

Machine Learning-Based Prediction of Acute Coronary Syndrome Using Only the Pre-Hospital 12-Lead Electrocardiogram

By

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ABSTRACT

Prompt identification of acute coronary syndrome is a pivotal challenge in clinical practice. The 12-lead ECG is readily available during initial patient evaluation, but current rule-based interpretation approaches lack sufficient accuracy. Here we report machine learning-based methods for the prediction of underlying acute myocardial ischemia in patients with chest pain. Using 554 temporal-spatial features of 12-lead ECG, we train and test multiple classifiers on two independent prospective patient cohorts ($n = 1244$). While maintaining higher negative predictive value, our final fusion model achieves 52% gain in sensitivity compared to commercial interpretation software and 37% gain in sensitivity compared to experienced clinicians. Such ultra-early, ECG-based clinical decision support tool, when combined with the judgment of trained emergency personnel, would be imperative for improving clinical outcomes and reducing unnecessary costs in patients with chest pain.

Key Words: machine learning; electrocardiogram; acute coronary syndrome; ischemia

SUMMARY OF WORK FOR DISPLAY ON HOMEPAGE

Diagnosing a heart attack requires excessive testing and prolonged observation, which usually requires hospital admission. Here we report an intelligent system based exclusively on EKG data that can help clinicians identify 37% more heart attacks during initial screening.

1 INTRODUCTION

2 Nearly seven million Americans visit the Emergency Department annually for a chief
3 complaint of chest pain. Approximately 10% of those patients have an acute disruption in blood
4 supply to the heart attributed to underlying atherosclerotic disease in the coronary arteries, a
5 life-threatening condition referred to as acute coronary syndrome (ACS).^{1,2} However, more than
6 50%–75% of the seven million patients with chest pain are admitted to the hospital because the
7 initial clinical evaluation is not sufficient to rule in or rule out ACS. This problem results from the
8 low sensitivity of the electrocardiogram (ECG) and initial clinical data to predict the presence of
9 ongoing acute myocardial ischemia in those with ACS. As the first available clinical test, the
10 standard 10-second 12-lead ECG can identify a small subset of ACS patients that have ST
11 segment elevation on their ECG, hence the term ST elevation myocardial infarction (STEMI).
12 The majority (>50%) of ACS patients, however, have no such ST elevation on their ECG,³ and
13 thus require a time-consuming biomarker-driven approach and/or provocative testing to rule in
14 or out acute myocardial ischemia. More sensitive classification tools using the 12-lead ECG
15 could improve speed and accuracy of ACS detection.

16 Critical narrowing or occlusion of a coronary artery leads to myocardial ischemia in the
17 region supplied by that coronary artery.^{4,5} Regional ischemia leads to reduction in the duration,
18 resting potential, and propagation velocity of action potentials in the affected myocardium, which
19 leads to a wide variability in the conduction speeds between various myocardial regions.
20 Variability in conduction speeds between the epicardial and endocardial walls of the affected
21 region results in temporal changes in specific ECG leads facing that region (i.e., features of
22 waveform duration and amplitude in individual leads),⁶ whereas variability ischemic regions and
23 healthy myocardium results in spatial changes between orthogonal ECG leads (i.e., features of
24 global electrical heterogeneity).⁷ Thus, using both temporal and spatial features of the 12-lead
25 ECG would be more robust in detecting ACS than using temporal waveform features alone,

1 such as ST elevation. We have shown that mild-to-moderate ischemia distorts the temporal-
2 spatial features of the 12-lead ECG before ST changes evolve,^{8,9} suggesting that building
3 sensitive ACS classification algorithms using only the 12-lead ECG is plausible.

4 A single 10-second 12-lead ECG provides a large number of temporal-spatial features
5 and is thus a rich data platform to model and quantify the presence of ongoing myocardial
6 ischemia. Analysis of the high-dimensional, highly-correlated ECG features requires
7 sophisticated machine learning (ML) classifiers. A number of ML classifiers to predict ACS using
8 ECG data have been reported in the literature.¹⁰⁻¹⁸ However, most studies either used small
9 and limited public datasets (e.g., MIT-BIH, PTB, etc.) or used historical ECGs from hospital
10 records in a case-control approach. Other studies focused only on the identification of patients
11 with STEMI,¹⁸ while others used single-lead ECGs or individual heart beats for algorithm
12 development. All of these limitations diminished the clinical utility for predicting acute ischemia
13 in the general, non-selected chest pain populations seen at ED settings. Validation of
14 generalizable ML classifiers on real-world data from prospective cohort studies is needed.

15 Herein, we present ML-based methods for the prediction of underlying acute myocardial
16 ischemia in patients with chest pain using only the standard 12-lead ECG. We validate and test
17 this approach using two large prospective cohorts from three tertiary care hospitals in the United
18 States. Each cohort includes consecutive prehospital 12-lead ECGs obtained during first
19 medical contact. A key feature of our ML method is that it not only utilizes traditional ECG
20 features, but it also takes advantage of novel temporal-spatial features of the 12-lead ECG.¹⁹
21 Another key element of this method is that feature selection and data recoding is guided by
22 domain-specific knowledge of the pathological nature of acute myocardial ischemia. Using
23 different ML-based classifiers trained and tested on separate prospective cohorts, we arrive at a
24 model that compares to and outperforms clinicians in their expert ECG interpretations based on
25 current practice guidelines.

1 RESULTS

2 *Patient Characteristics*

3 The study population consisted of two patient cohorts from the ongoing EMPIRE study
4 (ECG Methods for the Prompt Identification of Coronary Events).²⁰ The first cohort (2013–2014)
5 included 745 patients and the second cohort (2014–2015) included 499 patients with
6 interpretable ECGs (i.e., no excessive noise or artifacts, not in ventricular tachycardia or
7 fibrillation). Paramedics enrolled consecutive patients in both cohorts and acquired 12-lead
8 ECGs in the field prior to any medical treatment. Paramedics transmitted ECGs to our medical
9 command center where they were stored for offline analysis. We collected clinical data for 30
10 days from the date of index encounter. The primary study outcome used to train classifiers and
11 to test performance was defined as any ACS event. In subsequent sensitivity analyses, we
12 excluded patients with confirmed STEMI on prehospital ECG who were sent to the
13 catheterization lab emergently. Finally, to optimize the clinical utility of our algorithms, we
14 included all ECGs in our analyses, including those with secondary repolarization changes (i.e.,
15 pacing, bundle branch block, and left ventricular hypertrophy). Table 1 summarizes the clinical
16 characteristics of each cohort.

