

1 **Machine learning to classify left ventricular hypertrophy using ECG feature**
2 **extraction by variational autoencoder**

3 Short title: ML models for LVH

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ABSTRACT

Background: Traditional ECG criteria for left ventricular hypertrophy (LVH) have low diagnostic yield. Machine learning (ML) can improve ECG classification.

Methods: ECG summary features (rate, intervals, axis), R-wave, S-wave and overall-QRS amplitudes, and QRS/QRST voltage-time integrals (VTIs) were extracted from 12-lead, vectorcardiographic X-Y-Z-lead, and root-mean-square (3D) representative-beat ECGs. Latent features were extracted by variational autoencoder from X-Y-Z and 3D representative-beat ECGs. Logistic regression, random forest, light gradient boosted machine (LGBM), residual network (ResNet) and multilayer perceptron network (MLP) models using ECG features and sex, and a convolutional neural network (CNN) using ECG signals, were trained to predict LVH (left ventricular mass indexed in women $>95 \text{ g/m}^2$, men $>115 \text{ g/m}^2$) on 225,333 adult ECG-echocardiogram (within 45 days) pairs. AUROCs for LVH classification were obtained in a separate test set for individual ECG variables, traditional criteria and ML models.

Results: In the test set (n=25,263), AUROC for LVH classification was higher for ML models using ECG features (LGBM 0.790, MLP 0.789, ResNet 0.788) as compared to the best individual variable ($\text{VTI}_{\text{QRS-3D}}$ 0.677), the best traditional criterion (Cornell voltage-duration product 0.647) and CNN using ECG signal (0.767). Among patients without LVH who had a follow-up echocardiogram >1 (closest to 5) years later, LGBM false positives, compared to true negatives, had a 2.63 (95% CI 2.01, 3.45)-fold higher risk for developing LVH ($p < 0.0001$).

Conclusions: ML models are superior to traditional ECG criteria to classify—and predict future—LVH. Models trained on extracted ECG features, including variational autoencoder latent variables, outperformed CNN directly trained on ECG signal.

47 **Keywords:** Left ventricular hypertrophy, LVH, machine learning, deep learning, artificial
48 intelligence, electrocardiogram, ECG, variational autoencoder.

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50 **Abbreviations:**

51 ECG, electrocardiogram

52 LVH, left ventricular hypertrophy

53 ML, machine learning

54 AI, artificial intelligence

55 MLP, multilayered perceptron

56 LGBM, light gradient-boosting machine

57 AUROC, area under the receiver operator characteristic curve

58 VAE, variational Autoencoder

59 LVMi, left ventricular mass indexed

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INTRODUCTION

Left ventricular hypertrophy (LVH) refers to increased left ventricular mass, characterized by an increase in left ventricular wall thickness and/or enlargement of the left ventricular cavity. This is often secondary to pathological or physiological stressors such as chronic hypertension, valvular heart disease, athletic training, or genetic conditions. LVH is associated with over a two-fold increase in cardiovascular morbidity and all-cause mortality (1). Early detection and initiation of pharmacological treatment, along with lifestyle modifications, have been associated with improved outcomes (2).

Transthoracic echocardiography is the standard-of-care for the diagnosis of LVH. However, despite its non-invasive nature and widespread utilization, universal screening for LVH using echocardiography even in high-risk groups, such as those with hypertension, is not cost-effective (3,4).

Electrocardiography (ECG) is an affordable, widely accessible, and frequently used diagnostic tool for cardiovascular screening. Often considered an extension of the cardiovascular physical examination, it is estimated that over 100-300 million ECGs are performed annually in the United States (5). Several criteria for 12-lead ECG diagnosis of LVH have been published over many decades, mainly based on the magnitude of QRS voltages in various—especially precordial—leads. However, these criteria have poor sensitivities in detecting LVH, making them unsuitable for standalone ECG screening (6-8). In a 2023 consensus statement, the International Society of Electrocardiology and the International Society for Holter Monitoring and Noninvasive Electrocardiology highlighted the need for a paradigm shift in ECG-based LVH diagnosis (9). The statement emphasized the limitations of traditional ECG criteria and discussed the potential of artificial intelligence (AI)-driven approaches for LVH detection.

83 Machine learning (ML) can reduce reliance on human interpretation and yet increase the
84 diagnostic accuracy of ECG (10,11). Several ECG-based ML models have been developed for
85 detecting LVH, with varying sensitivities and specificities (12). Many of these studies use
86 convolutional neural network (CNN) deep learning architecture to train models using ECG
87 signals often with fewer than 10,000 training ECGs. Given that each 12-lead 10-second ECG
88 signal at 500 Hz consists of 60,000 data points, using such a high-dimensionality input for ML
89 training with a limited number of samples can result in overfitting and reduced generalizability
90 (13-15). On the other hand, non-neural network ML architectures—such as logistic regression,
91 random forest, gradient boosted machine—are not suited to use high-dimensional ECG signal
92 data as input and are usually limited to using extracted ECG features with potential loss of
93 diagnostic information (15).

94 To mitigate these limitations—while preserving the advantages of deep learning—we
95 developed a variational autoencoder (VAE) that can encode 0.75-sec-representative-beat from
96 either X-Y-Z-lead or root-mean-squared ECG into 30 variables (15-17). These VAE latent
97 encodings retain the ECG morphological information and can reconstruct back the ECG signal
98 with high fidelity. In this study, we aimed to train and test different ML models using extracted
99 ECG features including the latent encodings or the ECG signal to classify LVH from the
100 representative-beat ECG.

