



OPEN ACCESS

EDITED BY

Evangelos Oikonomou,
National and Kapodistrian University of Athens,
Greece

REVIEWED BY

Emanuele Micaglio,
IRCCS San Donato Polyclinic, Italy
Giuseppe Mascia,
University of Genoa, Italy
Nicola Pierucci,
Sapienza University of Rome, Italy

*CORRESPONDENCE

Jian Zhou

✉ zj9758@126.com

Xiaorong Li

✉ jsxrl@yeah.net

†These authors have contributed equally to this work and share first authorship

RECEIVED 08 August 2024

ACCEPTED 28 November 2024

PUBLISHED 16 December 2024

CITATION

Zhang B, Xie X, Yu J, Wu Y, Zhou J, Li X and Yang B (2024) A new prediction model for sustained ventricular tachycardia in arrhythmogenic cardiomyopathy. *Front. Cardiovasc. Med.* 11:1477931. doi: 10.3389/fcvm.2024.1477931

COPYRIGHT

© 2024 Zhang, Xie, Yu, Wu, Zhou, Li and Yang. This is an open-access article distributed under the terms of the [Creative Commons Attribution License \(CC BY\)](https://creativecommons.org/licenses/by/4.0/). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

A new prediction model for sustained ventricular tachycardia in arrhythmogenic cardiomyopathy

Baowei Zhang^{1†}, Xin Xie^{1†}, Jinbo Yu¹, Yizhang Wu¹, Jian Zhou^{1*}, Xiaorong Li^{1,2*} and Bing Yang^{1,3}

¹Department of Cardiology, Shanghai East Hospital, Tongji University School of Medicine, Shanghai, China, ²Department of Cardiology, Ji'an Center People's Hospital, Ji'an, China, ³Department of Cardiology, The First Affiliated Hospital of Nanjing Medical University, Nanjing, China

Background: Arrhythmogenic cardiomyopathy (ACM) is an inherited cardiomyopathy characterized by high risks of sustained ventricular tachycardia (sVT) and sudden cardiac death. Identifying patients with high risk of sVT is crucial for the management of ACM.

Methods: A total of 147 ACM patients were retrospectively enrolled in the observational study and divided into training and validation groups. The least absolute shrinkage and selection operator (LASSO) regression model was employed to identify factors associated with sVT. Subsequently, a nomogram was constructed based on multivariable logistic regression analysis. The performance of the nomogram was evaluated using the area under the curve (AUC) of the receiver operating characteristic (ROC) curve and calibration curve. Decision curve analysis was conducted to assess the clinical utility of the nomogram.

Results: Seven parameters were incorporated into the nomogram: age, male sex, syncope, heart failure, T wave inversion in precordial leads, left ventricular ejection fraction (LVEF), SDNN level. The AUC of the nomogram to predict the probability of sVT was 0.867 (95% CI, 0.797–0.938) in the training group and 0.815 (95% CI, 0.673–0.958) in the validation group. The calibration curve demonstrated a good consistency between the actual clinical results and the predicted outcomes. Decision curve analysis indicated that the nomogram had higher overall net benefits in predicting sVT.

Conclusion: We have developed and internally validated a new prediction model for sVT in ACM. This model could serve as a valuable tool to accurately identify ACM patients with high risk of sVT.

KEYWORDS

arrhythmogenic cardiomyopathy, sustained ventricular tachycardia, sudden cardiac death, prediction model, nomogram

Introduction

Arrhythmogenic cardiomyopathy (ACM), an inherited cardiomyopathy characterized by high risks of ventricular arrhythmias and sudden cardiac death, is one of the leading causes of SCD in young people and athletes (1–3). Sustained ventricular tachycardia (sVT) constitutes a major contributor to SCD in ACM patients. Although frequency catheter ablation (RFCA) and anti-arrhythmic drugs have effectively reduced sVT incidence, the implanted cardioverter defibrillator (ICD) remains the most efficacious

therapy for preventing SCD in ACM patients (4). However, ACM often presents in adolescents or at young age, and ICD treatment is invasive, associated with various complications, and imposes a dual burden of physiological and psychological stress on patients. Therefore, patients should be thoroughly assessed before undergoing ICD treatment (5). ACM patients with a history of sVT or resuscitated SCD can benefit from secondary prevention with ICD. Nevertheless, for ACM patients without documented sVT, the primary goal in the management of ACM patients is to identify those at high risk of sVT and treated with prophylactic ICD implantation (6).

Previous studies have identified several risk factors for predicting sVT in ACM patients, including gender, proband status, cardiac syncope, degree of exercise restriction, premature ventricular complex (PVC) burden, non-sustained VT, genetic testing, and extent of myocardial involvement (7, 8). Two studies have constructed prediction models for sudden cardiac death and sustained ventricular tachycardia in ACM patients using these risk factors (9, 10). These prediction models could effectively predict the occurrence of sVT and sudden cardiac death, but their predictive value was limited in ACM patients without identified pathogenic gene variants and in those with left ventricular involvement (11–13). These two prediction models considered demographic factors, symptoms, PVCs burden, and extent of myocardial involvement, but they overlooked the pathophysiological mechanisms underlying the occurrence of sVT in ACM patients. Increased cardiac sympathetic nerve activity plays an important role in the occurrence of sVT in ACM patients (14). Our preliminary study found that SDNN, an indicator of cardiac sympathetic nerve activity, could effectively predict the occurrence of sVT in ACM patients (15). The predictive value of SDNN is based on the role of hyperactivity of cardiac sympathetic nerve system in the occurrence of sVT, a mechanism distinct from other traditional risk factors. Therefore, we hypothesized that incorporating SDNN in the prediction model could provide additional value in distinguishing ACM patients with high risk of sVT.

In the present study, a total of 147 ACM patients were retrospectively enrolled and divided into training and validation groups. A prediction model was built on the basis of multivariable logistic regression analysis, and performed well in identifying ACM patients with high risk of sVT in both the training and validation groups.

Methods

Patients

A total of 212 patients diagnosed with ACM were retrospectively enrolled from the First Affiliated Hospital of Nanjing Medical University and Shanghai East Hospital between January 2006 and October 2024. This study was conducted in accordance with the principles outlined in the Declaration of Helsinki (revised in 2013) and received approval from the institutional ethics committee board of the First Affiliated

Hospital of Nanjing Medical University (Approval No. 2011-SR-014). Given the retrospective nature of the study, individual consent was waived. Diagnosis of ACM was based on the updated diagnostic criteria (6, 16). Patients with incomplete data, a history of heart transplantation (having received a normal heart), or other comorbidities associated with a high risk of ventricular arrhythmias (e.g., LVEF <35%) were excluded from the study.

