



# A comparison of artificial intelligence-based algorithms for the identification of patients with depressed right ventricular function from 2-dimensional echocardiography parameters and clinical features

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**Background:** Recognizing low right ventricular (RV) function from 2-dimensional echocardiography (2D-ECHO) is challenging when parameters are contradictory. We aim to develop a model to predict low RV function integrating the various 2D-ECHO parameters in reference to cardiac magnetic resonance (CMR)—the gold standard.

**Methods:** We retrospectively identified patients who underwent a 2D-ECHO and a CMR within 3 months of each other at our institution (American University of Beirut Medical Center). We extracted three parameters (TAPSE, S' and FAC<sub>RV</sub>) that are classically used to assess RV function. We have assessed the ability of 2D-ECHO derived parameters and clinical features to predict RV function measured by the gold standard CMR. We compared outcomes from four machine learning algorithms, widely used in the biomedical community to solve classification problems.

**Results:** One hundred fifty-five patients were identified and included in our study. Average age was 43±17.1 years old and 52/156 (33.3%) were females. According to CMR, 21 patients were identified to have RV dysfunction, with an RVEF of 34.7%±6.4%, as opposed to 54.7%±6.7% in the normal RV population (P<0.0001). The Random Forest model was able to detect low RV function with an AUC =0.80, while general linear regression performed poorly in our population with an AUC of 0.62.

**Conclusions:** In this study, we trained and validated an ML-based algorithm that could detect low RV function from clinical and 2D-ECHO parameters. The algorithm has two advantages: first, it performed better than general linear regression, and second, it integrated the various 2D-ECHO parameters.

**Keywords:** RV function; 2D-ECHO; CMR; machine learning

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## Introduction

The cardiology community has historically neglected the right ventricle's (RV) role in the development and progression of heart failure (HF). However, with the recognition of RV function as an important prognostic factor in several diseases such as HF and pulmonary hypertension (PH) along with the need of accurate RV measurements for ventricular assist devices (VADs), assessment of the RV has become a cornerstone in cardiac evaluation (1-6). Unfortunately, the irregular shape of the RV and lack of normative data challenged the cardiologists when trying to integrate the assessment of the RV as part of the standard Echo evaluation and in clinical decision-making (7).

Currently, assessing RV function by 2-dimensional echocardiography (2D-ECHO) represents the first choice for clinicians (7). However, despite years of experience in measuring RV function, cardiologists still find this task difficult and frequently the RV function is misclassified. In one study assessing concordance levels between 2D-ECHO and CMR imaging there was fair agreement between the two modalities with the prevalence-adjusted bias-adjusted Kappa being 0.43 (range, 0.36–0.45), and poor inter-echocardiographer agreement with a Kappa of 0.12 (7). Simply, we cannot extrapolate data and long-term experience gained from conventional left ventricular (LV) evaluation to the RV. This could be attributed to the complex three-dimensional RV shape, poor definition of RV endocardial surface (LV is much thicker than RV), limited direct visualization due to the retrosternal position of RV, as well as uneven complex contraction-relaxation forces along the RV segment (8-11).

RV EF calculation from 2D-ECHO requires several geometric assumptions, and this has proven to be an inaccurate way for EF estimation. Another way is to derive RV systolic function from other measurements such as the RV fractional area-change ( $FAC_{RV}$ ) and lateral TA peak systolic excursion (TAPSE), which provide similar information to RVEF (8,12,13). Other quantitative values also include RV strain, peak S wave velocity of the lateral tricuspid annulus by tissue Doppler imaging (RV S') (9). The abnormality thresholds for S', TAPSE, RV Strain, and  $FAC_{RV}$  found in the literature are  $<9.5$  cm/s,  $<17$  mm,  $<-25\%$ , and  $<48.5\%$ , respectively (10,11,14,15). With TAPSE being the most sensitive (93%) and specific (100%) of all 4.

While 3D-ECHO is supposed to give a better RVEF

estimation in comparison to CMR-derived RV EF and negate the geometrical assumptions made with 2D-ECHO RVEF estimation, it has its own limitations such as being technically challenging, not being widely available yet, and the difficulty of tracing RV border (16,17). When the RV is dilated, such as in corrected Tetralogy of Fallot patients, identifying specific landmarks such as the true RV apex and the pulmonary valve becomes very difficult (18).

