

000 001 002 003 004 005 006 007 008 009 010 011 012 013 014 015 016 017 018 019 020 021 022 023 024 025 026 027 028 029 030 031 032 033 034 035 036 037 038 039 040 041 042 043 044 045 046 047 048 049 050 051 052 053 PHAROS+: NON-INVASIVE LONGITUDINAL MONITORING OF PULMONARY HYPERTENSION WITH UNIMODAL AND MULTIMODAL LEARNING

Anonymous authors

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ABSTRACT

Pulmonary hypertension is a disease characterized by elevated pressures in the blood vessels that supply the lungs. It is a progressive and incurable disease that can lead to right heart failure and premature death if improperly managed. Close monitoring plays an important role in management of patients with pulmonary hypertension, as it facilitates the timely detection of disease progression and enables the prompt administration of therapies that can alter the course of the disease. The gold standard for monitoring disease progression is a right heart catheterization (RHC) – a procedure that involves the insertion of a catheter, attached to a pressure transducer, into the pulmonary vasculature to measure the pulmonary pressures. This procedure is typically repeated several times during the course of a patient’s life to monitor the response to therapies designed to reduce pulmonary pressures. Although RHC is an important tool that can help guide the care of patients with pulmonary hypertension, the procedure itself entails some risk to the patient and can only be performed in hospitals that have the needed equipment and trained personnel. Prior attempts to develop non-invasive alternatives for measuring pulmonary pressures have primarily focused on the task of initial diagnosis, rather than long-term monitoring of patients that have already been diagnosed. In this work, we propose a novel deep learning paradigm for the non-invasive assessment of pulmonary artery pressures. The method leverages electrocardiographic signals and, when available, cardiac ultrasound data to enable long-term monitoring in these patients. We demonstrate that our method achieves strong performance on an internal dataset from one hospital and generalizes well to the MIMIC-III Waveform Database from a different hospital. Our approach provides a cheap and accessible method that can be used to monitor patients with pulmonary hypertension at home. To the best of our knowledge, this work is the first to address the task of longitudinal monitoring in patients with pulmonary hypertension.

1 INTRODUCTION

Pulmonary hypertension is a chronic and progressive disease estimated to affect around 1% of the global population (Hooper et al., 2016). It is characterized by high pressures in the pulmonary vasculature (blood vessels of the lungs) and is formally defined by a mean pulmonary artery pressure (mPAP) greater than 20 mmHg (Humbert et al., 2022). If the disease is not well-controlled by medications, it can lead to right heart failure and premature death. As a result, repeated hemodynamic evaluations (measuring pulmonary artery pressures) are crucial for the diagnosis and long-term monitoring and management. Currently, the gold standard for measuring these pressures is right heart catheterization (RHC), an invasive procedure that involves threading a catheter through a major vein in the body to the heart. This procedure entails risk, must be performed in a hospital setting, and is limited to hospitals that have a catheterization suite and trained personnel.

In recent years, deep learning approaches have shown promise in assessing cardiac hemodynamics from non-invasive data modalities. For instance, Tripathi et al. (2024) used cardiac magnetic resonance imaging and information from electronic health records (EHR) and Suvon et al. (2024) used electrocardiograms (ECG) and chest x-rays to detect elevated mean pulmonary capillary wedge pressures (mPCWP) for heart failure patients. For assessing pulmonary hypertension, Zhao et al.

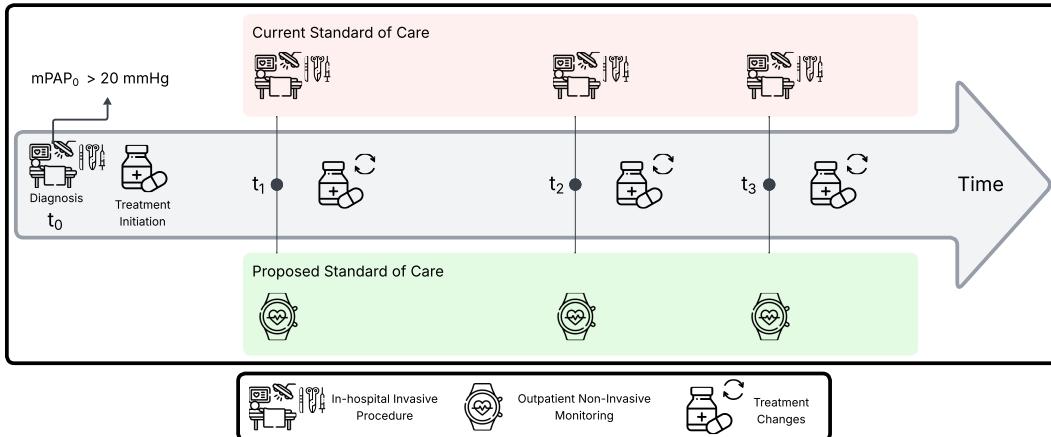


Figure 1: Clinical impact of our proposed work.

(2025) used a combination of tabular and textual data from EHRs along with chest x-rays while Liu et al. (2024b) used ECGs and chest x-rays to diagnose pulmonary hypertension. These methods, however, focus on identifying patients who have pulmonary hypertension (the diagnosis task) rather than longitudinally following patients who are known to have elevated pulmonary pressures.

In patients with pulmonary hypertension, the mPAP rarely normalizes, even with appropriate therapy. The goal in these patients is therefore to reduce the pulmonary pressure as much as possible in an attempt to prevent disease progression. Routine care therefore typically involves repeated RHCs to assess changes in the mPAP as therapy is adjusted (Figure 1). For effective longitudinal monitoring in patients with known pulmonary hypertension, rather than detecting elevated pressures, we propose a method to detect whether the mPAP has increased from its baseline value at any given point in time. As an increase in mPAP reflects disease progression, methods that identify such increases would directly influence therapy without the need for repeated invasive procedures. We envision the impact of our work to be as illustrated in Figure 1.

Our approach leverages ECGs to enable cheap, accessible, and outpatient or at-home monitoring, without the need for in-person hospital visits. In particular, we leverage single lead (Lead I) ECGs for its convenience and widespread availability in wearable and handheld electronic devices. We develop a flexible approach that can use ECGs as a single modality, when only ECGs are available, or use a combination of ECGs and cardiac ultrasound data, when both modalities are available.

