

1 **Yield of Molecular Autopsy in Sudden Cardiac Death in Athletes. Data from a Large**
2 **Registry in the United Kingdom**

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5 Gherardo Finocchiaro^a MD, PhD*, Davide Radaelli^{a,b} MD*, David Johnson^a PhD, Raghav T.
6 Bhatia^a MBBS, MRCP, Joseph Westaby^a MBBS, Stefano D'Errico^b, MD, PhD, Michael
7 Papadakis^a MBBS, FRCP, MD, Sanjay Sharma^a BSc, MBChB, FRCP, MD, Mary N.
8 Sheppard^a MBBCh, BAO, BSc, MD, FRCPath, Elijah R. Behr^a MA, MBBS, MD, FRCP

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10 **Institutions:**

11 ^a Cardiovascular Clinical Academic Group and Cardiology Research Section, St. George's,
12 University of London, St. George's University Hospitals NHS Foundation Trust, United
13 Kingdom

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15 ^b Department of Medicine, Surgery and Health, University of Trieste, Trieste, Italy

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21 * equally contributed as first author

22

23

24 **Author for correspondence:**

25 Professor Elijah R Behr,
26 Professor of Cardiology, St. George's University of London, Cardiovascular Sciences,
27 Cranmer Terrace, London, SW17 0RE, UK.
28 E-mail: ebehr@sgul.ac.uk

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Abstract

4 **Background:** Sudden cardiac death (SCD) may occur in apparently healthy individuals,
5 including athletes. We report the diagnostic role of post-mortem genetic testing, molecular
6 autopsy (MA), in elucidating the cause of SCD in athletes.

7 **Methods:** We reviewed a database of 6860 consecutive cases of SCD referred to our
8 specialist cardiac pathology centre. All cases underwent detailed cardiac autopsy and 748
9 were deemed to be athletes. Of these, 42 (6%) were investigated with MA (28 using a
10 targeted sequencing, 14 exome sequencing). Variants were classified manually as pathogenic
11 (P), likely pathogenic (LP), variant of unknown significance (VUS) using international
12 guidelines. Clinical information was obtained from referring coroners who completed a
13 detailed health questionnaire.

14 **Results:** Out of the 42 decedents (average age 35 years old, 98% males) who were
15 investigated with MA, the autopsy was in keeping with a structurally normal heart (sudden
16 arrhythmic death syndrome, SADS) in n=33 (78%) cases, followed by arrhythmogenic
17 cardiomyopathy (ACM) in 8 (19%) individuals and idiopathic left ventricular fibrosis in 1
18 (2%). Death occurred during exercise and at rest in 26 (62%) and 16 (38%) individuals
19 respectively. Variants that were adjudicated clinically actionable were present in 7 cases
20 (17%). There was concordance between the genetic and phenotypic findings in 2 cases of
21 ACM. None of the variants identified in SADS cases were previously linked to
22 channelopathies. Clinically actionable variants in cardiomyopathy-associated genes were
23 found in 5 cases of SADS.

24 **Conclusions:** The yield of MA in athletes who died suddenly is 17%. In SADS cases,
25 clinically actionable variants were found in cardiomyopathy-associated genes and not in

1 channelopathy-associated genes. ACM is a common cause of SCD in athletes and one in four
2 decedents had a clinically actionable variant in FLNC and TMEM43 genes.

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4 **Keywords:** Molecular autopsy, sudden cardiac death.

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7 Sudden cardiac death (SCD) may occur in apparently healthy individuals including athletes.

8 The yield of post-mortem genetic testing (molecular autopsy - MA) in athletes who died

9 suddenly is unknown¹. We aimed to assess the yield of pathogenic (P) and likely pathogenic

10 (LP) variant by MA in a cohort of athletes who died suddenly and underwent post-mortem

11 examination by a expert cardiac pathologists.