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METHODS

103 Patient selection and data retrieval: An automated retrospective retrieval of records was
104 performed from our clinical database at the University of Kansas Medical Center between May
105 2010 and Jan 2022 to search for ECG and echocardiogram performed on the same patient within

106 45 days of each other. Echocardiograms-ECG pairs with echocardiographic left ventricular mass
107 index (LVMI) $>95 \text{ g/m}^2$ for females and $>115 \text{ g/m}^2$ for males were labelled as ‘LVH’ while rest
108 of the pairs were assigned to the ‘no LVH’ group (15). The study was conducted under an
109 approval from the Institutional Review Board.

110 Data extraction: ECGs were acquired with Philips 12-lead ECG machines. The 12-lead
111 ECG 10-second and 1200-ms-representative-beat signals along with standard features like heart
112 rate, PR interval, etc. were exported to a research SQL data server. Echocardiograms were
113 standard clinical studies performed for clinical indications both as outpatient and inpatient
114 evaluations. Individual echocardiogram numeric variables including diastolic measurements of
115 left ventricular internal diameter (LVIDd), interventricular septum (IVSd) and posterior wall
116 (PWd) from 2D parasternal long-axis view were extracted using a backend query in HERON
117 (Healthcare Enterprise Repository for Ontological Narration), a search discovery tool that
118 facilitates searches on various hospital electronic data sources (18,19). The query results were
119 recombined using medical record number, encounter number and study date to generate back the
120 list of variables belonging to each echocardiogram study. Left ventricular mass was calculated
121 using the American Society of Echocardiography recommended formula: $0.8 \times 1.04[(\text{LVIDd} +$
122 $\text{IVSd} + \text{PWd})^3]$ and indexed to body surface area (20).

123 ECG processing: The details of ECG processing performed using Python are provided in
124 prior publications (15,21,22). In summary, vectorcardiographic X-Y-Z-lead ECGs were
125 constructed from 12-lead ECGs using Kors’ matrix (23). Using these orthogonal X, Y, Z leads,
126 the root-mean-square (RMS or 3D) ECG was constructed. Voltage-time integrals (VTIs) were
127 obtained by the integration of the instantaneous voltage over the duration of QRS (VTI_{QRS}) or
128 QRS-T (VTI_{QRST}).

129 Traditional Criteria and Univariable Models: Based on review of literature, we selected 5
130 widely used ECG-based LVH diagnostic criteria for comparison, i.e. Peguero-Lo Presti criteria
131 ($\max S + S_{v4}$), Cornell voltage ($R_{avL} + S_{v3}$), Cornell voltage-duration product (VDP), Sokolow-
132 Lyon criteria ($S_{V1} + \max R_{(V5 \text{ or } V6)}$), and Gubner-Ungerleider criteria ($R_I + S_{III}$). We also selected
133 3 ECG variables for comparison namely QRS duration, $\text{amplitude}_{\text{QRS-3D}}$, and $\text{VTI}_{\text{QRS-3D}}$
134 (21,22,24). The latter 2 were calculated off the QRS from the RMS/3D ECG.

135 Variational Autoencoder: We trained a variational autoencoder (VAE) on 1.18 million
136 unlabeled ECG signals to encode a 0.75-sec segment centered on the representative beat ECG
137 signal into 60 variables (30 variables for X, Y, Z leads and 30 for RMS of these leads). The VAE
138 has a dual neural network architecture with the encoder taking the ECG input and outputting 30
139 latent variables, and the decoder inputting the 30 latent variables and outputting the ECG signal.
140 The network is rewarded in training to encode the signal such as to learn accurate reconstruction
141 of the original signal from the latent variables alone. Our VAEs are able to reconstruct the
142 original signal back from the latent variables with high fidelity (16,17,25). The X-Y-Z-lead and
143 RMS/3D representative-beat ECGs included in this study were processed using these 2 VAEs to
144 generate latent encodings or variables.

145 ECG Features: The following features were available for ML model training:

- 146 • Summary features like heart rate, PR interval, QRS duration, corrected QT interval (26),
147 frontal plane QRS axis, etc.
- 148 • From 16 leads—each of 12-leads, 3 X-Y-Z-leads and 1 RMS ECG—we obtained QRS
149 amplitudes, VTI_{QRS} , VTI_{QRST} , R-wave amplitudes, S-wave amplitudes.
- 150 • 30 latent variables each from VAEs trained to reconstruct the X-Y-Z-lead and RMS
151 representative-lead ECGs.