The main contributions of this work are:

- We propose an ECG-based method, which we refer to as PHAROS (Pulmonary Hypertension Assessment fROm ECG Signals), and a multi-modal ECG and echocardiography-based method, PHAROS+, for detecting pulmonary hypertension progression. To the best of our knowledge, our work is the first to address the task of longitudinal monitoring in patients with pulmonary hypertension.
- We evaluate the model on two independent datasets and demonstrate that strong performance is maintained despite a dataset shift.

2 RELATED WORK

Data modalities for ML-based non-invasive pulmonary hypertension assessment. Prior studies suggest that echocardiograms (echo) can be used to estimate pulmonary artery pressure (Yock & Popp, 1984), and hence it can be used to follow patients with pulmonary hypertension (Humbert et al., 2022). However, measurements from cardiac ultrasound require a trained sonographer, and can vary depending on their skill level as well as the quality of the echocardiographic images obtained. As a result, estimates of pulmonary artery pressure from echos have large errors (Fisher et al., 2009; D’Alto et al., 2013; Slegg et al., 2021; Janda et al., 2011), making them unreliable

for long-term monitoring and therapeutic decision-making. Some machine learning methods have been introduced as attempts to improve the accuracy of echocardiography-guided estimates of pulmonary artery pressures (Anand et al., 2024; Salehi et al., 2025; Leha et al., 2019), while others have leveraged other data modalities, including cardiac magnetic resonance imaging (Cheng et al., 2025; Swift et al., 2020), computed tomography pulmonary angiography (Xie et al., 2025; Sharkey et al., 2022), and chest x-rays (Huang et al., 2025). Combinations of multiple modalities have also been explored, such as tabular, textual, and x-ray (Zhao et al., 2025) or ECG and x-ray (Liu et al., 2024b). Disadvantages of these methods are that they primarily address the diagnosis task and they require patients to make an in-person visit to a hospital or imaging center to obtain the scans. Other methods have used only readily available data such as electronic health records (Kogan et al., 2023), phonocardiograms (Guo et al., 2025), and ECGs (DuBrock et al., 2024; Raghu et al., 2023a; Aras et al., 2023; McLean et al., 2025; Suvon et al., 2025) to diagnose pulmonary hypertension. The goal of our work is also to use readily available data, primarily ECGs, to enable outpatient or at-home monitoring. Our method is flexible in the sense that it can be used when only an ECG available and when both ECG and echocardiographic measurements are available. Moreover, instead of addressing the diagnosis task as done by prior work, our proposed method aims to identify disease progression.

Non-invasive hemodynamics assessment from ECG. ECGs have been widely used in non-invasive, deep learning-based health assessment, including detecting demographics and health conditions (Abbaspourazad et al., 2024), sleep stages and disorders (Thapa et al., 2024), cardiac arrhythmias (Hannun et al., 2019; Liu et al., 2024a), heart attacks (Acharya et al., 2017), heart failure (Acharya et al., 2019), and hemodynamic abnormalities in patients with heart failure (Schlesinger et al., 2022; 2025; Raghu et al., 2023b) or pulmonary hypertension (Schlesinger et al., 2022; DuBrock et al., 2024; Aras et al., 2023; McLean et al., 2025; Suvon et al., 2025; Sadrawi et al., 2021). The hemodynamics assessment methods all aim to distinguish elevated from non-elevated pressures according to a predefined pressure threshold. The performance of these methods for estimating mPAP vary widely and, more importantly, these methods may not be useful for longitudinal monitoring as patients with pulmonary hypertension rarely have pressures that fall within the normal range. Instead of diagnosis, a few other methods have introduced hemodynamics estimation as regression tasks using physiologic signals as input (Klein et al., 2025; Jeong et al., 2023; Sadrawi et al., 2021). Klein et al. (2025) and Jeong et al. (2023) present methods that regress mPCWP and Sadrawi et al. (2021) regress mPAP. However, regression errors exceed variability in the RHC (Melillo et al., 2020) and clinically significant changes in pressure, making these methods unreliable for long-term monitoring. In contrast to prior work, we use ECGs to classify disease progression in pulmonary hypertension for the purpose of long-term monitoring and management.

3 METHOD

A RHC that definitively measures an elevated mPAP is required to make the diagnosis of pulmonary hypertension. Consequently, virtually all patients who have this diagnosis have had at least one RHC (Humbert et al., 2022). Using a dataset consisting of paired ECG and mPAP measurements from a RHC, our task is therefore to develop a method that uses ECGs to non-invasively determine whether there has been an increase in mPAP over time, thereby circumventing the need for repeated invasive procedures (Figure 1).

3.1 PROBLEM FORMULATION

We formulate the pulmonary hypertension monitoring problem as an optimization problem where the goal is to estimate the probability distribution of disease progression at time t_j relative to previous time t_i . More precisely, we estimate $p(Y_{ij}|\mathbf{s}_i, \mathbf{s}_j, m_i, \Delta t, \mathbf{f}_j) \in [0, 1]$, where $Y_{ij} \in \{0, 1\}$ is a binary label denoting whether the disease has progressed (i.e., the mPAP has increased) between times t_i and t_j . We define disease progression as an increase in mPAP that is larger than a certain threshold. That is, the ground truth label $Y_{ij}^{gt} = 1$ when $\Delta mPAP = m_j - m_i > \text{threshold}$

The time of the initial RHC and ECG is t_i ; $\Delta t = t_j - t_i > 0$, is the time interval between the initial ECG at time t_i , and the current ECG at time t_j ; \mathbf{s}_i is the initial ECG obtained at time t_i ; \mathbf{s}_j is the current ECG obtained at time t_j ; m_i is the known mPAP acquired via RHC at the time t_i ; m_j is the