Data collection

All enrolled ACM patients underwent retrospective evaluation of elementary clinical characteristics, as well as electrocardiographic and echocardiographic features. Elementary clinical data encompassed age, sex, age at diagnosis, prior syncope episodes, family history, comorbidities, and both pharmacological and non-pharmacological therapies. Surface electrocardiography (ECG), 24-h Holter monitoring, and echocardiography were meticulously reviewed by two independent physicians to identify potential predictors for risk stratification in ACM. The standard deviation of all normal-normal (NN) intervals (SDNN) was computed as the dispersion of all NN intervals over a 24-h period of Holter monitoring (17). Furthermore, all patients were meticulously assessed by two independent physicians to detect evidence of sVT. sVT was defined as follows: (1) recorded spontaneous persistent ventricular tachycardia (lasting for ≥ 30 s at ≥ 100 beats per minute, or with unstable hemodynamics requiring cardioversion), (2) ventricular fibrillation/flutter, or (3) appropriate ICD intervention in patients with such devices.

Statistical analysis

For continuous variables, the Kolmogorov-Smirnov test was employed to assess the distribution of the data. Variables exhibiting a normal distribution are expressed as mean \pm standard deviation (SD), with differences between groups investigated using the Student's *t*-test. Variables not conforming to a normal distribution are presented with interquartile range and median, and differences were examined using the Mann-Whitney test. Categorical data were summarized using percentages, and differences between groups were evaluated using the χ^2 or Fisher's exact test, as appropriate.

The initial population was randomized into training and validation groups at a ratio of 7:3, which were subsequently utilized for model development and validation, respectively. The least absolute shrinkage and selection operator (LASSO) logistic regression model was employed to identify factors with significant predictive value for the occurrence of sVT from the data in the training group. Factors exhibiting nonzero coefficients in the LASSO regression model were retained. Subsequently, a multivariable logistic regression analysis, incorporating factors selected from the LASSO regression as well as other clinically significant variables, was conducted to construct a predictive model. To maximize the inclusion of effective predictive factors,

variables with a two-sided p -value ≤ 0.1 were incorporated into the prediction model, while factors of evident clinical significance were directly included. To provide quantitative predictions of sVT probability for individual patients, a nomogram was constructed based on the multivariable logistic regression analysis conducted in the training group.

To evaluate the discriminatory capability of the nomogram, the area under the curve (AUC) of the receiver operating characteristic (ROC) curve and calibration curve analyses were conducted in the training group. These methods were also applied to the validation group to further validate the performance of the nomogram. Additionally, decision curve analyses (DCA) were performed to assess the clinical utility of the nomogram by quantifying the net benefits in both groups. All statistical analyses were conducted using R software (Version 4.2.0; <https://www.r-project.org>).

Results

Baseline characteristics

After excluding 65 patients for various reasons, a total of 147 ACM patients were ultimately enrolled in this study (Supplementary Figure S1). Among them, 60.5% of patients (89 individuals) had a history of sVT. Patients with and without sVT exhibited similar characteristics in terms of age, history duration, age at diagnosis, family history of ACM, comorbidities, and pharmacological therapies. However, patients with sVT demonstrated a higher proportion of males (87.6% vs. 56.9%, $p < 0.001$), smokers (29.2% vs. 6.9%, $p = 0.002$), alcohol consumers (15.7% vs. 3.4%, $p = 0.04$), users of class III anti-arrhythmic drugs (49.4% vs. 19.0%, $p < 0.001$), recipients of ICD devices (34.8% vs. 10.3%, $p = 0.002$), individuals who underwent RFCA (34.8% vs. 12.1%, $p = 0.004$), those with T wave inversion in anterior leads (73.0% vs. 48.3%, $p = 0.004$), right ventricular (RV) enlargement (68.5% vs. 32.8%, $p < 0.001$), and lower levels of SDNN [113.0 (89.0, 135.0) vs. 134.5 (113.0, 150.5) ms, $p < 0.001$], compared with patients without sVT (Table 1).

All 147 ACM patients were subsequently randomly divided into the training group ($n = 104$, 70.7%), with the remaining patients allocated to the validation group ($n = 43$, 29.3%). The incidence of sVT was found to be similar between the training and validation groups (56.7% vs. 69.8%, $p = 0.20$). Moreover, demographic characteristics, comorbidities, pharmacological therapies, electrocardiographic, and echocardiographic features were well-balanced between the training and validation groups (Supplementary Table S1).

Feature selection and nomogram construction

In the training group, patients with sVT exhibited a higher proportion of males (93.2% vs. 55.6%, $p < 0.001$), smokers (32.2% vs. 4.4%, $p = 0.001$), alcohol consumers (18.6% vs. 0, $p = 0.006$), recipients of ICD devices (32.2% vs. 11.1%, $p = 0.02$), individuals

TABLE 1 Baseline characteristics of patients enrolled.

Variables	sVT group ($n = 89$, 60.5%)	Non-sVT group ($n = 58$, 39.5%)	P -value
Age (years)	45.9 \pm 15.3	45.0 \pm 15.8	0.72
Male (n , %)	78 (87.6)	33 (56.9)	<0.001
History (years)	3.0 (1.0, 10.0)	4.0 (1.0, 8.5)	0.91
Age of diagnosis (years)	42.0 (28.5, 48.0)	38.0 (28.5, 51.0)	0.93
Syncope (n , %)	39 (43.8)	21 (36.2)	0.46
ICD (n , %)	31 (34.8)	6 (10.3)	0.002
RFCA (n , %)	31 (34.8)	7 (12.1)	0.004
Family history (n , %)	8 (9.0)	3 (5.2)	0.59
Comorbidities			
Hypertension (n , %)	10 (11.2)	10 (17.2)	0.43
Diabetes (n , %)	0 (0)	3 (5.2)	0.12
Coronary artery disease (n , %)	2 (2.2)	0 (0)	0.67
Heart failure (n , %)	7 (7.9)	9 (15.5)	0.24
Smoking (n , %)	26 (29.2)	4 (6.9)	0.002
Alcohol (n , %)	14 (15.7)	2 (3.4)	0.04
Anti-arrhythmic drugs			
Class I (n , %)	15 (16.9)	7 (12.1)	0.58
Class II (n , %)	43 (48.3)	18 (31.0)	0.06
Class III (n , %)	44 (49.4)	11 (19.0)	<0.001
Class IV (n , %)	3 (3.4)	0 (0)	0.41
Characteristics on electrocardiography			
RBBB (n , %)	33 (37.1)	18 (31.0)	0.57
TWI-A (n , %)	65 (73.0)	28 (48.3)	0.004
TWI-I (n , %)	38 (42.7)	19 (32.8)	0.30
NSVT (n , %)	39 (43.8)	25 (43.1)	1.00
SDNN (ms)	113.0 (89.0, 135.0)	134.5 (113.0, 150.5)	<0.001
Characteristics on echocardiography			
RV enlargement (n , %)	61 (68.5)	19 (32.8)	<0.001
RVOT dyskinesia (n , %)	40 (44.9)	31 (53.4)	0.40
RV free wall dyskinesia (n , %)	63 (70.8)	34 (58.6)	0.18
PAH (n , %)	32 (36.0)	13 (22.4)	0.12
Tricuspid regurgitation (n , %)	63 (70.8)	32 (55.2)	0.08
Mitral regurgitation (n , %)	26 (29.2)	19 (32.8)	0.79
LVDD (mm)	47.0 (41.5, 50.0)	48.0 (45.0, 51.0)	0.09
LVEF (%)	61.0 (52.5, 65.8)	62.9 (55.0, 65.8)	0.59

sVT, sustained ventricular tachycardia; ICD, implantable cardiac defibrillator; RFCA, radiofrequency catheter ablation; RBBB, right bundle branch block; TWI-A, T wave inversion on anterior leads; TWI-I, T wave inversion on inferior leads; NSVT, non-sustained ventricular tachycardia; SDNN, Standard deviation of NN intervals; RV, right ventricle; PAH, pulmonary artery hypertension; LVDD, left ventricular end diastolic dimension; LVEF, left ventricular ejection fraction.

