Prediction of short-term atrial fibrillation risk using primary care electronic health records

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ABSTRACT

Objective  Atrial fibrillation (AF) screening by age achieves a low yield and misses younger individuals. We aimed to develop an algorithm in nationwide routinely collected primary care data to predict the risk of incident AF within 6 months (Future Innovations in Novel Detection of Atrial Fibrillation (FIND-AF)).

Methods  We used primary care electronic health record data from individuals aged ≥30 years without known AF in the UK Clinical Practice Research Datalink-GOLD dataset between 2 January 1998 and 30 November 2018, randomly divided into training (80%) and testing (20%) datasets. We trained a random forest classifier using age, sex, ethnicity and comorbidities. Prediction performance was evaluated in the testing dataset with internal bootstrap validation with 200 samples, and compared against the CHA2DS2-VASc (Congestive heart failure, Hypertension, Age ≥75 (2 points), Stroke/transient ischaemic attack/thromboembolism (2 points), Vascular disease, Age 65–74, Sex category) and C2HEST (Coronary artery disease/Chronic obstructive pulmonary disease (1 point each), Hypertension, Elderly (age ≥75, 2 points), Systolic heart failure, Thyroid disease (hyperthyroidism) scores. Cox proportional hazard models with competing risk of death were fit for incident longer-term AF between higher and lower FIND-AF-predicted risk.

Results  Of 2,081,139 individuals in the cohort, 7,386 developed AF within 6 months. FIND-AF could be applied to all records. In the testing dataset (n=416,228), discrimination performance was strongest for FIND-AF (area under the receiver operating characteristic curve 0.824, 95% CI 0.814 to 0.834) compared with CHA2DS2-VASc (0.784, 0.773 to 0.794) and C2HEST (0.757, 0.744 to 0.770), and robust by sex and ethnic group. The higher predicted risk cohort, compared with lower predicted risk, had a 20-fold higher 6-month incidence rate for AF and higher long-term hazard for AF (HR 8.75, 95% CI 8.44 to 9.06).

Conclusions  FIND-AF, a machine learning algorithm applicable at scale in routinely collected primary care data, identifies people at higher risk of short-term AF.

INTRODUCTION

Atrial fibrillation (AF) is a major public health issue. There are now more new cases of AF diagnosed each year in the English National Health Service (NHS) than the four most common causes of cancer combined. Moreover, it is estimated that up to 35% of disease burden remains undiagnosed, and 15% of strokes occur in the context of undiagnosed AF.
Early detection of AF may permit the initiation of oral anticoagulation to reduce embolic stroke risk, and early antiarrhythmic therapy to reduce the risk of death and stroke. Accordingly, early AF detection is a key cardiovascular priority in the UK NHS Long Term Plan, and the European Society of Cardiology recommends opportunistic screening by pulse palpation or ECG rhythm strip in persons aged ≥65 years and systematic ECG screening in those aged ≥75 years. However, there is an increasing cohort of individuals aged younger than 65 years who are being diagnosed with AF and are eligible for anticoagulation.

A large proportion of the population is registered in primary care with a routinely collected electronic health record (EHR). An algorithm that uses routinely collected EHR data to calculate AF risk could give a scalable, efficient and fair approach to targeting AF detection. However, previous algorithms tested in community-based EHRs have a number of shortcomings (online supplemental tables 1 and 2). First, many algorithms developed using traditional regression techniques show only moderate discriminative performance. Second, algorithm prediction using RF across different EHR datasets. RF is a machine learning method consisting of many algorithms, plus an updated literature review (online supplemental tables 4–6). Diagnostic code lists only included the primary care coding system (Read codes), ensuring that only information readily available within a primary care EHR could be incorporated within the algorithm. Concordantly, our entire analytical cohort had no missing data for any of the predictor variables and the algorithm could be applied to all records.

