







Family screening for hypertrophic cardiomyopathy: Is it time to change practice guidelines?

Myriam Lafreniere-Roula¹, Yoav Bolkier ², Laura Zahavich², Jacob Mathew³, Kristen George², Judith Wilson ², Elizabeth A. Stephenson², Leland N. Benson ², Cedric Manlhiot¹, and Seema Mital ^{2*}

¹Department of Surgery, Cardiovascular Data Management Centre, Hospital for Sick Children, 555 University Avenue, Toronto, Ontario M5G 1X8, Canada; ²Department of Pediatrics, Hospital for Sick Children, University of Toronto, 555 University Avenue, Toronto, Ontario M5G 1X8, Canada; and ³Department of Pediatrics, The Royal Children's Hospital, 50 Flemington Road, Parkville, Victoria 3052, Australia

Received 29 March 2019; revised 23 April 2019; editorial decision 22 May 2019; accepted 24 May 2019; online publish-ahead-of-print 6 June 2019

See page 3682 for the editorial comment on this article (doi: 10.1093/eurheartj/ehz487)

Aims

Current guidelines recommend initiating family screening for hypertrophic cardiomyopathy (HCM) after age 10 or 12 years unless early screening criteria are met. The aim was to evaluate if current screening guidelines miss early onset disease.

Methods and results

Children who underwent family screening for HCM before age 18 years were analysed. Major cardiac events (MaCEs) were defined as death, sudden cardiac death (SCD), or need for major cardiac interventions (myectomy, implantable cardioverter-defibrillator insertion, transplantation). Of 524 children screened, 331 were under 10 years of age, 9.9% had echocardiographic evidence of HCM, and 1.1% were symptomatic at first screening. The median (interquartile range) age at HCM onset was 8.9 (4.7–13.4) years, and at MaCE was 10.9 (8.5–14.3) years with a median time to MaCE from HCM onset of 1.5 (0.5–4.1) years. About 52.5% phenotype-positive children and 41% with MaCEs were <10 years old. Only 69% children with early HCM met early screening criteria. Cox regression identified male gender, family history of SCD, and pathogenic variants in *MYH7*/*MYBPC3* as a predictor of early onset HCM and MaCEs.

Conclusion

A third of children not eligible for early screening by current guidelines had phenotype-positive HCM. *MYH7* and *MYBPC3* mutation-positive patients were at highest risk for developing early HCM and experiencing an event or requiring a major intervention. Our findings suggest that younger family members should be considered for early clinical and genetic screening to identify the subset in need of closer monitoring and interventions.

Keywords

Hypertrophic cardiomyopathy • Family screening • Sudden cardiac death • Implantable cardioverter-defibrillator • Sarcomeric mutations

Introduction

The American College of Cardiology (ACC)/American Heart Association (AHA) consensus guidelines recommend 12 years as the starting age for family screening of first-degree relatives of affected probands with hypertrophic cardiomyopathy (HCM). Earlier

screening is recommended only in cases with an early growth spurt, family history of sudden cardiac death (SCD), and prior to competitive sports participation [Class I recommendation, level of evidence (LOE)-C].¹ The European Society of Cardiology (ESC) recommends that clinical and/or genetic screening be offered from age 10 years onwards (Class IIa, LOE-C) with earlier screening to be considered in

* Corresponding author. Tel: +1 416 813 7418, Fax: +1 416 813 5857, Email: seema.mital@sickkids.ca

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families with malignant early onset disease, presence of cardiac symptoms or are involvement in demanding physical activity (Class IIb, LOE-C).^{2–4} Furthermore, relying on symptoms to initiate screening is challenging in children who are often unable to verbalize symptoms like angina, palpitations, or exercise intolerance. Setting 10–12 years as the minimum age cut-off for screening involves a risk of missing earlier onset disease. The primary goal of this study was to evaluate if the current screening guidelines miss early onset disease which can impact timely interventions aimed at preventing adverse outcomes.

Methods

This was a single centre, retrospective cohort study of children who underwent echocardiography prior to 18 years of age as part of family screening for primary HCM between 2000 and 2018. As a tertiary care referral centre, families from across the province were followed at our institution. First-degree relatives of HCM probands were screened clinically independent of age, although echocardiography was sometimes delayed till 3 years of age to avoid sedation for echocardiography. Screening included a clinic visit, electrocardiogram (ECG), echocardiogram, and genetic counselling for the purpose of cascade genetic testing. Hypertrophic cardiomyopathy probands and families with secondary HCM e.g. syndromic, neuromuscular, metabolic, or mitochondrial disease were excluded. The study was approved by the Institutional Research Ethics Board of the Hospital for Sick Children and the requirement for informed consent was waived.

The medical records were reviewed for demographics, family history, baseline clinical characteristics, symptoms, serial echocardiographic data, Holter results, genetic test results, and clinical outcomes. Phenotype-positive status was defined as maximal left ventricular (LV) posterior wall diameter or interventricular septal (IVS) diameter z-score >2 on two-dimensional echocardiography. Major cardiac events (MaCEs) were defined as a composite of death, SCD, resuscitated SCD due to ventricular fibrillation or ventricular tachycardia, or need for major cardiac interventions including surgical myectomy, implantable cardioverter-defibrillator (ICD) implantation, or cardiac transplantation.