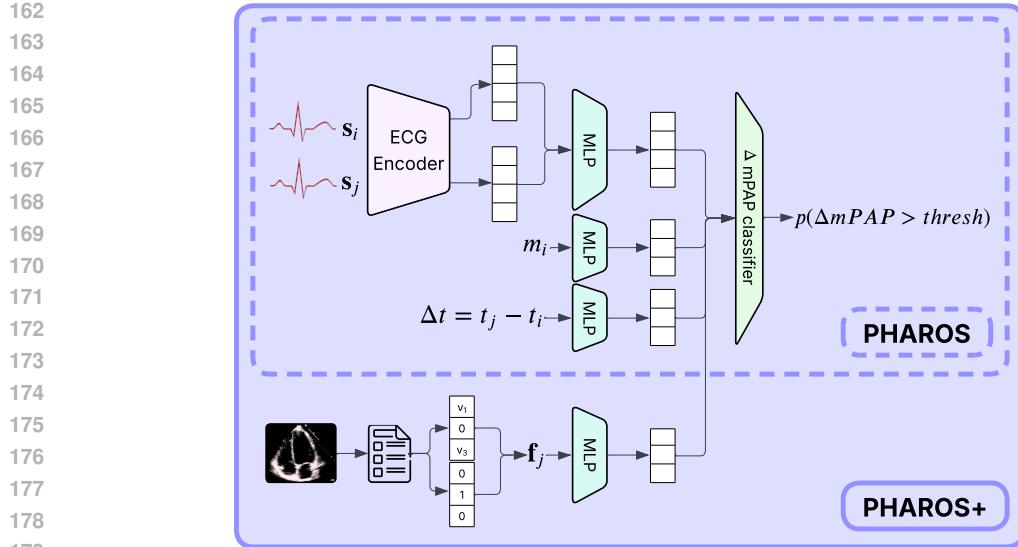


Figure 2: Overview of PHAROS (dashed box) and PHAROS+ (solid box). Merging arrows denote concatenation. Our models take two ECGs, s_i and s_j , corresponding to time points t_i and t_j , respectively, the baseline mPAP m_i corresponding to the first ECG, and the time gap $\Delta t = t_j - t_i > 0$. In PHAROS+, we also input a vector of echo features f_j corresponding to the most recent echo relative to t_j . The embeddings of all the inputs are concatenated and passed to a final MLP to output the final prediction of disease progression.

mPAP at time t_j ; f_j represents additional features of the patient corresponding to time t_j . In this work, f_j represents measurements from the echo most recent to time t_j if available for the patient.

3.2 MODEL ARCHITECTURE

Our model architecture consists of an encoder that individually encodes s_i and s_j , corresponding to two different time points, an encoder for the baseline mPAP m_i , and an encoder for the time gap between the ECGs Δt . We refer to this model as PHAROS. We additionally explore including echo measurements for patients who have echos, train a separate model, and compare performance. The model trained with echo parameters is referred to as PHAROS+. An overview of our architecture is shown in Figure 2.

Encoding ECGs. As input to our model, we use Lead I ECGs due to its convenience and availability on at-home and wearable health monitoring devices. Given two Lead I ECGs s_i and s_j from two different time points t_i and t_j , respectively, we encode each ECG using a ResNet-inspired (He et al., 2016) 1D convolutional neural network (CNN). We use the same encoder weights for both ECGs. The embeddings of s_i and s_j from the ECG encoder are concatenated and fed to a multi-layer perceptron (MLP) to fuse the embeddings.

Encoding the baseline mPAP m_i and time gap Δt . The baseline mPAP m_i and time gap Δt are each input to their own MLPs to obtain embeddings of each value. Prior to training, the mPAP values are normalized to have zero mean and unit variance in the training dataset. The values of Δt are scaled to be in units of months.

Encoding echo measurements as optional features. We train a version of our model, PHAROS+, that optionally takes echo measurements as input when they are available for a patient. Each feature type in the vector is normalized across the dataset to have zero mean and unit variance. Not all patients have echos, and not all measurements are taken for the patients that do have echos. As a result, there can be missing values. To account for this, we vectorize all possible features and fill in missing values with zeros. We concatenate the features with a mask that identifies which elements are missing. The result forms the echo input f_j , which is passed through a MLP to obtain an echo embedding.

216 **Classifying disease progression.** Finally, for determining disease progression, the fused ECG em-
 217 bedding, the embeddings of m_i and Δt , and the echo embedding (for PHAROS+) are concatenated
 218 and fed to a MLP that outputs the estimated probability of disease progression at t_j . For the training
 219 loss, we use the binary cross-entropy loss.
 220

221 4 EXPERIMENTS

223 4.1 DATASETS

225 To train our model, a large source of time-aligned Lead I ECGs and mPAP measurements is required.
 226 Although obtaining ECGs is typically routine procedure in preparation for RHCs, the ECG and RHC
 227 are not done at the exact same time and could even be weeks apart. This data would have noisy
 228 ground truth labels due to inherent variability in mPAP even within the same day (Rich et al., 1985)
 229 or hour (Melillo et al., 2020). Therefore, large quantites of such well-aligned data is difficult to
 230 obtain from the RHC procedures done in a catheterization lab of a hospital. Alternatively, patients
 231 are sometimes admitted to intensive care units (ICU) with a pulmonary artery catheter left inside
 232 the pulmonary artery to continuously monitor the mPAP in real-time. Essentially all ICU patients
 233 also have continuous ECG monitoring. We therefore take advantage of this ICU data to train our
 234 model. In our experiments, we verify that our model performance is maintained at larger time gaps
 235 by evaluating on the subsets of data corresponding to longer ICU stays.

236 **Massachusetts General Hospital (MGH) Dataset.** We train, validate, and test our model on a
 237 large private dataset from MGH. This dataset consists of waveform data recorded from ICU bedside
 238 monitors, including Lead I ECG and pulmonary artery pressure waveforms. There are also echo
 239 measurements for patients that had echos. More details about the echos are described in Appendix
 240 A. We randomly split the dataset by patient into 80%, 10%, and 10% splits for training, validation,
 241 and testing, respectively.

242 **MIMIC-III Waveform Database (Moody et al., 2020; Johnson et al., 2016).** We externally eval-
 243 uate our model on the publicly available MIMIC-III Waveform Database from PhysioNet (Goldberger
 244 et al., 2000). This dataset was collected from ICUs at Beth Israel Deaconess Medical Center. We
 245 use the subset of recordings that contain aligned Lead I ECG and pulmonary artery pressure wave-
 246 forms. There are no echo measurements in this dataset, so when evaluating PHAROS+, we input
 247 fully masked vectors.

