

# Clinical characteristics and follow-up of pediatric-onset arrhythmogenic right ventricular cardiomyopathy

**Short title:** Pediatric-onset arrhythmogenic right ventricular cardiomyopathy

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45

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47

48 **Structured Abstract**

49 **Objectives:** To describe characteristics, cascade screening results, and predictors of adverse  
50 outcome in pediatric-onset arrhythmogenic right ventricular cardiomyopathy (ARVC).

51 **Background:** While ARVC is increasingly recognized in children, pediatric ARVC cohorts  
52 remain underrepresented in the literature.

53 **Methods:** We included 12 probands with pediatric-onset ARVC (<18 years at diagnosis) and 68  
54 pediatric relatives (<18 years at first evaluation) referred for cascade screening. ARVC diagnosis  
55 was based on 2010 Task Force Criteria (TFC). Clinical presentation, diagnostic testing, and  
56 outcomes (sustained ventricular tachycardia [VT]; heart failure) were ascertained. Predictors of  
57 adverse outcome were determined by univariable logistic regression.

58 **Results:** Pediatric-onset ARVC was diagnosed in 12 probands and 12 (18%) relatives at age 16.6  
59 (interquartile range 13.8-17.4) years, while 12 (18%) relatives reached ARVC diagnosis as adults  
60 (age 22.0 [interquartile range 20.0-26.7] years). Sudden cardiac death/arrest was the first disease  
61 manifestation in 3 (25%) probands and 3 (4%) relatives. In patients without ARVC diagnosis at  
62 presentation (n=61), ECG and Holter monitoring abnormalities occurred prior to development of  
63 imaging TFC ( $7.3\pm 5.0$  vs.  $8.4\pm 5.0$  years). Clinical course was characterized by sustained VT  
64 (91%) and heart failure (36%) in probands, which were rare in relatives (2% and 0%, respectively).  
65 Male sex ( $p<0.01$ ), T-wave inversion V1-3 ( $p<0.01$ ), premature ventricular complexes/runs  
66 ( $p\leq 0.01$ ) and decrease of biventricular ejection fraction ( $p\leq 0.01$ ) were associated with VT  
67 occurrence.

68 **Conclusions:** Pediatric ARVC carries high arrhythmic risk, particularly in probands. Disease  
69 progression is particularly observed on ECG or Holter monitoring. Arrhythmic events are

70 associated with male sex, T-wave inversions, premature ventricular complexes/runs and reduced  
71 biventricular ejection fraction.

72

73 **Keywords:**

74 Arrhythmogenic right ventricular cardiomyopathy, cascade screening, genetics, heart failure,  
75 pediatric-onset, sudden cardiac death, ventricular tachycardia.

76

77 **Condensed Abstract**

78 While ARVC is recognized in young subjects, pediatric ARVC remains underrepresented in the  
79 literature. In this multicenter study of 80 (12 probands, 68 relatives) pediatric ARVC patients,  
80 disease penetrance peaked during late adolescence (median age at diagnosis 16.6[range 9-17]  
81 years). Notably, sudden cardiac death/arrest was the presenting symptom in both probands  
82 (n=3/12; 25%) and relatives (n=3/68; 4%), while clinical course after diagnosis was more benign  
83 in relatives compared to probands (2% vs. 91% ventricular tachycardia during follow-up).  
84 Arrhythmic events associated with male sex ( $p<0.01$ ) ECG ( $p<0.01$ ), Holter monitoring  
85 abnormalities ( $p=0.01$ ) and reduced biventricular ejection fraction ( $p\leq 0.01$ ). These data may  
86 inform decision making in ARVC families.

87

88

89 **Abbreviations list:**

90 ARVC = arrhythmogenic right ventricular cardiomyopathy; CMR = cardiac magnetic resonance;  
91 ECG = electrocardiogram; ICD = implantable cardioverter defibrillator; LVEF = left ventricular  
92 ejection fraction; NSVT = non-sustained ventricular tachycardia; PVC = premature ventricular  
93 complex; RVEF = right ventricular ejection fraction; TFC = task force criteria; VT = ventricular  
94 tachycardia.

95 **Introduction**

96 Arrhythmogenic right ventricular cardiomyopathy (ARVC) is an inherited cardiomyopathy  
97 characterized by fibrofatty replacement of predominantly the right ventricular myocardium, which  
98 may result in potentially lethal ventricular arrhythmias and progressive ventricular dysfunction  
99 (1). Definite ARVC diagnosis requires an extensive diagnostic work-up as described in the so-  
100 called “Task Force” Criteria (TFC), which were based on an adult population (1-3). Furthermore,  
101 penetrance of ARVC is highly variable and age-dependent (4). Although disease penetrance peaks  
102 during young adulthood, rare but severe cases of pediatric-onset ARVC have been described (5,6).  
103 As such, a diagnosis of pediatric-onset ARVC has important implications, because these  
104 individuals are exposed to radical lifestyle modifications at young age, frequent cardiovascular re-  
105 evaluations to monitor disease progression, and decisions on implantable cardioverter defibrillator  
106 (ICD) implantation (7).

107 Pediatric-onset ARVC has increasingly been recognized during the past decade, and  
108 cascade screening guidelines now recommend to start screening at the age of 10 years (7-10).  
109 Nonetheless, pediatric ARVC patients are still underrepresented in the literature, and diagnostic as  
110 well as management recommendations are based on the adult disease phenotype (3). Likewise, the  
111 yield of cascade screening and predictors of adverse outcome in pediatric subjects remain  
112 unknown.

113 To address these shortcomings, we aim to describe the clinical presentation, phenotypic  
114 characteristics, and outcomes of patients diagnosed with pediatric-onset ARVC. As a secondary  
115 objective, we sought to determine predictors of disease expression and adverse outcome among  
116 pediatric relatives referred for cascade screening.

117

118 **Methods**

119 *Study Population*

120 The patient population was drawn from the Netherlands Arrhythmogenic Cardiomyopathy  
121 Registry ([www.acmregistry.nl](http://www.acmregistry.nl)) (11). This registry is a national, multicenter observational cohort  
122 that prospectively enrolls patients with ARVC and their relatives who are referred for ARVC  
123 cascade screening. For the purpose of this study, we included subjects who consented to our  
124 registry in the University Medical Center Utrecht, Amsterdam University Medical Center, or  
125 University Medical Center Groningen, and were evaluated for ARVC prior to the age of 18 years.  
126 The study was exempt from the Medical Research Involving Human Subjects Act as per judgement  
127 of the Medical Ethics Committee (18-126/C, Utrecht, the Netherlands).

128

129 *Clinical evaluation*

130 Patients were evaluated according the registry protocol (11). Detailed clinical information  
131 regarding demographics, presentation, and symptom onset was obtained for every participant using  
132 medical record review. Phenotypic characterization was performed upon discretion of the treating  
133 physician, and included 12-lead electrocardiograms (ECG), signal averaged electrocardiograms,  
134 exercise testing, Holter monitoring recordings, echocardiography, cardiovascular magnetic  
135 resonance (CMR), electrophysiological study and/or endomyocardial biopsy.