17 *Dataset Derivation and Preparation*

18 Figure 1 shows the stages of dataset derivation and preparation. First, all ECGs were
19 pre-processed using manufacturer-specific commercial software (Philips Healthcare, Andover,
20 MA) and then we manually inspected tracings for noise and artifact. After ectopic beats were
21 removed and median beats were computed, we extracted 554 temporal-spatial features from
22 each ECG using previously-validated, commercial algorithms (Philips Healthcare, Andover,
23 MA). Overall, less than 0.2% of values in all features were missing and these were imputed
24 using either the mean or the mode. We then tested a menu of various ML classifiers and
25 selected for further training and validation the three algorithms that had the best performance:

1 logistic regression (LR), gradient boosting machine (GBM), and artificial neural network (ANN).
2 We further tuned these three classifiers by excluding ECG features that were least likely to have
3 any mechanistic link to the pathogenesis of acute myocardial ischemia, yielding a reduced set of
4 65 clinically-important ECG features. Finally, we recoded the 65 continuous ECG features into
5 categories using previously published cutoff values of clinical significance, for instance, spatial
6 QRS-T angle was re-labeled as normal (0–49°), borderline (50–99°), and abnormal ($\geq 100^\circ$).
7 There was a total of 9 classifiers trained and tested in this paper.

8 ***Classification Performance Using Various ECG Feature Subsets***

9 Using 10-fold cross-validation approach, we first trained each ML classifier using all 554
10 features (LR₅₅₄, GBM₅₅₄, and ANN₅₅₄) on Cohort 1, and then tested performance on Cohort 2.
11 Figure 2A shows the ROC curves for each classifier. Although GBM₅₅₄ and ANN₅₅₄
12 outperformed LR₅₅₄ using all available ECG features, we observed a wide variability in
13 classifiers' performance with poor generalizability to testing set, reflecting low bias – high
14 variance tradeoff.

15 Next, we trained and tested the performance of each ML classifier using only the 65
16 ECG features deemed as clinically relevant to the pathogenesis of acute myocardial ischemia
17 (LR₆₅, GBM₆₅, and ANN₆₅). Figure 2B shows the ROC curves for these classifiers. Interestingly,
18 all three classifiers performed equally on the training set and they generalized well on the
19 independent testing set, reflecting low bias – low variance tradeoff.

20 Finally, we trained the three classifiers using the same 65 features after they were
21 relabeled using previously-validated cutoff thresholds of clinical significance (LR_{65+L}, GBM_{65+L},
22 and ANN_{65+L}). Figure 2C shows the ROC curves for these classifiers. Although these classifiers
23 performed well on the training set, we again observed a wide variability in classifiers'
24 performance with poor generalizability to testing set, reflecting low bias – low variance tradeoff.

1 **Comparing ML Classifiers to Reference Standard**

2 We used the ML classifiers with best low bias – low variance tradeoff to create a simple
3 fusion model. This fusion model was based on vote-count of classifiers built on the reduced
4 (LR₆₅, GBM₆₅, and ANN₆₅) and labeled (LR_{65+L}, GBM_{65+L}, and ANN_{65+L}) datasets. We used 3 or
5 more votes as the cutoff to compute diagnostic performance metrics for this model as compared
6 to two current ECG reference standards: (1) expert ECG read by clinicians and (2) automated
7 ECG reads by commercial rule-based software. To get these annotations, each 12-lead ECG
8 was annotated according to the fourth Universal Definition of Myocardial Infarction consensus
9 statement²¹ by two experienced clinicians who were blinded from study outcome. We used
10 Philips diagnostic 12/16 lead ECG analysis program (Philips Healthcare, Andover, MA) for
11 automated ECG read. Figure 3 and Table 2 show the ROC curves and diagnostic accuracy
12 metrics for the ML fusion model against the two ECG reference standards. To place these
13 comparisons in a context, we show the classification performance of the HEART score obtained
14 at the ED based on all available clinical, laboratory, and ECG data.

15 Figure 3A demonstrates that ML classifiers outperform expert clinicians and commercial
16 ECG algorithms in detecting ACS events. While maintaining higher negative predictive value,
17 our ML fusion model demonstrates 37% gain in sensitivity compared to experienced clinicians
18 and 52% gain compared to commercial ECG algorithms (Table 2), corresponding to a net
19 reclassification improvement of 0.19 (95% CI 0.06–0.31) and 0.30 (95% CI 0.19–0.41),
20 respectively. Furthermore, supplementing our ML algorithm with important patient history data
21 typically available during first medical contact did not result in any additional improvement in
22 classification performance.

23 Next, to explore the performance of our model in detecting non-ST elevation ACS
24 events, we removed the prehospital STEMI cases and repeated our analyses (Figure 3B). As
25 seen in this figure, our ML fusion model still outperforms experienced clinicians and commercial

1 ECG software in detecting the majority of ACS events (Table 2), with net reclassification
2 improvement of 0.28 (95% CI 0.13–0.43) against experienced clinicians and 0.37 (95% CI 0.26–
3 0.49) against commercial algorithms. This finding supports the notion that unlike current ECG
4 reference standards that are heavily geared toward evaluating ST amplitude changes, our ML
5 algorithm takes into account the subtle temporal-spatial signatures of ischemia, which explains
6 the large gain we observed in the diagnostic accuracy.

7 Finally, to identify potential room for improvement, we interrogated the sources of false
8 negatives in the classifications of our ML model. On independent testing, there were 21 patients
9 with ACS misclassified as no disease. We investigated the ECGs of these cases and identified
10 the following potential sources of error: excessive baseline wander (n=6), frequent PVCs (n=3),
11 tachycardia > 100 bpm (n=2), and left ventricular hypertrophy (n=2).

12 **DISCUSSION**

13 This study built and tested a classification algorithm that uses available ECG data from
14 first medical contact to predict ACS in consecutive, unselected patients presenting to ED with
15 chest pain. Using different ML-based classifiers trained and tested on separate prospective
16 cohorts, we arrived at a generalizable model that outperforms both commercial interpretation
17 software as well as experienced clinicians. Our findings show that, by incorporating existing
18 clinical knowledge in classification decisions, linear prediction models like LR can be equivalent
19 to complex and computationally-expensive algorithms like ANN and GBM. On the independent
20 test set, our final ML fusion classifier, while maintaining higher NPV, achieved 39% gain in
21 sensitivity compared to commercial interpretation algorithms and 24% gain in sensitivity
22 compared to experienced clinicians. To our knowledge, this is the first clinical study that
23 prospectively validated and tested the performance of ML-based models on two separate
24 cohorts to predict ACS using only the prehospital 12-lead ECG.