Statistical analysis

Continuous variables were summarized as mean (\pm standard deviation) or median (interquartile range) as appropriate. Dichotomous and polytomous variables were summarized as frequencies and proportions. Children were presumed phenotype-negative between birth and the first record of them being phenotype-positive. Age-dependent freedom from phenotype-positive status and from a MaCE was evaluated using Kaplan–Meier estimates. Incidence rate of outcomes was calculated as the ratio of events per 100 patient-years of follow-up. We assessed the proportion of children who became phenotype-positive and/or experienced a MaCE who did not meet early screening criteria of symptoms or family history of SCD. Univariable Cox regression models were used to identify factors associated with HCM at <10 years and MaCEs. Genetic risk factors were dichotomized to by the presence or absence of a pathogenic or likely pathogenic (P/LP) variant, variant in a sarcomeric gene, and multiple P/LP variants. Factors significant in the multivariable model were retained and forward selection was applied to the other candidate factors significant at the univariable level. Forward selection of additional factors was based on minimizing Akaike's information criteria (AIC). Models with clinical factors only were compared to models with clinical and genetic factors using the AIC, whereby a lower AIC indicates a better model. To quantify discriminative ability, the concordance statistic for Cox proportional hazard models was calculated.⁵

Table 1 Baseline characteristics (n = 524)

Variables	N	Statistics
Demographic characteristics		
Age at first echo (years)	524	7.7 (3.7–12.0)
Gender	524	
Female		235 (44.8%)
Reported race	157	
Afro-Caribbean		6 (3.8%)
Caucasian		131 (83.4%)
Mixed		7 (4.5%)
Other		13 (8.3%)
Phenotype-positive at first screening	524	52 (9.9%)
Symptomatic at screening ^a	523	6 (1.1%)
Screened before 12 years old	524	392 (74.8%)
Phenotype-positive at last follow-up	524	80 (15.3%)
Family history		
Family history of sudden cardiac death	524	198 (37.8%)
Family history of ICD	524	150 (28.6%)
Youngest family member with SCD	198	
>60 years old		11 (5.6%)
40–59 years old		52 (26.3%)
25–39 years old		74 (37.4%)
<25 years old		61 (30.8%)
Closest affected family member	522	
First degree (mother/father/sibling)		468 (89.7%)
Second degree (grand-parent, uncle/aunt)		47 (9.0%)
Third or fourth degree		7 (1.3%)

^aSymptoms included: palpitations (1), chest pain (1), murmur (2), syncope (1), syncope, and palpitations (1).

Results

Baseline characteristics

A total of 524 relatives (from 315 families) <18 years old at the time of screening echocardiography were eligible. The median number of at-risk children screened per family was 2 (range 1–6) (those screened due to a murmur were excluded). In total, 331 (63.2%) were screened before 10 years of age. *Table 1* describes the baseline characteristics of the cohort. Of these, 52 (9.9%) were phenotype-positive at first evaluation and an additional 28 (5.4%) became phenotype-positive during follow-up. About 198 (37.8%) had a positive family history of SCD. Only 6 (1.1%) children were symptomatic at first screening. Genetic characteristics are described in *Table 2*. Overall 215 families were genotype-positive. Of these, 120 children from these families had clinical genetic testing of which 86 (71.7%) were genotype-positive (95 families declined genetic testing). Variants in *MYH7* and *MYBPC3* accounted for over 60% of genotype-positive cases.

Outcomes

Hypertrophic cardiomyopathy onset

The median age at onset of HCM was 8.9 (4.7–13.4) years (*Table 3*). Freedom from phenotype-positive HCM at 10 years of age was

Table 2 Genetic characteristics (n = 524)

Variables	N	Statistics
Paediatric relative with genetic testing		
Subject had genetic testing	524	120 (22.9%)
Variant identified	120	
No		28 (23.3%)
Pathogenic		86 (71.7%)
Variant of uncertain significance		6 (5.0%)
Affected gene		
MYH7	86	27 (31.4%)
MYBPC3	86	43 (50.0%)
PRKAG2 + MYBPC3	86	1 (1.2%)
TPM1	86	8 (9.3%)
TNNI/TNNT	86	3 (3.5%)
MYL2	86	2 (2.4%)
TMPO	86	1 (1.2%)
Unknown/Missing	86	1 (1.2%)
Proband with genetic testing		
Variant identified in the proband		
Pathogenic		95 (23.5%)
Variant of uncertain significance		30 (7.4%)
Affected gene		
Unknown	95	15 (15.8%)
MYH7	95	27 (28.4%)
MYBPC3	95	41 (43.1%)
TNN13	95	5 (5.3%)
TPM1	95	4 (4.2%)
TNNT2	95	2 (2.1%)
PRKAG2	95	1 (1.1%)

91.0% (88.0–93.8) (Figure 1) and freedom from moderate-severe hypertrophy defined as IVS z-score >5 was 95.2% (92.8–96.8). Of the 80 phenotype-positive children, 42 (52.5%) became phenotype-positive before 10 years of age (Figure 2A).

Major cardiac events

Seventeen (3.2%) children experienced a MaCE during follow-up (Table 4). Seven of 17 (41%) children experienced it before 10 years of age (Figure 2B). Freedom from a MaCE at 10 years of age was 98.4% (96.7–99.2) in the overall cohort. The median age at MaCE was 10.9 (8.5–14.3) years. There were a total of 21 events in 17 children. The first event included three deaths (two from heart failure and one SCD), two aborted SCD, five surgical myectomies, and seven ICDs for primary prevention (including two with appropriate shocks) (Figure 2C). With 2206 patient-years of follow-up, incidence rate of MaCE was 0.95 per 100 patient-years. All events occurred in phenotype-positive children. The median follow-up from first echocardiographic evaluation to MaCE was 2.8 (0.5–4.5) years, and from onset of HCM to MaCE was 1.5 (0.5–4.1) years. The cumulative proportion of MaCE in phenotype-positive children is shown in Figure 3A.