who underwent RFCA (39.0% vs. 8.9%, $p = 0.001$), users of class III anti-arrhythmic drugs (AAD) therapy (54.2% vs. 24.4%, $p = 0.004$), those with T wave inversion in anterior leads (74.6% vs. 51.1%, $p = 0.02$), and RV enlargement (71.2% vs. 37.8%, $p = 0.001$), but exhibited lower levels of SDNN [114.0 (88.0, 136.0) vs. 132.0 (115.0, 151.0) ms, $p = 0.003$], consistent with findings observed in the entire cohort (Supplementary Table S2). The LASSO logistic regression model identified 11 features as potential predictors of sVT in the training group (Figure 1). Subsequently, a multivariable logistic regression

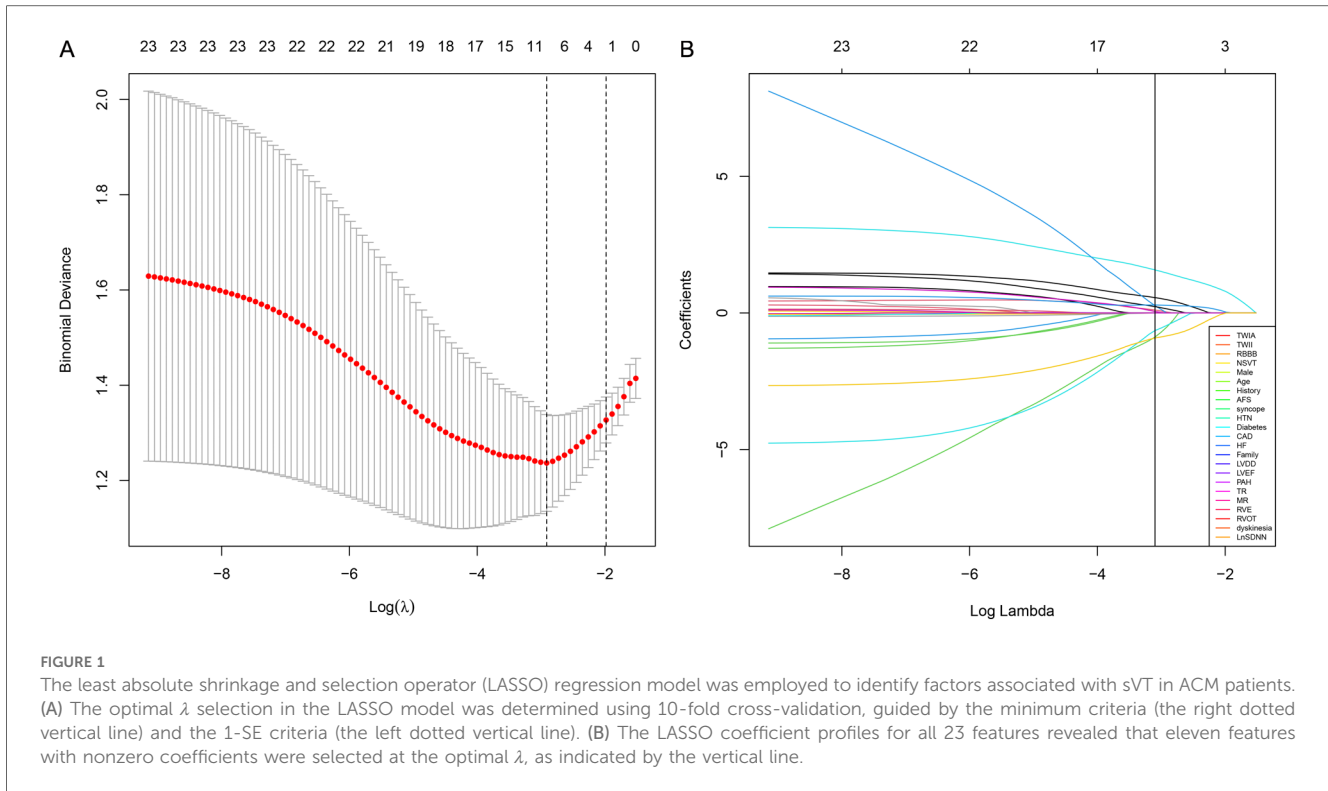


TABLE 2 Predictors for sVT in patients with ACM.

Variable	Prediction model		
	β	Odds ratio (95% CI)	P-value
TWI-A	1.38	3.97 (1.148–15.647)	0.04
LnSDNN	-2.19	0.11 (0.011–0.813)	0.04
Male	2.81	16.62 (3.613–108.370)	<0.001
Heart failure	-4.43	0.01 (0.001–0.155)	0.002
LVEF	-0.11	0.90 (0.818–0.978)	0.02
Syncope	1.01	2.73 (0.698–11.527)	0.15
Age	0.02	1.02 (0.970–1.071)	0.44

sVT, sustained ventricular tachycardia; ACM, arrhythmic cardiomyopathy; CI, confidence interval; TWI-A, T wave inversion on anterior leads; TWI-I, T wave inversion on inferior leads; NSVT, non-sustained ventricular tachycardia; LVDD, left ventricular end diastolic dimension; LVEF, left ventricular ejection fraction; RV, right ventricle; PAH, pulmonary artery hypertension; NSVT, non-sustained ventricular tachycardia; SDNN, Standard deviation of NN intervals; NA, not available.

analysis based on the factors selected in the LASSO regression was conducted to construct the prediction model. Additionally, for the comprehensive predictive value of sVT in ACM patients, syncope and left ventricular ejection fraction (LVEF) were also included. Ultimately, age, syncope, LVEF, male sex, heart failure, T wave inversion in anterior leads, and SDNN were incorporated into the prediction model (Table 2).

Based on the predictors identified by the LASSO logistic regression model in the training group, a nomogram was constructed in the training group for identifying ACM patients at high risk of sustained ventricular tachycardia (sVT). The nomogram included five significant predictive features and two canonical predictors (Figure 2). Each factor in the nomogram

was assigned an individual score based on its value, and a total score was calculated by summing the scores of all factors. The final score derived from the nomogram could be utilized to estimate the risk of sVT for a given patient (Figure 2).