152 • Sex

153 Model Training and Testing: Approximately 10% of the medical record numbers in the
154 dataset were withheld as the testing set, and remainder used for model training (**Figure 1**). We
155 trained the following ML architectures on the training set – logistic regression, random forests,
156 light gradient boosted machine (LGBM), residual neural network (ResNet), multilayered
157 perceptron (MLP) and CNN. The CNN was trained on the representative-beat X-Y-Z-lead ECG
158 signal, and the other 5 ML models trained on the extracted ECG features (as above) plus sex. Sex
159 was provided to the models as the definition of LVH is sex specific. The results are reported
160 from the performance of the trained models in the holdout test set. We also report the models’
161 performance in 4 subgroups based on intraventricular conduction – QRS duration <120 ms,
162 typical right bundle branch block (RBBB, QRS duration ≥ 120), typical left bundle branch block
163 (LBBB, QRS duration ≥ 120 ms), and interventricular conduction delay (IVCD, QRS duration \geq
164 120 ms but not meeting either RBBB or LBBB criteria). American Heart Association-American
165 College of Cardiology Foundation-Heart Rhythm Society criteria for bundle branch blocks were
166 used (27).

167 Statistical analysis: Continuous variables are reported as mean \pm standard deviation, and
168 categorical variables as percentages. Comparisons were made using Student’s t-test for
169 continuous variables and χ^2 -test for categorical variables. Statistical analysis was conducted in
170 Python version 3.12.7 and 2-tailed p-value of less than 0.05 was considered statistically
171 significant.

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175 **RESULTS**

176 Patient characteristics: A total of 250,596 ECG-echocardiogram pairs were included, with
177 149,612 (59.7%) pairs belonging to females. The mean age of the overall population of ECG-
178 echocardiogram samples was 63.8 ± 15.3 years. In the training sets, 40,839 (28.2%) of the
179 female samples and 23,309 (24.3%) male samples had LVH on echocardiography. The testing
180 set consisted of 25,263 ECG-echocardiogram pairs. In the testing set, 4470 (27.8%) female
181 samples and 2672 (24.6%) male samples had LVH. The detailed distributions of the ECG and
182 echocardiographic variables in the testing set are shown in **Table 1** and for the training set in
183 **Supplementary Table 1**. The testing samples were divided into 4 subgroups i.e. narrow QRS
184 <120 ms ($n=215,228$), typical RBBB ($n=24,800$), typical LBBB ($n=13,893$), and IVCD
185 ($n=13,714$).

186 LVH classification models: The testing set performance of the 3 univariable models, 5
187 traditional criteria and the 6 ML models is summarized in **Table 2** and **Supplementary Table**
188 **2A-D**.

189 Univariable models: Amongst the linear univariable models, VTI_{QRS-3D} was the best
190 predictor of LVH in the overall population, with an AUROC 0.677. Further, VTI_{QRS-3D}
191 performed the best in all subgroups except in typical LBBB (narrow QRS 0.659, RBBB 0.674,
192 LBBB 0.585, IVCD 0.578). In typical LBBB, $amplitude_{QRS-3D}$ performed the best, with an
193 AUROC 0.590.

194 Traditional criteria: Overall, the performance of traditional ECG criteria for predicting
195 LVH was poor, with AUROCs ranging from 0.507 to 0.647. Cornell VDP was the best
196 performing criteria overall and in narrow QRS subgroup (overall 0.647; narrow QRS 0.643). In

197 other subgroups, Peguero-Lo Presti criteria performed the best (RBBB 0.598, LBBB 0.572,
198 IVCD 0.578). In general, these criteria performed better in females as compared to males.

199 ML Models: All ML models outperformed the traditional criteria and univariate models.
200 LGBM (AUROC 0.790), MLP (0.789) and ResNet (0.788), which were trained on ECG features
201 including VAE latent encodings and sex, were the best performing models in the overall
202 population. The CNN model, which was trained on the raw ECG signal alone, demonstrated an
203 AUROC 0.767. The ROC curves, separately for females and males, for the top 4 ML models vis-
204 à-vis the best univariable and best traditional criteria are plotted in **Figure 2**.

205 When evaluated in the 4 ECG subgroups by intraventricular conduction, models with
206 highest AUROCs were LGBM in narrow QRS (0.785), MLP in RBBB (0.778) and LBBB
207 (0.698) and ResNet in IVCD (0.720). The ROC curves of the best model each amongst
208 univariable, traditional criteria and ML for each of the 4 subgroups separately for females and
209 males is shown in **Figure 3 and 4**.

210 Linear analysis of LGBM prediction probabilities: LVMi was plotted against the
211 prediction probabilities output generated by LGBM model for females and males as shown in
212 **Figure 5**. A strong linear trend between prediction probabilities and LVMi can be noted for both
213 females and males (respectively R^2 0.851 and 0.833, or correlation coefficient ρ 0.922 and 0.913).

214 Longitudinal analysis of LVH negatives: Among false positives and true negatives
215 produced by the LGBM model in the testing set, we searched for the ECG-echocardiogram pairs
216 where a follow-up echocardiogram >1 year and closest to 5 years later was available for further
217 analysis. We used a 2x2 table to compare the development of LVH in 161 false-positive as
218 compared to the 1,019 true-negative samples. On mean follow-up of 3.9 ± 1.8 years, 54/161
219 (33.5%) patients in false-positive group, and 130/1019 (12.8%) patients in true-negative group

220 developed LVH. The risk ratio for development of LVH was 2.63 (95% CI 2.01, 3.45) in false-
221 positives compared to true-negatives from the LGBM model (**Table 3**).

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DISCUSSION

224 To the best of our knowledge, this is the largest evaluation of ECG criteria and ML models for
225 predicting LVH till date. We have applied the innovative framework of using DL-based latent
226 space ECG encodings for building ML models, which allows simpler models to make accurate
227 predictions without overfitting.