12

13 We reviewed a database of 6860 consecutive cases of SCD referred to our specialist cardiac

14 pathology centre at St George's, University of London between 1994 and 2020. SCD was

15 defined as death from a cardiovascular cause within one hour of onset of symptoms if

16 witnessed, or within 12 hours of onset if unwitnessed. Clinical information was obtained

17 from referring coroners who were asked to complete a detailed health questionnaire. We

18 arbitrarily defined athletes as individuals that engaged in at least 5 hours of organised

19 exercise activity per week. All cases underwent detailed post-mortem evaluation of the heart,

20 including histological analysis, by expert cardiac pathologists (MNS, JW). A minimum of 10

21 blocks of tissue were taken for histological analysis and cardiomyopathy was defined as

22 reported previously^{1,2}. Death during exercise was defined as occurring while the individual

23 was engaging in exercise, as opposed to death during daily activities or rest. MA was

24 performed with targeted panel and whole exome sequencing focusing on a broad panel of

25 genes implicated in channelopathy and cardiomyopathy as previously described³. Variants

26 were classified manually as P, LP or a variant of unknown significance (VUS) using the

27 American College of Medical Genetics and Genomics (ACMG) consensus statement

1 guidelines⁴. Ethical and research governance approval have been granted for this study
2 (10/H0724/38). Results are expressed as mean \pm standard deviation (SD) for continuous
3 variables or as number of cases and percentage for categorical variables.

4

5 Out of the total cohort, 748 individuals were athletes. A minority of consecutive athletes with
6 DNA available (n=42, 6%) were investigated with MA: 28 individuals with targeted panel
7 sequencing and 14 with whole exome sequencing. The average age was 35 ± 11 years old and
8 98% (n=41) were male. A structurally normal heart at the post-mortem examination with
9 negative toxicology (sudden arrhythmic death syndrome - SADS) was found in 33 (78%)
10 decedents. Arrhythmogenic cardiomyopathy (ACM) was observed in 8 (19%) cases and
11 idiopathic left ventricular fibrosis in 1 (2%). Death occurred during exercise in 26 (62%)
12 cases and at rest in 16 (38%), including 5 (12%) cases where death occurred during sleep.
13 MA showed P/LP variants in 9 (21%) individuals. Variants that were adjudicated as the likely
14 cause of death were present in 7 cases (17%). Two cases had a P/LP variant in *HFE* in
15 heterozygous state unlikely to cause haemochromatosis. There was concordance between the
16 genetic and phenotypic findings in two cases of ACM (Table 1). No P/LP variants linked to
17 channelopathies were identified in the SADS cases (Table 1).

18

19 Athletes often appear to epitomize health but SCD may occur in apparently healthy
20 individuals^{1,5}. Post-mortem examination is a crucial diagnostic step in establishing the cause
21 of SCD and in guiding the clinical evaluation of surviving relatives^{6,7,8,9}. The interpretation of
22 the post-mortem results, however, can be challenging especially when the heart is structurally
23 normal or when abnormalities of uncertain significance are found¹⁰. MA has the potential to
24 establish the cause of death^{11,12}. Indeed, a study from our group on a large cohort of SADS
25 decedents investigated with MA found a clinically actionable pathogenic or likely pathogenic

1 variants in 13% of the cases, mostly associated with channelopathy³. Our study comprised 42
2 athletes where the post-mortem examination performed by expert cardiac pathologists was
3 mostly in keeping with a structurally normal heart and ACM. Roughly one fifth of athletes
4 had a pathogenic or likely pathogenic variant. Clinically actionable variants were found in
5 17% of cases. Interestingly, none of the athletes with SADS were found to have P/LP variants
6 associated with channelopathies. In all cases P/LP variants were identified in
7 cardiomyopathy-associated genes. This suggest that cardiomyopathy, even when ‘concealed’
8 and not detected at expert cardiac autopsy, may predispose to SCD in young male athletes.
9 This is in line with a recent study on 91 autopsy-inconclusive SCD cases where
10 cardiomyopathy-associated genes harboured 70% of clinically actionable variants¹³. Genetic
11 findings correlated with the phenotype in only two cases, both with ACM; *FLNC* and
12 *TMEM43* were the genes involved, which emphasizes their arrhythmic risk of these specific
13 disorders^{14,15}. Although our cohort is large, not all cases of SCD are referred to our centre
14 and this introduces a bias. Further, only 6% of athletes who died suddenly and were referred
15 to our centre, were investigated with MA. This implies potential selection bias which should
16 be taken in account when interpreting the results.