Performance of the nomogram

We firstly validated the predictive value of the nomogram for identifying ACM patients at high risk of sVT in the training group by means of the ROC curve. The AUC of the nomogram for predicting the occurrence of sVT in ACM patients was 0.867 (95% CI, 0.797–0.938) (Figure 3A). The calibration curve demonstrated that the predictive nomogram for the occurrence of sVT provided excellent estimations of actual probabilities (Figure 3B).

Subsequently, the performance of the prediction model was validated in the validation group. Among the 43 patients allocated to the validation group, 30 had a history of sVT (69.8%). Patients with sVT in the validation exhibited similar proportions of males, smokers, recipients of ICD devices, individuals who underwent RFCA, and those with T-wave inversion in anterior leads, which was different from patients in the training group (Supplementary Table S3). However, the AUC of the prediction model to forecast the incidence of sVT was 0.815 (95% CI, 0.673–0.958) in the validation group (Figure 3C), indicating that the prediction model demonstrated excellent performance in identifying ACM patients at high risk of sVT. Furthermore, the calibration curve also illustrated good consistency between the actual incidence of sVT and the predicted probability in the validation group (Figure 3D).

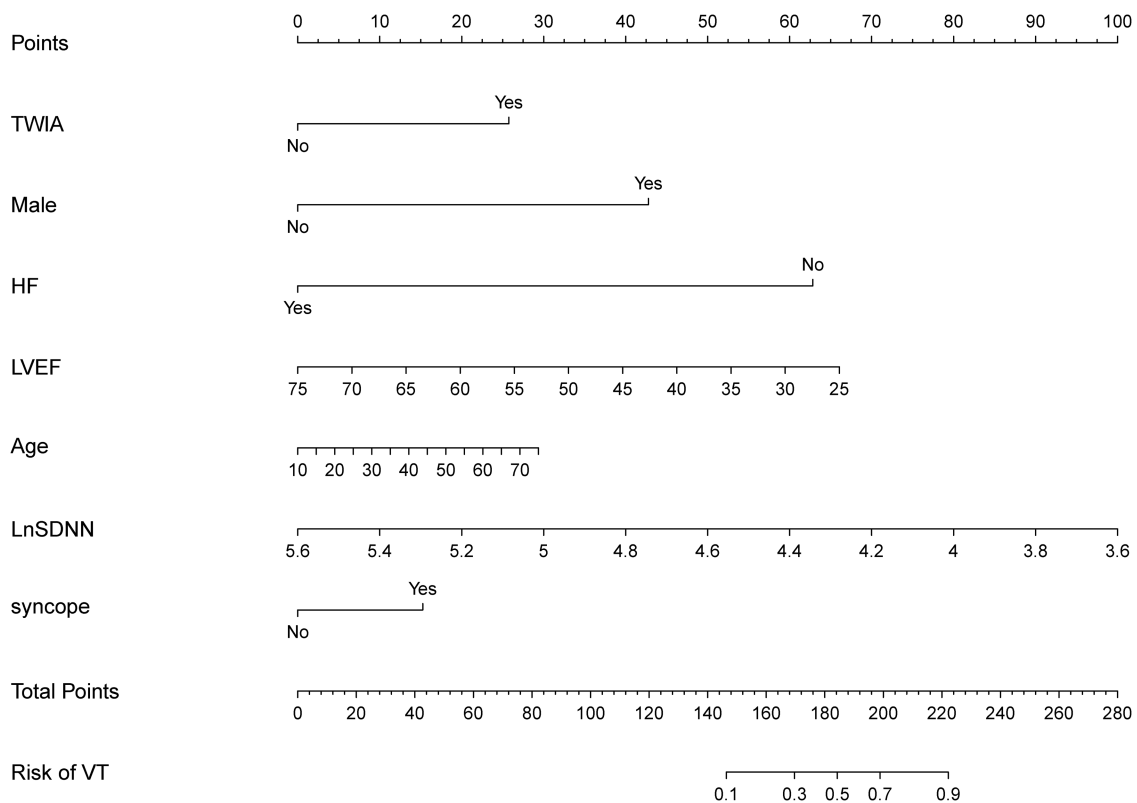


FIGURE 2

The nomogram was constructed in the training group for identifying ACM patients at high risk of sVT. Each factor in the nomogram was assigned an individual score based on its value, and the scores of all factors were summarized to a total score, which was used to estimate the risk of sVT for a given patient.

Clinical utility

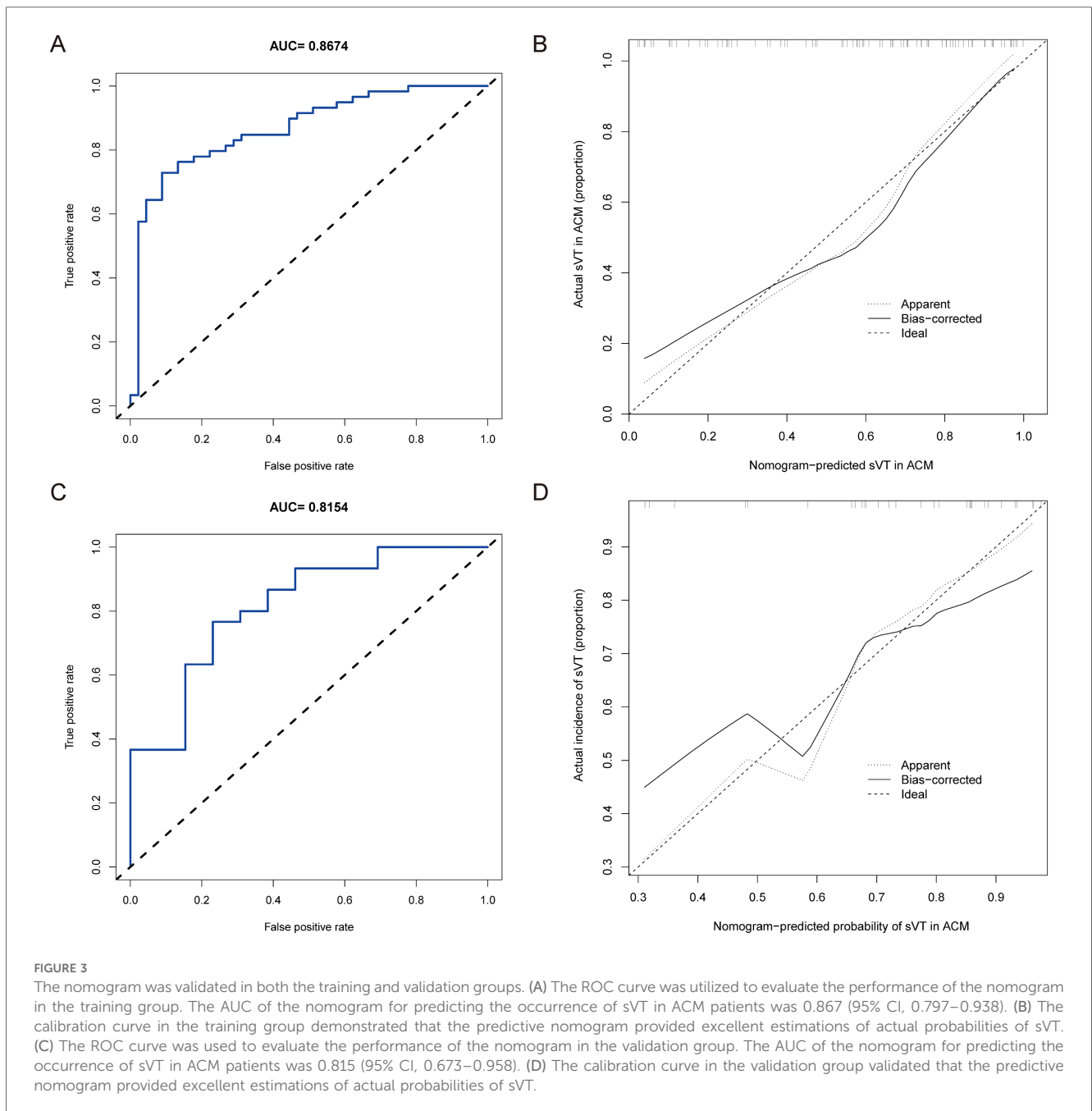
To explore the clinical implications of our predictive model, decision curve analyses were conducted in both the training group and the validation group. As depicted in Figure 4, the results of DCA illustrated that the predictive model provided excellent overall net benefits in predicting the incidence of sVT in ACM patients.