228 ***Salient findings:*** First, traditional ECG-based criteria demonstrate suboptimal
229 performance in diagnosing LVH, with the Cornell VDP showing the highest accuracy among
230 them (AUROC 0.647). Second, univariable models including QRS duration, $\text{amplitude}_{\text{QRS-3D}}$,
231 and $\text{VTI}_{\text{QRS-3D}}$ were at par or better than traditional criteria for the diagnosis of LVH, with
232 $\text{VTI}_{\text{QRS-3D}}$ achieving the best overall results (AUROC 0.677). Third, ML models outperform both
233 traditional and univariable models, with LGBM models demonstrating the highest performance
234 in our study (overall AUROC 0.790). Last, the performance of traditional, univariable, and ML
235 models vary across sex and QRS morphologies. Further, the LGBM model trained on ECG latent
236 encodings and features successfully captured the underlying trend of LVM_i , showing strong
237 correlation and predicting future development of LVH.

238 ***Univariable models:*** Previous studies have demonstrated the utility of linear univariable
239 predictors of LVH, such as QRS duration and QRS-VTIs (22,31). In our analysis, we evaluated
240 QRS duration, $\text{amplitude}_{\text{QRS-3D}}$, and $\text{VTI}_{\text{QRS-3D}}$ for predicting LVH across various subgroups. Our
241 findings indicate that these measures generally outperform traditional LVH criteria. Among them,
242 $\text{VTI}_{\text{QRS-3D}}$ emerged as the best overall criteria, except in the typical LBBB subgroup, where

243 amplitude_{QRS-3D} was superior. Similar to Cornell VDP, VTI_{QRS-3D} incorporates both QRS voltage
244 and duration. Since VTI_{QRS-3D} is calculated from the reconstructed 3D-orthogonal leads,
245 ostensibly, it captures the QRS complex more comprehensively as compared to Cornell VDP,
246 which uses information from a pair of 2-D leads (V3 and aVL).

247 ***Traditional ECG criteria:*** As demonstrated in previous studies, our analysis reaffirmed
248 the poor discrimination of LVH offered by standard electrocardiographic criteria using a large
249 dataset (28,29). Unlike other voltage-based rules, Cornell VDP, which emerged as the best
250 overall criterion, accounts for both QRS voltage and duration in its calculation. Both of these
251 parameters are affected in LVH (30). In the subset of ECGs with conduction abnormalities
252 (RBBB, LBBB, and IVCD), Peguero-Lo Presti criteria performed better than Cornell VDP.
253 Although the difference in performance was marginal, if this trend is real, it could be explained
254 by obfuscation of LVH-related changes in QRS duration due to QRS prolongation inherent to
255 conduction delays. However, this cannot be verified in our study. Notably, compared to the
256 combined population, individual criteria generally performed better in females and males
257 separately. This underscores the importance of using different cut-off values for females and
258 males, recognizing the sex-based differences in ECGs and definition of LVH. (28,29).

259 ***ML models:*** We tested several ML architectures for LVH prediction, including simple
260 models (LR), tree-based models (RF, LGBM), and neural networks (ResNet, MLP, and CNN).
261 The LGBM model demonstrated the best overall performance (AUROC 0.790), with AUROCs
262 comparable to those of the MLP (0.789) and ResNet (0.788) models. The performance of all the
263 models was worse in the subgroups with conduction abnormalities. MLP was the best
264 performing model in typical RBBB and LBBB subgroups (0.778 and 0.698) while ResNet

265 performed the best in the IVCD subgroup (0.720). Nevertheless, it is important to note that the
266 differences in the performance these models were only marginal.

267 We further evaluated the interpretability and physiological relevance of the LGBM model.
268 First, we plotted the prediction probabilities from this model against LVMi, which showed a
269 strong linear positive correlation, suggesting that the model captures meaningful physiological
270 patterns rather than artificial class boundaries. Second, we analyzed the false positives produced
271 by this model for future development of LVH, finding that the false positives were more than 2.5
272 times as likely to develop LVH in the future compared to true negatives. This indicates that the
273 model captures underlying ECG abnormalities even before patients meet the criteria for overt
274 LVH diagnosis.

275 ***Previous literature:*** In a recently published study from China, Zhu et al. used a large
276 dataset comprising of over 90,000 ECGs to create deep learning multilabel classifier algorithms.
277 They achieved AUROCs ranging from 0.78-0.92 using their 12-lead model, and showed that a
278 reduced 4-lead model using lead I, aVR, V1 and V5 had equivalent performance (32). In a
279 Taiwanese study, Liu et al. developed a deep learning model for predicting LVH using
280 approximately 23,000 training samples (33). They achieved high AUROCs ranging from 0.83-
281 0.89 across different testing sets. However, the definition of LVH used in this study was different,
282 using LV mass >186 g for females and >258 g for males. In a South Korean study, Kwon et al.
283 developed an ensemble deep neural network + CNN model using approximately 36,000 training
284 samples, combining information from ECG signal, ECG features, and patient demographics (34).
285 While using higher cut-off values for LVMi (109 g/m² females and 132 g/m² males), their model
286 achieved AUROCs ranging from 0.87-0.88 in testing sets.