Machine learning develops a prediction model using data, algorithms and computing power. The use of logistic regression models to predict complex biological relationships may be limited in some contexts, partly because many biological relationships are non-linear. In the realm of ML and imaging, the use of artificial intelligence approaches (19,20), has enabled automated detection of LV and left atrial endocardial boundaries throughout the cardiac cycle from 3DE data sets, allowing accurate measurements of LV and left atrial volumes and EF (21-25).

The aim of our study is to develop models that can predict low RV function by integrating several 2D-ECHO derived parameters in reference to cardiac magnetic resonance, and then compare these models to single-parameter assessment of RV function. We present the following article in accordance with the TRIPOD checklist. Available at <http://dx.doi.org/10.21037/cdt-20-471>.

## Methods

The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by the Institutional Review Board of the American University of Beirut (Registration Number: IRB00003225). Patient consent was waived due to the retrospective nature of the research. Patient characteristics were extracted by one investigator (AA) while ECHO parameters were extracted and validated by 2 different investigators (ZI and LM). CMR parameters were extracted by an experienced operator (HI), blinded to the clinical and ECHO parameters.

### *Selection of subjects*

We retrospectively identified consecutive patients older than 18 years of age who underwent a 2D-ECHO and a CMR within 3 months of each other at our institution (American University of Beirut Medical Center, Beirut, Lebanon) from 2012 to 2013. Indications for the standard Two-Dimensional Echocardiography (2D-ECHO) were various and no exclusion criteria were applied. From the ECHO,

**Table 1** Data used for predictive modelling

Continuous variables	Binary variables
“Age”	“Dyslipidemia Yes/No”
“RV FAC”	“Coronary artery disease Yes/No”
“TAPSE”	“Smoking Yes/No”
“S”	“Diabetes type2 Yes/No”
	“Hypertension Yes/No”
	“Gender Male/Female”

FAC, fractional area change; TAPSE, tricuspid annular plane systolic excursion; S, S-wave velocity.

LV EF, TAPSE, S' and FAC<sub>RV</sub> were extracted from four standard views [Apical Four Chamber (A4C), Parasternal Long axis (PLAX), Parasternal Short axis: Aortic Valve level (PSAX), and RV inflow view]. These views are done as per the standard 2D-ECHO protocol in our institution. We then used the CMR determined RV function as our reference standard. In our population, the CMR was mostly ordered to assess the LV (function, scar burden and/or viability) (25%), to rule out myocarditis (13%), or to rule out Arrhythmogenic Right Ventricular Disease (ARVD) (13%). Other indications included cardiac sarcoidosis evaluation, ruling out hypertrophic cardiomyopathy and others. As per recent data from the UK Biobank population cohort, we classified CMR RVEF values below 45% as abnormal (26). Furthermore, patient related characterizes (age, sex, cardiovascular risk factors) at the time of the studies were also extracted from their records.

### Statistical analyses

All statistical analyses and data curation were performed using the R programming language. The data was cleaned by excluding patients with missing values (especially outcome values). Cohort-specific categorical and continuous data as well as corresponding inferential and descriptive statistics are shown in *Table 1*. Continuous variables distributed normally were expressed as the mean  $\pm$  standard deviation, categorical variables were expressed as frequency (percentage). Enrolled patients were divided into two groups—those with RV Dysfunction (RVEF <45%) and those without RV dysfunction (RVEF >45%). For between-group comparisons, unpaired *t*-test was used for normally distributed continuous variables, Mann-Whitney U test for non-normally distributed variables, and  $\chi^2$  test (and Fisher's

exact test) for categorical variables. For all tests, a P value <0.05 was considered statistically significant. Accuracy refers to  $[\text{true pos}/\text{pos} + \text{true neg}/\text{neg}]/2$ .

### Machine learning pipeline

After splitting the data into training and validation sets, we randomly replicate samples from the minority class (abnormal RV) in the training set to deal with data imbalance (over sampling technique) in the subsequent cross-validation step. To test the ability of 2D-ECHO and laboratory records to predict RV dysfunction, we implemented four machine learning methods with varying complexity using the CARET package in R: a generalized linear model (GLM) (logistic regression), a support vector machine (SVM) using a radial basis function (RBF), a Random Forest (RF) which employs an ensemble of decision trees to construct a predictive model and Elastic net regularization, a regression method that combines the penalties of lasso and ridge regressions (Rpart).