**METHODS**

**Study design and population**

In this population-based study, we used primary care EHRs from the UK Clinical Practice Research Datalink (CPRD)-GOLD dataset. CPRD is one of the largest databases of longitudinal medical records from primary care worldwide and contains anonymised patient data from approximately 7% of the UK population. CPRD-GOLD represents the UK population in terms of age, sex and ethnicity, and has been used to develop algorithms for predicting AF. Data collection happens as part of routine clinical care in participating practices and patients are included in the primary care dataset from their first until their last contact with a participating practice. Diagnostic coding for AF in CPRD has been shown to be consistent and valid, with a positive predictive value (PPV) of 98%. All individuals in the CPRD dataset were linked to Hospital Episode Statistics (HES) Admitted Patient Care (APC) records to obtain comprehensive coverage of AF cases diagnosed in secondary care. We included all adults registered at practices within CPRD who were ≥30 years of age at entry with no history of AF from either data source and at least 1-year follow-up between 2 January 1998 and 30 November 2018. Individuals were censored to a diagnosis of AF (or atrial flutter (AFL)), since it has similar thromboembolic risk and anticoagulation guidelines, withdrawal from CPRD or 6 months, whichever came first. Diagnoses of AF or AFL in primary care were identified using Read codes in CPRD and in secondary care with the 10th revision of the International Statistical Classification of Diseases and Related Health Problems codes in HES-APC (online supplemental table 3). Individuals were randomly split 4:1 to establish a training dataset (80%) and a testing dataset (20%) using the Merseme twist pseudorandom number generator.

We followed the Transparent Reporting of a Multivariable Prediction Model for Individual Prognosis or Diagnosis reporting guideline and the CODE-EHR best-practice framework for using structured electronic healthcare records in clinical research.

**FIND-AF algorithm development**

A random forest (RF) classifier was trained to predict AF at 6 months. Our systematic review evidenced strong discriminative performance for AF prediction using RF across different EHR datasets. RF is a machine learning method consisting of many individual decision trees that operate as an ensemble. The FIND-AF was trained using 10-fold cross-validation on the full training set (full details available in online supplemental methods).

To create an algorithm that could be implemented at scale in national primary care EHRs, we restricted candidate variables to age, sex, comorbidities (72 binary variables, indicating presence or absence of recorded diagnosis) and ethnicity (six categories; online supplemental table 6). Observations and laboratory results were not included. Ethnicity information is routinely collected in the UK NHS and so has increasingly high completeness, and we included an 'ethnicity unrecorded' category where it was unavailable because missingness was considered to be informative. Predictor variables were selected a priori from systematic review of variables included in previous AF risk prediction algorithms, plus an updated literature review (online supplemental tables 4–6). Diagnostic code lists only included the primary care coding system (Read codes), ensuring that only information readily available within a primary care EHR could be incorporated within the algorithm. Concordantly, our entire analytical cohort had no missing data for any of the predictor variables and the algorithm could be applied to all records.

**Statistical analyses**

The baseline characteristics are summarised by incident AF status. Continuous variables were reported as mean ± SD. Categorical variables were reported as frequencies with corresponding percentages.

The degree of variation of each feature in FIND-AF to classification was calculated using the mean decrease in the Gini coefficient, a measure of how each variable contributes to the homogeneity of nodes and leaves in the resulting RF.