Table 3 Outcomes (frequency and incidence rate) (n = 524)

Variables	Statistics** (incidence rate)
Age at last echocardiogram (years)	13.4 (8.6–16.6)
Length of follow-up (years)	3.2 (0.0–6.4)
Maximal LV wall or IVS z-score >2	80 (3.62)
Primary outcome—earliest event	
Major cardiac event	17 (0.77)
Death	3 (0.14)
Heart transplant	0 (0.00)
Aborted sudden cardiac death	2 (0.09)
ICD implantation (all primary indication)	7 (0.32)
Myectomy	5 (0.23)
Other outcomes	
Abnormal blood pressure response on exercise test	16 (0.73)
Non-sustained ventricular tachycardia on Holter	7 (0.32)
Coronary unroofing	3 (0.14)
Maximal LV wall or septal Z-score >5	49 (2.22)
Maximal LV wall or septal Z-score >10	13 (0.59)
LV outflow gradient >30 mmHg	13 (0.59)
LV outflow gradient >50 mmHg	10 (0.45)
LV outflow gradient >100 mmHg	9 (0.41)

Major cardiac events in early onset hypertrophic cardiomyopathy

Thirteen of 17 (76.5%) children with a MaCE had early onset HCM. The cumulative proportion (95% confidence interval) of MaCEs at 5 years after HCM onset was 28.9% (15.9–48.7) in those who became phenotype-positive before 10 years of age and 19.1% (8.3–40.6) in those who became phenotype-positive after 10 years of age. For those who became phenotype-positive before 10 years of age, the median age at MaCE was 8.9 (5.9–11.2) years and median time from HCM onset to MaCE was 3.5 (0.9–4.9) years. For subjects who became phenotype-positive after 10 years of age, the median age at MaCE was 14.3 (14.27–14.59) years and the median time from HCM onset to MaCE was 0.6 (0.2–1.3) years (although the latter entails a shorter duration of follow-up) (Figure 3B). The freedom from death/aborted SCD was 99.6% (98.5–99.9) at 5 years, 99.6% (98.5–99.9) at 10 years, 99.3% (97.8–99.8) at 12 years, and 98.5% (96.2–99.4) at 15 years. The freedom from interventions (myectomy, ICD) was 100% at 5 years, 98.8% (97.1–99.5) at 10 years, 98.5% (96.7–99.3) at 12 years, and 96.9 (94.2–98.4) at 15 years. This suggests that childhood onset HCM can be progressive with a subset experiencing adverse events and major interventions early after diagnosis.

Eligibility for early screening

Overall, 201 (38.4%) children fulfilled criteria for early screening based on symptoms or family history of SCD. However, among the 42 children with HCM before 10 years of age, only 29 (69%) met

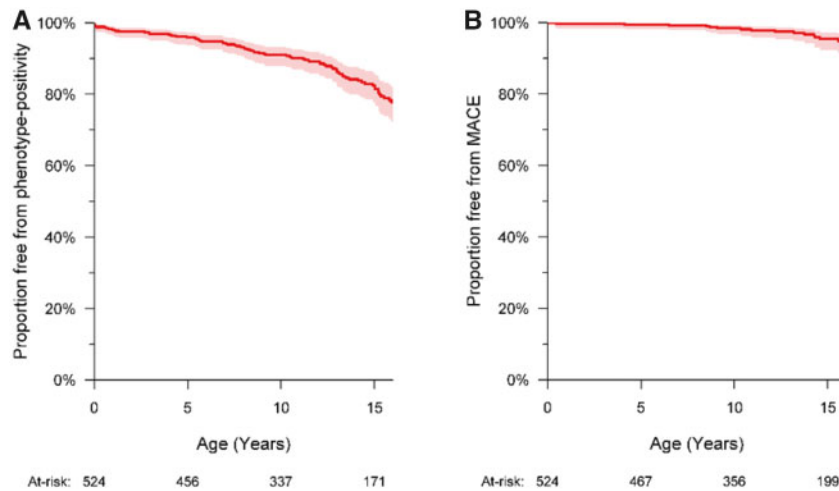


Figure 1 Freedom from hypertrophic cardiomyopathy and major cardiac event ($n = 524$). (A) Freedom from phenotype-positive hypertrophic cardiomyopathy (95% confidence interval) was 95.9% (93.7–97.3) at 5 years, 91.0% (88.0–93.3) at 10 years, 89.2% (85.9–91.8) at 12 years, and 82.4% (77.9–86.1) at 15 years. (B) Freedom from a major cardiac event was 99.4% (98.2–99.8) at 5 years, 98.4% (96.7–99.2) at 10 years, 97.8% (95.8–98.9), and 95.4% (92.3–97.3) at 15 years. MaCE, major cardiac event.

criteria for early screening. The criteria in these 29 cases included symptoms in two and a family history of SCD in 28 cases (with one overlapping case). The freedom from HCM at 10 years of age was 84.3% (77.8–89.0) in subjects eligible for early screening compared to 94.9% (91.6–96.9) in subjects not eligible for early screening ($P = 0.0003$) (Figure 4). Also, of the seven children who had MaCE before 10 years of age, one (11.1%) was not eligible for early screening. This patient had *MYH7* variant-positive HCM and received a primary prevention ICD for extreme septal hypertrophy and abnormal blood pressure response on exercise.