For all ML models, seed values were fixed, tuning and hyper-parameters were set to default as implemented in CARET. We randomly partitioned the data into the ratio of 60:20:20 for training, validation and testing respectively. Then, we opted for the repeated train/validation split or leave group out cross-validation strategy and the ratio of 3:1 to split the training set into train/validation splits; this step is repeated 100 times and is coupled, within the cross-validation, with oversampling for class imbalance in CARET (since out data is skewed towards normal RV).

Each of the ML methods has a built-in feature selection scheme and ranks variables by importance. The top four most important variables were retained with regard to each of the algorithms. In addition, the Boruta algorithm implemented in R was also independently used for feature selection (27). It is built around the random forest classification algorithm and tries to capture all the significant features in a dataset. Subsequently, several ML models are built with either the whole set of variables or selected features from Boruta. A decision tree was built using the best algorithm with the version with the lowest out-of-bag (OOB) error.

Finally, the model with the optimal receiver operating characteristic (ROC) curve was selected to further investigate its corresponding performance on the test set (20% of the population). Furthermore, and to confirm that RF outperformed other methods in terms of AUROC, we used a pairwise t-test with Bonferroni correction to compare

Table 2 Baseline characteristics

	RV function		P value
	RV EF $\geq$ 45%, N=134	RV EF <45%, N=21	
Clinical parameters			
Age (years)	42.2 $\pm$ 16.9	49.7 $\pm$ 17.4	0.07
Females, n (%)	46 (34.3)	6 (28.5)	0.6
Hypertension, n (%)	43 (32.0)	11 (32.4)	0.07
Dyslipidemia, n (%)	41 (30.6)	6 (28.6)	0.85
Type 2 diabetes, n (%)	17 (12.7)	7 (33.3)	0.015
Smoking exposure, n (%)	56 (41.8)	7 (33.3)	0.45
Coronary artery disease, n (%)	25 (18.7)	7 (33.3)	0.12
Imaging parameters			
RV EF (%)	54.7 $\pm$ 6.7	34.7 $\pm$ 6.4	<0.0001
LV EF (%)	52.7 $\pm$ 11.8	34.2 $\pm$ 13.5	<0.0001
FAC (%)	43.6	34.8	0.0002
S' (cm/s)	12.9	11.3	0.01
TAPSE (mm)	21	17.8	0.04

RV, right ventricle; LV, left ventricle; EF, ejection fraction; FAC, fractional area change; TAPSE, tricuspid annular plane systolic excursion; S, S-wave velocity.

the mean of AUROC values, across the same resampling results, from the leave group out cross-validation folds extracted from all ML methods. To compare resampling results across ML methods, we matched the same indices in the R object `trainObject$control$index`.

## Results

One-hundred ninety-six [196] patients were eligible to be included in our study. Twenty-five were excluded because of missing outcomes data and 16 with missing ECHO data. One-hundred fifty-five [155] patients were included in our analysis. Average age was 43 $\pm$ 17.1 years and 52/155 (33.5%) were females. Baseline characteristics are summarized in Table 2, patients with abnormal RV function had higher diabetes (33.3% vs. 12.7%; P=0.02) and lower LV EF (34.2 vs. 52.7; P<0.0001). According to CMR, 21 patients were identified to have RV dysfunction, with an RVEF of 34.7% $\pm$ 6.4%, as opposed to 54.7% $\pm$ 6.7% in the normal RV population (P<0.0001).

All four ML algorithms optimized the ROC and generated best models, from either all variables (Table 1) or selected features from Boruta algorithm (Table 3), which

were evaluated on the test set (20% of the population). Representative confusion matrices for all variables and selected top variables are shown in Tables 4,5, respectively. Clearly, the Random Forest (RF) model (OOB error of 3.24%, 500 trees, and a mtry of 2) yielded the highest accuracy and the best balance between sensitivity and specificity, especially when using Boruta features: age, FAC<sub>RV</sub>, hypertension, TAPSE and S'. Its AUC reached 0.82 when trained with these features, higher than other ML models and general linear regression analysis (AUC =0.71). When comparing ROC curves to confirm that RF outperformed all other methods (Table 6), there was no significant difference in area under ROC between RF and ElasticNet when the same cross-validation folds were compared (P=1). However, ElasticNet seems to overfit; thus, did not generalize well on the test dataset. However, and as predicted, the accuracy drops when we generated the MCC for our models. However, when comparing MCCs; RF (MCC<sub>RF</sub> =0.52) was significantly a better choice for classification in our dataset when compared to other algorithms (MCC<sub>SVM</sub> =0.3) and general linear regression (MCC<sub>GLM</sub> =0.31) (P=0.002).

