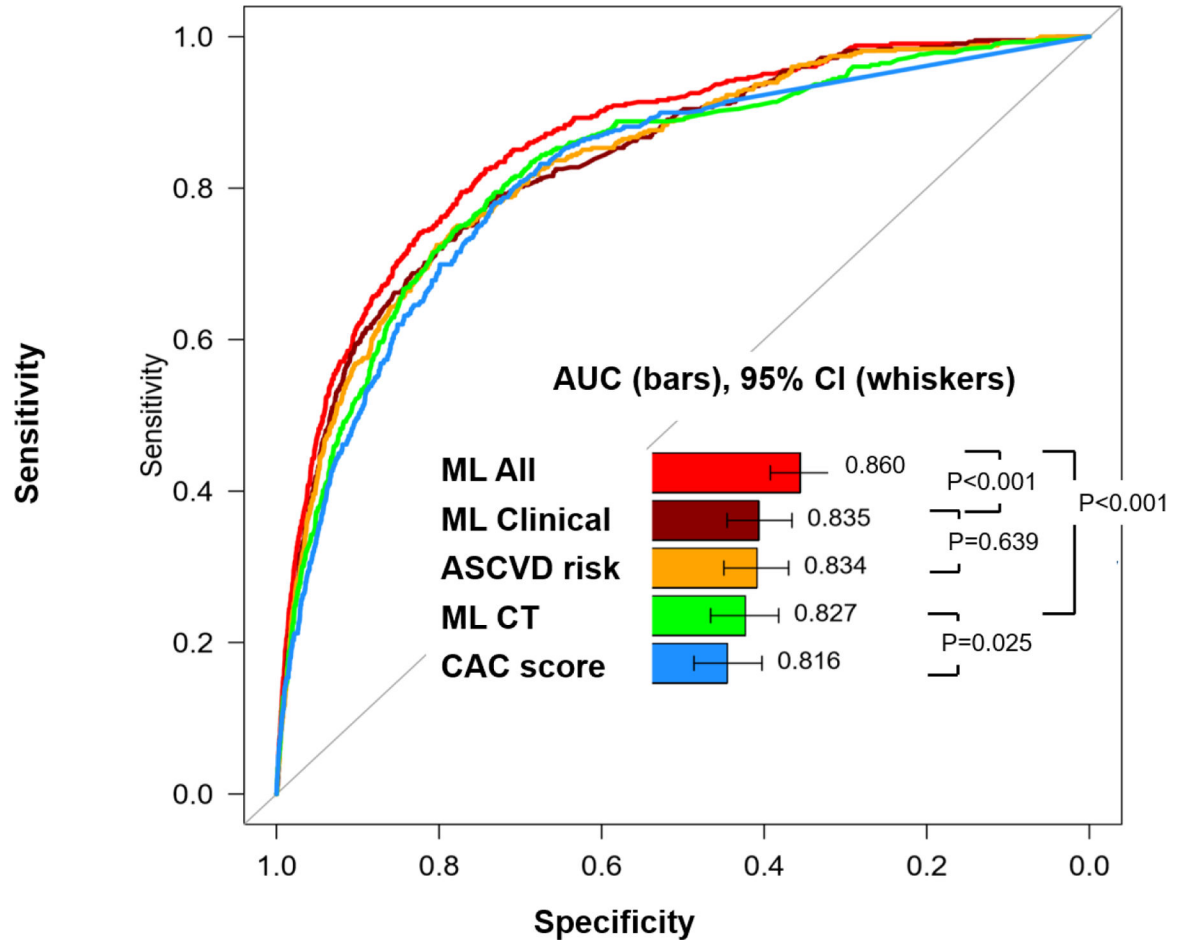
Prediction of 10 year CVD death (772/66636 events)



Central illustration-a.

Receiver operating characteristic curves for prediction of CVD death

Prediction of 10 year CHD death (429/66636 events)



Central illustration-b.

Receiver operating characteristic curves for prediction of CHD death

Table 1.