Factors associated with early hypertrophic cardiomyopathy and major cardiac events

On multivariable Cox regression, male gender, family history of SCD/ICD, and P/LP variant in *MYH7* or *MYBPC3* in the child or family were associated with early onset HCM. Family history of SCD prior to 25 years of age, symptoms at screening and presence of a P/LP *MYH7* or *MYBPC3* variant were associated with early occurrence of a MaCE (Table 5 and Figure 5).

Modelling

Factors associated with early onset HCM and MaCEs were included in the final model. Models with clinical factors only were compared to models with clinical and genetic factors using AIC whereby a lower AIC indicates a better model. For both outcomes, i.e. freedom from early onset HCM and freedom from early MaCEs, the combined clinical and genetic model had a lower AIC (i.e. performed better) than the model with only clinical factors (Table 6). The concordance statistic of the final combined Cox proportional hazard model for freedom from early onset HCM was 0.71 and for MaCE was 0.85 indicating good discriminative ability.

Discussion

Our study has several important findings. First, 9.9% children screened for HCM were already phenotype-positive at first evaluation with MaCEs occurrence at a median follow-up of 1.5 (0.5–4.1) years from onset of HCM. 52.5% children with clinical HCM became phenotype-positive before 10 years of age and 41% MaCEs occurred in children before 10 years of age. The short interval to a MaCE suggests that many children already had a well-established phenotype by the time they were screened with a relatively rapid progression on follow-up. Second, a third of early HCM cases did not fulfil criteria for early screening like symptoms or family history of SCD. Missing data precluded our ability to include early onset HCM in the family in the screening criteria. Although the children who met early screening criteria had a lower freedom from early onset HCM and MaCEs, 31% children with early onset HCM did not meet eligibility for early screening and would potentially have been missed. Thirdly, we identified male gender, family history of SCD and presence of P/LP variants in *MYH7* or *MYBPC3* as predictors of early onset HCM and of MaCEs. Several published studies have reported an association of genetic factors with phenotype and outcomes in children and adults explaining some of the variability in disease onset both within and between families with variants in the same gene.^{2–4}

As an autosomal dominant disorder, every first-degree relative from a HCM family regardless of age has a 50% chance of inheriting the disease. However, the guidelines for when to offer screening differ by age likely based on previous studies with an underrepresentation of children and pre-adolescents that reported that most HCM does not manifest until adolescence or adulthood.^{6–10} Our findings and the recent report by Norrish *et al.*¹¹ which include large patient cohorts suggest that disease prevalence in children and pre-adolescents is not lower than in adolescents. They found that in 1198 consecutive children aged ≤ 18 years from 594 families who