Interestingly, the five important variables, flagged with

**Table 3** Top five important variables from built-in feature selection with different models

Method	1	2	3	4	5
SVM	Age	FAC	TAPSE	S'	HTN
GLM	Age	FAC	DL	DM2	HTN
RF	Age	FAC	TAPSE	S'	HTN
ELASTICNET	DL	FAC	Smoker	DM2	HTN

A Seed of [123] was fixed for all methods with set. Seed function in R. SVM, support vector machine; GLM, generalized linear method; RF, random forest; HTN, hypertension; DM2, diabetes mellitus type 2; DL, dyslipidemia; FAC, fractional area change, TAPSE, tricuspid annular plane systolic excursion, S, S-wave velocity.

**Table 4** Confusion matrices for the test set

Method	Sens (%)	Spec (%)	Accuracy (%)	AUC (ROC)	MCC
RF	25	100	90	0.80	0.47 (P=0.001)
ELASTICNET	50	81	77	0.69	0.25 (P=0.024)
Rpart	50	77	73	0.65	0.21 (P=0.063)
SVM	25	92	83	0.69	0.2 (P=0.036)
GLM	50	73	70	0.62	0.17 (P=0.061)

All attributes were used to train the ML models. SVM, support vector machine; GLM, generalized linear method; RF, random forest.

**Table 5** Confusion matrices for the test set using important attributes (age, S, TAPSE, FAC, HTN), from the Boruta Feature selection algorithm, were used to train the ML models

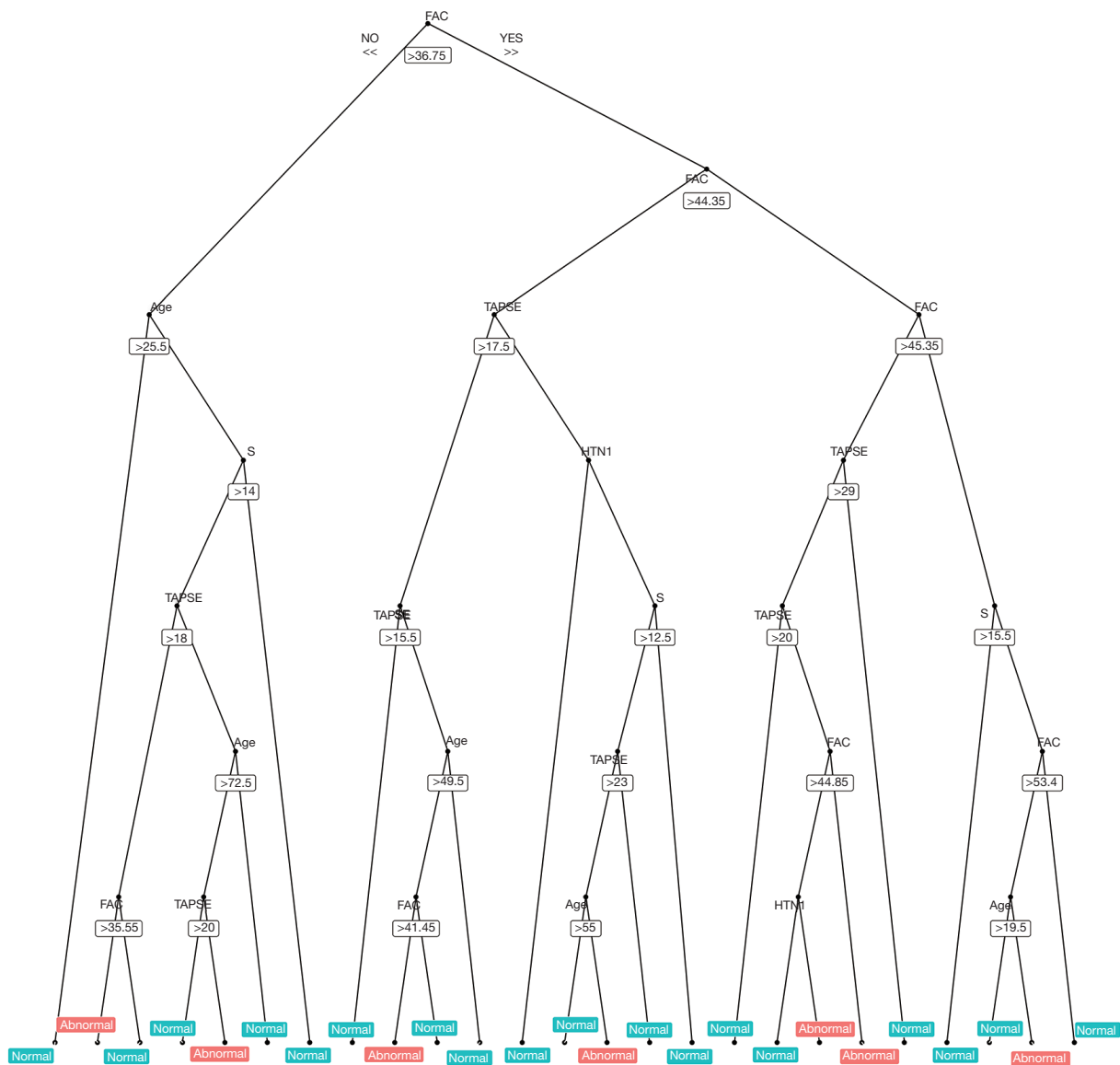
Method	Sens (%)	Spec (%)	Accuracy (%)	AUC (ROC)	MCC
RF	50	96	90	0.86	0.52 (P=0.002)
ELASTICNET	75	81	80	0.75	0.43 (P=0.002)
GLM	75	70	70	0.71	0.31 (P=0.012)
SVM	50	85	80	0.82	0.3 (P=0.012)
Rpart	50	77	73	0.65	0.21 (P=0.063)