248 4.2 DATA PRE-PROCESSING

249 We prepare the data by splitting up aligned continuous Lead I ECG and pulmonary artery pressure
 250 waveforms into 10-second segments, the standard duration of outpatient ECGs (Sattar & Chhabra,
 251 2023). The aligned waveforms are then filtered to ensure they have good signal quality and align-
 252 ment, and poor quality waveforms are discarded. Further details are discussed in Appendix B. The
 253 ECGs in the MGH dataset have a frequency of 120 Hz or 240 Hz and those in the MIMIC-III dataset
 254 have a frequency of 125 Hz. Prior to inputting ECGs to our model, they are upsampled such that
 255 $s_i, s_j \in \mathbb{R}^{4096}$. The statistics of both datasets after pre-processing are shown in Table 1.

256 For validation and testing on the MGH dataset and for external evaluation on MIMIC-III, the pairs
 257 of ECGs, s_i and s_j , are randomly sampled offline and saved to ensure we evaluate on the exact
 258 same pairs each time. During training, we randomly sample different pairs in an online fashion,
 259 allowing pairs to vary at each epoch to make full use of the large training data. We validate every
 260 approximately 5.4 million pairs sampled across all ICU stays and refer to this as one epoch. For
 261 all data splits, ECGs in a sampled pair always come from the same ICU stay and the number of
 262 sampled pairs per ICU stay is proportional to its duration. We compute the mPAPs m_i and m_j from
 263 the aligned pulmonary artery pressure waveform corresponding to s_i and s_j , respectively.

264 4.3 IMPLEMENTATION, TRAINING, AND EVALUATION DETAILS

265 **Implementation.** The ECG encoder is a 1D CNN that begins with a convolutional layer of size (1,
 266 64) followed by a residual block of size (64, 256) and a global pooling layer, leading to embeddings
 267 of s_i and s_j that have a size of 256 each. The two embeddings are concatenated and a two-layer

| | | MGH Dataset | MIMIC-III |
|-----|--|--------------|-------------|
| 272 | # patients | 7894 | N/A |
| 273 | # ICU stays | 8213 | 295 |
| 274 | # 10-sec aligned Lead I ECG and mPAP | 63.3 million | 1.1 million |
| 275 | Average aligned waveform duration per ICU stay | 1.7 days | 0.7 days |
| 276 | # patients with echos | 3885 | N/A |
| 277 | # ICU stays with echos | 4121 | N/A |
| 278 | # echos | 7993 | N/A |
| 279 | Age (years, mean \pm std) | 64 \pm 14 | N/A |
| 280 | Sex (female, %) | 30.3% | N/A |

282 Table 1: Statistics of the parts of the MGH and MIMIC-III Waveform (Moody et al., 2020) datasets
 283 that contain aligned Lead I ECG and pulmonary artery pressure waveforms. Numbers shown corre-
 284 spond to the data after pre-processing.

285
 286 MLP with layer sizes of (512, 256) is used to combine them into an embedding of size 256. The
 287 MLPs for m_i and Δt have the same architecture. Both are two-layer MLPs with layer sizes of (32,
 288 32) that map their respective scalar inputs to embeddings of size 32. The echo inputs consist of 57
 289 measurements and a mask of the same size to indicate missing values. They are concatenated to
 290 give $\mathbf{f}_j \in \mathbb{R}^{114}$. The MLP that embeds \mathbf{f}_j consists of two layers with sizes of (64, 32), yielding an
 291 embedding of size 32. Finally, all the embeddings are concatenated and fed to a three-layer MLP
 292 with layer sizes (256, 256, 1) to obtain the final classification. More details about the architecture
 293 are presented in Appendix C.

294 Measurements of mPAP are generally presumed to have inherent variability, so small increases in
 295 mPAP should not be considered disease progression. In fact, Melillo et al. (2020) have shown that
 296 the standard deviation of differences between two mPAP measurements in a single RHC procedure
 297 is 3.9 mmHg. Considering this, we use a $\Delta mPAP$ threshold of 4 mmHg to define the ground truth
 298 positive class. That is, for every ECG pair corresponding to $\Delta mPAP > 4$ the ground truth label
 299 is $Y_{ij}^{gt} = 1$. This threshold results in a positive class prevalence of 0.1990 in the MGH dataset and
 300 0.1841 in MIMIC-III.

301 **Training.** We train our models using binary cross-entropy loss and the AdamW optimizer
 302 (Loshchilov & Hutter, 2019) with a batch size of 256, learning rate of 1×10^{-6} , and weight
 303 decay of 0.01. Each model is trained for 20 epochs or until the validation loss converged as defined by
 304 early stopping after at least 5 epochs. Models were trained on a single NVIDIA RTX A6000 GPU,
 305 with training time taking around 32 hours. We use the validation AUC to select the best model and
 306 report results on the test split of the MGH dataset and externally evaluate on MIMIC-III.

307 **Evaluation.** We evaluate our models on several common binary classification metrics, includ-
 308 ing Area Under the Receiver Operating Characteristic Curve (AUC), Brier score, Area Under the
 309 Precision-Recall Curve (AUPRC), positive predictive value (PPV, also known as precision), and
 310 negative predictive value (NPV). The probability threshold used to binarize model outputs for the
 311 PPV and NPV was the threshold corresponding to a sensitivity (also known as recall or true positive
 312 rate) of 80%, as commonly done in related work (Schlesinger et al., 2022; Raghu et al., 2023b). To
 313 calculate statistical measures of uncertainty for all metrics, sets of inputs were randomly drawn with
 314 replacement from the test set to produce 10 bootstrapped datasets with equal size to the original test
 315 dataset. The reported errors represent one standard deviation computed using the results across the
 316 10 bootstraps.