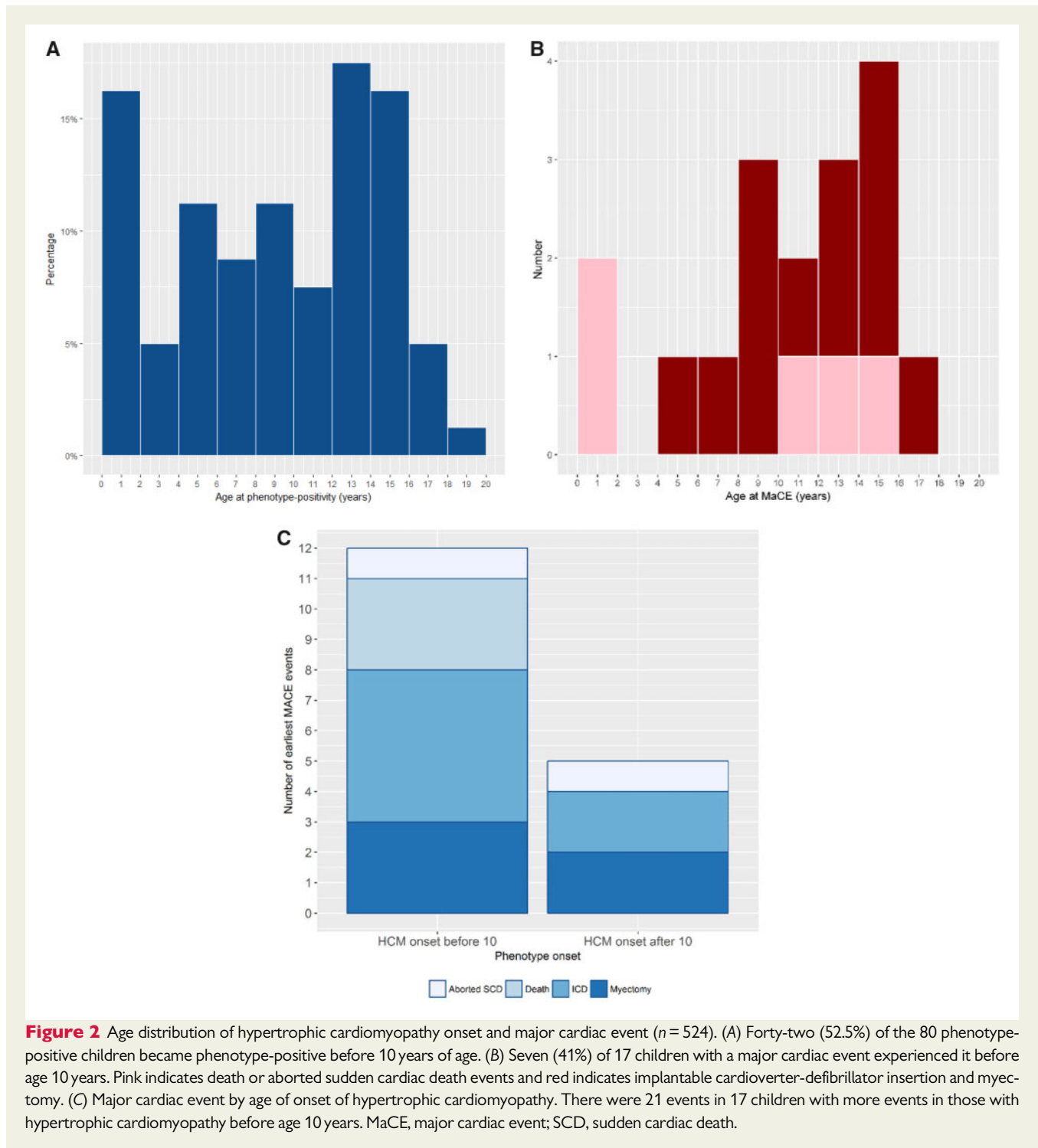


Figure 2 Age distribution of hypertrophic cardiomyopathy onset and major cardiac event ($n = 524$). (A) Forty-two (52.5%) of the 80 phenotype-positive children became phenotype-positive before 10 years of age. (B) Seven (41%) of 17 children with a major cardiac event experienced it before age 10 years. Pink indicates death or aborted sudden cardiac death events and red indicates implantable cardioverter-defibrillator insertion and myectomy. (C) Major cardiac event by age of onset of hypertrophic cardiomyopathy. There were 21 events in 17 children with more events in those with hypertrophic cardiomyopathy before age 10 years. MaCE, major cardiac event; SCD, sudden cardiac death.

underwent serial evaluation, 72% of diagnoses were made in pre-adolescents with over a third of patients requiring medical, surgical or device therapy before 18 years of age. They concluded that the phenotype of familial HCM in childhood is varied and includes severe disease, suggesting that clinical screening should commence at a younger age. At the other end of the spectrum, Maurizi et al.¹² recently published a report that showed that genotype-positive

individuals who remain phenotype-negative by 18 years have a low frequency (10%) of phenotype conversion. Together, this suggests that disease prevalence and penetrance in children is not lower than in adolescents or even adults.

Secondly, the event rates in phenotype-positive children are not different than in adults. The risk of a SCD event in an adult diagnosed with clinical HCM is 1–2% per year.¹³ Our previous report on 98

Table 4 Major cardiac events in the overall cohort

Age at diagnosis (years)	Age at MaCE (years)	Genetic test results (P/LP variant)	Major cardiac event	Cause of death/indication for ICD
0 (foetal)	0.4	None	Death (cardiac)	Advanced heart failure
0 (foetal)	0.5	MYBPC3 (homozygous)	Death (cardiac)	Advanced heart failure awaiting transplant
0 (foetal)	12.2	MYH7	ICD—primary prevention	Family history of SCD, severe septal hypertrophy, LGE on CMR
1.1	4.1	MYBPC3	Surgical myectomy	Symptomatic with LVOT gradient >100 mmHg
2.7	13.7	MYH7	Aborted SCD; ICD—secondary prevention	Secondary prevention
4.2	8.7	MYH7	ICD—primary prevention	Family history of SCD, extreme septal hypertrophy, and abnormal BP response
4.4	8.5	MYBPC3 (compound heterozygote)	ICD—primary prevention	Family history of SCD, abnormal BP response (two appropriate shocks)
5.4	6.5	MYBPC3	Surgical myectomy	Symptomatic with severe LVOTO on optimal medical therapy
7	10.9	MYBPC3 (compound heterozygote)	Death (SCD)	SCD
7.6	15.6	MYH7	ICD—primary prevention	Family h/o SCD, severe septal hypertrophy, and palpitations
8.8	10	MYH7	ICD—primary prevention	Extreme septal hypertrophy, abnormal BP response
8.8	14.3	None	Aborted SCD; ICD—secondary prevention	Secondary prevention
9.1	9.1	None	Surgical myectomy (+ primary prevention ICD 3 years later)	Family history of SCD, severe LVOTO, and NSVT
11.5	14.3	MYH7	Surgical myectomy	Symptomatic with severe LVOTO on optimal medical therapy
13	13.2	MYBPC3 (compound heterozygote)	ICD—primary prevention	Severe septal hypertrophy, NSVT, and abnormal BP response (one appropriate shock)
14.5	14.6	None	Surgical myectomy (+ primary prevention ICD)	Severe LVOTO, NSVT, and abnormal BP response on exercise
15.9	16.5	TNNT2	ICD—primary prevention	Unexplained syncope, severe septal hypertrophy, and abnormal ECG on exercise

CMR, cardiac magnetic resonance; LGE, late gadolinium enhancement; LVOTO, LV outflow tract obstruction; MaCE, major cardiac event; NSVT, non-sustained ventricular tachycardia; P/LP, pathogenic or likely pathogenic.

genotype-positive probands with HCM <18 years old revealed a 10% event rate over a median follow-up of 1.4 years (0.4–3.0). These included 7% SCD events i.e. one SCD, three resuscitated SCD, and three appropriate shocks post primary prevention ICD.³ Another recent publication on outcomes in 446 children and adolescents with ICDs showed that appropriate shocks occurred in 15.7% of patients with primary prevention ICDs.¹⁴