SVM, support vector machine; GLM, generalized linear method; RF, random forest; FAC, fractional area change; TAPSE, tricuspid annular plane systolic excursion; S, S-wave velocity; HTN, hypertension.

**Table 6** Comparison of area under ROC curves among ML methods using the same folds from the leave group out cross-validation

	SVM	ELASTICNET	GLM	RF	Rpart
SVM	–	–0.013778	0.022889	–0.001333	0.061861
ELASTICNET	1	–	0.036667	0.012444	0.075639
GLM	0.27595	5.651E-08	–	–0.024222	0.038972
RF	1	1	0.18639	–	0.063194
Rpart	2.444E-06	1.68E-07	0.02722	1.18E-08	–

Upper diagonal: estimates of the difference from unpaired *t*-test. Lower diagonal: P values. SVM, support vector machine; GLM, generalized linear method; RF, random forest.



**Figure 1** Decision Tree built based on the random forest machine learning algorithm. FAC, fractional area change; TAPSE, tricuspid annular plane systolic excursion; S, S-wave velocity.

Boruta, were also concordantly selected as the top five most important features only from RF and SVM built-in feature selection functions. All models have an improved performance when trained with the top variables from Boruta except in the ElasticNet model, where it was unchanged.

Also, from the top five important variables extracted from the built-in feature selection functions (Table 3),  $FAC_{RV}$  and hypertension were featured by the different models as top predictors of RV dysfunction. TAPSE and

S' were emphasized by two methods; SVM and random forest but not GLM and Elastic Net. While age was not correlated with the outcome (spearman rho =0.056, P=0.48), it featured in all models except Elastic Net. Diabetes type 2 status showed up only in the regression-based models (GLM and Elastic Net), while smoking exposure was chosen in all models but SVM.

The tree (Figure 1) from the random forest model which classifies best the training data (mtry: 3, OOB error: 4.63%, number of trees: 500). The tree has the maximum number

of nodes (25 terminal nodes corresponding to predictions, either normal or abnormal). Finally, an online calculator was constructed using both RF and Elastic-Net Models and is using this link: [https://nelhachem.shinyapps.io/ShinyApp\\_ML/](https://nelhachem.shinyapps.io/ShinyApp_ML/).

## Discussion

In this paper, we describe a new ML-based algorithm based on a random forest plot analysis that provides reasonable accuracy in detecting low RV function using clinical characteristics and 2D-ECHO derived parameters. The top predictors of RV dysfunction vary between ML-models used, but age,  $FAC_{RV}$ , and HTN were recurring selections. TAPSE and S' appeared in the two most accurate models.