Model performance of FIND-AF was determined using the full holdout test set with internal bootstrap validation with 200 samples and compared with a multivariable logistic regression (MLR) model developed with backward model selection with Akaike information criterion. Performance was compared with the CHA2DS2-VASC (Congestive heart failure, Hypertension, Age >75 (2 points), Stroke/transient ischaemic attack/thromboembolism (2 points), Vascular disease, Age 65–74, Sex category) and C,H,HEST (Coronary artery disease/Chronic obstructive pulmonary disease (1 point each), Hypertension, Elderly (age ≥75, 2 points), Systolic heart failure, Thyroid disease (hyperthyroidism)) scores. The CHA2DS2-VASC score was originally developed to predict stroke risk in individuals with AF, and the C,H,HEST score for Asian people without structural heart disease. These algorithms are robust to missing data in routinely collected primary care EHRs and have been tested for AF risk prediction in European cohorts (online supplemental table 2). Other algorithms that can only be applied to a minority of European primary care
EHRs (Pfizer-AI, CHARGe-AF) were not considered. The area under the receiver operating characteristic (AUROC) curve was used to evaluate predictive ability (concordance index) with 95% CIs calculated using the DeLong method. Youden Index was established for the outcome measure as a method of empirically identifying the optimal dichotomous cut-off to assess sensitivity, specificity, PPV and negative predictive value (NPV). Youden Index was calculated and optimised for each test set for each score to derive the optimal cut-off threshold. Calibration was assessed by plotting predicted AF risk against observed AF incidence and by the calibration slope. We calculated the Brier score, a measure of both discrimination and calibration, by taking the mean squared difference between predicted probabilities and the observed outcome. To assess the clinical impact of using FIND-AF as opposed to other risk prediction scores, we calculated the net reclassification index at 0.4% AF risk threshold (the average 6-month incidence rate in the cohort) and conducted a decision curve analysis.

We investigated the performance of FIND-AF, CHA2DS2-VAsC and C_HEST within relevant subgroups defined by sex, ethnicity (white vs black vs Asian vs other non-white ethnic minorities) and age (≥65 years and ≥75 years). We plotted Kaplan-Meier plots for individuals identified as higher and lower FIND-AF-predicted risk of AF to assess the event rate for AF censored at 10 years, and calculated the HR for AF between higher and lower FIND-AF-predicted risk of AF using the Cox proportional hazard model with adjustment for the competing risk of death. We used R V.4.1.0 for all analyses.

Patient and public involvement
The Arrhythmia Alliance, an AF association, provided input on the FIND-AF scientific advisory board. The FIND-AF patient and public involvement group have given input to reporting and dissemination plans of the research.

RESULTS
Patient population
There were 2 081 139 individuals registered in our UK primary care cohort (1 664 911 in the training dataset, 416 228 in testing dataset), with average age 49.9 years (SD 15.4), 50.7% women and 86.7% white. Baseline characteristics and clinical outcomes were similar in the training and testing datasets (online supplemental table 7). Within 6 months, 7386 individuals (0.4%) were diagnosed with AF. Those who developed AF were older and had a higher prevalence of baseline comorbidities than individuals who did not develop AF (table 1). Of new cases, 1546 (20.9%) were younger than 65 years old.

Prediction factors and model accuracy
According to mean decrease in the Gini coefficient, age contributed the most to the prediction, followed by ethnicity and history of heart failure (figure 1). AF discrimination and accuracy of predictions, by AUROC and Brier scores, were better using FIND-AF than the MLR, CHA2DS2-VaSc and C_HEST algorithms (table 2 and figure 2). Sensitivity was highest for the CHA2DS2-VaSc algorithm, but specificity lowest.

According to the Youden Index, the optimal cut-off was 0.0032, leading to a sensitivity of 78% and a specificity of 73%, with a PPV of 2.5% and NPV of 99.8%. The low incidence of AF over 6 months led to similar values for PPV and NPV across the algorithms. Of the algorithms, FIND-AF was the best calibrated (calibration slope 0.782 (95% CI 0.743 to 0.824), table 2 and online supplemental figure 1), yet showed underestimation of risk in the mid-risk strata and overestimation in the highest risk strata.

Risk classification
Of the 416 228 individuals in the testing set, 82 942 (19.9%) were classified as higher risk using FIND-AF, 84 282 (20.2%) using the CHA2DS2-VaSc score and 84 542 (20.3%) using the C_HEST score, respectively. Net reclassification analyses at the 0.4% risk threshold demonstrated modestly favourable reclassification using FIND-AF as opposed to using CHA2DS2-VaSc (net reclassification 0.032, 95% CI 0.029 to 0.051) and strong favourable reclassification using FIND-AF as opposed to using C_HEST (net reclassification 0.113, 95% CI 0.098 to 0.135; online supplemental table 8). In a decision curve analysis, FIND-AF had a superior net benefit compared with the CHA2DS2-VaSc and C_HEST risk scores across all threshold probabilities (online supplemental figure 2).