Our study provided additional new insights compared to the Norrish study by evaluating the yield of clinical testing in patients who met early screening criteria vs. those who did not and also by evaluating the importance of early genetic screening to help predict early onset HCM. Despite good discriminative ability, the combined clinical and genetic models were not perfect in their predictions. This may be related to the availability of a genetic diagnosis in only a subset of the cohort since many families either decline genetic testing or

remain genotype-elusive. This suggests that until better validated prediction models of early onset disease are available, clinical and genetic screening should not be delayed till 10 or 12 years of age since studies report effective coping with a positive result and no difference in quality of life between gene-positive children and peers.^{15,16} We recognize that there are costs associated with early screening. However, even if event rates are low, there are significant costs (emotional and financial) to families and to society associated with missed diagnoses and the preventable loss of a young life.^{17,18}

Limitations

A retrospective study can entail a survivor bias and underestimate out of hospital deaths and early onset HCM since those who were not screened until 12 years of age may have had unrecognized disease prior to their first screening. The prediction model for early onset

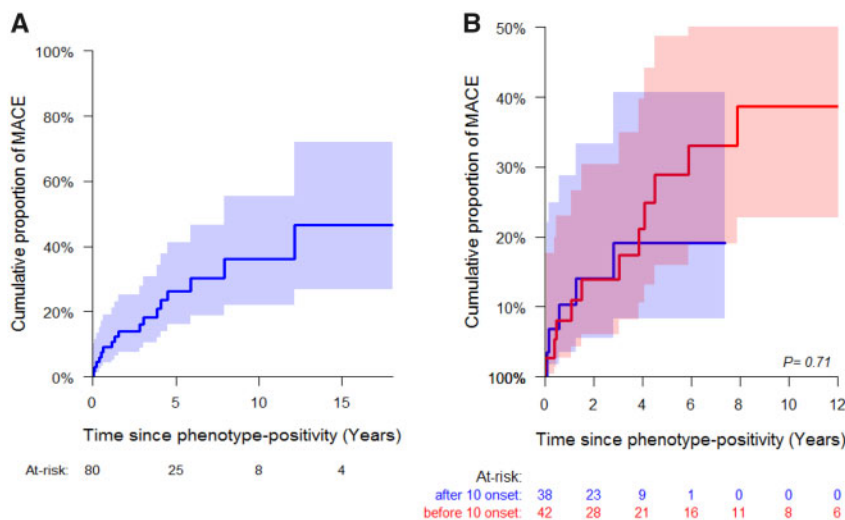


Figure 3 Major cardiac event in phenotype-positive children (n = 80). The x-axis shows the time between phenotype-positivity and major cardiac event in years. (A) The cumulative proportion (95% CI) of major cardiac event (A) at 5 years was 26.4% (58.7–83.9) in the overall cohort, and (B) at 10 years was 28.9% (15.9–48.7) in subjects with hypertrophic cardiomyopathy before 10 years of age and 19.1% (8.3–40.6) in subjects with hypertrophic cardiomyopathy after 10 years of age. MaCE, major cardiac event.

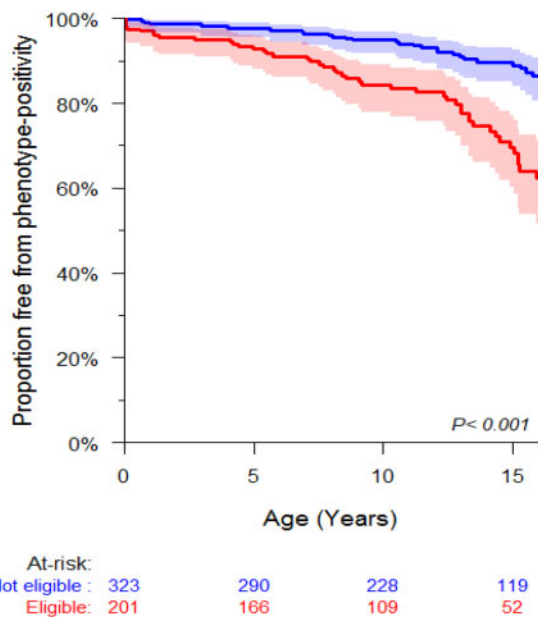


Figure 4 Freedom from hypertrophic cardiomyopathy stratified by eligibility for early screening. Freedom from hypertrophic cardiomyopathy in those eligible for early screening (red) was 92.8% (88.1–95.7) at 5 years, 84.3% (77.8–89.0) at 10 years, and 82.6% (75.7–87.7) at 12 years. Freedom from clinical hypertrophic cardiomyopathy in the remaining cohort (blue) was 97.8% (95.4–98.9) at 5 years, 94.9% (91.6–96.9) at 10 years, and 93.1% (89.3–95.6) at 12 years.

HCM and MaCEs was not validated in an independent external cohort. Also, a majority of events were cardiac interventions like myectomy and ICD which can involve practitioner bias.

In summary, the results of our study coupled with other recent evidence of the yield on clinical screening in young family members indicate that disease prevalence and penetrance as well as event rates are not lower in children compared to adolescents (and adults). Although annual event rate in children who develop HCM is low, given the potential for rapid progression, early detection of clinical HCM is important in order to facilitate close follow-up and timely interventions to prevent SCD, and to alleviate symptoms of worsening LV outflow obstruction before adverse events occur. In conclusion, younger family members should be considered for early clinical and genetic screening to identify the subset in need of close monitoring and interventions.

Funding

This work was supported by the Ted Rogers Centre for Heart Research and the Heart and Stroke Foundation Chair funds (to S.M.).

Conflict of interest: none declared.