287 In a study from Massachusetts General Hospital, Haimovich et al. create ML models for
288 predicting LVH in specific disease populations like cardiac amyloidosis, hypertrophic
289 cardiomyopathy, aortic stenosis, and others using a total of 34,258 training samples (35). Similar
290 to our approach, they used a pretrained deep learning model to produce latent encodings and
291 trained a simpler classifier for LVH classification although they used full 10-second ECG signal
292 instead of representative beat ECG. Their model achieved AUROCs ranging from 0.69 to 0.96 in
293 various subgroups. Khurshid et al. used data from the UK Biobank to create a CNN model
294 trained on 32,000 samples and achieved AUROCs ranging from 0.62 to 0.65 in predicting LVH.
295 Owing to heterogeneity in study populations, data structures, and labels for LVH, it is difficult to
296 evaluate the performance of models across studies. Nonetheless, the AUROCs attained by ML
297 models in our study are comparable to previous work.

298 Limitations: Our work is best understood in the context of its limitations. Both training
299 and testing sets for the models were from a single center, and these models might have sub-
300 optimal performance when generalized to other datasets. Further, since the median beat ECGs
301 were derived from a proprietary system, additional steps may be required in processing ECGs
302 from other systems. Additionally, to calculate ECG parameters for traditional criteria and
303 univariate models, automated feature extraction was done, which might not be as accurate as
304 expert-created labels.

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306 CONCLUSIONS

307 Traditional voltage-based criteria for ECG diagnosis have poor diagnostic performance. Simple
308 univariable models, especially VTI_{QRS-3D} , perform better than the traditional criteria. ML
309 techniques can significantly enhance the accuracy of ECG-based diagnosis of LVH over both

310 traditional voltage-based criteria and univariable models. Dimensionality reduction of ECG
311 using variational autoencoder can facilitate utilization of non-deep learning ML architectures,
312 which may otherwise struggle with high dimensionality of ECG data. Further external testing
313 and testing is needed for clinical utilization of these ML models.

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TABLE LEGENDS

- Table 1.** Patient characteristics of the testing set.
- Table 2.** Model performance for LVH prediction in the entire testing set. Area under receiver-operating characteristic curve (AUROC) and sensitivity at specificity fixed at 0.75 are provided.
- Table 3.** Comparison between presence of LVH on subsequent echocardiogram (>1 year and closest to 5 years after index echocardiogram) in false positives versus true negatives of LVH LGBM model in testing set

FIGURE LEGENDS

- Figure 1.** Data pipeline for model training and testing
- Figure 2.** ROC curves from the entire testing set for males (*left panel*) and females (*right panel*).
- Figure 3.** ROC curves for subgroups of testing set in females, narrow QRS (*top left*), typical right bundle branch block (RBBB, *top right*), typical left bundle branch block (LBBB, *bottom left*), intraventricular conduction delay (IVCD, *bottom right*).
- Figure 4.** ROC curves for subgroups of testing set in males, narrow QRS (*top left*), typical right bundle branch block (RBBB, *top right*), typical left bundle branch block (LBBB, *bottom left*), intraventricular conduction delay (IVCD, *bottom right*)
- Figure 5.** Scatterplots of echocardiographic left ventricular mass indexed (LVMI) plotted against prediction probabilities from the LGBM model for females (*left panel*) and males (*right panel*).

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- 459
- 460

461 **Table 1.** Patient characteristics of the testing set.

	Females			Males		
	No LVH	LVH	p	No LVH	LVH	p
n (%)	10782 (71.6%)	4268 (28.4%)		7466 (73.1%)	2747 (26.9%)	
Age (years), mean±S.D.	61.6 ± 16.6	67.4 ± 16.1	< 0.0001	63.2 ± 14.9	67.0 ± 13.6	< 0.0001
QRS duration (ms), mean±S.D.	96.1 ± 23.3	108.6 ± 27.8	< 0.0001	104.8 ± 25.0	123.6 ± 33.2	< 0.0001
Frontal plane QRS axis (°), mean±S.D.	28.5 ± 53.1	20.3 ± 66.3	< 0.0001	24.2 ± 58.7	24.9 ± 82.3	< 0.0001
Amplitude _{QRS-3D} (µV), mean±S.D.	938.7 ± 405.5	1128.9 ± 525.5	< 0.0001	975.1 ± 416.6	1147.5 ± 525.1	< 0.0001
VTI _{QRS-3D} (nVs), mean±S.D.	30507.8 ± 16684.7	42976.0 ± 24444.6	< 0.0001	34575.5 ± 17143.0	49621.2 ± 25866.2	< 0.0001
QTc (ms), mean±S.D.	431.6 ± 40.1	450.0 ± 45.5	< 0.0001	433.5 ± 43.2	458.9 ± 49.5	< 0.0001
LVEF (%), mean±S.D.	57.7 ± 10.3	50.9 ± 15.6	< 0.0001	54.4 ± 12.4	43.7 ± 17.5	< 0.0001
LV mass index (g/m ²), mean±S.D.	70.2 ± 14.2	122.6 ± 224.6	< 0.0001	83.2 ± 17.7	150.2 ± 192.3	< 0.0001
ECG-defined subgroups			<0.0001			<0.0001
Narrow QRS ^a	9279 (86.1%)	3091 (72.7%)		5836 (78.5%)	1556 (56.8%)	
Typical RBBB ^b	664 (6.2%)	439 (10.3%)		902 (12.1%)	530 (19.4%)	
Typical LBBB ^b	434 (4.0%)	375 (8.8%)		253 (3.4%)	296 (10.8%)	
IVCD ^b	404 (3.7%)	346 (8.1%)		448 (6.0%)	356 (13.0%)	

462
463 RBBB, Right Bundle Branch Block; LBBB, Left Bundle Branch Block; IVCD, Intraventricular
464 Conduction Delay
465 ^aQRS duration <120 ms, ^bQRS duration ≥120 ms
466

467 **Table 2.** Model performance for LVH prediction in the entire testing set. Area under receiver-
 468 operating characteristic curve (AUROC) and sensitivity at specificity fixed at 0.75 are provided.