136

137 *Genetic evaluation*

138 Probands were defined as the first affected patient in a family in whom definite ARVC diagnosis  
139 was confirmed. Relatives were defined as family members of a proband with definite ARVC, as  
140 done previously (see **Supplementary Table 1** (5,6,11)). All probands underwent genetic testing

141 for the ARVC-associated genes, while relatives were only tested for the variant identified in their  
142 proband. Pathogenicity of variants was evaluated based on the guidelines developed by the  
143 American College of Medical Genetics and Genomics (12,13). Subjects with an autopsy diagnosis  
144 of ARVC who either did not undergo genetic screening or were obligate carrier were considered  
145 to carry the same variant as identified in their first-degree relatives.

146

#### 147 *Arrhythmogenic right ventricular cardiomyopathy diagnosis*

148 Diagnosis of ARVC was based on the revised 2010 TFC (3). Definite ARVC was defined as either  
149  $\geq 2$  major, 1 major and  $\geq 2$  minor or  $\geq 4$  minor TFC. Pediatric-onset ARVC was defined as a definite  
150 ARVC diagnosis prior to the age of 18 years. Precordial T-wave inversions were excluded from  
151 the TFC for patients prior to the age of 14 years (3,14,15). Age-appropriate criteria were used to  
152 assess late potentials on signal averaged electrocardiogram recordings, as published previously  
153 (16).

154

#### 155 *Follow-up and outcome definitions*

156 Clinical management was performed upon discretion of the treating (pediatric) cardiologist.  
157 Follow-up was determined from the date of first evaluation until the most recent clinical  
158 evaluation. In case of cardiac transplantation or death, the dates at which these occurred were  
159 considered to be the dates of last follow-up.

160 For cascade screening analyses, we evaluated the presence or absence of disease  
161 progression, which was defined as the development of a new diagnostic 2010 TFC and/or  
162 progression from a minor to a major criterion at last follow-up, that was absent at first presentation.



163 Our primary outcome was the occurrence of potentially life-threatening arrhythmic events,  
164 which was defined as a composite of sustained ventricular tachycardia (VT), ventricular  
165 fibrillation, resuscitated sudden cardiac arrest, sudden cardiac death, or appropriate ICD  
166 intervention, as done previously (17). Secondary outcomes included symptomatic heart failure,  
167 cardiac transplantation, and death. Definitions are provided in **Supplementary Table 1**.

168

### 169 *Statistical Analysis*

170 All continuous data were presented as mean  $\pm$  standard deviation or median (interquartile range),  
171 while categorical variables were presented as numbers (percentages). Continuous variables were  
172 compared using the independent Student *t*-test or Mann Whitney *U* test, and categorical data using  
173 Chi-square or Fisher's exact tests, as appropriate. By definition, serial cascade screening  
174 constitutes observed time, which was visually evaluated by Kaplan Meier curves to determine  
175 differences in survival free from disease progression. However, due to heterogeneity in the  
176 performed diagnostic modalities and/or baseline evaluation, comparison using log-rank tests was  
177 considered inappropriate. Data were therefore presented as incidence rates with 95% confidence  
178 intervals. Predictors of ventricular arrhythmias were determined by univariable logistic regression,  
179 which was chosen over Cox regression since many subjects presented with an event (i.e. the  
180 absence of observed time until the arrhythmic outcome limited our ability to perform survival  
181 analysis). A p-value of  $<0.05$  was considered statistically significant. Data was analyzed using  
182 SPSS version 25.0 for Windows (SPSS Inc., Chicago, USA) and GraphPad Prism version 8.3 (La  
183 Jolla, California, USA).

184 **Results**

185 *Study Population*

186 **Figure 1** shows a flowchart of our study population. As of July 1<sup>st</sup>, 2020 the Netherlands  
187 arrhythmogenic cardiomyopathy registry includes a total of 1109 patients, of whom 80 (7%)  
188 patients were referred for genetic and clinical evaluation prior to the age of 18 years in the  
189 predefined enrolling centers (11). Their clinical characteristics at first presentation are presented  
190 in **Table 1**.

191

192 *Characteristics of Proband*s

193 A total of 12 pediatric probands (67% male, 16.8 [13.7-17.1] years old at first evaluation) were  
194 identified (**Figure 1 and Table 1**). One (8%) proband (a 13-year old female) presented with sudden  
195 cardiac death and died upon presentation (**Supplementary Table 2**). Since no clinical evaluation  
196 was available prior to presentation, she was excluded from further analyses. The remaining  
197 pediatric probands most commonly presented with sustained VT (n=7, 58%), followed by  
198 recurrent syncope (n=2, 17%) and resuscitated sudden cardiac arrest (n=2, 17%). Among 11  
199 pediatric probands who underwent genetic testing, 9/11 (82%) carried a pathogenic/likely  
200 pathogenic genetic variant. Variants in *PKP2* (n=5, 42%) were most prevalent, followed by digenic  
201 variants in *PKP2* and *DSG2* (n=2, 17%) and compound heterozygosity (n=2, 17%) (**Figure 2 and**  
202 **Supplementary Table 3**).

203 Clinical evaluation was available in 11 probands who presented alive. An abnormal ECG  
204 (n=10, 91%) was the most frequently observed abnormality, with T-wave inversion in V1-3 among  
205 those >14 years of age as most common finding (n=9, 82%). Almost half (n=5, 46%) of probands  
206 had abnormal cardiac imaging, with a median right ventricular ejection fraction (RVEF) and left

207 ventricular ejection fraction (LVEF) of 33% [25-45%] and 49% [39-56%], respectively. A definite  
208 diagnosis of ARVC was made at presentation in 10/11 (91%) pediatric probands. Detailed  
209 information regarding evaluation and management of probands is described on a case-by-case  
210 basis in **Supplementary Table 4**.

211

### 212 *Characteristics of Relatives*

213 Overall, 68 pediatric relatives (44% male, 13.2 [10.1-15.3] years old at first evaluation) were  
214 identified (**Figure 1 and Table 1**). Two (3%) relatives (a 17-year old male and a 9-year old female)  
215 presented with sudden cardiac death and died upon presentation prior to cascade screening (see  
216 **Supplementary Table 2**). Both cases were asymptomatic prior to their death and were not  
217 recognized as possible ARVC cases during autopsy. Since no clinical evaluation was available  
218 prior to presentation, these subjects were excluded from further analyses. The remaining relatives  
219 were most commonly referred for clinical evaluation because of cascade screening (n=61, 90%),  
220 followed by (near)-syncope (n=2, 3%), palpitations (n=1, 2%), sustained VT (n=1, 2%), and  
221 resuscitated sudden cardiac arrest (n=1, 2%). A total of 19 (29%) relatives did not (yet) participate  
222 in genetic evaluation. Among 47 pediatric relatives who underwent genetic evaluation, 44 (94%)  
223 carried a pathogenic/likely pathogenic genetic variant. Variants in *PKP2* (n=34, 72%) and *PLN*  
224 (n=8, 17%) were most prevalent (see **Figure 2 and Supplementary Table 3**). Differences between  
225 the identified genetic variants of probands and relatives occurred because we included all subjects  
226 who were evaluated prior to the age of 18 years. This included relatives whose respective proband  
227 presented at adult age, and whose proband was excluded for this study.