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Table 5 Factors associated with early onset hypertrophic cardiomyopathy and major cardiac event (n = 524)

	Univariable HR (95% CI)	P-value	Multivariable HR (95% CI)	P-value
Early phenotype-positivity				
Male gender (ref = female)	2.048 (1.099–3.816)	0.024	2.028 (1.081–3.807)	0.028
Family history of SCD	2.618 (1.474–4.650)	0.001	1.935 (1.071–3.497)	0.029
Family history of SCD before age 25 years	1.615 (0.756–3.452)	0.22		
First-degree phenotype or genotype-positive relative	1.695 (0.527–5.456)	0.38		
Family history of ICD	2.842 (1.612–5.010)	<0.001	2.212 (1.218–4.019)	0.009
Family history of transplant	1.240 (0.385–43.990)	0.72		
Symptomatic at screening	4.130 (1.002–17.026)	0.050	—	—
P/LP MYH7 variant	2.995 (1.526–5.877)	0.001	3.247 (1.555–6.782)	0.002
P/LP MYBPC3 variant	2.451 (1.331–4.513)	0.004	2.319 (1.173–4.587)	0.016
P/LP variant in other gene	0.385 (0.053–2.787)	0.34	—	—
P/LP variants in more than one gene	3.234 (0.446–23.445)	0.25	—	—
MaCE ^a				
Male gender (ref = female)	1.176 (0.448–3.090)	0.74	—	—
Family history of SCD	3.516 (1.299–9.513)	0.013	1.177 (0.325–4.260)	0.80
Family history of SCD before age 25 years	5.829 (2.217–15.326)	<0.001	2.621 (0.687–9.990)	0.158
Family history of ICD	3.497 (1.347–9.076)	0.010	2.869 (0.903, 9.119)	0.074
Family history of transplant	1.195 (0.158–9.018)	0.86		
Symptomatic at screening	15.400 (4.408–53.805)	<0.001	57.611 (9.549–347.595)	<0.001
P/LP MYH7 variant	6.068 (2.234–16.481)	<0.001	18.574 (4.183–82.475)	<0.001
P/LP MYBPC3 variant	2.884 (1.06–7.802)	0.037	9.340 (2.123–41.190)	0.003
P/LP variant in other gene	1.126 (0.149–8.491)	0.91	—	—

Bold values significant on multivariable analysis.

CI, confidence interval; HR, hazard ratio; ICD, implantable cardioverter-defibrillator; LP, likely pathogenic; MaCE, major cardiac event; P, Pathogenic; SCD, sudden cardiac death.

^aFirst degree affected relative and presence of multiple P/LP variants were not included in the models for MaCE because HR was unestimable as the events did not occur in one of the two groups.