Of the 82 942 individuals identified as higher risk by FIND-AF, 3483 were <65 years of age, of whom 3448 had a CHA2DS2-VaSc score of at least 1. The incidence rate of AF in routine clinical practice at 6 months was 20-fold higher among individuals identified as a higher predicted risk of AF by FIND-AF compared with individuals identified as lower risk (2.0% vs 0.1%). In routine clinical practice, 1 in every 71 individuals aged ≥65 years were diagnosed with AF within 6 months, 1 in every 58 individuals aged ≥75 years and 1 in every 40 individuals identified at higher predicted AF risk.

Higher predicted AF risk was also associated with increased long-term AF occurrence. Within 5 and 10 years, respectively, 5.1% and 11.9% of the higher predicted risk cohort had been diagnosed with AF, with an 8.75-fold increased hazard (95% CI 8.44 to 9.06) relative to individuals at lower predicted risk (figure 3).

Table 1 Baseline characteristics of analytical cohort with and without atrial fibrillation (AF)

<table>
<thead>
<tr>
<th>Incident AF</th>
<th>No AF</th>
<th>AF</th>
</tr>
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<tbody>
<tr>
<td>n (%)</td>
<td>n (%)</td>
<td></td>
</tr>
<tr>
<td>Demographics</td>
<td></td>
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<tr>
<td>Demographics</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age, years</td>
<td>49.82</td>
<td>73.72</td>
</tr>
<tr>
<td>Sex (women)</td>
<td>1 051</td>
<td>3619</td>
</tr>
<tr>
<td>Comorbidity</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Diabetes mellitus</td>
<td>71 966</td>
<td>815</td>
</tr>
<tr>
<td>Stroke or TIA</td>
<td>37 773</td>
<td>892</td>
</tr>
<tr>
<td>Ischaemic heart disease</td>
<td>77 060</td>
<td>1542</td>
</tr>
<tr>
<td>Hypertension</td>
<td>247 436</td>
<td>2887</td>
</tr>
<tr>
<td>Heart failure</td>
<td>13 717</td>
<td>650</td>
</tr>
<tr>
<td>Dyslipidaemia</td>
<td>60 357</td>
<td>532</td>
</tr>
<tr>
<td>Hyperthyroidism</td>
<td>16 147</td>
<td>155</td>
</tr>
<tr>
<td>Chronic kidney disease</td>
<td>29 359</td>
<td>449</td>
</tr>
<tr>
<td>Anaemia</td>
<td>66 844</td>
<td>501</td>
</tr>
<tr>
<td>Cancer</td>
<td>72 621</td>
<td>887</td>
</tr>
<tr>
<td>Valvular heart disease</td>
<td>9 497</td>
<td>376</td>
</tr>
<tr>
<td>Mean CHA2DS2-VaSc score (SD)</td>
<td>0.97</td>
<td>2.72</td>
</tr>
</tbody>
</table>

| CHA2DS2-VaSc, Congestive heart failure, Hypertension, Stroke/transient ischaemic attack/thromboembolism (2 points), Vascular disease, Age 65–74 years, Sex category; COPD, chronic obstructive pulmonary disease; TIA, transient ischaemic attack.
Arrhythmias and sudden death

Model performance in clinically relevant subgroups
FIND-AF discrimination performance remained strong in both sexes, whereas for the CHA2DS2-VASc and C_HEST scores, performance was better in men than women (table 3). The scores performed differently across ethnic groups. In black individuals, AF discrimination was highest for CHA2DS2-VASc, and in white and Asian individuals, FIND-AF had the strongest discrimination performance.

DISCUSSION
In this population-based study, we trained a machine learning algorithm (FIND-AF) on more than 1.5 million individuals registered in UK primary care to predict the risk of incident AF within the next 6 months (figure 4). When tested in over 400,000 individuals, FIND-AF demonstrated good predictive accuracy, which was superior to other risk scores and robust in both sexes and across ethnic groups. FIND-AF identified a cohort of younger people at higher risk of AF and more efficiently identified individuals diagnosed with AF within 6 months compared with age-based risk stratification. Finally, short-term predicted AF risk also translated to long-term AF occurrence.