228 Clinical evaluation was available in 66 relatives who presented alive (**Table 1**). An  
229 abnormal ECG was observed at first evaluation in 8 (12%) relatives. Abnormal imaging was

230 observed in 3 (5%) relatives, with a median RVEF and LVEF of 53% [46-57%] and 60% (57-  
231 64%), respectively. A definite diagnosis of ARVC was made at presentation in 6 (9%) living  
232 relatives. There was no difference in age ( $p=0.11$ ) and sex ( $p=0.90$ ) among relatives with and  
233 without a definite ARVC diagnosis at presentation. Detailed information regarding evaluation and  
234 management of relatives is described on a case-by-case basis in **Supplementary Table 5**.

235

### 236 *Yield of Serial Evaluation in Subjects without Diagnosis at Presentation*

237 Overall, one proband (male, 17 years old, who presented with palpitations) and 60 relatives (46%  
238 male, 12.6 [9.8–15.3] years old) did not fulfill ARVC diagnosis at time of first evaluation.  
239 Phenotypic development during follow-up in these individuals is shown in **Table 2** and **Figure 3**.

240 During 8.5 [5.2–12.4] years of follow-up, disease progression (defined as the presence of  
241 an abnormal TFC at last follow-up, which was absent at enrollment) was observed in 30/61 (49%)  
242 subjects (43% males, median age at first evaluation 13.8 [11.4–16.1] years old). Progression was  
243 most frequently observed on ECG ( $n=14/56$ , 25%), followed by CMR ( $n=12/59$ , 20%), and Holter  
244 monitoring ( $n=11/56$ , 20%). **Figure 4** shows progression on ECG, Holter monitoring and imaging  
245 during follow-up. Time to progression was  $7.3\pm 5.0$  years for ECG/Holter monitoring  
246 abnormalities and  $8.4\pm 5.0$  years for imaging changes (CMR and/or echocardiography). This  
247 corresponded to a 1-year incidence rate of 0.054 (95% confidence interval 0.035-0.081) for  
248 ECG/Holter monitoring, which was almost double the 1-year incidence rate for newly abnormal  
249 imaging (incidence rate 0.029, 95% confidence interval 0.017–0.048). Among ECG and Holter  
250 monitoring criteria, a newly prolonged terminal activation duration ( $n=10/56$ , 18%), premature  
251 ventricular complex (PVC) count  $>500/24\text{hrs}$  ( $n=8/56$ , 14%) and non-sustained ventricular  
252 tachycardia (NSVT,  $n=7/56$ , 13%) were the most prevalent findings. Regarding imaging criteria,

253 new CMR abnormalities (n=12/59, 20%) were more prevalent than new echocardiography criteria  
254 (n=6/59, 10%). None of the tests that were abnormal at presentation reverted to normal during  
255 follow-up.

256 Nineteen (1 proband and 18 relatives) subjects were diagnosed with ARVC during follow-  
257 up (31% of those without diagnosis at presentation). This specific proband was referred for  
258 evaluation of recurrent syncope and palpitations. While he did not meet the 2010 diagnostic TFC  
259 at first evaluation, he experienced disease progression and a definite ARVC diagnosis was made  
260 before 18 years of age. This made him eligible for inclusion as a pediatric proband in this study.  
261 The majority of these individuals were females (n=11, 58%) who reached diagnosis in young  
262 adulthood (median age at diagnosis 19.9 [16.2–22.6] years). There was no significant difference  
263 in duration of follow-up between those with and without definite ARVC diagnosis (median 8.6  
264 [7.6–13.5] years vs. 8.1 [4.6–11.4] years, respectively, p=0.14).

265

### 266 *Clinical Characteristics associated with Arrhythmic Events*

267 The remaining analyses were performed among all 77 patients who presented alive. Overall, 12  
268 (16%) subjects (10 probands and 2 relatives) experienced ventricular arrhythmias after median 1.8  
269 [0.3 – 1.6] years of follow-up, which included 10 probands who had a first event at presentation  
270 but experienced a second event during follow-up (**Figure 3**). Heart failure developed in four (36%)  
271 probands after median 17.7 [6.0 – 24.1] years of follow-up, of whom one underwent cardiac  
272 transplantation (at age 47 years) and two died (at age 21 and 37 years) due to therapy-resistant  
273 congestive heart failure (**Figure 3**). None of the relatives experienced heart failure or death. As  
274 shown in **Figure 3**, first sustained VTs typically occurred during late adolescence (median age  
275 17.9 [17.2–18.8] years), whereas heart failure was only observed during adulthood (median age

276 35.6 [23.1–38.2] years). Overall, 20 (11 probands and 9 relatives) subjects had transvenous ICDs  
277 implanted, all of whom had a definite diagnosis at time of ICD implantation. ICD follow-up was  
278 complicated by lead revision in two patients, which included lead dysfunction in one patient and  
279 upgrade to a two-chamber system because of pacemaker syndrome in the other patient.  
280 Furthermore, one patient had two episodes of inappropriate shocks due to atrial tachycardia and  
281 atrial fibrillation. In addition, 20 patients (11 probands and 9 relatives) received antiarrhythmic  
282 medication: sotalol (n=12), betablockers (n=7) or flecainide (n=1). A total of five probands  
283 underwent VT ablation for recurrent sustained VT.

284 **Table 3** shows the association of clinical characteristics with arrhythmic outcome in  
285 pediatric-onset ARVC. As expected, probands were more likely to experience ventricular  
286 arrhythmia compared to relatives ( $p<0.01$ ). In addition, males had significantly increased risk of  
287 events compared to females ( $p<0.01$ ), while the identification of a pathogenic/likely pathogenic  
288 genetic variant associated with arrhythmic occurrence with borderline significance ( $p=0.06$ ). With  
289 regards to clinical evaluation, T-wave inversions in V1-3 ( $p<0.01$ ), documentation of NSVT  
290 ( $p<0.01$ ) and a higher PVC count on Holter monitoring ( $p=0.01$ ) were associated with ventricular  
291 arrhythmias. In addition, both LVEF ( $p=0.01$ ) and RVEF ( $p<0.01$ ) were lower in subjects with  
292 arrhythmic events.

293 **Discussion**

294 This study reports the clinical characteristics of subjects with pediatric-onset ARVC, describes  
295 testing results during serial evaluation, and determines predictors of adverse clinical outcome. The  
296 main findings of the study are: 1) sudden cardiac death may be the first manifestation of ARVC in  
297 both pediatric probands and relatives; 2) definite ARVC diagnosis before adolescence is rare but  
298 should not be overlooked, as potentially lethal complications may occur; 3) ECG, Holter  
299 monitoring and CMR most frequently identify disease progression during follow-up; 4) T-wave  
300 inversions, frequent PVCs including NSVT, and reduced biventricular ejection fraction are  
301 associated with arrhythmic events.