Accurate evaluation of RV function is crucial as a diagnostic and prognostic factor of several diseases. This has led to appreciation of its complex 3D shape and underscored the limit of 2D-ECHO images in assessing this function. However, 3D-echocardiography requires expertise and is not yet widely available. In some cases of multiple ECHO-derived parameter RV assessment, contradictory interpretation of the measurements might occur, often leading to more confusion. Hence using clinical features in addition to echo data would be able to guide physician effort to detect impaired RV function. Therefore, the use of parameters other than  $FAC_{RV}$ , TAPSE, and S' should be investigated.

Since TAPSE and S' are inherently similar and usually correlate (28), the use of another widely studied RV 2D-ECHO derived parameter, such as RV strain, might provide an additional benefit in detection of low RV function when added to the production of the accuracy of our models. In fact, RV strain was shown to have a good correlation with RV function in multiple diseases such as COPD, PH and certain types of HF (15,28-31). Furthermore, our group recently showed that using a speckle-based strain analysis could help differentiate between types of cardiomyopathy in an artificial convoluted network, a commonly used ML method (32). Finally, the use of automatic-border detection (ABD) for strain analysis could be used to eliminate intra-operator variability and to lessen the need for heavy post-processing of images as shown in two recent studies which used ABD and ML-based algorithm to assess RV function from 3D-ECHO (33,34). Another study used ABD for volumetric analysis of the RV, an important factor when assessing RV function (35). Recently, the use of ABD in the estimation of RVEF with

3D-ECHO was shown to be superior to manually drawn border, reducing inter- and intra-observer variability of measurements (36-40).

Another easily accessible and cheap method that could be used in detecting low ventricular function is an ECG. In a study published in nature, investigators were able to produce an AI that would detect low LV function (<35%) with an AUC of 0.93 (41). Replicating the model with a large dataset to create an AI to detect RV function or to incorporate ECG with other parameters to help detect low RV function should be studied.

Alternatively, perhaps there would be value in building predictive models that prioritize either specificity or sensitivity. In our study, models were designed to maximize overall predictive accuracy (F1 score), but possibly in this context a useful model might be one prioritizing high sensitivity for identifying low RV function so we can identify patients in need for CMR.

Finally, future investigations will include ABD along with automatic detection of TAPSE + S' from captured ECHO clips, spontaneously inserting them into the model and hence getting an instant RV assessment of RV function. Several studies have previously applied ABD technology to extract RV parameters. This cutting-edge technology is still limited by availability of the advanced imaging machines.

## Limitations

A major limitation of our study is our small population. Using a larger population would make the algorithm more reliable/replicable and would allow the use of additional data features, such as ECG waveforms. In addition, our population represents a negatively balanced dataset, where dysfunctional RV represented only 14% of our total population. Negatively balanced datasets are a notorious limitation for machine learning algorithm. This is the reason we calculated the Matthews correlation coefficient, which is regarded as a balanced measure in that can be used when classes are of different sizes, i.e., unbalanced datasets. Another limitation is that our study is a single center study, with a majority of subjects from a Middle Eastern ethnic background. Thus, more studies should be conducted to validate our model in other populations.

Another limitation is that the study was retrospective, and thus some limitations in extracting parameters from the ECHO windows, since the studies were not done to assess RV function solely. This has led to limitations in computing RV strain, which could have been used in our algorithm as

another feature, as discussed above. The studies included in this analysis were mostly done between 2012 and 2013, before the acquisition of newer higher resolution echocardiography machines. The new machine would have made it easier to acquire images of the RV despite its retrosternal, angle-dependent, respiratory cycle-dependent, position.

Furthermore, the use of RV strain and automated 3D-ECHO-derived RVEF measurements from ECHO images could have provided an added value to our model. However, these measurements pose a technical difficulty outside specialized tertiary medical centers and are not widely used yet, therefore our model uses common 2D-ECHO measurements. Finally, the use of raw ECHO DICOM images in a convolutional neural network-based algorithm might have had an incremental value to the diagnosis of RV dysfunction; however, convolutional neural networks require a significantly larger data set than what is available for training, validating, and testing the models.

## Conclusions

Non-invasive assessment of RV function is a technically challenging task and clinicians usually rely on visual assessment of RV size and contractility. In this study, we trained and validated an ML-based algorithm that could detect low RV function from clinical and 2D-ECHO parameters. The algorithm performed better than common statistical methods.

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## Footnote

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