Current approaches to targeting investigation for undiagnosed AF are based on age.7 Our analysis demonstrated that one-fifth of newly detected AF cases within 6 months occur in people aged ≤65 years, emphasising the opportunity lost when enhanced AF investigation is restricted to older populations. ECGs can be used to accurately predict AF risk,20 but they are not widely available in the community, whereas 98% of the UK population are registered in primary care with an accompanying EHR.7 Our meta-analysis of AF prediction algorithms using EHRs demonstrated that algorithms developed using traditional regression techniques provided only moderate discrimination performance.10 In our study, a machine learning prediction algorithm (FIND-AF) outperformed the C_HEST and CHA2DS2-VASc scores.

For a machine learning prediction algorithm to be useful in clinical practice, it must be implementable within the clinical workflow, provide prediction that meaningfully informs decision-making and engender confidence in how outputs were interpreted.

Table 2 Performance for 6-month incident AF with optimal threshold determined by Youden Index

<table>
<thead>
<tr>
<th>Algorithm</th>
<th>FIND-AF</th>
<th>MLR</th>
<th>CHA2DS2-VASc</th>
<th>C_HEST</th>
</tr>
</thead>
<tbody>
<tr>
<td>AUROC (95% CI)</td>
<td>0.824 (0.814 to 0.834)</td>
<td>0.765 (0.755 to 0.769)</td>
<td>0.784 (0.773 to 0.794)</td>
<td>0.757 (0.744 to 0.770)</td>
</tr>
<tr>
<td>Sensitivity (95% CI)</td>
<td>0.781 (0.731 to 0.829)</td>
<td>0.760 (0.653 to 0.814)</td>
<td>0.847 (0.829 to 0.866)</td>
<td>0.642 (0.619 to 0.791)</td>
</tr>
<tr>
<td>Specificity (95% CI)</td>
<td>0.731 (0.693 to 0.771)</td>
<td>0.679 (0.635 to 0.776)</td>
<td>0.611 (0.608 to 0.612)</td>
<td>0.790 (0.622 to 0.792)</td>
</tr>
<tr>
<td>PPV (% (95% CI))</td>
<td>2.5% (2.3 to 2.7)</td>
<td>2.0% (1.8 to 2.6)</td>
<td>2.2% (2.1 to 2.3)</td>
<td>2.0% (1.5 to 2.2)</td>
</tr>
<tr>
<td>NPV (% (95% CI))</td>
<td>99.8% (99.8 to 99.9)</td>
<td>99.7% (99.6 to 99.7)</td>
<td>99.8% (99.8 to 99.8)</td>
<td>99.7% (99.7 to 99.8)</td>
</tr>
<tr>
<td>Calibration slope* (95% CI)</td>
<td>0.782 (0.743 to 0.824)</td>
<td>0.698 (0.654 to 0.735)</td>
<td>0.621 (0.589 to 0.652)</td>
<td>0.608 (0.576 to 0.648)</td>
</tr>
</tbody>
</table>

*Calibration slope was derived from linear regression models by forcing the intercept through origin (0, 0).

AF, atrial fibrillation; AUROC, area under the receiver operating characteristic; CHA2DS2-VASc, Congestive heart failure, Hypertension, Age >75 (2 points), Stroke/transient ischaemic attack/thromboembolism (2 points), Vascular disease, Age 65–74, Sex category; C_HEST, Coronary artery disease/Chronic obstructive pulmonary disease (1 point each), Hypertension, Elderly (age ≥75, 2 points), Systolic heart failure, Thyroid disease (hyperthyroidism); FIND-AF, Future Innovations in Novel Detection of Atrial Fibrillation; MLR, multivariable logistic regression; NPV, negative predictive value; PPV, positive predictive value.
Figure 2  Receiver operating characteristic curves for FIND-AF, the multivariable logistic regression (MLR), CHA₂DS₂-VASc and C.HEST algorithm.