Discussion

In this study, we developed and internally validated a prediction model for identifying ACM patients at high risk of sVT. All features incorporated into the prediction model consisted of elementary clinical information readily available in clinical practice, thus rendering the prediction model suitable for routine use in clinical settings. Predicted risks were consistent with the actual risks of sVT in both the training group [AUC 0.867 (95% CI, 0.797–0.938)] and the validation group [AUC 0.815 (95% CI, 0.673–0.958)]. Furthermore, decision curve analysis also demonstrated that the model was associated with a high net benefit in predicting the incidence of sVT in ACM patients.

High risk of sVT is the prominent clinical feature in patients with ACM. Previous studies have shown that patients with ACM

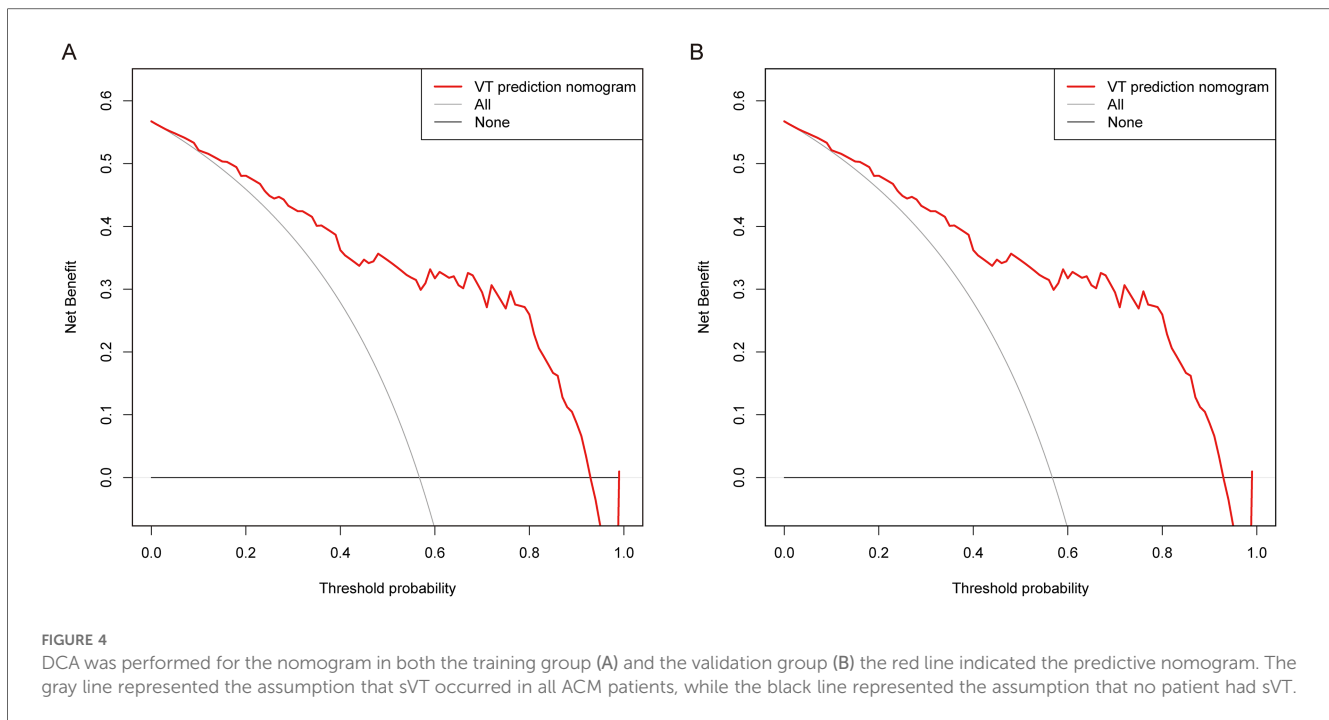
exhibit annual sVT rates ranging from 2% to 10%, which are notably higher compared to patients with other non-ischemic cardiomyopathies (18–20). Substitution of the ventricular myocardium with fibrous or fibrofatty tissue represents a distinctive pathological feature in patients with ACM (21). The infiltration of fibrous or fibrofatty tissue disrupts the conduction velocity and direction of cardiac electrical activity, predisposing individuals to the formation of reentrant circuits and thereby increasing the risk of sVT (22). Premature ventricular beats and sVT exhibiting a left bundle branch block (LBBB) morphology with a non-inferior axis pattern in ACM patients indicate the “triangle of dysplasia” as the origin of ventricular arrhythmias, thereby supporting cardiac fibrosis as an important mechanism of sVT in ACM (23). However, clinical studies have shown that VAs could manifest before identifiable cardiac structural changes occur, suggesting the involvement of other mechanisms contributing to ventricular arrhythmias in ACM. Subsequent studies identified impairment of action potential upstroke velocity and propagation of action potential in cardiomyocytes from ACM mice, attributed to the dislocation of connexin 43 (Cx43) and dysfunction of sodium channel Nav1.5 (24–26). Furthermore, dysregulation of calcium (Ca^{2+}) handling has been observed in cardiomyocytes from ACM mice (27). This disruption of Ca^{2+} homeostasis predisposes to sarcoplasmic reticulum spontaneous Ca^{2+} release and early afterdepolarizations, ultimately leading to increased cardiomyocyte



excitability and the onset of ventricular arrhythmogenesis (28). Consequently, a combination of cardiac fibrosis, dislocation of Cx43 and Nav1.5, and disturbance of Ca^{2+} homeostasis is postulated to contribute to arrhythmogenesis in ACM.