1 Our findings have several important clinical implications. First, using an ultra-early, ECG-
2 based clinical decision support tool, when combined with the judgment of trained emergency
3 personnel, could be imperative for improving outcomes in patients with chest pain. Our gain in
4 net classification improvement implies that 37%–59% of patients with ACS could be better
5 targeted for transfer to appropriate destinations (e.g., centers with advanced cardiac care units
6 or PCI capabilities) or for the initiation of guideline-recommended anti-ischemic therapies. Our
7 high NPV suggest that our algorithm could also be useful for the early and safe discharge of
8 chest pain patients at low risk of ACS. This could save substantial time and cost compared to
9 traditional chest pain evaluations reliant on biomarkers and provocative testing performed over
10 a 24-hour hospital observation. Second, a strength of our predictive model lies in its real-time
11 applicability and scalability since it could be automated and directly integrated into existing ECG
12 machines without the need to input additional clinical data into the model. This means that our
13 model can be very useful in non-tertiary care settings where more invasive diagnostics might
14 not be readily available. Third, the real-time clinical decision support that could be provided by
15 our model is specifically useful to non-specialists and nurses or prehospital personnel with
16 limited experience in ECG interpretation. The classification performance of our model not only
17 outperformed rule-based predictions by standard commercial software, but also met and
18 outperformed the expert ECG interpretation by trained physicians. This means that our
19 algorithm can be used by non-specialized emergency personnel to screen patients and identify
20 the subset of patients whose ECGs need to be further evaluated by offsite experts, a strategy
21 that has long been shown to improve outcomes in those with confirmed ACS.²² Finally, our
22 algorithm can be used to detect ACS in patients whose ECG is confounded by baseline
23 abnormalities such as pacing and bundle branch blocks. Current clinical sensitivity in classifying
24 these patients is low, and our algorithm’s ability to triage these vulnerable patients would
25 significantly enhance the generalizability of our approach to real world clinical settings.

1 Applying ML algorithms to predict ACS has been widely described in literature. A
2 challenge in the development of such models using ECG data is the absence of relevant
3 datasets for training and validation. Most prior algorithms²³⁻³⁴ have used the open-source
4 Physikalisch-Technische Bundesanstalt (PTB) diagnostic ECG database. This highly-selected
5 dataset contains the 12-lead ECGs of only 200 subjects (148 ACS and 52 healthy controls).
6 Although most of these ML classifiers report accuracy that ranges from 93.5% to 98.8%, the
7 generalizability of such models to real-world clinical settings remain questionable, and it is likely
8 these algorithm were overfitting the data contained in the PTB dataset

9 On the other hand, there were few studies that used clinical datasets to build ML
10 classifiers. However, most of these studies combined classical ECG features (e.g., diagnostic
11 ST-T amplitude changes) with a full range of other clinical data elements (e.g., patient history,
12 physical exam abnormalities, laboratory values, and / or diagnostic tests).^{10,15,17,35} Despite the
13 high accuracy achieved by these models (≥ 0.90), classifiers that incorporate such extensive
14 findings from patient clinical profiles have limited utility during early patient triage decisions.

15 There are two prior studies that used only ECG data to build ML-based classifiers.
16 Forberg et al.³⁶ trained and cross-validated ANN and LR classifiers on a dataset of 861 patients
17 with chest pain (ACS = 344, 40%). Using 228 ECG features (i.e., 19 temporal measures from
18 each of the 12 leads), their classifiers achieved AUC of 0.86 and 0.88, respectively, providing
19 nearly 13% gain in sensitivity compared to expert clinicians and 20% gain compared to
20 automated interpretation software. Although LR classifier developed by Forberg et al. (AUC =
21 0.88) performed better compared to ours (AUC = 0.79), it is worth noting that their approach
22 selectively recruited positive cases to enrich their dataset for ACS, making their classes
23 artificially better balanced compared to ours (ACS prevalence = 40% vs. 18%). In addition, their
24 classifier was not evaluated on an independent test set, suggesting that their classifier yielded a
25 too optimistic estimate of the ROC performance. Another clinical study that is relevant to our

1 current work is the study by Green et al.¹³ that trained ANN and LR classifiers on 643
2 consecutive patients with chest pain (ACS = 130, 20%). Using a set of 16 ECG features (i.e.,
3 duration, amplitude, area, and slope measures of QRS and ST segment) that were reduced
4 using PCA from nearly 72 features, their classifiers achieved an AUC of 0.80 and 0.71,
5 respectively. However, these results were not evaluated on an independent test set, again
6 raising concerns about the generalizability of their model.

7 How to select the most appropriate ML approach in clinical applications is a debatable
8 issue. Prior studies have generally showed that ANN outperforms LR classifiers in the task of
9 ACS prediction. Green et al.¹³ specifically compared the performance of ANN against LR, taking
10 into account computational data reduction using PCA, and found that the earlier provides
11 significant clinical advantage in risk stratification. Our data supported this notion that non-linear
12 models like ANN and GBM are more powerful tools to handle the high-dimensional, highly-
13 correlated nature of excessive ECG features. However, our data interestingly showed that
14 feature selection and annotation based on existing clinical knowledge can boost the
15 classification performance of linear models like LR. This is reasonable given that data reduction
16 and labeling could reduce the dimensionality and complexity in the data. Although this
17 significant improvement in LR classifier is yet to be compared against other data reduction
18 techniques in subsequent methodological studies, it has important technical implications. First, if
19 it is confirmed that simple linear classifiers can be equivalent to complex non-linear models such
20 as ANN, then future applications can focus on less computationally-exhaustive models like LR
21 classifiers. Second, and more importantly, LR has the appealing property of being fully
22 interpretable by clinicians, which can improve the clinical utility by removing the black-box
23 stigma of ML. Identifying a subset of features that are prevailing in the prediction of ACS can
24 shed the light on some hidden important disease pathways in our current understanding of the
25 electrocardiographic presentation of acute myocardial ischemia.

1 Our study has several strengths that addressed some of the existing gaps in the
2 literature. First, unlike previous studies, we did not exclude ECGs confounded by baseline
3 abnormalities such as pacing and bundle branch blocks, which would significantly enhance the
4 generalizability of our approach to real world clinical settings. This was evident by the
5 performance of our model on the external validation prospective cohort that had ACS
6 prevalence similar to what is seen in real-world patient populations. Second, our dataset was
7 unique in that it used the prehospital 12-lead ECG rather than the initial ECG from the
8 emergency department. Both have been previously shown to be dissimilar;³⁷ the prehospital
9 ECG could capture the subtle and transient acute cardiac ischemia during its ongoing evolution,
10 changes which could be easily masked on ECGs obtained at the emergency department
11 following early treatment. To our knowledge, this is the first study to build a generalizable ML
12 classifier to predict the likelihood of ACS using only the prehospital ECG data.