C.HEST, Coronary artery disease/Chronic obstructive pulmonary disease (1 point each), Hypertension, Elderly (age ≥75, 2 points), Systolic heart failure, Thyroid disease (hyperthyroidism); CHA₂DS₂-VASc, Congestive heart failure, Hypertension, Age >75 (2 points), Stroke/transient ischaemic attack/thromboembolism (2 points), Vascular disease, Age 65–74, Sex category.

Figure 3  Kaplan-Meier plots for AF occurrence, by predicted risk from FIND-AF. AF, atrial fibrillation; FIND-AF, Future Innovations in Novel Detection of Atrial Fibrillation.
monitoring. Finally, the most important predictors in FIND-AF are already well-recognised risk factors for AF (for example, age, heart failure, valvular heart disease), which provide reassurance in the associations being made by the algorithm.7

Fairness is a critical characteristic when considering the impact of prediction algorithms in healthcare. The CHARGE-AF and PuLSE-AI algorithms have strong AF prediction performance,9 11 yet incorporate variables that are frequently missing (height, weight and systolic and diastolic blood pressure).10 Consequently, their applicability is limited to 17% and 35% of primary care EHRs, respectively.9 11 Often, health data poverty disproportionately affects individuals from minority ethnicities and deprived backgrounds, so the application of these algorithms could reinforce health inequities.22 Furthermore, whether their performance varies by sex and in minority ethnic groups in European populations is unknown. In our study, the C2HEST and CHA2DS2-VASc scores were less accurate in women compared with men, and their performance varied substantially across different ethnic groups. FIND-AF’s design enabled its application to every single patient record in a nationally representative

Table 3  Discrimination performance of FIND-AF, CHA2DS2-VASc and C2HEST by sex, age and ethnicity

<table>
<thead>
<tr>
<th></th>
<th>FIND-AF AUROC (95% CI)</th>
<th>CHA2DS2-VASc AUROC (95% CI)</th>
<th>C2HEST AUROC (95% CI)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Overall</td>
<td>0.824 (0.814 to 0.834)</td>
<td>0.784 (0.773 to 0.794)</td>
<td>0.757 (0.744 to 0.770)</td>
</tr>
<tr>
<td>Sex</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Men</td>
<td>0.819 (0.809 to 0.829)</td>
<td>0.807 (0.793 to 0.821)</td>
<td>0.793 (0.777 to 0.810)</td>
</tr>
<tr>
<td>Women</td>
<td>0.821 (0.810 to 0.831)</td>
<td>0.776 (0.760 to 0.793)</td>
<td>0.746 (0.727 to 0.765)</td>
</tr>
<tr>
<td>Age</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>≥65 years</td>
<td>0.712 (0.698 to 0.727)</td>
<td>0.669 (0.654 to 0.684)</td>
<td>0.675 (0.661 to 0.690)</td>
</tr>
<tr>
<td>≥75 years</td>
<td>0.657 (0.638 to 0.675)</td>
<td>0.612 (0.593 to 0.632)</td>
<td>0.589 (0.570 to 0.608)</td>
</tr>
<tr>
<td>Ethnicity</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>White</td>
<td>0.810 (0.799 to 0.821)</td>
<td>0.781 (0.769 to 0.792)</td>
<td>0.756 (0.743 to 0.770)</td>
</tr>
<tr>
<td>Asian</td>
<td>0.796 (0.693 to 0.899)</td>
<td>0.758 (0.639 to 0.876)</td>
<td>0.731 (0.611 to 0.850)</td>
</tr>
<tr>
<td>Black</td>
<td>0.801 (0.680 to 0.923)</td>
<td>0.843 (0.764 to 0.923)</td>
<td>0.707 (0.511 to 0.902)</td>
</tr>
<tr>
<td>Other non-white ethnic minority</td>
<td>0.805 (0.765 to 0.845)</td>
<td>0.768 (0.729 to 0.807)</td>
<td>0.805 (0.765 to 0.846)</td>
</tr>
<tr>
<td>Ethnicity unrecorded</td>
<td>0.823 (0.770 to 0.875)</td>
<td>0.838 (0.777 to 0.900)</td>
<td>0.788 (0.705 to 0.870)</td>
</tr>
</tbody>
</table>