317 **Baseline Comparison.** We compare PHAROS and PHAROS+ to the current non-invasive standard
 318 for longitudinal monitoring – pressure estimation using echo data (Humbert et al., 2022). In an
 319 echo, the peak velocity of tricuspid regurgitation (the abnormal backflow of blood from the right
 320 ventricle to the right atrium) can be often be measured in patients with pulmonary hypertension.
 321 This velocity is used to compute the pressure gradient between the right atrium and ventricle using
 322 $g_i = 4v_i^2$, where g_i is the pressure gradient and v_i is the peak tricuspid regurgitation velocity from
 323 the echo at time t_i (Yock & Popp, 1984). From this, the pulmonary artery systolic pressure (PASP),
 which is approximately equal to the right ventricular systolic pressure (RVSP), can be computed as

324 $PASP_i \approx RVSP_i = g_i + RAP$, where RAP is the pressure in the right atrium. RAP cannot be
 325 measured on an echo, so a constant value is commonly assumed and simply added to g to estimate
 326 $RVSP$. Likewise, we assume that RAP is some constant and does not change in time. Therefore,
 327 given two velocities v_i and v_j from t_i and t_j , respectively, we can compute the change in $PASP$
 328 as $\Delta PASP = PASP_j - PASP_i = (g_j + RAP_j) - (g_i + RAP_i) = g_j - g_i$. To convert
 329 $\Delta PASP$ to $\Delta mPAP$ we use the widely accepted equation proposed by Chemla et al. (2004) to
 330 get $\Delta mPAP = 0.61 \times \Delta PASP$. To compute the binary classification metrics described above,
 331 we use $\text{sigmoid}(\Delta mPAP - \text{thresh})$ to convert our computations to probabilities, where
 332 $\text{thresh}=4$ is the $\Delta mPAP$ threshold used to define the positive class.

333

334 5 RESULTS AND DISCUSSION

335

336 5.1 EVALUATION ON THE MGH DATASET

337

338 Our results on the MGH dataset are shown in Table 2. PHAROS achieves a strong performance,
 339 significantly outperforming the echo baseline on all metrics. In PHAROS+, by including echo data
 340 when available, the AUC, Brier score, AUPRC, and PPV improve. This shows that echo measure-
 341 ments can help the model achieve better performance when it is available. The NPV of PHAROS+
 342 decreases slightly compared to PHAROS, but remains significantly higher than the echo baseline.
 343 Overall, our method significantly outperforms the current standard for non-invasive longitudinal
 344 monitoring. We also note that the performance of PHAROS+ when only using ECG data (i.e.,
 345 masking out echo data for all inputs) is similar to the performance of PHAROS which was trained
 346 without echo.

| Model | Eval Inputs | AUC (\uparrow) | Brier (\downarrow) | AUPRC (\uparrow) | PPV (\uparrow) | NPV (\uparrow) |
|---------------|-------------|---------------------------------------|---------------------------------------|---------------------------------------|---------------------------------------|---------------------|
| Echo Baseline | – | 0.5147 ± 0.0789 | 0.2523 ± 0.0256 | 0.1932 ± 0.0676 | 0.1765 ± 0.0390 | 0.8750 ± 0.0720 |
| PHAROS | ECG | 0.7803 ± 0.0010 | 0.1312 ± 0.0003 | 0.4932 ± 0.0023 | 0.3289 ± 0.0010 | 0.9236 ± 0.0006 |
| PHAROS+ | ECG only | 0.7809 ± 0.0009 | 0.1307 ± 0.0002 | 0.4895 ± 0.0021 | 0.3267 ± 0.0010 | 0.9235 ± 0.0005 |
| PHAROS+ | ECG + Echo | 0.7827 ± 0.0009 | 0.1296 ± 0.0002 | 0.4979 ± 0.0019 | 0.3305 ± 0.0009 | 0.9211 ± 0.0004 |

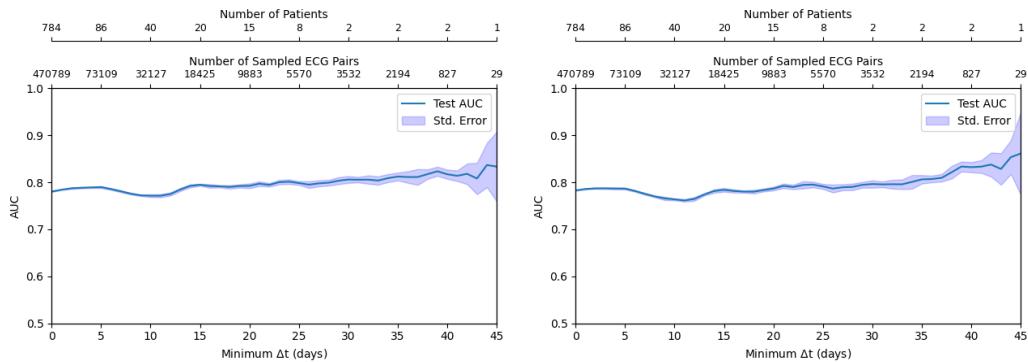
352 Table 2: Performance of PHAROS and PHAROS+ on the MGH dataset. (\uparrow indicates higher is better
 353 and \downarrow indicates lower is better.)

354

355 5.2 EVALUATION ON THE MGH DATASET FOR LARGER TIME GAPS

356

357 We examine the effect of large time gaps on performance. Plots for PHAROS and PHAROS+ are
 358 shown in Figure 3. In general, the models maintain strong performance and slightly improves, albeit
 359 with larger standard error, as the time gap increases up to a duration of 45 days. This is a promising
 360 finding, as it indicates the potential for PHAROS and PHAROS+ to perform well on larger time
 361 gaps beyond the span of typical ICU lengths of stay.