Patient characteristics (n= 66,636)

Age (years)	54±11
Male (n, %)	44,633 (67)
White/Black/Hispanic/Others (%)	89/2/3/6
BMI (kg/m ²)	27.5±5.3
Hypertension (n, %)	20,624 (31)
Diabetes (n, %)	4,503 (7)
Hyperlipidemia (n, %)	36,227 (54)
Current smoking (n, %)	6,400 (10)
Family history of CHD (n, %)	30,720 (46)
ASCVD risk score (mean ± SD)	7.4 ± 8.9%
CAC score (n, %) (n=66,636)	
CAC 0	29,757 (45)
CAC 1–99	20,534 (30)
CAC 100–399	7,341 (14)
CAC 400	9,004 (11)
TAC score (n, %) (n=41,066)	
TAC 0	19,476 (48)
TAC 1–99	11,927 (29)
TAC 100–399	5,415 (13)
TAC 400	4,248 (10)
AVC score (n, %) (n=10,007)	
AVC 0	8,610 (86)
AVC 1–99	876 (9)
AVC 100–399	352 (3)
AVC 400	169 (2)
MVC score (n, %) (n=10,008)	
MVC 0	9,416 (94)
MVC 1–99	283 (3)
MVC 100–399	150 (1)
MVC 400	159 (2)

Abbreviations: BMI- Body mass index, CHD- Coronary heart disease, ASCVD- Atherosclerotic cardiovascular disease, CAC- Coronary artery calcium, TAC- Thoracic aortic calcification, AVC- Aortic valve calcification, MVC- Mitral valve calcification

Table 2.

Clinical and CT variables

Clinical variables		CT variables	
		CAC variables	Non-CAC variables
ASCVD risk score	Diabetes	Total CAC score	Presence of TAC
Age	Oral diabetic medications	LM CAC score	TAC score
Sex	Insulin	LAD CAC score	TAC volume score
Race	Glucose	LCx CAC score	Presence of AVC
Height	Current smoker	RCA CAC score	AVC score
Weight	Past smoker	Total volume score	AVC volume score
Obese	Smoking years	LM volume score	Presence of MVC
Body mass index	Smoking pack years	LAD volume score	MVC score
Heart rate	Smoking packs	LCx volume score	MVC volume score
Menopause	Digoxin	RCA volume score	Descending aorta diameter
Site	Statins	Total CAC lesions	Ascending aorta diameter
Fasting	Nitrates	LM CAC lesions	
Hypertension	Angiotensin-converting enzyme (ACE) inhibitors	LAD CAC lesions	
Hypertension medications	Beta blockers	LCx CAC lesions	
Systolic blood pressure	Vitamin C	RCA CAC lesions	
Diastolic blood pressure	Aspirin	Total CAC mean density	
Dyslipidemia	Niacin	LM CAC mean density	
Dyslipidemia medications	Calcium blockers	LAD CAC mean density	
Low density lipoprotein	Blood thinner medications	LCx CAC mean density	
Cholesterol	Stroke	RCA CAC mean density	
High density lipoprotein	Peripheral vascular disease		
Triglyceride	Kidney disease		
	Lung disease		
	Family history		

Abbreviations: CT-Computed tomography, ASCVD-Atherosclerotic cardiovascular disease, CAC-Coronary artery calcium, LM-Left main, LAD-Left anterior descending artery, LCx-Left circumflex, RCA-Right coronary artery, TAC-Total aorta calcium, AVC-Aortic valve calcium, MVC-Mitral valve calcium

Table 3.

Optimal cut-points and Youden index for each model in 10-year CHD deaths

Model	Optimal cut-point	Sensitivity	Specificity	Youden index
ML All	0.005	0.825	0.745	0.570
ML Clinical	0.007	0.720	0.802	0.522
ASCVD risk	0.114	0.746	0.783	0.529
ML CT	0.007	0.748	0.778	0.526
CAC score	76.80	0.781	0.732	0.512

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Table 4.

Optimal cut-points and Youden index for each model in 10-year CVD deaths

Model	Optimal cut-point	Sensitivity	Specificity	Youden index
ML All	0.011	0.737	0.803	0.540
ML Clinical	0.010	0.742	0.761	0.504
ASCVD risk	0.113	0.720	0.780	0.500
ML CT	0.013	0.731	0.765	0.496
CAC score	56.0	0.751	0.702	0.453

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