In addition, the abnormal activity of sympathetic nerve system contributes to the ventricular arrhythmogenesis in ACM (29). Our clinical studies, along with others, have identified the overactivation of the cardiac sympathetic system in patients with ACM (14, 15, 30). Stimulation of myocardial β_1 adrenergic receptors can increase cellular Ca^{2+} loading through the cyclic adenosine monophosphate (cAMP)–protein kinase A (PKA) pathway, resulting in an augmentation in the L-type calcium current, as well as through SERCA2-mediated reduction of Ca^{2+} re-uptake. The resultant

elevation in diastolic Ca^{2+} levels within the cardiomyocyte may heighten the likelihood of spontaneous sarcoplasmic reticulum calcium release, subsequently leading to the induction of afterdepolarization and extra stimulus, which could serve as triggers for inducing sVT (31). Additionally, the sprouting and remodeling of sympathetic nerve fibers secondary to sympathetic nerve fiber necrosis result in the heterogeneous reinnervation of cardiomyocytes, potentially promoting the dispersion of action potential duration and providing a suitable substrate for the maintenance of sVT (32). Therefore, the overactivity of the cardiac sympathetic nervous system not only provides triggers but also a suitable substrate for sVT in ACM patients.



Sustained ventricular tachycardia in ACM patients is associated with abnormal electroactivity of cardiomyocytes, cardiac fibrosis, and overactivation of the cardiac sympathetic system. Therefore, clinical parameters associated with these factors may serve as predictors of sVT in ACM patients. Previous studies have identified several predictors of sVT in ACM patients, including recent syncope, younger age at presentation, male sex, decreased left ventricular ejection fraction (LVEF), non-sustained ventricular tachycardia (NSVT), PVC burden, RV dysfunction, and leads with T-wave inversion (33–36). Syncope is a common initial symptom of ACM and often indicates that the patient has experienced hemodynamically unstable sVT, although non-cardiogenic causes of syncope should also be considered in some cases (37). Therefore, recent syncope is regarded as a significant indicator for identifying high-risk patients with sVT in ACM. NSVT and PVC burden may reflect the abnormal electroactivity of cardiomyocytes. Decreased LVEF and RV dysfunction may indicate more severe cardiac fibrosis and scar formation in both ventricles (38). Additionally, the 12-lead ECG plays a crucial role in the diagnosis of ACM, localization of ventricular arrhythmias, and identification of high-risk patients for sVT (39). T-wave inversion is not only one of the ECG criteria for diagnosing ACM, but it is also an independent factor in predicting the occurrence of sVT in patients with ACM (20, 40). This study also identified that T-wave inversion in the anterior leads was independently associated with sVT in ACM patients. A recent study integrated these predictors and developed a prediction model for sVT in ACM patients (10). However, subsequent study showed that this prediction model had good performance in distinguishing patients with and without sVT in ACM patients with pathogenic mutant *PKP2* (C-index 0.77), but had limited value in predicting the risk of sVT in ACM patients with

unknown pathogenic gene mutations (9). In addition, this predictive model only incorporated parameters related to the abnormal electroactivity of cardiomyocytes and cardiac fibrosis. We hypothesized that incorporation parameters associated with the overactivity of the cardiac sympathetic system might enhance the value of the prediction model.

SDNN, a parameter of heart rate variability, is regarded as an indicator of cardiac sympathetic system activity. Our previous study revealed that ACM patients exhibited decreased SDNN levels, and SDNN level emerged as an independent predictor of sVT in ACM patients (15). The present study also demonstrated that the SDNN levels in the sVT group were significantly lower than those in the control group. Further analysis using LASSO regression and logistic regression indicated that SDNN levels were independently associated with the occurrence of sVT in patients with ACM. The prediction model incorporating SDNN and these canonical parameters effectively identifies high-risk patients for sVT in individuals with ACM. Furthermore, calibration plots exhibited good agreement between predicted and observed sVT in ACM patients. Therefore, the predictive model in this study may more effectively identify high-risk patients for sVT among individuals with ACM.

The significant clinical benefit of our prediction model lies in its precise quantification of arrhythmic risk tailored to individual ACM patients, offering prognostic insights crucial for guiding clinical decisions regarding prophylactic ICD implantation. Additionally, all parameters included in the prediction model are readily accessible clinical variables, rendering the model user-friendly. It is important to acknowledge that all these parameters in the prediction model might change as ACM progresses. Thus, ACM patients should undergo periodic re-stratification as recommended in a recent expert consensus document.

Limitations

One potential limitation is that all patients in this study were from tertiary medical centers in China and had relatively high incidences of sVT within this cohort (60.5%). The bias in patient selection could potentially impact the overall accuracy of this prediction model when applied to other ACM patients. Secondly, the relatively small sample size of patients used to develop a new prediction model might be associated with selection bias, reduce the statistical power, increase the risk of type II errors, and affect the reliability of the conclusions drawn. Although we have tried to minimize the impact of a small sample size by adding a validation cohort, caution is still required when interpreting and applying the results of our study. Additionally, due to its retrospective design, our study possesses certain drawbacks compared to a prospective study. Moreover, genetics play important roles in the diagnosis of ACM and sVT risk evaluation in ACM (41, 42). However, only a small portion of patients in this study underwent genetic testing, so genetics were not included in the construction of the prediction model. Finally, some *p*-values in this study were close to the significance threshold, raising the potential for type I error. Therefore, further large prospective cohort studies are needed to assess the accuracy and clinical utility of the prediction model in risk stratification for sVT in ACM patients.

In summary, we developed and validated a new prediction model to generate individualized estimates of the risk of incident sVT in ACM patients. Utilizing easily accessible clinical parameters, this model has the potential to accurately identify high-risk patients with sVT, thereby offering significant value for guiding prophylactic ICD implantation in ACM patients.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Ethics statement

The studies involving humans were approved by the institutional Ethics Committee board of the First Affiliated Hospital of Nanjing Medical University. The studies were conducted in accordance with the local legislation and institutional requirements. The ethics committee/institutional review board waived the requirement of written informed consent for participation from the participants or the participants' legal guardians/next of kin because Given the retrospective nature of the study, individual consent was waived.

Author contributions

BZ: Writing – original draft, Writing – review & editing. XX: Writing – original draft. JY: Methodology, Supervision, Writing – review & editing. YW: Data curation, Formal Analysis, Project administration, Validation, Writing – original draft. JZ: Conceptualization, Formal Analysis, Methodology, Project administration, Resources, Supervision, Validation, Visualization, Writing – review & editing. XL: Conceptualization, Data curation, Formal Analysis, Funding acquisition, Investigation, Resources, Software, Visualization, Writing – review & editing. BY: Conceptualization, Funding acquisition, Resources, Supervision, Writing – review & editing.