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18 In conclusion, in a small cohort of athletes who died suddenly and who were investigate with
19 post-mortem genetic test, the yield of MA was 17%. In SADS cases P/LP variants were
20 found in cardiomyopathy-associated genes and not in channelopathy-associated genes.
21 Genetic panels should include assessment of genes implicated in cardiomyopathy even when
22 a clear phenotype is not identified through post-mortem examination. One in four decedents
23 with arrhythmogenic cardiomyopathy were identified with P/LP variants.

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7 **Conflict of interest:** none declared.

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9 **Data Availability Statement:** Data supporting this study are available from CRY Centre for
10 Cardiac Pathology in St. George's University, London. Access to the data is subject to
11 approval and a data sharing agreement due to ethical reason.

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1 **Table 1.** Genetic test results in athletes who died suddenly. All variants were heterozygous. The first seven cases (light grey shading) are

2 deemed to harbour an actionable variant.

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Case	Gene	Variant details	p. annotations	ACMG Interpretation	ClinGen	Genecards	ClinVar	PM diagnosis	Concordance
1	FLNC	c.4718T>A	p.Leu1573Ter	Pathogenic	HCM – Definitive (AD) DCM- Definitive (AD) Myofibrillar Myopathy		HCM-definitive (AD) DCM-definitive (AD) Myofibrillar Myopathy	Arrhythmogenic cardiomyopathy involving both ventricles	YES
2	TMEM43	c.1073C>T	p.Ser358Leu	Pathogenic	AC – Definitive (AD)		AC	Arrhythmogenic Cardiomyopathy	YES
3	CDH2	c.2060A>G	p.Asn687Ser	Likely Pathogenic	AC – Limited (AD)	Agenesis of corpus callosum, cardiac, ocular, and genital syndrome	Not present specific variant	SADS	NO
4	CDH2	c.586A>G	p.Ser196Gly	Likely Pathogenic	AC – Limited (AD)	Agenesis of corpus callosum, cardiac, ocular, and	Not present specific variant	SADS	Disputed – Slight fibrosis of RV interpreted as not significant.

						genital syndrome			
5	TTN	c.9577C>T	p.Arg3193Ter	Pathogenic	DCM – Definitive (AD) TTN-related myopathy – Definitive (AR) ACM – Limited (AD) HCM – Limited (AD)		DCM - Conflicting interpretations of pathogenicity: Likely pathogenic; Uncertain significance	SADS	NO – No evidence of cardiomyopathy
6	CRYAB	c.32G>A	p.Arg11His	Likely Pathogenic	/	Myofibrillar myopathy, DCM	DCM – Conflicting - VUS 2 submitters – LP 1 submitter	SADS	NO –No evidence of cardiomyopathy
7	SOS1	c.3392G>A	p.Arg1131Lys	Pathogenic	Noonan Syndrome– Definitive (AD) Costello Syndrome– Disputed (AD) Cardiofaciocutaneous Syndrome - Disputed (AD)	Noonan Syndrome, Gingival fibromatosis	RASopathy (Uncertain significance) Noonan syndrome (Uncertain significance)	SADS	NO –No evidence of cardiomyopathy

8	HFE	c.845G>A	p.Cys282Tyr	Pathogenic	Type 1 hemochromatosis (Gene Associated with Autosomal Recessive Phenotype)	Type 1 hemochromatosis, Microvascular complications of diabetes	Cardiomyopathy (NOS)	SADS	NO
9	HFE	c.845G>A	p.Cys282Tyr	Pathogenic	Type 1 hemochromatosis (Gene Associated with Autosomal Recessive Phenotype)	Hemochromatosis, Type 1 Microvascular Complications of Diabetes	Cardiomyopathy (NOS)	Arrhythmic cardiomyopathy (predominant RV involvement)	NO