302

303 *Pediatric-onset ARVC in the literature*

304 Studies evaluating pediatric-onset ARVC are hampered by small sample sizes (**Table 4**).  
305 Regardless, the available data suggests that the majority of pediatric-onset ARVC cases present  
306 during puberty and adolescence, which was also observed in our cohort. Most prior studies focused  
307 on diagnostic evaluation of pediatric ARVC cases, specifically investigating the diagnostic tests  
308 and criteria of the TFC (18-21). We now extend prior findings in one of the largest cohorts of  
309 pediatric-onset ARVC subjects, by evaluating the performance of diagnostic testing during  
310 cascade screening of relatives and focusing on pediatric-specific markers of arrhythmic risk.

311 *Sudden cardiac death as mode of presentation in pediatric-onset ARVC*

312 In our study, sudden cardiac death (n=3) or sudden cardiac arrest (n=3) was the mode of  
313 presentation in 6/80 (8%) of pediatric cases. While this percentage is much higher than in adult  
314 patients with ARVC (22) and in children with dilated cardiomyopathy (23), it is similar to high-  
315 risk pediatric hypertrophic cardiomyopathy cohorts (24). Of note, none of the subjects  
316 experiencing sudden cardiac death or sudden cardiac arrest were medically evaluated before their  
317 event, which suggests the absence of symptoms and/or short interval between symptom onset  
318 and arrhythmic events in these children (**Supplementary Table 2**) (25). While these events will  
319 therefore be difficult to prevent in probands, vigilance in relatives should be high. Indeed, we  
320 observed one relative with sudden cardiac arrest without prior participation in cascade screening.  
321 Importantly, none of the subjects experienced sudden cardiac death during follow-up after  
322 medical evaluation. This suggest that proper management by monitoring of disease progression,  
323 restriction of participation in sports, and ICD implantation may indeed prevent potentially lethal  
324 complications.

325

326 *Penetrance of disease before and during adolescence*

327 The majority of definite ARVC diagnoses occurred during adolescence or young adulthood, while  
328 the youngest proband with definite ARVC diagnosis was 9 years old. This age distribution is  
329 similar to observations in prior studies (**Table 4**) (8,10,18,19). In this context, it is important to  
330 note that the diagnostic TFC for ARVC were derived from a predominantly adult population and  
331 lack both sensitive and specific criteria for pediatric-onset of disease (3). For example, the cutoffs  
332 used for RV dilatation and dysfunction are likely to overdiagnose disease in pediatric patients (26).  
333 Development of pediatric-specific criteria may enhance the diagnostic performance of the TFC in



334 general and of specific modalities in particular, which however likely requires a large dataset  
335 derived from multiple longitudinal ARVC cohorts (17,19-22). Furthermore, future studies may  
336 elucidate the relation between environmental factors such as exercise and (epi)-genetic modifiers  
337 associated with early disease manifestation of ARVC (10,27).

338

### 339 *Diagnostic testing during cascade screening*

340 Our study showed that approximately one in three pediatric relatives develop definite ARVC  
341 during follow-up. It is important to note that the presence of multiple pathogenic variants plays a  
342 pivotal role in the early expression of an ARVC phenotype among pediatric probands and relatives,  
343 as has been suggested previously (10). This is also in line with our findings, which underline the  
344 role of multiple genetic variants influencing early expression of disease in very young ARVC  
345 patients. Of note, our study supports the current recommendation to start cascade screening at the  
346 age of 10 years old, since the youngest relative fulfilling ARVC diagnosis in our study was 11  
347 years old. Furthermore, our findings confirm the suggested approach to perform periodic screening  
348 with ECG, Holter recording, exercise testing and cardiac imaging (7). However, the guidelines do  
349 not frame recommendations on the optimal screening interval and best combination of tests during  
350 cascade screening.

351 Our study suggests that ARVC is a slowly progressive disease, and that ECG criteria, a  
352 higher PVC count during Holter recording, and CMR imaging criteria are associated with disease  
353 progression during follow-up. This may aid the development of follow-up recommendations in  
354 this overwhelmingly young population. Although ECG parameters were important for diagnosis,  
355 only one case of epsilon waves was confirmed in this cohort. This may be explained by both 1)  
356 variability in the identification of epsilon waves in general; and 2) the rather significant amount of

357 activation delay which should be present for an epsilon wave to appear on the ECG, which is likely  
358 to be absent in our young pediatric population (28). In addition to the parameters tested in this  
359 study, newer techniques can aid early disease detection. One of those is echocardiographic  
360 deformation imaging, which was shown to have diagnostic and prognostic potential during cascade  
361 screening of adult patients. Future studies should evaluate the role of echocardiographic  
362 deformation imaging during cascade screening of pediatric patients, define criteria to aid ARVC  
363 diagnosis in children, and define pediatric-specific criteria for disease progression (29-31).

364

#### 365 *Predictors of ventricular arrhythmia*

366 It has been suggested that arrhythmogenicity in ARVC depends on several coexisting mechanisms:  
367 triggered activity related to sympathetic activity, reentrant mechanisms due to myocardial fibrosis,  
368 myocardial inflammation and/or ion channel dysfunction (32,33). These factors may explain the  
369 observation of both fast, unstable rhythms such as polymorphic VT or ventricular fibrillation, and  
370 hemodynamically stable monomorphic VT in ARVC patients. In our cohort, T-wave inversion in  
371 the right precordial leads, NSVT, higher PVC count and reduced biventricular ejection fraction  
372 were associated with ventricular arrhythmias, similar to risk factors recently described in a risk  
373 prediction model for adult ARVC patients ([www.arvcrisk.com](http://www.arvcrisk.com)) (17,22). Based on our results, risk  
374 stratification of pediatric patients evaluated for ARVC should be based on a minimal clinical  
375 evaluation including an ECG, Holter monitoring, exercise testing and imaging with either  
376 echocardiography and/or CMR. Future studies with a larger sample size should validate this risk  
377 prediction model in pediatric subjects before widespread use can be advocated in these young  
378 patients.

379

380 *Limitations*

381 The retrospective nature of this study and the small sample size may compromise generalizability  
382 of our results. Ascertainment bias may have occurred due to referral to our tertiary cardiogenetic  
383 clinics. The age cutoff of <18 years to define pediatric-onset of disease may be considered  
384 arbitrary. Although this age cutoff is frequently used in (Dutch) clinical practice to distinguish  
385 pediatric from adult cases, disease progression and severity should be assessed on a continuum.  
386 Participation in endurance exercise has long been suggested as risk factor for early ARVC  
387 occurrence; however, reliable exercise data was lacking in this cohort. Furthermore, based on the  
388 best available evidence, the 2010 TFC were used to define ARVC diagnosis. However, these  
389 criteria have not been validated in pediatric subjects. Since family history and genetic screening  
390 results play an important role in the 2010 TFC for relatives, a less severe phenotype may be  
391 observed in relatives who fulfill diagnostic TFC.