Funding

The author(s) declare financial support was received for the research, authorship, and/or publication of this article. This work was supported by grants from the Key Specialty Construction Project of Shanghai Pudong New Area Health Commission (PWZzk2022-03), the Top-level Clinical Discipline Project of Shanghai Pudong District Grant/Award (PWYgf2021- 01), the Health Commission of Shanghai Pudong District (PW2023A-17), and the Science and Technology Plan Project of Health Commission of Jiangxi Province (202310098).

Acknowledgments

We thank all participants in this study and all work staff in the department of cardiology, the First Affiliated Hospital of Nanjing Medical University and Shanghai East Hospital.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fcvm.2024.1477931/full#supplementary-material>

References

- Corrado D, Link MS, Calkins H. Arrhythmogenic right ventricular cardiomyopathy. *N Engl J Med*. (2017) 376:61–72. doi: 10.1056/NEJMra1509267
- Finocchiaro G, Radaelli D, D'Errico S, Papadakis M, Behr ER, Sharma S, et al. Sudden cardiac death among adolescents in the United Kingdom. *J Am Coll Cardiol*. (2023) 81:1007–17. doi: 10.1016/j.jacc.2023.01.041
- Zhang B, Wu Y, Zhou C, Xie J, Zhang Y, Yang X, et al. Hyperactivation of ATF4/TGF- β 1 signaling contributes to the progressive cardiac fibrosis in arrhythmogenic cardiomyopathy caused by DSG2 variant. *BMC Med*. (2024) 22:361. doi: 10.1186/s12916-024-03593-8
- Zeppenfeld K, Tfelt-Hansen J, de Riva M, Winkel BG, Behr ER, Blom NA, et al. 2022 ESC guidelines for the management of patients with ventricular arrhythmias and the prevention of sudden cardiac death. *Eur Heart J*. (2022) 43:3997–4126. doi: 10.1093/eurheartj/ehac262
- Mariani MV, Pierucci N, Fanasio F, Laviola D, Silvetti G, Piro A, et al. Inherited arrhythmias in the pediatric population: an updated overview. *Medicina (Kaunas)*. (2024) 60:94. doi: 10.3390/medicina6010094
- Towbin JA, McKenna WJ, Abrams DJ, Ackerman MJ, Calkins H, Darrioux FCC, et al. 2019 HRS expert consensus statement on evaluation, risk stratification, and management of arrhythmogenic cardiomyopathy. *Heart Rhythm*. (2019) 16:e301–72. doi: 10.1016/j.hrthm.2019.05.007
- Wallace R, Calkins H. Risk stratification in arrhythmogenic right ventricular cardiomyopathy. *Arrhythm Electrophysiol Rev*. (2021) 10:26–32. doi: 10.15420/aer.2020.39
- Bosman LP, Te Riele A. Arrhythmogenic right ventricular cardiomyopathy: a focused update on diagnosis and risk stratification. *Heart*. (2022) 108:90–7. doi: 10.1136/heartjnl-2021-319113
- Cadrin-Tourigny J, Bosman LP, Nozza A, Wang W, Tadros R, Bhonsale A, et al. A new prediction model for ventricular arrhythmias in arrhythmogenic right ventricular cardiomyopathy. *Eur Heart J*. (2019) 40:1850–8. doi: 10.1093/eurheartj/ehz103
- Cadrin-Tourigny J, Bosman LP, Nozza A, Wang W, Tadros R, Bhonsale A, et al. A new prediction model for ventricular arrhythmias in arrhythmogenic right ventricular cardiomyopathy. *Eur Heart J*. (2022) 43:e1–9. doi: 10.1093/eurheartj/ehac180
- Protonotarios A, Bariani R, Cappelletto C, Pavlou M, Garcia-Garcia A, Cipriani A, et al. Importance of genotype for risk stratification in arrhythmogenic right ventricular cardiomyopathy using the 2019 ARVC risk calculator. *Eur Heart J*. (2022) 43:3053–67. doi: 10.1093/eurheartj/ehac235
- Aquaro GD, De Luca A, Cappelletto C, Raimondi F, Bianco F, Botto N, et al. Prognostic value of magnetic resonance phenotype in patients with arrhythmogenic right ventricular cardiomyopathy. *J Am Coll Cardiol*. (2020) 75:2753–65. doi: 10.1016/j.jacc.2020.04.023
- Casella M, Gasperetti A, Gaetano F, Busana M, Sommariva E, Catto V, et al. Long-term follow-up analysis of a highly characterized arrhythmogenic cardiomyopathy cohort with classical and non-classical phenotypes—a real-world assessment of a novel prediction model: does the subtype really matter. *Europace*. (2020) 22:797–805. doi: 10.1093/europace/euz352
- Assis FR, Krishnan A, Zhou X, James CA, Murray B, Tichnell C, et al. Cardiac sympathectomy for refractory ventricular tachycardia in arrhythmogenic right ventricular cardiomyopathy. *Heart Rhythm*. (2019) 16:1003–10. doi: 10.1016/j.hrthm.2019.01.019
- Zhang B, Zhou C, Liu J, Zhang J, Jiang C, Tang M, et al. Impaired heart rate variability in patients with arrhythmogenic cardiomyopathy: a multicenter retrospective study in China. *Front Cardiovasc Med*. (2022) 9:1044797. doi: 10.3389/fcvm.2022.1044797
- Marcus FI, McKenna WJ, Sherrill D, Basso C, Baucé B, Bluemke DA, et al. Diagnosis of arrhythmogenic right ventricular cardiomyopathy/dysplasia: proposed modification of the task force criteria. *Eur Heart J*. (2010) 31:806–14. doi: 10.1093/eurheartj/ehq025
- Zhang B, Yu J, Wu Y, Li X, Xie X, Tao A, et al. The significance of heart rate variability in patients with frequent premature ventricular complex originating from the ventricular outflow tract. *Clin Cardiol*. (2024) 47:e24174. doi: 10.1002/clc.24174
- Bhonsale A, James CA, Tichnell C, Murray B, Gagari D, Philips B, et al. Incidence and predictors of implantable cardioverter-defibrillator therapy in patients with arrhythmogenic right ventricular dysplasia/cardiomyopathy undergoing implantable cardioverter-defibrillator implantation for primary prevention. *J Am Coll Cardiol*. (2011) 58:1485–96. doi: 10.1016/j.jacc.2011.06.043
- Folino AF, Buja G, Baucé B, Thiene G, dalla Volta S, Nava A. Heart rate variability in arrhythmogenic right ventricular cardiomyopathy correlation with clinical and prognostic features. *Pacing Clin Electrophysiol*. (2002) 25:1285–92. doi: 10.1046/j.1460-9592.2002.01285.x