376 Figure 3: AUC for $0 < \min(\Delta t) \leq 45$ on the MGH dataset for PHAROS (left) and PHAROS+
 377 (right).

378 5.3 EXTERNAL EVALUATION ON THE MIMIC-III WAVEFORM DATABASE
379

380 In our previous analysis (Table 2), we found that when given ECGs only, PHAROS+ has a similar
381 performance to PHAROS on our internal dataset, and when echo data is available, PHAROS+ per-
382 forms better. PHAROS+ would therefore be the model to use in practice, so we focus on this model
383 for external evaluation. We use the MIMIC-III Waveform Database (Moody et al., 2020), which was
384 collected from a different hospital, to verify that performance translates to other patient populations.
385 The waveforms in MIMIC-III do not have associated echo measurements, so we can only evaluate
386 PHAROS+ using ECGs and need to fully mask the input vector f_j . Our results are shown in Table 3.
387 We find that despite a performance gap, our model still performs well with an AUC above 0.7 and a
388 slight increase in NPV. These results demonstrate good generalization to unseen data distributions.
389

| Dataset | Eval Inputs | AUC (\uparrow) | Brier (\downarrow) | AUPRC (\uparrow) | PPV (\uparrow) | NPV (\uparrow) |
|-----------|-------------|---------------------|------------------------|----------------------|---------------------|---------------------|
| MGH | ECG only | 0.7809 ± 0.0009 | 0.1307 ± 0.0002 | 0.4895 ± 0.0021 | 0.3267 ± 0.0010 | 0.9235 ± 0.0005 |
| MIMIC-III | ECG only | 0.7052 ± 0.0007 | 0.1649 ± 0.0002 | 0.3145 ± 0.0011 | 0.2375 ± 0.0006 | 0.9550 ± 0.0004 |

393 Table 3: Performance of PHAROS+ on MIMIC-III compared to MGH. (\uparrow indicates higher is better
394 and \downarrow indicates lower is better.)
395

396 6 CONCLUSION
397

400 In this work, we proposed a multi-modal approach that uses ECGs and echos for addressing the novel
401 task of non-invasively identifying pulmonary hypertension progression, defined by an increase in
402 mPAP between two timepoints of $\Delta mPAP > 4$ mmHg. The performance of PHAROS, trained us-
403 ing only ECGs, was improved upon by PHAROS+, which used ECGs and was also able to incorpo-
404 rate echo data when available to boost performance. Our models demonstrated strong performance
405 across various metrics, even for time gaps of up to 45 days. The discriminatory ability of PHAROS
406 and PHAROS+ is significantly above what is obtained used echocardiographic measurements alone.
407 Indeed, the AUC for the estimating pulmonary artery pressures using echocardiography is close to
408 random (0.5), suggesting that this is not reliable method for estimating changes in pulmonary artery
409 pressures. By contrast, our results argue that more reliable estimates of pulmonary artery pressure
410 changes can be obtained with single-lead ECG signals, which do not require a highly trained opera-
411 tor for data acquisition. The inadequacy of echocardiographic information for assessing changes in
412 pulmonary artery pressures is further highlighted by the fact that the PHAROS+, which uses ECG
413 and echo data, has only modest improvement over PHAROS, which only uses ECG data.

414 External validation of PHAROS+ on the MIMIC-III Waveform Database demonstrated good dis-
415 criminatory ability, albeit worse than what was observed in the MGH dataset. Nevertheless, the
416 NPV (at a threshold corresponding to 80% recall) remains strong ($> 95\%$) in this cohort. The high
417 NPVs in both the MGH and MIMIC datasets argue that a negative result from PHAROS/PHAROS+
418 does not suggest that corresponding pulmonary artery pressure is elevated, at a threshold that ensures
419 a true positive rate of 80%.

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614 A ECHO PARAMETERS

615 To train PHAROS+, we incorporated the echo parameters shown in Table 4. This table also shows
 616 the number of times each feature appeared in our dataset. When a parameter had more than one
 617 value recorded within a single echo procedure, the mean of the values was used as input.

620 B FILTERING ECGS AND PULMONARY ARTERY PRESSURE WAVEFORMS 621 FOR QUALITY AND ALIGNMENT

622 Both the MGH and MIMIC-III datasets consisted of raw waveforms obtained from ICU bedside
 623 monitors. These waveforms can be noisy due to external factors such as motion from the patient
 624 or other artifacts. Sometimes patients could be disconnected from the monitor resulting in a flat or
 625 nonsensical signal. As a result, after separating the continuous waveforms into 10-second segments,
 626 we processed the data as follows:

- 627 • We removed all segments where the magnitude of the ECG voltage exceeded 5 mV or the
 628 max - min voltage was less than 0.1 mV.
- 629 • We removed all segments where the mPAP computed from the pulmonary artery pressure
 630 (PAP) waveform exceeded 80 mmHg, the minimum of the waveform was less than -10
 631 mmHg, the mPAP was less than 0, the pulse pressure (max - min) was greater than 80, or
 632 the pulse pressure was less than 2 mmHg.
- 633 • We removed all segments where the number of peaks in the ECG did not match that of the
 634 PAP waveform.
 - 635 1. We performed R-peak detection on the ECG segment using NeuroKit2, specifically
 636 the `ecg_clean` and `ecg_peaks` functions with the default keyword arguments.
 - 637 2. For the PAP waveform, we used the function `scipy.signal.find_peaks()`
 638 with keyword arguments `prominence=10` and `distance=sampling_rate / 4`, where `sampling_rate` is the signal frequency in each dataset and 4 is an estimate
 639 of the maximum possible beats per second obtained from 240 beats per minute divided
 640 by 60 seconds per minute.
 - 641 3. If the difference in number of peaks between the ECG and PAP waves exceeded 1, we
 642 discarded the segment.
- 643 • We removed all segments where the mPAP range was too large.
 - 644 1. We found the R-peaks in the ECG as previously described.