392

393 **Conclusion**

394 To the best of our knowledge, this is the largest study describing the clinical characteristics and  
395 outcomes of pediatric ARVC cases. Pediatric-onset ARVC is particularly recognized during  
396 adolescence, and is characterized by frequent ventricular arrhythmias in probands. However,  
397 definite ARVC diagnosis before adolescence should not be overlooked, because sudden cardiac  
398 death may be the first manifestation of disease in both probands and relatives. Disease progression  
399 during cascade screening is most frequently identified by ECG, Holter monitoring and CMR.  
400 Arrhythmic events are associated with male sex, the number of T-wave inversions in the precordial  
401 leads, ventricular ectopy on Holter monitoring, and reduced biventricular ejection fraction on  
402 cardiac imaging.

403 **Clinical competencies**

404 ARVC can also be a disease of adolescents, and is characterized by frequent ventricular  
405 arrhythmias particularly in probands. Disease in relatives is most commonly identified by cascade  
406 screening and has a more benign clinical course. Cascade screening and arrhythmic risk prediction  
407 should focus on ECG, Holter monitoring and CMR imaging.

408

409 **Translational outlook**

410 Future studies should establish pediatric-specific diagnostic criteria for ARVC, validate the  
411 performance of a prognostic risk prediction model (arvcrisk.com) in pediatric subjects, and address  
412 which (epi)genetic and environmental modifiers influence early disease manifestation in ARVC.

413

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- 529

530 **Figure legends**

531 **Central Illustration. Cascade Screening and Follow-up in Pediatric-onset ARVC**

532 The results of follow-up in probands and cascade screening in at-risk relatives are shown in the  
533 upper panel. Factors associated with sustained VT and disease progression during follow-up are  
534 shown in the lower panels. Abbreviations as in manuscript.

535

536 **Figure 1. Flowchart of the Study Population**

537 Flowchart of the included cohort from our ARVC registry, the selection of pediatric population,  
538 and clinical course of patients included in the study. Symptomatic was defined as (near)-syncope  
539 or palpitations. \* = population stratified to enrolled centers as described in Methods: University  
540 Medical Center Utrecht, Amsterdam University Medical Center, or University Medical Center  
541 Groningen. \*\* = Cause of death was therapy-resistant congestive heart failure after a long history  
542 of VTs and heart failure. Abbreviations: Dx= definite ARVC diagnosis; HF= congestive heart  
543 failure; other abbreviations as in manuscript.

544

545 **Figure 2. Molecular Genetic Analysis in the Study Population**

546 Pathogenic variants identified during molecular genetic testing, stratified by probands and relatives  
547 and presented as percentages of tested individuals. Abbreviations: *PKP2*= Plakophilin-2; *DSG2*=  
548 Desmoglein-2; *PLN*= Phospholamban; *TTN*= Titin; Mult. PV.= multiple pathogenic variants. See  
549 **Supplementary Table 3** for a detailed summary of the identified variants.

550

551 **Figure 3. Clinical Course and Follow-up among Patients Presenting Alive**

552 Clinical course and follow up stratified by probands, relatives with definite ARVC diagnosis and  
553 relatives without definite ARVC diagnosis. Abbreviations: HF = congestive heart failure; other  
554 abbreviations as in manuscript.

555

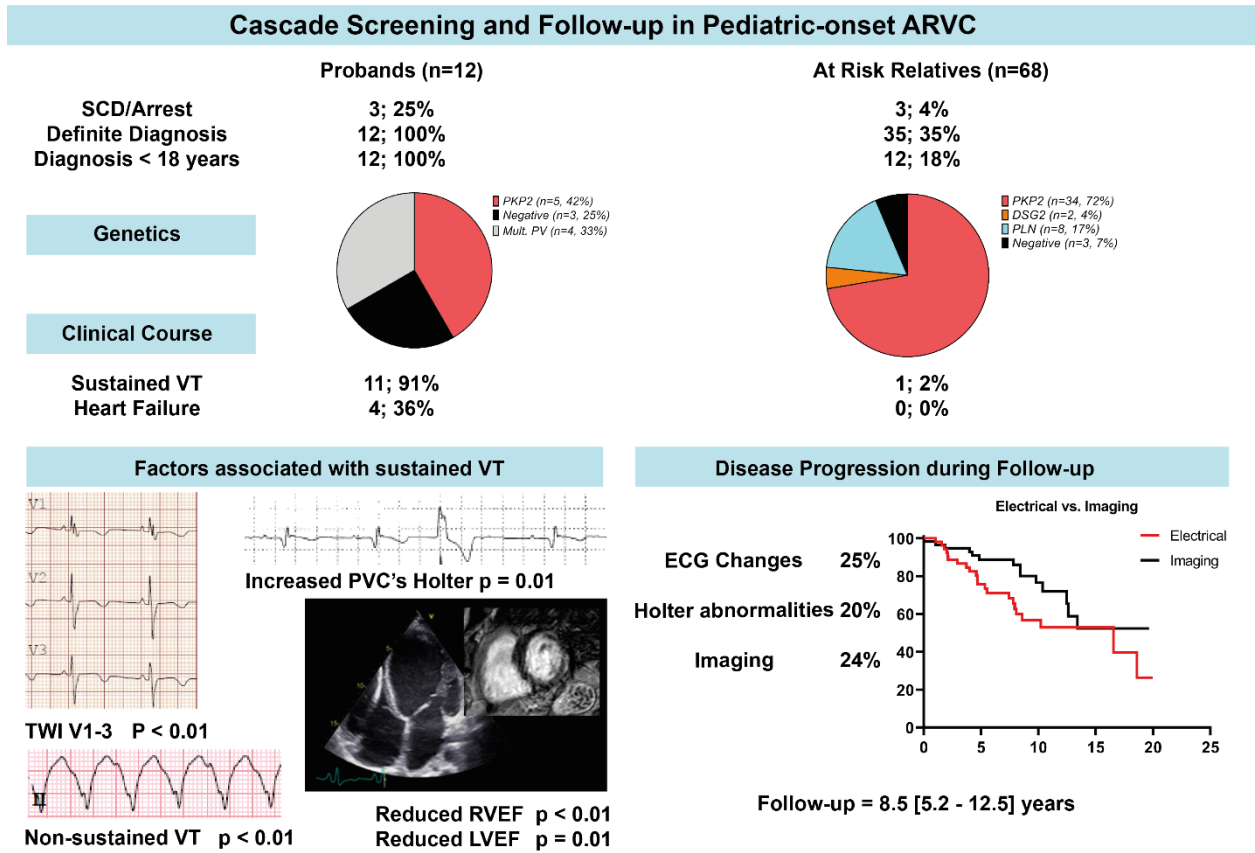
556 **Figure 4. Serial Evaluation of Patients without Definite Diagnosis at Presentation.**

557 Panels A-C show Kaplan Meier curves per diagnostic modality depicting survival free from a new  
558 diagnostic criterion during follow-up, which was absent at presentation. Per diagnostic test, only  
559 patients with a normal evaluation of that specific modality at presentation were included in the  
560 analysis. A 1-year incidence rate with 95% confidence interval was presented per diagnostic  
561 modality. (A) New “electrical” (ECG, Holter monitoring) criteria versus new “imaging” (CMR  
562 and echocardiogram) criteria. (B) “Electrical” criteria: new ECG versus new Holter monitoring  
563 criteria. The difference between the number of included ECG and Holter monitoring cases is  
564 explained by including only cases with normal evaluation of that specific test at baseline. (C)  
565 “Imaging” criteria: new CMR versus new echocardiographic criteria. Abbreviations: CI =  
566 confidence interval; IR = incidence rate.