13 Nevertheless, our study had some limitations. First, the number of the abstracted ECG
14 features was not proportional to the size of the dataset, which might have affected the
15 performance of our classifiers. Increasing the number of patients would probably lead to
16 increased performance. Second, although manual feature selection had a positive effect on the
17 performance of our ML classifiers, further data-driven techniques for feature selection need to
18 be further investigated. Third, the majority of commercial ECG software are designed based on
19 strict criteria geared toward rule in STEMI, which explain the very low sensitivity for NSTEMI-ACS
20 detection observed in this study. As such, the emphasis of our approach should focus on
21 comparing the performance of our ML classifiers against experienced clinicians. Finally,
22 although not widely used in the U.S., high sensitivity cardiac troponin I has been shown as an
23 indispensable rule out tool in patients with suspected ACS in the Emergency Department.³⁸ On
24 contrast, our algorithm has been shown to improve the net gain classification performance of
25 patients with ACS (rule in), plus it can aid decisions at the prehospital setting. A recent study

1 has shown that point-of-care troponin assays has a sensitivity of only 27% when used during
2 ambulance transport.³⁹

3 In conclusion, using features extracted from only the prehospital 12-lead ECG, we
4 arrived at a generalizable ML model that outperforms both commercial interpretation software
5 and experienced clinician interpretation. Such ultra-early, ECG-based clinical decision support
6 tool, when combined with the judgment of trained emergency personnel, could be imperative for
7 improving clinical outcomes and reducing unnecessary costs in patients with chest pain.
8 Furthermore, domain-specific knowledge can boost the classification performance of linear
9 models like LR, which has important implications for building user-friendly and acceptable
10 decision support tools for wider clinical use.

11 **METHODS**

12 ***Design & Settings***

13 The data set used in this paper was obtained from the EMPIRE study. EMPIRE is a
14 prospective observational cohort study that recruited consecutive, non-traumatic chest pain
15 patients transported by emergency medical services to one of three UPMC-affiliated tertiary
16 care hospitals (UPMC Presbyterian, Mercy, and Shadyside). As per prehospital medical
17 protocols, standard 10-second 12-lead ECGs are obtained on all patients with suspected ACS
18 during first medical contact. If the initial patient evaluation by paramedics was judged to be
19 highly suspicious for cardiac ischemia, then the ECG was transmitted to UPMC medical
20 command, where the raw digital ECG data are permanently stored. In the EMPIRE study, we
21 recruited all consecutive chest pain patients with transmitted ECG data. Data was collected
22 under a waiver of informed consent because there was no direct contact with patients, and
23 follow up data was collected after all routine medical care was completed. This study was
24 approved by the Institutional Review Board of University of Pittsburgh, and all relevant ethical
25 regulations on human experiments, including the declaration of Helsinki, have been followed.

1 The current data set consisted of 1251 patients from the EMPIRE study from which 30-
2 day follow up outcome data were available. To estimate the minimum sample size required for
3 adequate AUC analysis of new diagnostic tests, we used the methods described by Hajian-
4 Tilaki.⁴⁰ So that the maximum marginal error of estimates (precision) does not exceed 5% with
5 95% confidence level, at desired validation values of sensitivity and specificity of 90%, the
6 minimum sample size required for ACS detection given a prevalence of at least 15% is 927.
7 Moreover, given that ML does not follow the same statistical rules for sample size estimation,
8 we assessed the adequacy of sample size for our ML classifiers by evaluating models for
9 overfitting (common with inadequate sample size). In our analysis, our algorithms generalized
10 well from Cohort 1 to Cohort 2, suggesting that data were not over-fitted and the sample size
11 was adequate.

12 ***Study Outcome***

13 The primary outcome of the study was the presence of ACS (myocardial infarction or
14 unstable angina) during the primary indexed admission, defined according to the 4th Universal
15 Definition of MI guidelines as the presence of symptoms of ischemia (i.e. diffuse discomfort in
16 the chest, upper extremity, jaw, or epigastric area for more than 20 minutes) and at least one of
17 the following criteria: (1) elevation of cardiac troponin I (>99th percentile) with or without
18 subsequent development of diagnostic ischemic ECG changes during hospitalization, (2)
19 imaging evidence of new loss of viable myocardium or new regional wall motion abnormalities,
20 or (3) coronary angiography or nuclear imaging demonstrating >70% stenosis of a major
21 coronary artery with or without treatment.²¹ Two independent reviewers annotated available
22 medical data and adjudicated this outcome based on serial ECGs, results of cardiac diagnostic
23 tests (e.g. echocardiography, angiography, biomarkers lab test) and other pertinent information
24 (e.g. past medical record, prescribed medications). Patients discharged from the emergency
25 department were classified as negative for ACS if they had no 30-day adverse events.

1 Disagreements were resolved by a third reviewer. Finally, in subsequent sensitivity analyses,
2 we tested the performance of our algorithms in detecting patients with non-ST elevation ACS
3 (NSTEMI-ACS) after excluding patients with confirmed STEMI on their prehospital ECG and who
4 were sent to the catheterization lab emergently.

5 ***ECG Reference Standard***

6 Two independent physicians who were blinded from the study outcome evaluated the
7 12-lead ECG image of each patient. All ECGs were de-identified, labeled with study ID, and
8 were stored on a secure server. First, ECGs with excessive noise or ventricular tachycardia or
9 fibrillation were excluded from this analysis (Cohort 1 n=5/750; Cohort 2 n=2/501). All other
10 available ECGs, including those with pacing, bundle branch blocks, atrial fibrillation, or left
11 ventricular hypertrophy were included in the analysis. Second, each reviewer labeled diagnostic
12 ECG changes according to the fourth Universal Definition of Myocardial Infarction consensus
13 statement²¹ as two contiguous leads with (1) ST elevation in V2–V3 ≥ 2 mm in men ≥ 40 years,
14 ≥ 2.5 mm in men < 40 years, or ≥ 1.5 mm in women; or ST elevation ≥ 1 mm in other leads; (2)
15 new horizontal or downsloping ST depression ≥ 0.5 mm; or (3) T wave inversion > 1 mm in
16 leads with prominent R wave or R/S ratio > 1 . Finally, taking into account the prior criteria for
17 ST-T changes and all other ECG findings suspicious for ischemia (i.e., contiguous territorial
18 involvement, evidence of reciprocal changes, changes beyond those caused by secondary
19 repolarization, and lack of ECG evidence of non-ischemic chest pain etiologies), each reviewer
20 made a final determination about the likelihood of underlying ACS (yes / no). Disagreements
21 were resolved by a board-certified cardiologist.

22 Furthermore, to place the comparisons between our ML classifiers and ECG reference
23 standards in a context, we used the HEART score obtained at the ED as a reference to the
24 classification performance achieved in current clinical practice. We computed HEART score
25 (History, ECG, Age, Risk factors, Troponin) as described in details elsewhere.⁴¹

1

2 **ECG Data**

3 All digital ECG files were acquired using HeartStart MRX monitor-defibrillator at 500
4 samples/second (Philips Healthcare). Standard ECG signal pre-processing was completed
5 using manufacture-specific commercial software at the Philips Healthcare Advanced Algorithm
6 Research Center (Andover, MA). The raw digital ECG signals were first decompressed and the
7 ECG leads were extracted. Noise, artifact, and ectopic beats were removed, and the
8 representative average beat for each ECG lead was computed to eliminate residual baseline
9 noise and artifacts, yielding a high signal-to-noise ratio and stable average waveform signal for
10 each of the 12 leads. Feature extraction was performed on these representative beats.