Testing set	Combined (n=25,263)		Females (n=15,050)		Males (n=10,213)	
	AUROC	Sensitivity*	AUROC	Sensitivity*	AUROC	Sensitivity*
<i>Univariable models^a</i>						
▪ QRS duration	0.648	0.462	0.651	0.452	0.667	0.46
▪ Amplitude _{QRS-3D}	0.601	0.389	0.602	0.395	0.603	0.388
▪ VTI _{QRS-3D}	0.677	0.515	0.671	0.504	0.699	0.543
<i>Traditional criteria</i>						
▪ Peguero-Lo Presti voltage (max S + S _{V4})	0.626	0.384	0.634	0.392	0.623	0.377
▪ Cornell voltage (R _{avL} + S _{V3})	0.617	0.388	0.631	0.413	0.603	0.352
▪ Cornell VDP	0.647	0.419	0.657	0.425	0.644	0.401
▪ Sokolow-Lyon voltage (S _{V1} + max R (V5 or V6))	0.507	0.283	0.514	0.31	0.502	0.264
▪ Gubner-Ungerleider voltage (R _I + S _{III})	0.541	0.317	0.560	0.334	0.511	0.295
<i>ML models</i>						
▪ Logistic regression ^b	0.764	0.647	0.759	0.634	0.771	0.656
▪ Random forest ^b	0.772	0.654	0.766	0.646	0.783	0.679
▪ Light gradient boosted machine ^b	0.790	0.687	0.787	0.677	0.796	0.703
▪ Residual network ^b	0.788	0.684	0.783	0.674	0.796	0.693
▪ Multilayered perceptron network ^b	0.789	0.686	0.784	0.676	0.796	0.702
▪ Convolutional neural network ^c	0.767	0.638	0.773	0.646	0.769	0.644

469 * At specificity 0.75

470 ^a Logistic regressions

471 ^b Input of 117 ECG statistics like QRS duration, heart rate etc. and 60 variational autoencoder latent
 472 variables from ECG representative beat and sex

473 ^c Input of representative-beat ECG signal (X, Y, Z leads)

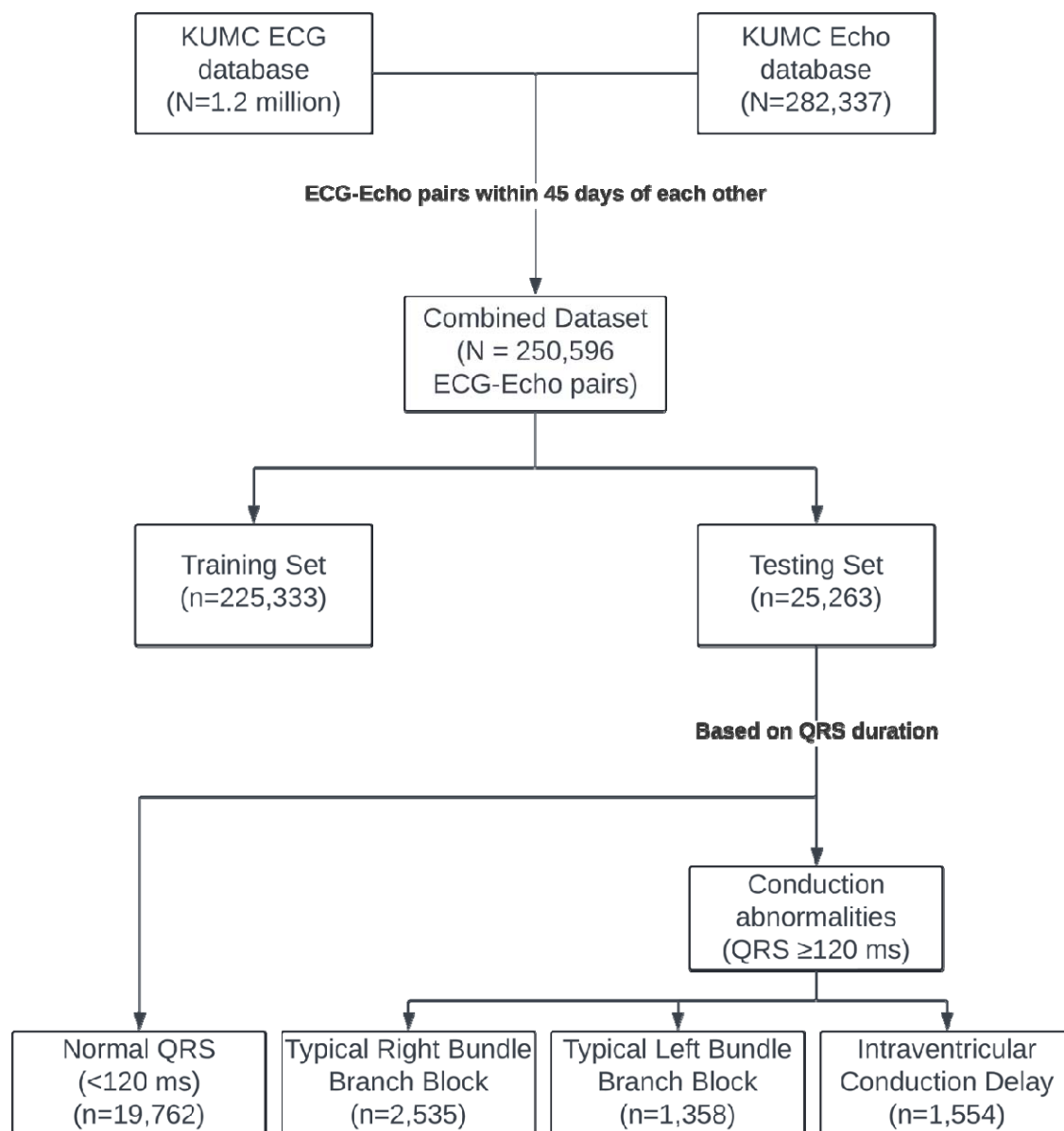
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475 **Table 3.** Comparison between presence of LVH on subsequent echocardiogram (>1 year and
476 closest to 5 years after index echocardiogram) in false positives versus true negatives of LVH
477 LGBM model in testing set