567

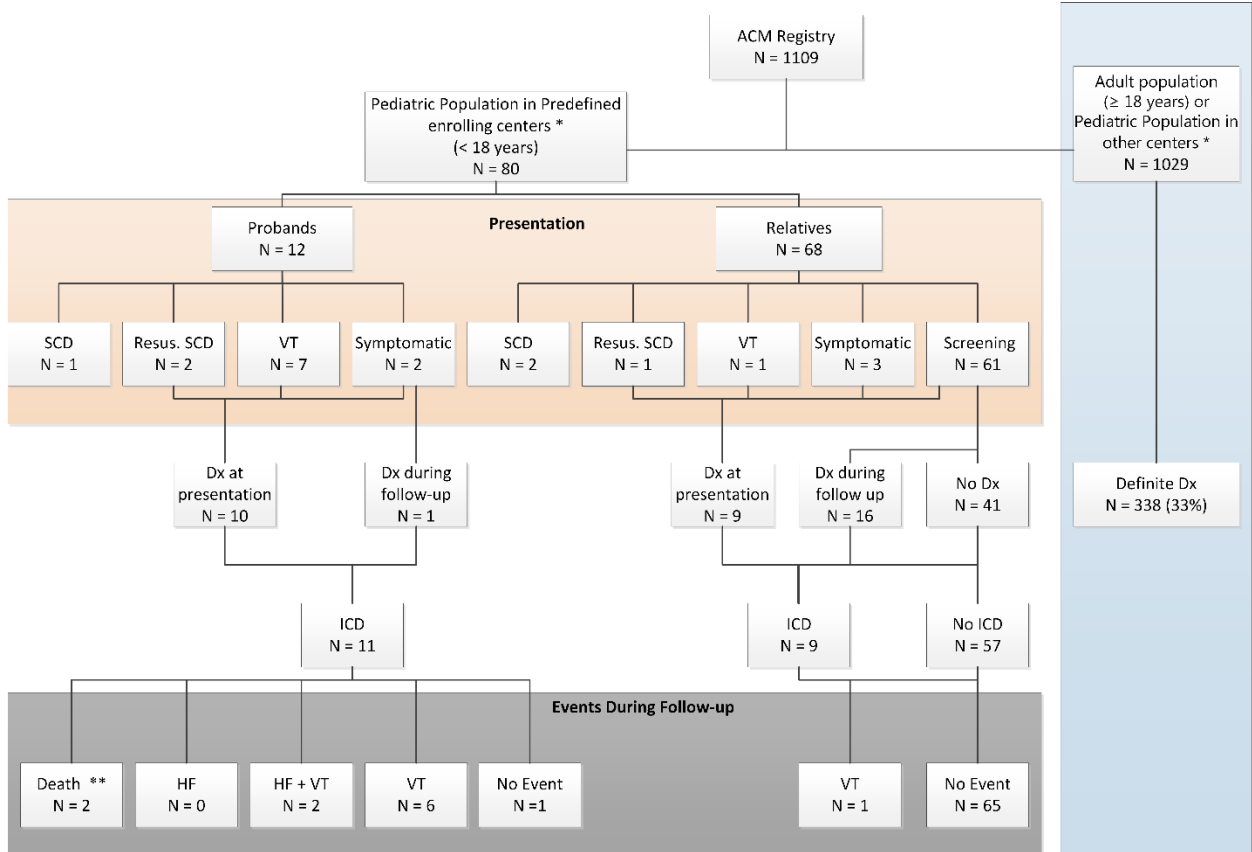
568 **Figures**

569 **Central Illustration**



570

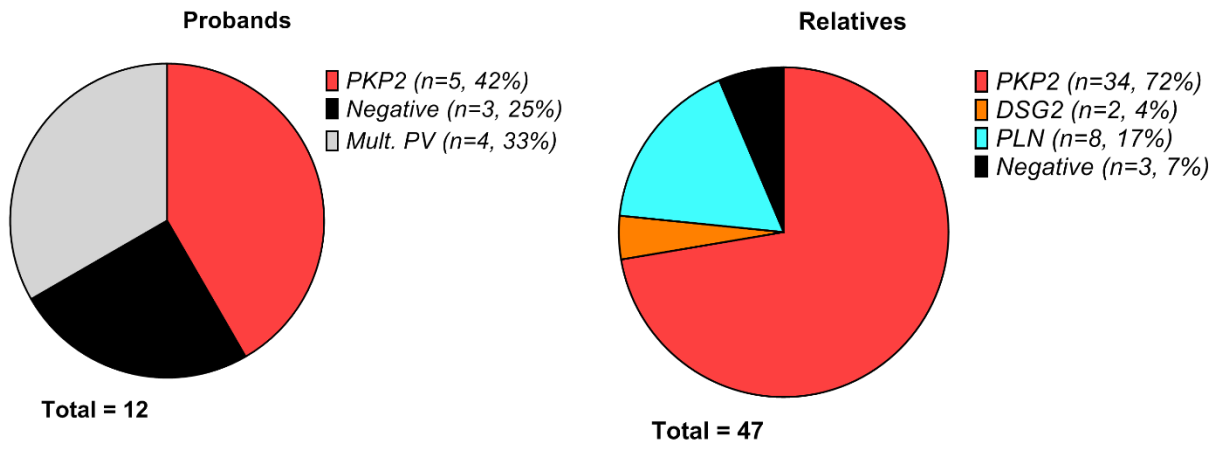
571 **Figure 1.**



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573

574 **Figure 2.**