11 First, from each of the 12 leads, the amplitude, duration, and/or area measures of the P
12 wave, Q wave, R wave, S wave, qR wave, rS wave, QRS complex, QRS peak, ST segment, T
13 wave, STT wave, QT interval, PP interval, RP interval, and SP interval were computed (k=384).
14 In addition, the amplitude of ST onset, ST peak, ST offset, and J+80, as well as ST slope were
15 computed from each lead (k=60). This yielded a total of 444 temporal ECG features. Then, all
16 representative beats were aligned and a set of global measures were obtained. Extracted
17 features included QRS, $J_{T_{end}}$, $J_{T_{peak}}$, $T_{peak-end}$, and QT interval measures (k=6); QRS and T axes
18 from the frontal, horizontal, and XYZ planes (k=16); spatial angle between QRS and T
19 waveforms (k=6); inflection, amplitude, and slope of global QT, QRS, and T wave in frontal and
20 horizontal planes (k=56); ratios between PCA eigenvalues of QRS, STT, J, and T subintervals
21 (k=13); T wave morphology and loop (k=7); signal noise values (k=6); regional MI scar using
22 Selvester score (k=19); and injury vector gradient and amplitude (k=14). This yielded a total of
23 143 spatial ECG features. All extracted ECG features were then z-score normalized.

24 We had a total of 587 temporal-spatial ECG features extracted from each 12-lead ECG.
25 First, to safeguard against systematically missing data due to noise and artifact, which are

1 usually common in prehospital setting (e.g., unsticking of electrodes), we manually evaluated
2 each record to exclude ECGs of poor quality or with failed leads. We found that a very small
3 subset of patients had uninterpretable ECGs due to excessive noise (< 3%). We speculate that
4 this low rate is because we enrolled only patients with ECG transmitted to medical command
5 center. Paramedics routinely repeat poor or problematic ECGs in the field before they transmit
6 to a command physician for medical consultations. After excluding these poor ECGs from
7 further analysis, subsequent dataset preparation identified a subset of features ($n = 33$) that
8 were completed unbalanced (i.e., <5% with non-zero values). Upon further evaluation by clinical
9 experts, we found out that zero values were the normal variant on most of these ECG features.
10 For instance, there are usually no S waves in leads II, aVL, V5, and V6, and most leads have no
11 Q waves, which means it is acceptable to see “zero” values for these features. After removing
12 these features, the final dataset included 554 features available for training ML classifiers. Any
13 residual missing values at random were imputed using the mean or mode.

14 Next, to evaluate the complexity of the non-linear correlations in ECG features
15 evaluated, we used recursive feature elimination technique to identify the most important
16 features nested in the developed ML classifiers. Figure 4 shows some selected features with the
17 2-dimensional display scatterplot matrices. As expected, linear correlations failed to separate
18 patients with or without the disease, computationally favoring non-linear classifiers like GBM
19 and ANN over linear classifiers like LR. As such, two expert clinician-scientists reviewed the
20 important ECG features and identified the features that were clinically relevant to the
21 pathogenesis of myocardial ischemia. The following 65 features were identified: (1) amplitude of
22 J+80 and T wave from each of the 12 leads ($k=24$); (2) QRS, $J_{T_{end}}$, $J_{T_{peak}}$, $T_{peak-end}$, and QT
23 interval measures ($k=6$); (3) QRS and T axis in the frontal plane ($k=2$); (4) spatial angle between
24 QRS and T waveforms ($k=6$); (5) inflection, amplitude, and slope of T wave in frontal plane

1 (k=5); (6) ratios between PCA eigenvalues of QRS, STT, J, and T subintervals (k=13); (7) T
2 wave morphology and loop (k=7); and signal noise values (k=2).

3 Finally, we annotated this reduced subset of 65 features to denote normal vs. abnormal
4 thresholds based on published cutoff clinical values. Symbolic dynamics is a mathematical
5 modeling approach that aims to convert infinite dynamical systems into discrete intervals each
6 of which denotes a particular state, with the discrete labels (dynamics) given by the shift
7 operator. This step seemed necessary given that many ECG features are non-linear. For
8 example, both a T wave amplitude $< 0 \text{ mV}$ or $> 1 \text{ mV}$ usually indicate a change of state (i.e.,
9 myocardial ischemia) along the continuum of T wave amplitude value. This clinically-annotated
10 and reduced subset was used during the final stage of model development.

11 ***Clinical Data***

12 Although the ECG is the only diagnostic tool available during early triage, age and sex
13 are important predictors of ACS during first medical contact. In fact, ECG data is always age
14 and sex normalized, that is users have to enter these values into the ECG machine before ECG
15 acquisition; and these entries are used in the machine-provided, rule-based interpretations. The
16 guideline recommendations used by clinicians were age- and sex-specific as well.²¹ Given that
17 we aimed to compare our ML classifiers to the reference standard by expert clinicians and
18 available commercial software, we therefore decided to keep the age and sex as input features
19 in all of our ML classifiers as we tuned our models.

20 ***Machine Learning Classifiers***

21 All analyses were performed using Matlab R2013a. We performed supervised learning
22 using a menu of different ML classifiers and selected for further training and validation the
23 algorithms that had the best performance: LR, GBM, and ANN. Other ML algorithms explored
24 but did not seem to perform as the selected ones include: SVM, Naïve Bayes, random forest,

1 etc. Regardless of the tuning of their parameters, these latter methods performed poorly and
2 were therefore excluded from the study.

3 The selected classifiers provide complementary values for predicting and classifying dire
4 outcomes in clinical research. LR classifiers simplify the relationship between the input and the
5 output, but becomes computationally expensive and limited when the relationship in the model
6 is complex. In contrast, GBM builds an efficient classifier by using regression decision trees as
7 weak learners, and combining them into a single stronger learner. ANN models complex
8 relationships between the input and the output thanks to its hidden layers, the activation
9 functions of its neurons, and the back propagation method updating its unit weights.

10 The ANN implemented in this study had one hidden layer with a number of hidden units
11 adjusted to each version of the data set. The network used the rectified linear unit activation
12 function for the forward propagation and the Adam solver to perform the back propagation and
13 update the units' weights. These parameters were determined via a grid search, trying to
14 optimize the mean AUC on the test splits of the 10-fold cross-validation. In the same way, the
15 parameters of LR and GBM were determined using the same grid search optimization
16 approach. The LR model used limited-memory BFGS as optimization solver, the value of the
17 regularization parameter was adjusted on each version of the data set, and dedicated weights
18 were attributed to each sample in order to account for the class imbalance. The GBM model
19 used the deviance as loss function, and the values of the learning rate and the number of
20 boosting iterations were adjusted on each version of the data set.