	Follow-up echocardiogram		Total	Risk	Risk Ratio	p
	No LVH	LVH				
True negatives	889	130	1,019	0.13	Ref.	-
False positives	107	54	161	0.34	2.63 (2.01-3.45)	<0.0001
Total	996	184	1,180	-	-	-

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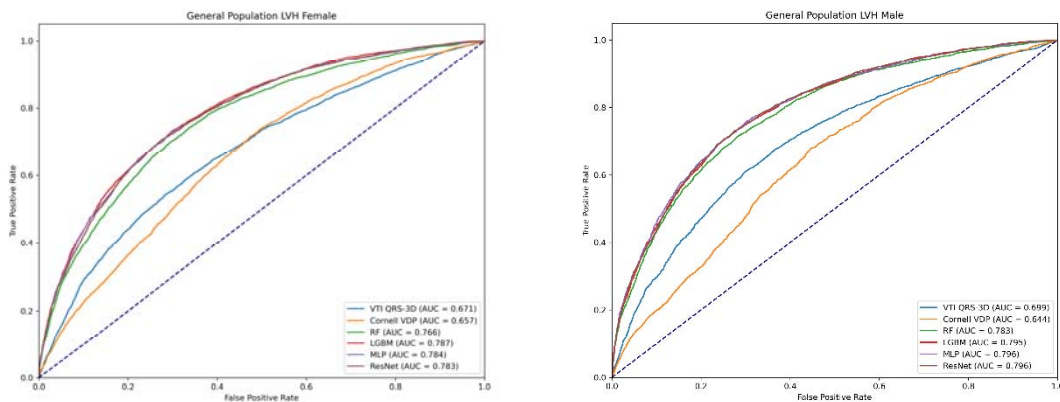
480 **Figure 1.** Data pipeline for model training and testing



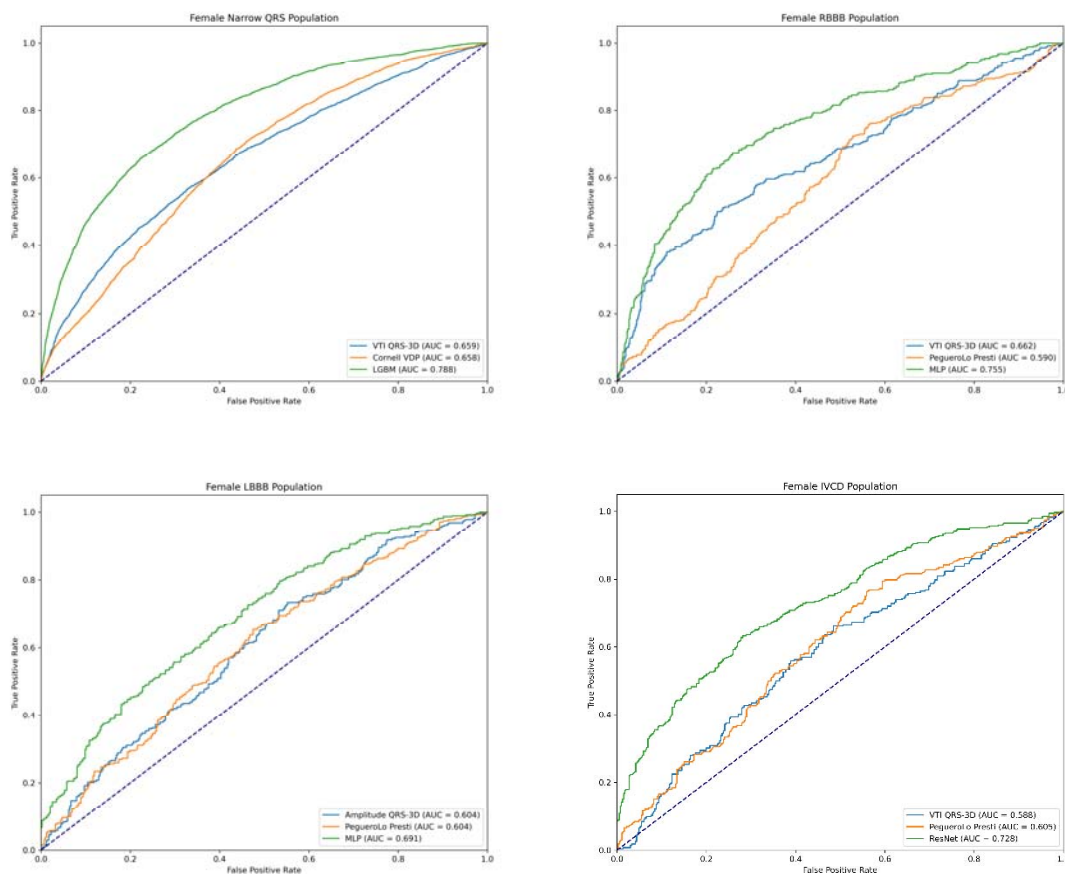
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483 **Figure 2.** ROC curves from the entire testing set for males (*left panel*) and females (*right panel*).