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2 **Abbreviations:** ACM: arrhythmic cardiomyopathy, AD: autosomal dominant, AR, autosomal recessive, DCM: dilated cardiomyopathy,

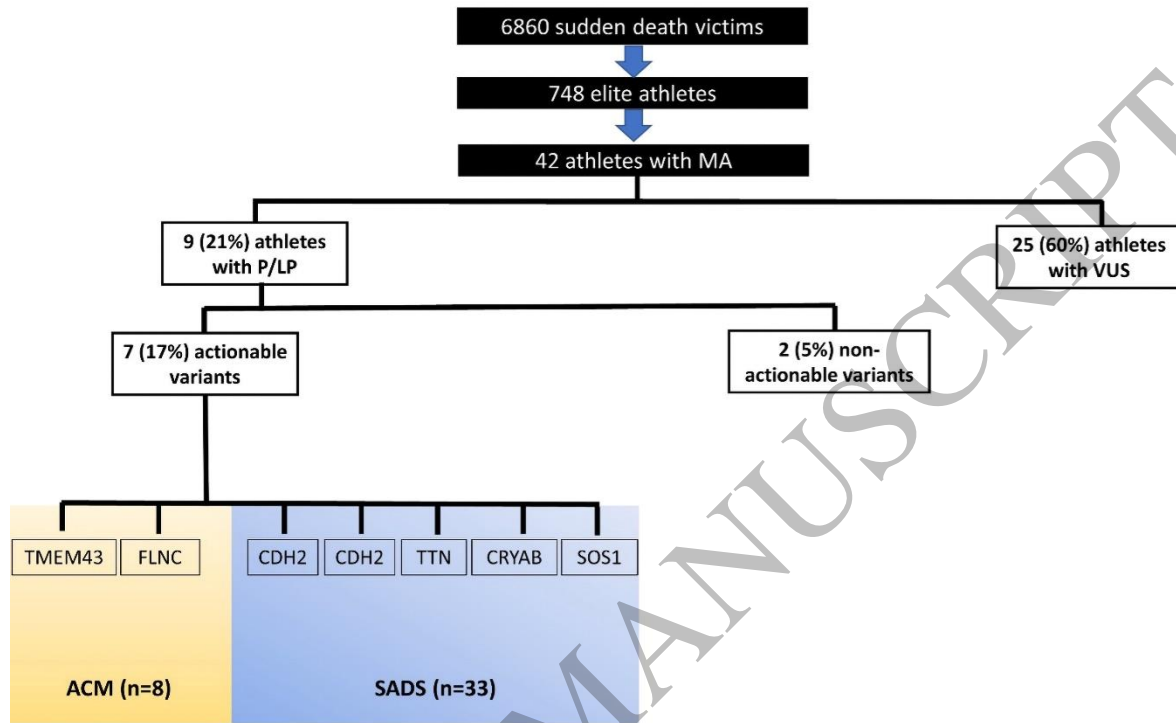
3 HCM: hypertrophic cardiomyopathy; PM: post-mortem; RV: right ventricle, SADS: sudden arrhythmic death syndrome, VUS: variant of

4 uncertain significance; NOS: not otherwise specified; LP: likely pathogenic; RV: right ventricle; ACMG: American College of Medical

5 Genetics; ClinGen: Clinical Genome Resource <https://clinicalgenome.org/>; ClinVar: NCBI database <https://www.ncbi.nlm.nih.gov/clinvar/>. The

6 “Concordance” column indicates whether there is concordance between the genetic data and the morphological findings.

1 **Figure 1.** Molecular autopsy in sudden cardiac death in athletes. Genes involved are
 2 described. Seven actionable variants have been found in 7 athletes.



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 4
 5
 6 *Figure 1*
 7 201x121 mm (x DPI)

8 **Abbreviations:** MA: Molecular Autopsy; ACM: arrhythmogenic cardiomyopathy; LP: likely
 9 pathogenic variant; P: pathogenic variant; VUS: variant of uncertain significance; SADS:
 10 sudden arrhythmic death syndrome.