21 To train and validate the models and account for the imbalance of the output classes,
22 the training data (n=745) was split into stratified 10-fold cross-validation sets, hence, the classes
23 were equally distributed from one split to the other. We then tested the performance of each
24 classifier on the independent test set (n=499). The model development and testing went into
25 three stages: (1) Using all extracted ECG features as input (LR₅₅₄, GBM₅₅₄, and ANN₅₅₄); (2)

1 Using only clinically-relevant EC features as input (LR₆₅, GBM₆₅, and ANN₆₅); and (3) Using the
2 clinically-relevant ECG features labeled according to published cutoff standards as input
3 (LR_{554+L}, GBM_{554+L}, and ANN_{554+L}). Finally, we selected the six classifiers with the best low bias –
4 low variance tradeoff and created a simple hybrid / fusion model based on vote count.
5 Classifications assigned by each model on the test set (disease vs. no-disease) were used as
6 votes. We considered 3 or more votes as the threshold to compute diagnostic performance
7 values for this fusion model.

8 ***Algorithm Performance & Statistical Testing***

9 The different performance metrics we refer to in this paper are the AUC, sensitivity,
10 specificity, and positive and negative predictive values (PPV and NPV) evaluated for a certain
11 classification cutoff value. We used 10-fold cross validation in Cohort 1 to estimate the statistical
12 uncertainty around ROC curves. To incorporate the uncertainty in the selection of cutoffs, we
13 first calculated the mean ROC of the 10 folds, then selected the ROC coordinate point that
14 maximized the sensitivity and kept the specificity above the arbitrary minimum level of 70%.
15 This selection aimed toward creating a good rule in model but with acceptable specificity. The
16 ROC cutoff value corresponding to this optimum was thus determined for each classifier studied
17 on the training set (Cohort 1), and the corresponding diagnostic metrics were computed on the
18 test set (Cohort 2). This independent testing supported whether or not the trained models
19 behaved consistently on new unseen patients. Lastly, we compared the performance of our final
20 ML classifier against standard reference on the test set. We used DeLong's non-parametric
21 approach to compare two ROC curves derived from related sample, with a p-value of < 0.05
22 (two-sided) indicating a significant difference between two diagnostic tests.⁴² We then computed
23 the net reclassification improvement (NRI) for our ML algorithm against the standard reference
24 using previously published methods.⁴³

25

1 **DATA AVAILABILITY**

2 The data that support the findings of this study are available on request from the
3 corresponding author S.S.A. The data are not publicly available due to intellectual
4 property claims under U.S. Patent and Trademark Office. However, source data for
5 figure 3 and Table 2 are provided with the paper.

6

7 **CODE AVAILABILITY**

8 The algorithms developed in this study are associated with a pending patent application
9 with IP rights under University of Pittsburgh, Pennsylvania, United States.

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END NOTES

AUTHOR CONTRIBUTION

S.S.A., C.M.G., C.C., and E.S. conceived the idea and supervised the research. Z.F. and S.F. supervised the dataset creation and annotation. S.S.A. and R.G. supervised electrocardiogram signal processing and feature extraction. Z.F., C.M.G, and C.C. supervised clinical outcomes adjudication. Z.F. and S.S. supervised the electrocardiograms annotation and reference standard adjudication. L.B., Z.B. and S.S.A., E.S. performed features engineering and developed the machine learning algorithms. S.S.A, Z.B., and E.S. performed statistical analyses and interpreted findings. The manuscript was written by S.S.A, L.B., Z.B., and E.S., with input from Z.F. and S.F., and comments from all other coauthors.

COMPETING INTERESTS

The algorithms developed in this study are associated with a pending PCT patent application # WO2016168181A1 invented by S.S.A, E.S., and C.C. The remaining authors declare no competing interests.

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1 **FIGURE LEGENDS**

2 **Figure 1:** *Stages of dataset derivation and preparation prior to developing the ML classifiers*

3 We used all available ECG features (k=554), selected ECG features (k=65), and
4 selected and relabeled ECG features (k=65+L) to train and test our machine learning
5 (ML) classifiers: logistic regression (LR), gradient boosting machine (GBM), and artificial
6 neural networks (ANN). Cohort 1 was used for training and Cohort 2 was used for
7 independent testing. The primary study outcome was acute coronary syndrome (ACS).

8 **Figure 2:** *Classification performance using Machine Learning Classifiers*

9 This figure shows the ROC curves of logistic regression (LR), gradient boosting machine
10 (GBM), and artificial neural network (ANN) classifiers using (A) all available ECG
11 features (k=554), (B) selected ECG features (k=65), and (C) selected and relabeled
12 ECG features (k=65+L). Cohort 1 was used for training with 10-fold cross validation, the
13 figure shows mean ROC curve with ± 2 standard errors. Cohort 2 was used for
14 independent testing using the algorithm trained on Cohort 1.

15 **Figure 3:** *Classification Performance of Final Model on Testing Set (n=499)*

16 This figure compares the ROC between our machine learning (ML) fusion model against
17 experienced clinicians and against rule-based commercial interpretation software for
18 detecting (A) any acute coronary syndrome (ACS) event, and (B) non-ST elevation acute
19 coronary syndrome events (NSTE-ACS). (***) $p < 0.001$ using two-sided DeLong's non-
20 parametric approach.

21 **Figure 4:** *Scatterplot matrix of some selected features using recursive feature elimination*

22 This figure shows selected ECG features with the 2-dimensional display scatterplot
23 matrices. These plots show how linear correlations fail to separate patients with or
24 without acute coronary syndrome (ACS), which explains why non-linear classifiers were
25 computationally favored over linear classifiers in our study.