648

649

650

| Measurement | Count |
|---|-------|
| AORTIC SINUS INDEX POST 1 | 11946 |
| RIGHT ATRIUM INDEX MEDIAL-LATERAL (1) | 8795 |
| BODY SURFACE AREA (1) | 7528 |
| EJECTION FRACTION (1) | 7499 |
| LEFT VENTRICLE INTERNAL DIAMETER END DIASTOLE (1) | 7447 |
| LEFT VENTRICLE INTERNAL DIAMETER END SYSTOLE (1) | 7390 |
| LEFT VENTRICLE EJECTION FRACTION - QUINES RAW (1) | 7389 |
| INTERVENTRICULAR SEPTUM THICKNESS (1) MM | 7260 |
| LEFT VENTRICLE POSTERIOR WALL THICKNESS (1) MM | 7259 |
| LEFT ATRIUM DIMENSION ANTERIOR-POSTERIOR (1) | 6640 |
| LEFT VENTRICLE APICAL CONTRIBUTION (1) | 5949 |
| WEIGHT | 5929 |
| HEIGHT | 5914 |
| PHS CV ECHO AV SINUS INDEX 1 | 5806 |
| ASCENDING AORTA DIAMETER (1) MM | 5173 |
| RIGHT ATRIUM PRESSURE ESTIMATED (1) | 5096 |
| RIGHT VENTRICLE TO RIGHT ATRIUM PRESSURE GRADIENT (1) | 4995 |
| TRICUSPID VALVE PEAK VELOCITY (1) | 4995 |
| RIGHT VENTRICLE PEAK SYSTOLIC PRESSURE (1) | 4966 |
| ASCENDING AORTA INDEX 1 | 4902 |
| ASCENDING AORTA INDEX POST 1 | 4902 |
| LEFT ATRIUM DIMENSION SUPERIOR-INFERIOR (1) MM | 4567 |
| LEFT ATRIUM DIMENSION MEDIAL-LATERAL (1) MM | 4537 |
| RIGHT ATRIUM DIMENSION SUPERIOR-INFERIOR (1) | 4533 |
| RIGHT ATRIUM INDEX SUPERIOR-INFERIOR POST | 4340 |
| RIGHT ATRIUM INDEX SUPERIOR-INFERIOR (1) | 4340 |
| RIGHT ATRIUM INDEX MEDIAL-LATERAL POST | 4301 |
| RIGHT VENTRICLE LINEAR DIMENSION (1) MM | 3451 |
| LEFT ATRIAL VOLUME (1) | 2652 |
| LEFT ATRIAL VOLUME INDEX (1) | 2632 |
| INFERIOR VENA CAVA DIAMETER (1) MM | 1877 |
| LEFT VENTRICULAR OUTFLOW TRACT VELOCITY (1) | 1300 |
| RIGHT VENTRICLE PULSE DOPPLER S WAVE (1) | 1284 |
| RIGHT VENTRICLE TAPSE (1) | 1251 |
| AORTIC VALVE PEAK GRADIENT (1) | 810 |
| AORTIC VALVE MEAN GRADIENT (1) | 780 |
| MITRAL VALVE PEAK GRADIENT (1) | 737 |
| MITRAL VALVE MEAN GRADIENT (1) | 734 |
| AORTIC VALVE PROSTHETIC PEAK GRADIENT (1) | 640 |
| AORTIC VALVE PROSTHETIC MEAN GRADIENT (1) | 635 |
| MITRAL VALVE GRADIENT HR (1) | 614 |
| LEFT VENTRICULAR OUTFLOW TRACT DIAMETER (1) | 359 |
| AORTIC VALVE AREA (1) | 320 |
| AORTIC VALVE AREA INDEX (1) | 298 |
| RIGHT VENTRICLE BASAL DIAMETER (1) | 281 |
| MITRAL VALVE PROSTHETIC MEAN GRADIENT (1) | 280 |
| MITRAL VALVE PROSTHETIC PEAK GRADIENT (1) | 280 |
| RIGHT VENTRICLE FRACTIONAL AREA CHANGE (FAC) (1) | 271 |
| PERICARDIUM EFFUSION DIMENSION 1 (1) MM | 261 |
| AORTIC VALVE DISTANCE TO CANNULA TIP MM | 249 |
| TRICUSPID VALVE PROSTHETIC MEAN GRADIENT (1) | 207 |
| TRICUSPID VALVE PROSTHETIC PEAK GRADIENT (1) | 207 |
| LEFT VENTRICULAR OUTFLOW TRACT TIME VELOCITY INTEGRAL (1) | 156 |
| PULMONARY ARTERY MAIN - DIMENSION (1) | 137 |
| PERICARDIUM EFFUSION DIMENSION 2 (1) MM | 126 |
| PULMONARY ARTERY RIGHT - DIMENSION (1) | 121 |
| PULMONARY ARTERY LEFT - DIMENSION (1) | 108 |

699

700

701

Table 4: Echo measurements used in PHAROS+ and the number of measurements for each feature across the full dataset.

702 2. Using the indices of a consecutive pair of R-peaks, we found the corresponding sec-
 703 tions in the PAP waveform. Each section corresponds to the PAP wave for one heart
 704 beat.
 705 3. Using the PAP waveform sections, we computed the mPAP for each heart beat.
 706 4. If any of the mPAPs were outside the range of $\pm 3\sigma_{mPAP}$, where σ_{mPAP} is the stan-
 707 dard deviation, we discarded the segment.
 708 • We removed all segments where the systolic PAP, diastolic PAP, and mPAP were inconsis-
 709 tent.
 710 1. We found the R-peaks in the ECG as previously described.
 711 2. Using the indices of a consecutive pair of R-peaks, we found the corresponding sec-
 712 tions in the PAP waveform. Each section corresponds to the PAP wave for one heart
 713 beat.
 714 3. Using the PAP waveform sections, we computed the mPAP and systolic and diastolic
 715 PAPs for each heart beat.
 716 4. If any heart beat had a corresponding systolic PAP less than or equal to the mPAP or
 717 diastolic PAP greater than or equal to the mPAP, the segment was discarded.
 718