The total number of individuals in each subgroup and number of incident AF cases are as follows: men (n=211 378, AF=720), women (n=204 850, AF=753), age ≥65 years (n=81 258, AF=1168), age ≥75 years (n=36 358, AF=796), white (n=279 027, AF=1301), Asian (n=8422, AF=16), black (n=6478, AF=11), other non-white ethnic minority (n=28 303, AF=96), ethnicity unrecorded (n=93 998, AF=49).

Figure 4  Summary of the study and main findings. Hitherto implementation of screening for atrial fibrillation (AF) has been targeted to older persons in the general population, but this may miss one-fifth of new cases. A machine learning algorithm using routinely collected data in primary care electronic health records in the UK can accurately predict short-term risk of AF in persons aged ≥30 years. This may be a more efficient method for guiding AF screening. C2HEST, Coronary artery disease/Chronic obstructive pulmonary disease (1 point each), Hypertension, Elderly (age ≥75, 2 points), Systolic heart failure, Thyroid disease (hyperthyroidism); FIND-AF, Future Innovations in Novel Detection of Atrial Fibrillation.
Arrhythmias and sudden death

dataset of routinely collected primary care EHRs; and performance was robust in both sexes and across minority ethnic groups.

Three barriers need to be overcome for FIND-AF to be accepted into clinical practice. First, it requires external validation, which is currently underway using The Phoenix Partnership UK primary care EHR system (ResearchOne) and the Israeli Clalit Health Services. Second, prospective validation of FIND-AF is critical before implementation into clinical practice. We are launching a pilot implementation study across primary care sites where individuals identified at higher risk will be offered rhythm monitoring (The BHF Bristol Myers Squibb Cardiovascular Catalyst Award—CC/22/250026). Third, a cost utility analysis and budget impact analysis of the use of FIND-AF will need to be conducted.

Primary care EHRs in the UK are nationwide and held centrally, so FIND-AF could be activated at scale across geographically disparate sites to identify a subpopulation at elevated AF risk. The cohort identified as higher risk in this study included younger people who would currently be excluded from screening pathways, and higher predicted AF risk was associated with elevated AF occurrence both in the short and long term. Therefore, FIND-AF could facilitate efficient population-based AF screening or comprehensive programmes designed to improve risk factor profiles (including targeted weight loss and optimisation of blood pressure control).23

Screening for AF would adhere to many of the Wilson and Junger principles for a screening programme.24 Opportunistic screening guided by age has not been demonstrated to increase AF detection rates,25 but this may change in a more precisely defined higher-risk cohort. Systematic screening of older patients with intermittent or continuous (invasive or non-invasive) rhythm monitors is associated with increased AF detection rates, compared with routine care.24 However, the yield of new AF detection rates,25 but this may change in a more precisely
detected AF of clinic

FIND-

study included younger people who would currently be excluded geographically disparate sites to identify a subpopulation at

CONCLUSIONS

We trained and tested a novel machine learning algorithm (FIND-AF) that was applicable at scale within a nationwide routinely collected primary care EHR dataset. FIND-AF was able to accurately predict AF risk within 6 months and identify a cohort at elevated risk of AF in the longer term.

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Contributors RN, JW and CPG conceived the idea of the study. JW undertook data extraction and statistical analysis. RN verified the underlying data. RN, JW and CPG interpreted the findings and RN drafted the manuscript. CB, JP, DH, CC, KR, YMN, KN, DZ, RA, MH, JW and CPG critically reviewed the manuscript and RN revised the manuscript for final submission. All authors had full access to all data in the study and accept responsibility to submit for publication. RN is the guarantor.

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Patient consent for publication Not required.

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Data availability statement Data may be obtained from a third party and are not publicly available. Data used in this study can be accessed through CPRD subject to protocol approval. The algorithm can be shared with researchers who agree to use it only for research purposes with a data sharing agreement.

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