1 **Table 1: Baseline Patient Characteristics**

	Cohort 1 (N=745) (Training and testing)	Cohort 2 (N=499) (External validation)
<u>Demographics</u>		
Age in years	59 ± 17	59 ± 16
Sex (Female)	317 (42%)	243 (49%)
Race (Black)	301 (40%)	202 (40%)
<u>Past Medical History</u>		
Hypertension	519 (69%)	329 (66%)
Diabetes mellitus	196 (26%)	132 (26%)
Old myocardial infarction	205 (27%)	122 (24%)
Known CAD	248 (33%)	179 (36%)
Known heart failure	130 (17%)	74 (15%)
Prior PCI / CABG	207 (28%)	124 (25%)
<u>Presenting Chief Complaint</u>		
Chest Pain	665 (89%)	454 (91%)
Shortness of Breathing	250 (34%)	234 (47%)
Indigestion, Nausea, or Vomiting	117 (16%)	109 (22%)
Dizziness or Syncope	106 (14%)	79 (16%)
Palpitation	96 (13%)	62 (12%)
Other Atypical Symptoms	54 (7%)	37 (7%)
<u>Baseline ECG Rhythm</u>		
Normal Sinus Rhythm	648 (87%)	442 (88%)
Atrial Fibrillation	71 (9%)	46 (9%)
Pacing	26 (4%)	8 (2%)
Right Bundle Branch Block	31 (4%)	27 (5%)
Left Bundle Branch Block	19 (3%)	16 (3%)
Left Ventricular Hypertrophy	37 (5%)	24 (5%)
<u>Primary Study Outcome</u>		
Any ACS Event	114 (15.3%)	92 (18.4%)
Prehospital STEMI	31 (4.2%)	18 (3.6%)
NSTE-ACS	83 (11.1%)	74 (14.8%)
<u>Course of Hospitalization</u>		
Length of Stay (median [IQR])	2.3 [1.0–3.0]	1.2 [0.6-2.5]
Stress Testing with SPECT	180 (24%)	115 (23%)
Treated by Primary PCI / CABG	74 (10%)	65 (13%)
30-Day Cardiovascular Death	33 (4.4%)	24 (4.8%)

1 **Table 2:** Diagnostic Accuracy Measures on Testing set (n=499)

PREDICTING ANY ACS EVENT

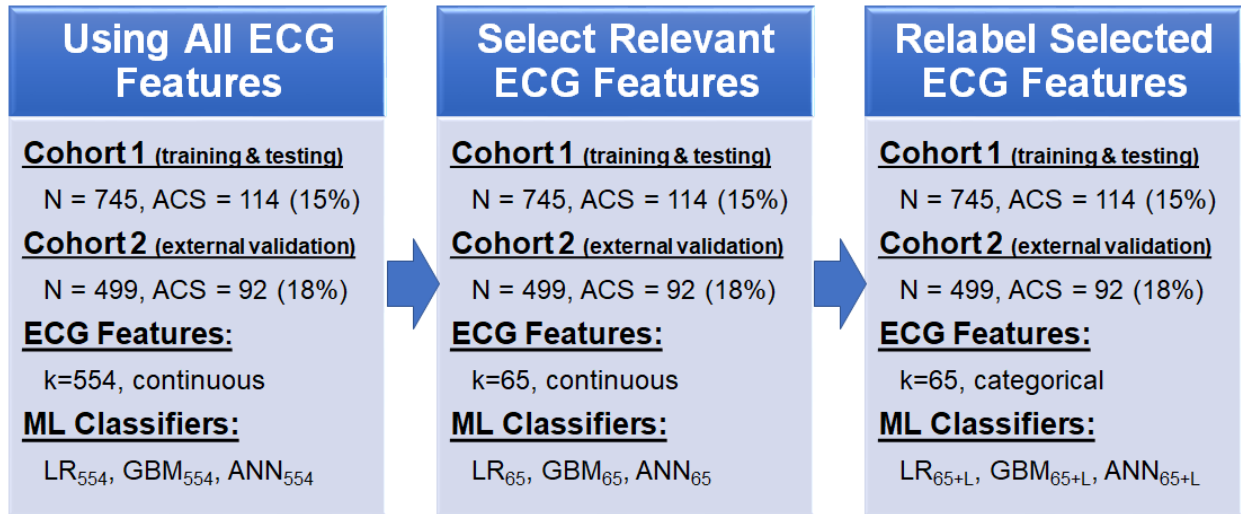
	ML FUSION MODEL	EXPERT ECG READ	AUTOMATED ECG READ
SENSITIVITY	0.77 (0.67 – 0.85)	0.40 (0.30 – 0.51)	0.25 (0.17 – 0.35)
SPECIFICITY	0.76 (0.72 – 0.81)	0.94 (0.92 – 0.96)	0.98 (0.97 – 0.99)
PPV	0.43 (0.38 – 0.48)	0.63 (0.51 – 0.73)	0.79 (0.62 – 0.90)
NPV	0.94 (0.91 – 0.96)	0.87 (0.86 – 0.89)	0.85 (0.82 – 0.88)

PREDICTING NSTE-ACS EVENTS

	ML FUSION MODEL	EXPERT ECG READ	AUTOMATED ECG READ
SENSITIVITY	0.72 (0.60 – 0.81)	0.26 (0.16 – 0.37)	0.12 (0.06 – 0.22)
SPECIFICITY	0.76 (0.72 – 0.80)	0.94 (0.92 – 0.96)	0.98 (0.97 – 0.99)
PPV	0.36 (0.31 – 0.41)	0.46 (0.33 – 0.60)	0.60 (0.35 – 0.80)
NPV	0.94 (0.91 – 0.93)	0.87 (0.85 – 0.89)	0.86 (0.85 – 0.87)

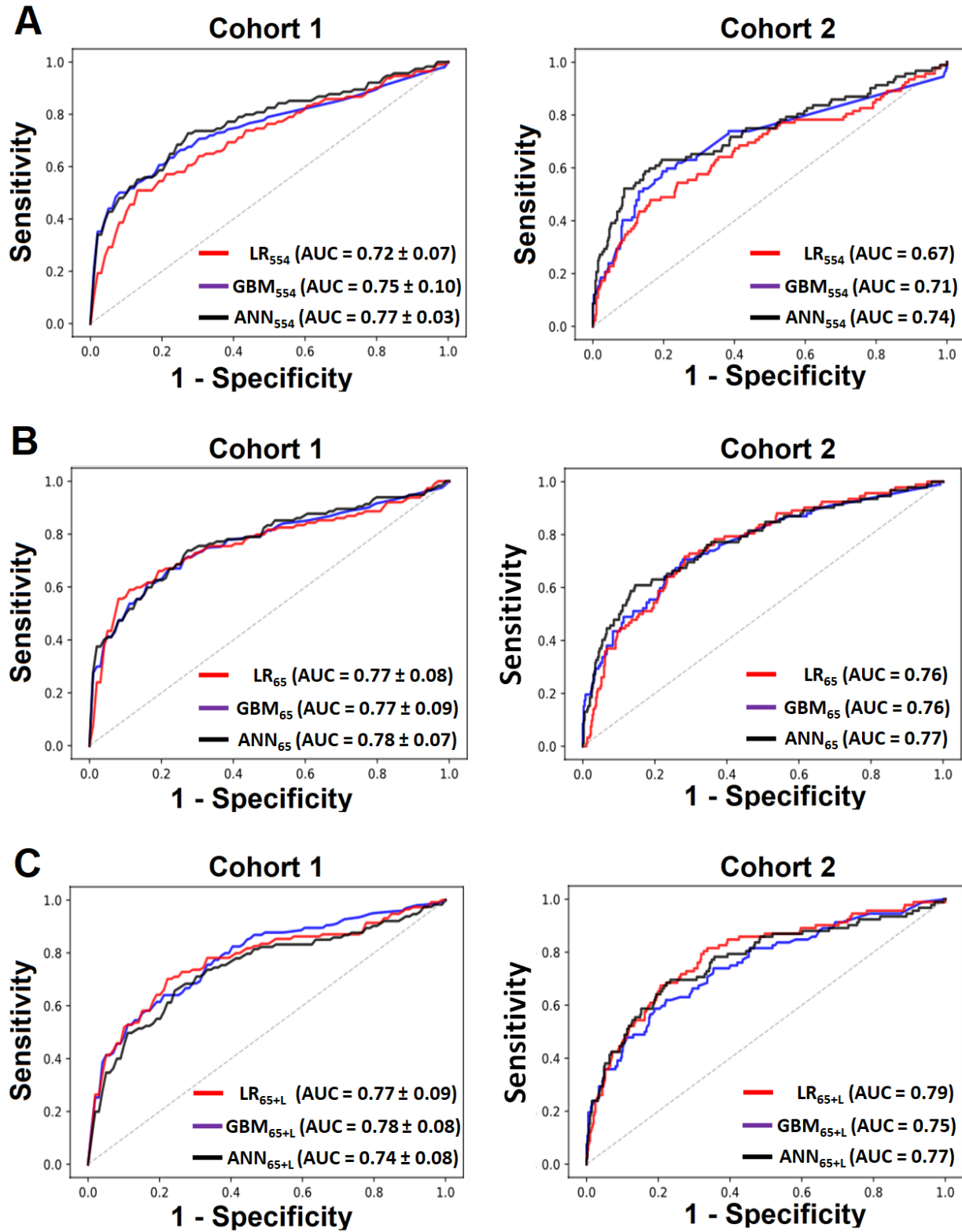
2 PPV: positive predictive value; NPV: negative predictive value