719 **C MODEL ARCHITECTURE**

720 The detailed architecture of PHAROS+ is shown below. PHAROS follows the same architecture, but
 721 without the MLP encoding the echo parameters, and with a MLP decoder that takes in 320 features
 722 instead of 352.

```

725 1 PHAROSPlus (
726 2     (encoder): ECGPairPAPDeltaEncoder (
727 3         (ecg_encoder): CNN1D (
728 4             (conv1): Conv1d(1, 64, kernel_size=(17,), stride=(1,), padding=same
729 5                 , bias=False)
730 6             (bn1): BatchNorm1d(64, eps=1e-05, momentum=0.1, affine=True,
731 7                 track_running_stats=True)
732 8             (res_blocks): ModuleList (
733 9                 (0): ResidualBlock (
734 10                     (conv1): Conv1d(64, 256, kernel_size=(17,), stride=(1,),
735 11                         padding=same, bias=False)
736 12                     (bn1): BatchNorm1d(256, eps=1e-05, momentum=0.1, affine=True,
737 13                         track_running_stats=True)
738 14                     (conv2): Conv1d(256, 256, kernel_size=(17,), stride=(12,), bias
739 15                         =False)
740 16                     (bn2): BatchNorm1d(256, eps=1e-05, momentum=0.1, affine=True,
741 17                         track_running_stats=True)
742 18                     (pool): MaxPool1d(kernel_size=12, stride=12, padding=0,
743 19                         dilation=1, ceil_mode=False)
744 20                     (identity_layer): Conv1d(64, 256, kernel_size=(1,), stride=(1,),
745 21                         , bias=False)
746 22             )
747 23         )
748 24         (global_avg_pool): AdaptiveAvgPool1d(output_size=1)
749 25     )
750 26     (ecg_fuser): Sequential (
751 27         (0): Linear(in_features=512, out_features=512, bias=True)
752 28         (1): ReLU()
753 29         (2): Dropout(p=0.4, inplace=False)
754 30         (3): Linear(in_features=512, out_features=256, bias=True)
755 31     )
756     (hemodynamics_encoder): Sequential (
757 32         (0): Linear(in_features=1, out_features=32, bias=True)
758 33         (1): ReLU()
759 34         (2): Dropout(p=0.4, inplace=False)
760 35         (3): Linear(in_features=32, out_features=32, bias=True)
761     )
762     (time_encoder): Sequential (

```

```

756     31     (0): Linear(in_features=1, out_features=32, bias=True)
757     32     (1): ReLU()
758     33     (2): Dropout(p=0.4, inplace=False)
759     34     (3): Linear(in_features=32, out_features=32, bias=True)
760     35   )
761     36   (echo_parameters_encoder): Sequential(
762     37     (0): Linear(in_features=114, out_features=64, bias=True)
763     38     (1): ReLU()
764     39     (2): Dropout(p=0.4, inplace=False)
765     40     (3): Linear(in_features=64, out_features=32, bias=True)
766     41   )
767     42   (decoder): MLP(
768     43     (layers): Sequential(
769     44       (0): Linear(in_features=352, out_features=256, bias=True)
770     45       (1): ReLU()
771     46       (2): Dropout(p=0.4, inplace=False)
772     47       (3): Linear(in_features=256, out_features=256, bias=True)
773     48       (4): ReLU()
774     49       (5): Dropout(p=0.4, inplace=False)
775     50       (6): Linear(in_features=256, out_features=1, bias=True)
776     51     )
777     52   )
778     53 )
779

```

D MODEL PERFORMANCE AT DIFFERENT $\Delta mPAP$ THRESHOLDS

In this work, we define disease progression as an increase in mPAP greater than 4 mmHg and trained our models with this threshold. In practice, however, other thresholds can be informative as well. For instance, in some cases, an increase in mPAP can be expected, such as when withdrawing medication (e.g., due to adverse events). A model trained with a larger threshold could be useful for ruling out increases in mPAP beyond a certain value. We studied the impact of the $\Delta mPAP$ classification threshold on performance by training different PHAROS+ models with thresholds ranging from 0 to 7 mmHg. The results are plotted in Figure 4. As the threshold increases, AUC increases. This is likely because larger changes in mPAP are more distinguishable in the ECGs, leading to better performance.

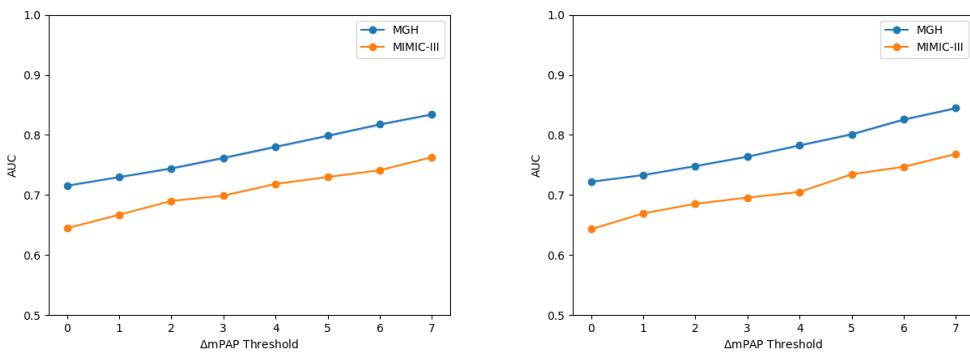


Figure 4: AUC of PHAROS (left) and PHAROS+ (right) when trained at different $\Delta mPAP$ thresholds. Error bars are present but are too small to see.

E ABLATION STUDY OF Δt AND m_i

An ablation study of Δt and the baseline mPAP m_i was conducted. The results are presented in Table 5. We see that both Δt and m_i improved AUC, with m_i contributing much more to the results

than Δt . This makes sense because the dataset we used to train our models consisted of ICU stays with short time scales, an average duration of 1.7 days, as shown in Table 1. Within such close time intervals, the baseline m_i would be a stronger predictor of changes in mPAP than Δt .

| Δt | m_i | AUC (\uparrow) | Brier (\downarrow) | AUPRC (\uparrow) | PPV (\uparrow) | NPV (\uparrow) |
|--------------|--------------|---------------------|------------------------|----------------------|---------------------|---------------------|
| \times | \times | 0.6081 ± 0.0007 | 0.1567 ± 0.0003 | 0.2704 ± 0.0010 | 0.2281 ± 0.0005 | 0.8671 ± 0.0008 |
| \checkmark | \times | 0.6101 ± 0.0009 | 0.1565 ± 0.0004 | 0.2720 ± 0.0015 | 0.2297 ± 0.0009 | 0.8681 ± 0.0006 |
| \times | \checkmark | 0.7812 ± 0.0004 | 0.1299 ± 0.0003 | 0.5025 ± 0.0014 | 0.3315 ± 0.0012 | 0.9219 ± 0.0005 |
| \checkmark | \checkmark | 0.7827 ± 0.0009 | 0.1296 ± 0.0002 | 0.4979 ± 0.0019 | 0.3305 ± 0.0009 | 0.9211 ± 0.0004 |

Table 5: Ablation study of the Δt and baseline mPAP (m_i) model inputs in PHAROS+. The results shown are for the MGH dataset. (\uparrow indicates higher is better and \downarrow indicates lower is better.)