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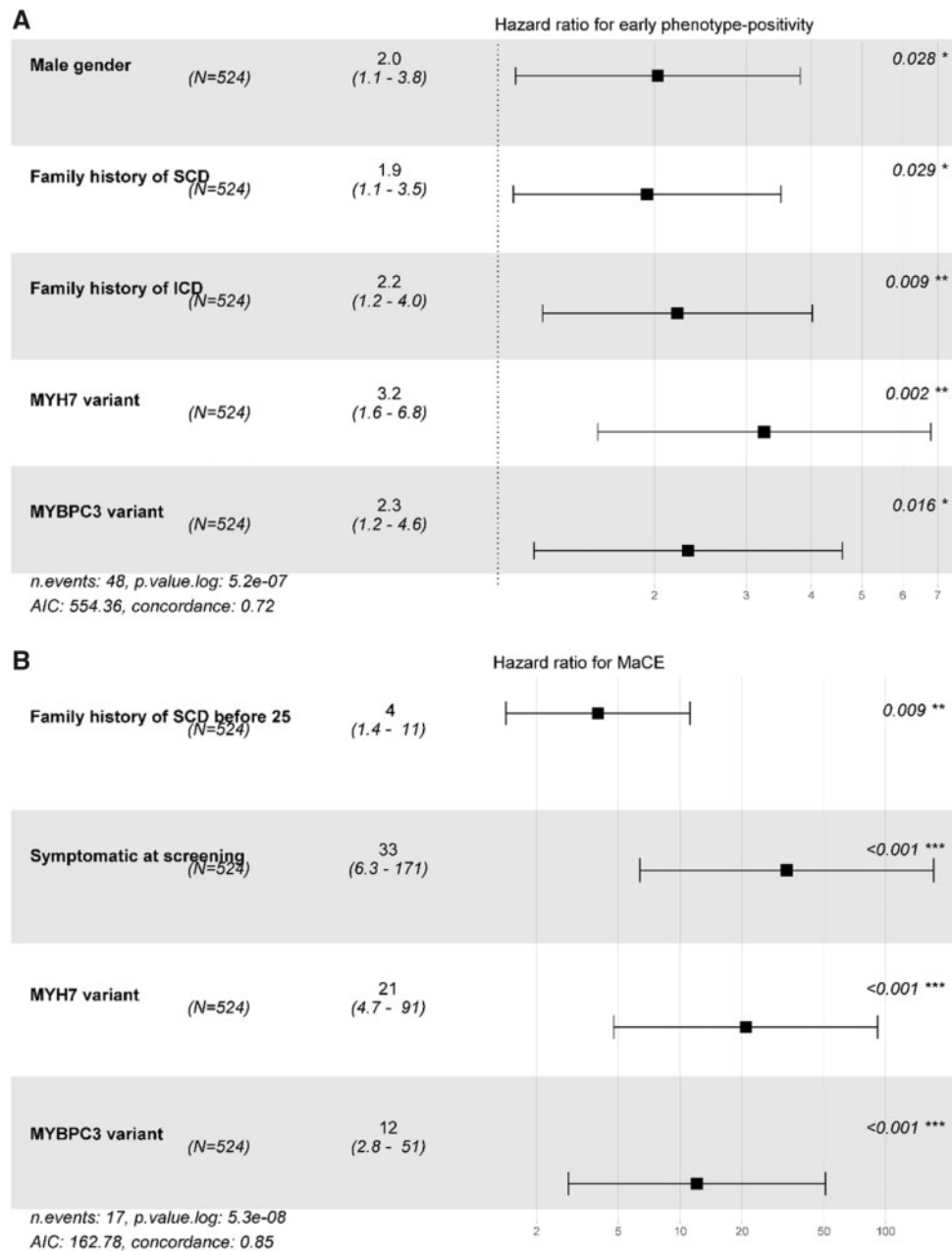
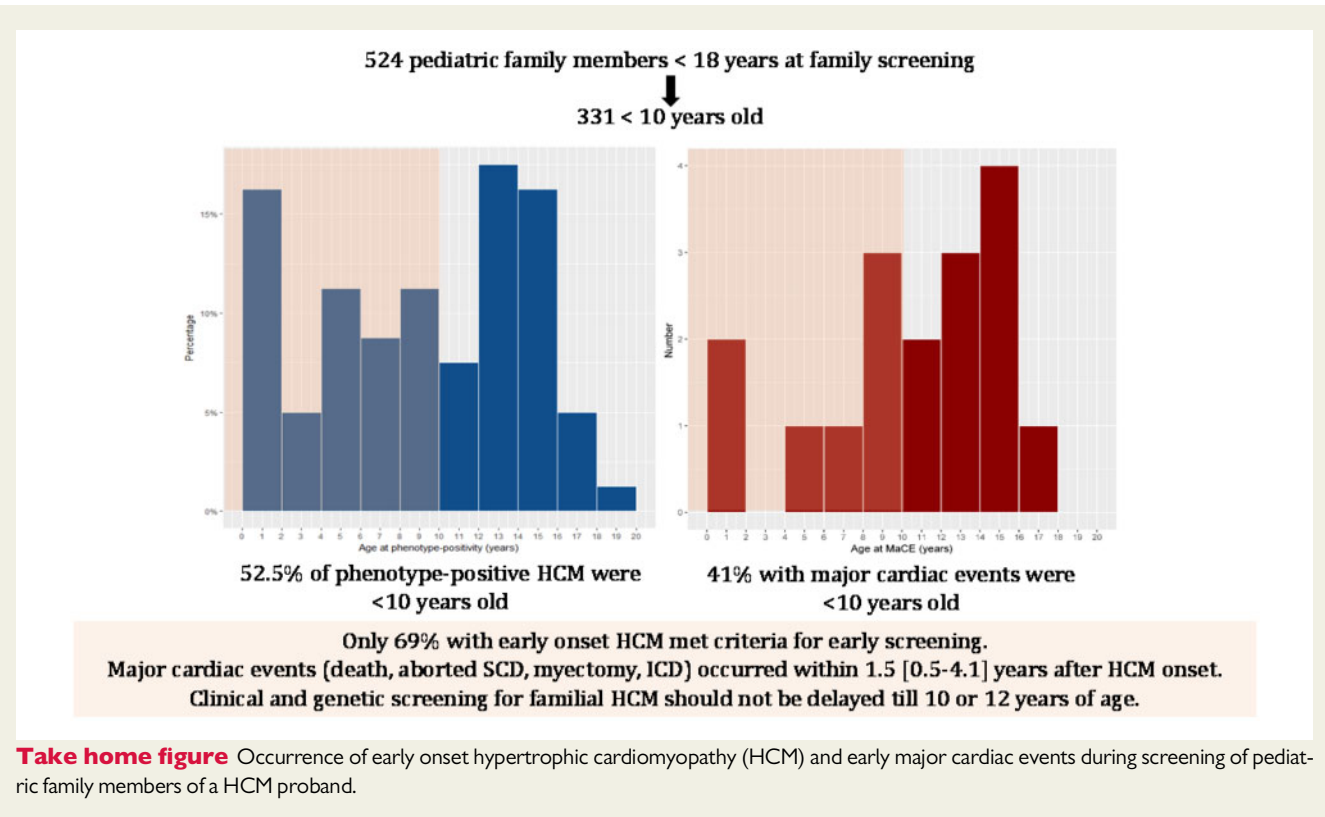


Figure 5 Factors associated with major cardiac event and early onset hypertrophic cardiomyopathy (n = 524). Forest plot of hazard ratios and 95% confidence intervals for multivariable model of freedom from (A) early onset hypertrophic cardiomyopathy and (B) major cardiac event. MaCE, major cardiac event.

Table 6 Clinical and genetic models

Model	Factors included	AIC	Concordance
Early onset HCM			
Clinical factors	Gender Family history of SCD Family history of ICD	565	0.64
Clinical + genetic factors	Gender Family history of SCD Family history of ICD Pathogenic MYH7 variant Pathogenic MYBPC3 variant	556	0.71
MaCE			
Clinical factors	Family history of SCD before 25 years Symptomatic at screening	182	0.72
Clinical + genetic factors	Family history of SCD before 25 years Symptomatic at screening Pathogenic MYH7 variant Pathogenic MYBPC3 variant	163	0.85

AIC, Akaike's information criterion; MaCE, major cardiac event; SCD, sudden cardiac death.



Take home figure Occurrence of early onset hypertrophic cardiomyopathy (HCM) and early major cardiac events during screening of pediatric family members of a HCM proband.

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