1 **Figure 1:** Stages of dataset derivation and preparation prior to developing the ML classifiers



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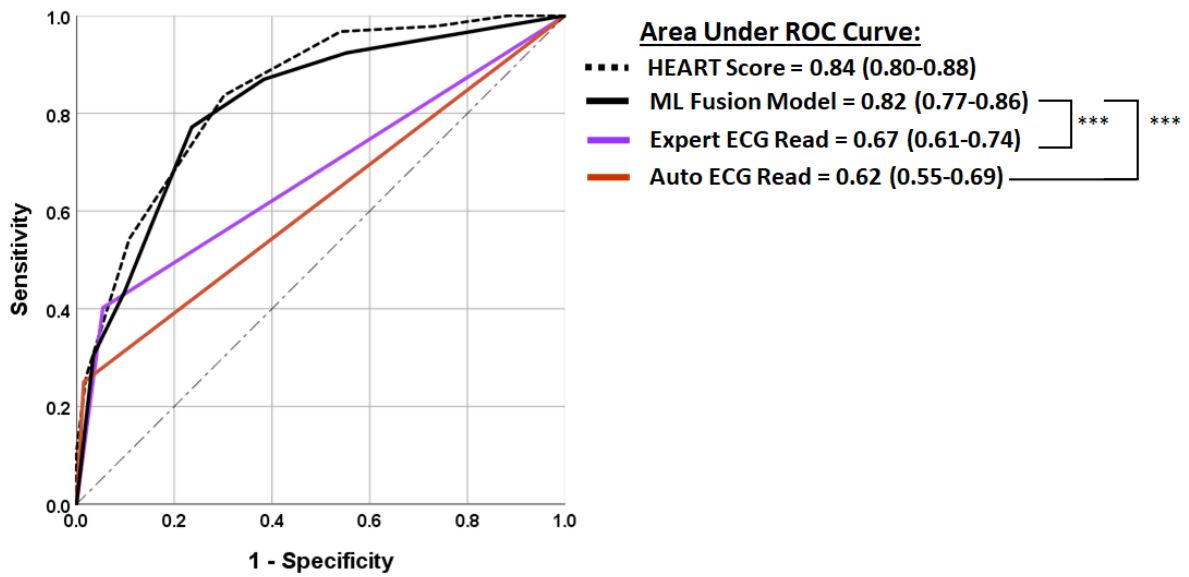
1 **Figure 2: Classification performance using Machine Learning Classifiers**



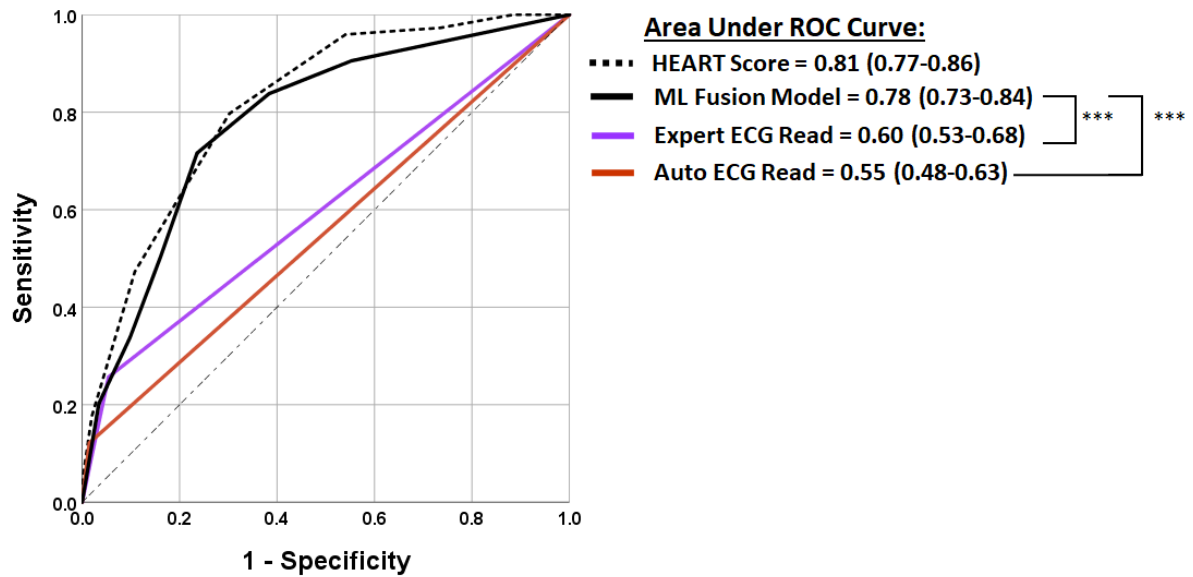
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1 **Figure 3: Classification Performance of Final Model on Testing Set (n=499)**

A Predicting ACS Events (n=92, 18%)



B Predicting NSTEMI—ACS Events (n=74, 15%)



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1 **Figure 4:** Scatterplot matrix of some selected features using recursive feature elimination