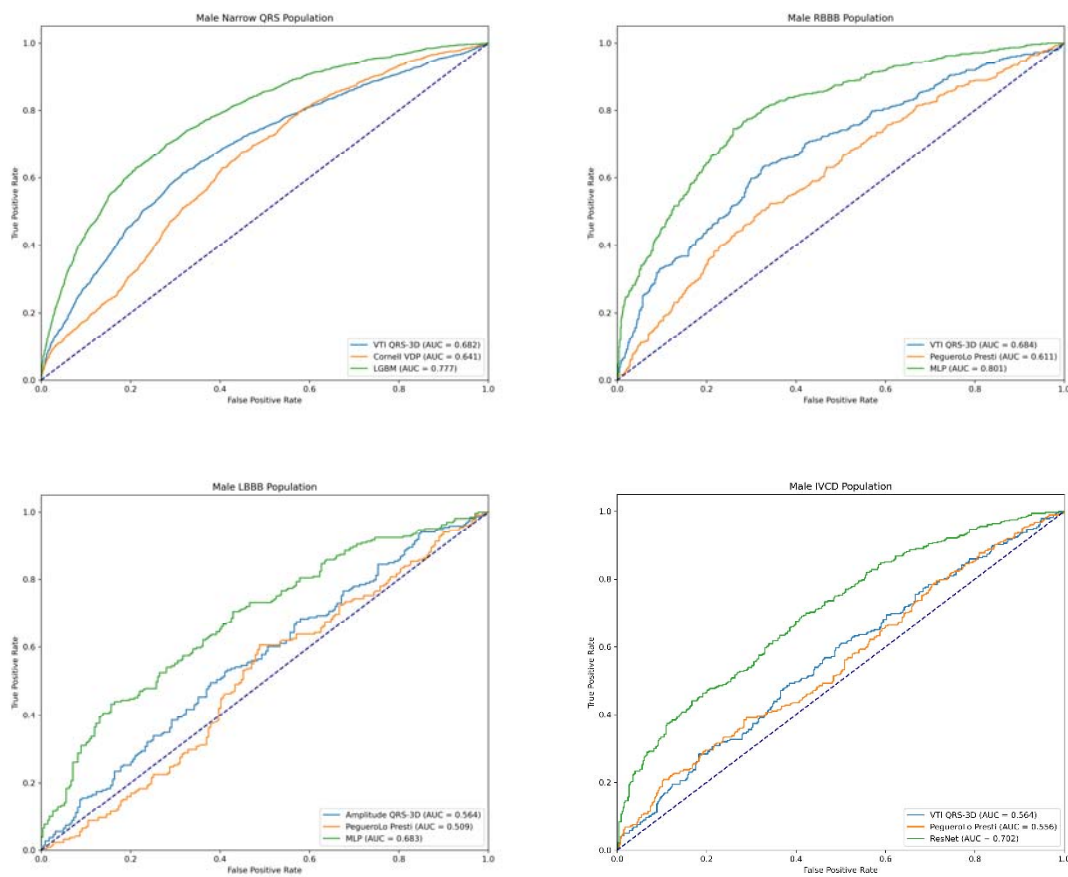
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486 **Figure 3.** ROC curves for subgroups of females in testing set, narrow QRS (*top left*), typical
487 right bundle branch block (RBBB, *top right*), typical left bundle branch block (LBBB, *bottom*
488 *left*), intraventricular conduction delay (IVCD, *bottom right*).
489
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491 **Figure 4.** ROC curves for subgroups of males in testing set, narrow QRS (*top left*), typical right
492 bundle branch block (RBBB, *top right*), typical left bundle branch block (LBBB, *bottom left*),
493 intraventricular conduction delay (IVCD, *bottom right*)
494



495 **Figure 5.** Scatterplots of echocardiographic left ventricular mass indexed (LVMI) plotted against
496 prediction probabilities from the LGBM model for females (*left panel*) and males (*right panel*).
497