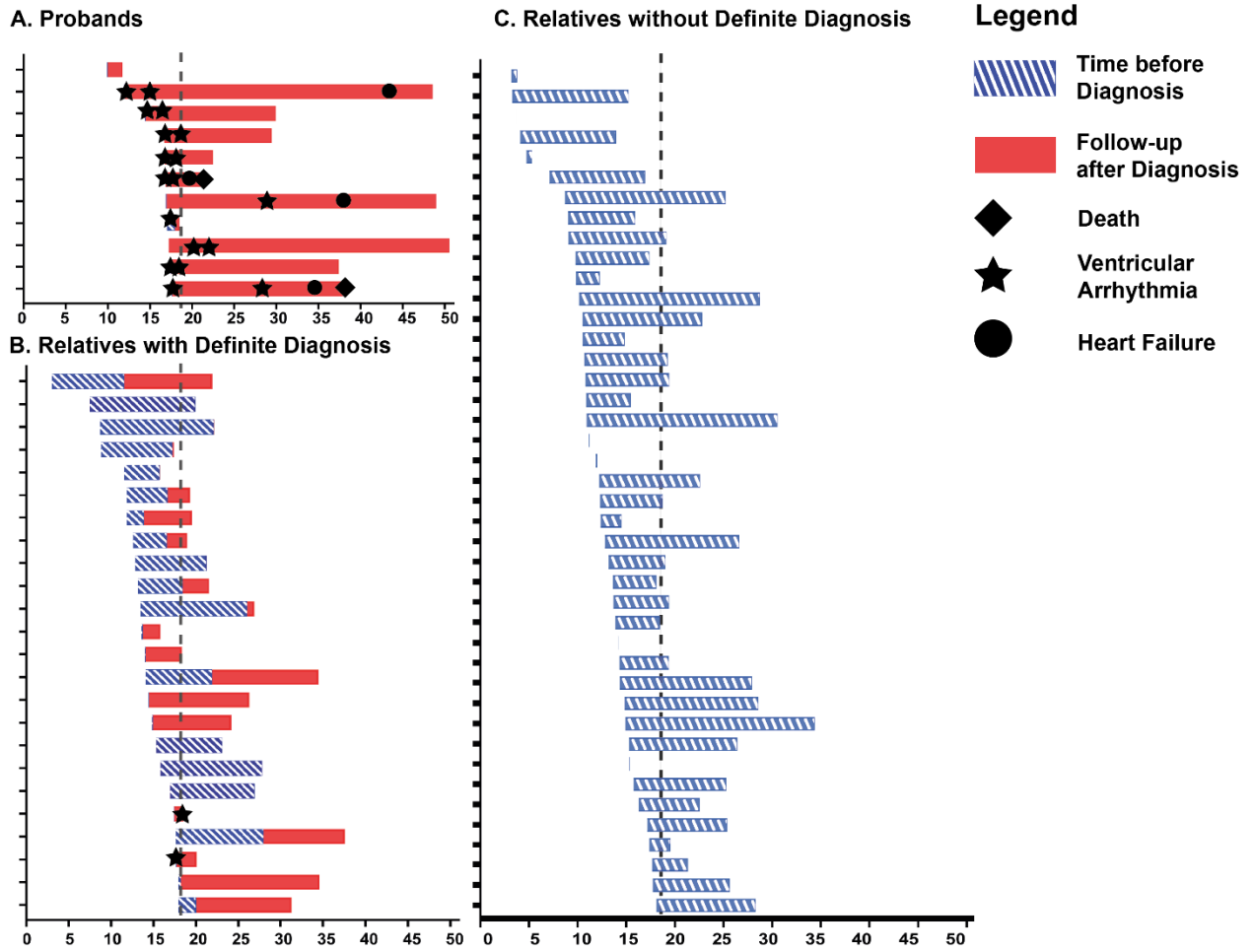


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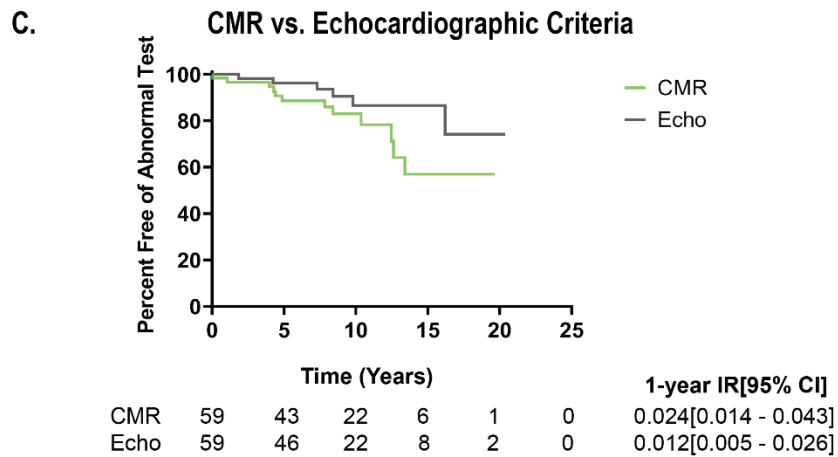
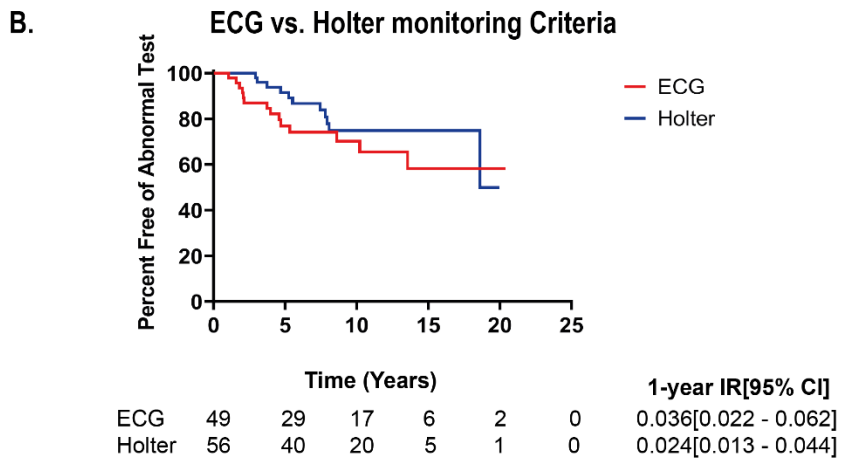
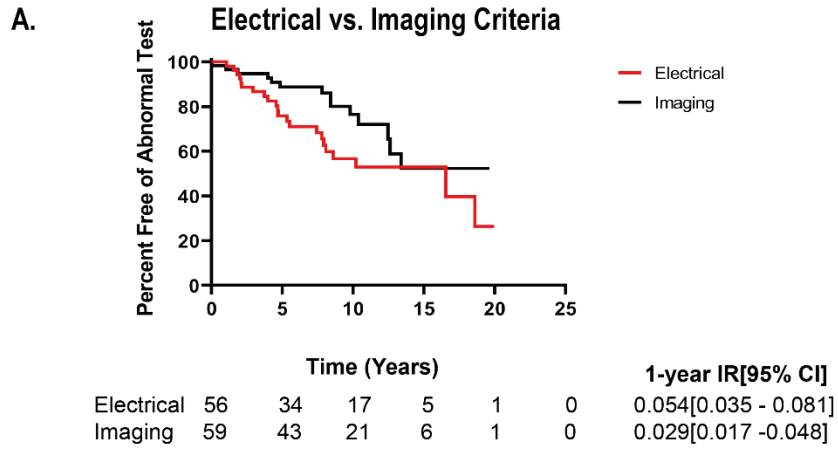
576