- Calkins H, Corrado D, Marcus F. Risk stratification in arrhythmogenic right ventricular cardiomyopathy. *Circulation*. (2017) 136:2068–82. doi: 10.1161/CIRCULATIONAHA.117.030792
- Austin KM, Trembley MA, Chandler SF, Sanders SP, Saffitz JE, Abrams DJ, et al. Molecular mechanisms of arrhythmogenic cardiomyopathy. *Nat Rev Cardiol*. (2019) 16:519–37. doi: 10.1038/s41569-019-0200-7
- Segura-Rodríguez D, Bermúdez-Jiménez FJ, Carriel V, López-Fernández S, González-Molina M, Oyonarte Ramírez JM, et al. Myocardial fibrosis in arrhythmogenic cardiomyopathy: a genotype-phenotype correlation study. *Eur Heart J Cardiovasc Imaging*. (2020) 21:378–86. doi: 10.1093/ehjci/jez277
- van der Voorn SM, Te Riele A, Basso C, Calkins H, Remme CA, van Veen TAB. Arrhythmogenic cardiomyopathy: pathogenesis, pro-arrhythmic remodeling, and novel approaches for risk stratification and therapy. *Cardiovasc Res*. (2020) 116:1571–84. doi: 10.1093/cvr/cvaa084
- Noorman M, Hakim S, Kessler E, Groeneweg JA, Cox MG, Asimaki A, et al. Remodeling of the cardiac sodium channel, connexin43, and plakoglobin at the intercalated disk in patients with arrhythmogenic cardiomyopathy. *Heart Rhythm*. (2013) 10:412–9. doi: 10.1016/j.hrthm.2012.11.018
- Zhang B, Wu Y, Yang X, Xiang Y, Yang B. Molecular insight into arrhythmogenic cardiomyopathy caused by DSG2 mutations. *Biomed Pharmacother*. (2023) 167:115448. doi: 10.1016/j.biopha.2023.115448
- Krahn AD, Wilde AAM, Calkins H, La Gerche A, Cadrin-Tourigny J, Roberts JD, et al. Arrhythmogenic right ventricular Cardiomyopathy. *JACC Clin Electrophysiol*. (2022) 8:533–53. doi: 10.1016/j.jacep.2021.12.002
- Cerrone M, Montnach J, Lin X, Zhao YT, Zhang M, Agullo-Pascual E, et al. Plakophilin-2 is required for transcription of genes that control calcium cycling and cardiac rhythm. *Nat Commun*. (2017) 8:106. doi: 10.1038/s41467-017-00127-0
- Kim JC, Perez-Hernandez M, Alvarado FJ, Maurya SR, Montnach J, Yin Y, et al. Disruption of $ca(2+)$ homeostasis and connexin 43 hemichannel function in the right ventricle precedes overt arrhythmogenic cardiomyopathy in plakophilin-2-deficient mice. *Circulation*. (2019) 140:1015–30. doi: 10.1161/CIRCULATIONAHA.119.039710
- Dusi V, De Ferrari GM. The sympathetic nervous system and arrhythmogenic right ventricular cardiomyopathy: further evidence of a strong tie. *Heart Rhythm*. (2019) 16:1011–2. doi: 10.1016/j.hrthm.2019.02.004
- Wichter T, Hindricks G, Lerch H, Bartenstein P, Borggrefe M, Schober O, et al. Regional myocardial sympathetic dysinnervation in arrhythmogenic right ventricular cardiomyopathy. An analysis using 123I-meta-iodobenzylguanidine scintigraphy. *Circulation*. (1994) 89:667–83. doi: 10.1161/01.CIR.89.2.667
- Maack C, Eschenhagen T, Hamdani N, Heinzel FR, Lyon AR, Manstein DJ, et al. Treatments targeting inotropy. *Eur Heart J*. (2019) 40:3626–44. doi: 10.1093/eurheartj/ehy600
- Herring N, Kalla M, Paterson DJ. The autonomic nervous system and cardiac arrhythmias: current concepts and emerging therapies. *Nat Rev Cardiol*. (2019) 16:707–26. doi: 10.1038/s41569-019-0221-2
- Link MS, Laidlaw D, Polonsky B, Zareba W, McNitt S, Gear K, et al. Ventricular arrhythmias in the north American multidisciplinary study of ARVC: predictors, characteristics, and treatment. *J Am Coll Cardiol*. (2014) 64:119–25. doi: 10.1016/j.jacc.2014.04.035
- Mazzanti A, Ng K, Faragli A, Maragna R, Chiodaroli E, Orphanou N, et al. Arrhythmogenic right ventricular cardiomyopathy: clinical course and predictors of arrhythmic risk. *J Am Coll Cardiol*. (2016) 68:2540–50. doi: 10.1016/j.jacc.2016.09.951
- Corrado D, Calkins H, Link MS, Leoni L, Favale S, Bevilacqua M, et al. Prophylactic implantable defibrillator in patients with arrhythmogenic right ventricular cardiomyopathy/dysplasia and no prior ventricular fibrillation or sustained ventricular tachycardia. *Circulation*. (2010) 122:1144–52. doi: 10.1161/CIRCULATIONAHA.109.913871
- Corrado D, Leoni L, Link MS, Della Bella P, Gaita F, Curnis A, et al. Implantable cardioverter-defibrillator therapy for prevention of sudden death in patients with arrhythmogenic right ventricular cardiomyopathy/dysplasia. *Circulation*. (2003) 108:3084–91. doi: 10.1161/01.CIR.0000103130.33451.D2
- Castillo E. Recurrent syncope in a patient with arrhythmogenic right ventricular cardiomyopathy. *Cureus*. (2023) 15:e45850. doi: 10.7759/cureus.45850
- Corrado D, Anastakis A, Basso C, Baucé B, Blomstrom-Lundqvist C, Bucciarelli-Ducci C, et al. Proposed diagnostic criteria for arrhythmogenic cardiomyopathy: european task force consensus report. *Int J Cardiol*. (2024) 395:131447. doi: 10.1016/j.ijcard.2023.131447
- Efremidis M, Vlachos K, Kyriakopoulou M, Mililis P, Martin CA, Bazoukis G, et al. The RV(1)-V(3) transition ratio: a novel electrocardiographic criterion for the differentiation of right versus left outflow tract premature ventricular complexes. *Heart Rhythm O2*. (2021) 2:521–8. doi: 10.1016/j.hroo.2021.07.009
- Laredo M, Tovia-Brodie O, Milman A, Michowitz Y, Roudijk RW, Peretto G, et al. Electrocardiographic findings in patients with arrhythmogenic cardiomyopathy and right bundle branch block ventricular tachycardia. *Europace*. (2023) 25:1025–34. doi: 10.1093/europace/euac267
- Casian M, Papadakis M, Jurcut R. Arrhythmogenic right ventricular cardiomyopathies (ARVC): diagnostic challenges from imaging to genetics. *Kardiol Pol*. (2024) 82:1059–70. doi: 10.33963/v.phj.102391
- Hoorntje ET, Burns C, Marsili L, Corden B, Parikh VN, Te Meerman GJ, et al. Variant location is a novel risk factor for individuals with arrhythmogenic cardiomyopathy due to a desmoplakin (DSP) truncating variant. *Circ Genom Precis Med*. (2023) 16:e003672. doi: 10.1161/CIRCGEN.121.003672