577 **Figure 3.**

578



580 **Figure 4.**



581

582



584 Table 1. Clinical characteristics at presentation stratified by probands and relatives.

	Overall	Probands	Relatives
<b>Demographics of all patients</b>	<b>n=80</b>	<b>n=12</b>	<b>n=68</b>
Male	38 (48)	8 (67)	30 (44)
Age at presentation	13.5[10.5 - 16.7]	16.8[13.7 - 17.1]	13.2[10.1 - 15.3]
<b>Type of presentation</b>			
Ventricular tachycardia	8 (10)	7 (58)	1 (2)
Resuscitated sudden cardiac arrest	3 (4)	2 (17)	1 (2)
Sudden cardiac death	3 (4)	1 (8)	2 (3)
Symptomatic	5 (6)	2 (17)	3 (4)
Cascade screening	61 (76)	0	61 (90)
<b>Molecular Genetic Analysis</b>			
(Likely)Pathogenic variant *	53/58 (91)	9/11 (82)	44/47 (94)
<b>Clinical evaluation in alive patients</b>	<b>n=77</b>	<b>n=11</b>	<b>n=66</b>
<b>Electrocardiogram</b>			
T-wave inversion V1-V3	11 (14)	9 (82)	2 (3)
T-wave inversion V1-V2	5 (7)	1 (8)	4 (6)
T-wave inversion V4-V6	1 (1)	0	1 (2)
Epsilon wave	1 (1)	0	1 (2)
Prolonged terminal activation duration	10 (13)	2 (18)	8 (12)
Any 12-lead ECG abnormality	18 (23)	10 (91)	8 (12)
<b>Electrophysiology</b>			
Late potentials on signal averaged electrocardiogram	2/6 (33)	0	2/6 (33)
>500 PVCs/24h	7/48 (15)	2/4 (50)	5/44 (11)
Number of PVCs/24h	0 [0 - 58]	890 [23 - 11933]	0 [0 - 34]
NSVT	10/74 (14)	6/10 (60)	4/64 (6)
LBBB VT superior axis	3 (4)	2 (18)	1 (2)
LBBB VT other axis	7 (9)	6 (55)	1 (2)
<b>Imaging</b>			
RVEF (%)	52 [44 - 55]	33 [25 - 45]	53 [46 - 57]
LVEF (%)	59 [56 - 64]	49 [39 - 56]	60 [57 - 64]
Late gadolinium enhancement on CMR †	7/22 (32)	3/4 (75)	4/18 (22)
Any imaging test fulfilling TFC	8 (10)	5 (46)	3 (5)

<b>Diagnosis</b>			
Definite diagnosis at presentation	16/67 (24)	10/11 (91)	6/66 (9)
Median TFC score at presentation	2 [2 - 3]	6 [5 - 7]	2 [2 - 2]

585

586 Values are n (%), mean  $\pm$  SD or median [interquartile range], as appropriate. Abbreviations as in  
587 manuscript. \* class 4 or 5 likely pathogenic and pathogenic variants (12), see **Figure 2** and  
588 **Supplementary Table 3** for a detailed description. † all right ventricular associated late  
589 gadolinium enhancement in probands, one case of left ventricular associated and three cases of  
590 right ventricular associated late gadolinium enhancement in relatives. Abbreviations: LBBB = left  
591 bundle branch block, rest as in manuscript.

592

593 **Table 2. Serial Evaluation among Subjects without Definite Diagnosis at Presentation.**

	Overall (n=61)
<b>Clinical evaluation during follow-up</b>	
<b>ECG progression*</b>	<b>14/56 (25)</b>
T-Wave Inversion V1-3	5 (9)
T-Wave Inversion V1-2	4 (7)
T-Wave Inversion V4-6	1 (2)
Prolonged terminal activation duration	10 (18)
<b>Holter monitoring progression*</b>	<b>11/56 (20)</b>
>500PVC/24h	8 (14)
Number of PVC/24h	2 [0-191]
NSVT	7 (13)
<b>Echocardiography progression*</b>	<b>6/59 (10)</b>
Major echocardiographic criterion	4 (7)
Minor echocardiographic criterion	2 (3)
<b>CMR progression*</b>	<b>12/59 (20)</b>
Major CMR criterion	11 (19)
Minor CMR criterion	1 (2)
<b>Disease progression on any test*</b>	<b>30/61 (49)</b>
<b>Events during follow-up</b>	
Ventricular tachycardia	1 (2)
Heart Failure	0 (0)
Death	0 (0)
Cardiac transplantation	0 (0)

594

595 Values are n (%), mean ± SD or median [interquartile range]. \* = Progression is defined as the  
 596 occurrence of a new diagnostic TFC criterion during follow-up which was absent at first  
 597 presentation. Abbreviations as in manuscript.

598

599

600 **Table 3. Characteristics of patients presenting alive, stratified by any VT event**

	<b>VT (n=12)</b>	<b>No VT (n=65)</b>	<b>p-value</b>
<b>Demographics</b>			
Proband status	10 (83)	1 (2)	<0.01
Male sex	10 (83)	27 (42)	<0.01
Pathogenic genetic variant	9/12 (75)	42/44 (96)	0.06
<b>ECG and Holter monitoring</b>			
T-wave inversion in V1-3	9 (75)	2 (3)	<0.01
Number of PVC/24h	3241 [413 – 6892]	3 [1 – 285]	0.01
NSVT	5/11 (46)	5/63 (8)	<0.01
<b>Imaging</b>			
LVEF	54 [49 – 59]	60 [57 – 64]	0.01
RVEF	32 [24 – 33]	53 [47 – 57]	<0.01
Late gadolinium enhancement on CMR	3/4 (75)	4/18 (22)	0.09

601

602 Values are n (%), mean ± SD or median [interquartile range]. Abbreviations as in manuscript.

603 **Table 4. Literature review of pediatric-onset ARVC**

First Author	Year	Definite ARVC cases	Mean age (yrs)	Sex (% males)	Evaluation	Findings	Ref.
Daliento	1995	17	14.9	59	Natural history	Based on 1994 TFC. High incidence sudden cardiac death.	(34)
Bauce	2011	14	15.0	79	Genotype-phenotype correlation	2010 TFC were superior over 1994 diagnostic criteria. High proportion of pediatric pathogenic variant carriers reached ARVC diagnosis.	(19)
Te Riele	2015	75	15.3	55	Natural history	Similar course of disease between adult- and pediatric-onset ARVC.	(8)
Etoom	2015	23	11.8	74	Diagnostic evaluation of CMR	Poor agreement between CMR and echocardiography 2010 TFC criteria.	(20)
Despande	2016	16	12.6	63	Histopathology-phenotype correlation	Only half of ARVC patients met histopathology TFC criterion.	(35)
Chungsom prasang	2017	24	15.2	56	Diagnostic evaluation of CMR	Association between right ventricular remodelling and left ventricular involvement.	(21)
Steinmetz	2018	12	14.0	66	Diagnostic evaluation of 2010 TFC	Genetic testing, CMR and ECG contributed to pediatric-onset ARVC diagnosis.	(18)
Sreetharan	2018	26	12.6	63	Histopathology-phenotype correlation	Poor agreement between phenotype and endomyocardial biopsy criteria.	(36)
Martins	2018	6	9.2	83	Diagnostic evaluation of CMR	Suggesting a role of myocarditis in pediatric-onset disease.	(37)
DeWitt	2019	32	15	56	Genotype-phenotype correlation	Association between specific pathogenic variants and right-dominant, left-dominant or biventricular disease.	(10)
Slesnink	2019	3	12	100	Histopathology-phenotype correlation	Correlation between biventricular fibrofatty replacement and CMR findings.	(38)
Pieles	2019	38	15	76	Diagnostic evaluation of echocardiography	Echocardiographic criteria, indexed to body surface area, did not differentiate between ARVC and controls. Reduced RV strain was associated with ARVC diagnosis.	(39)

