



# Indications and utility of cardiac genetic testing in athletes

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Sports Cardiology practice commonly involves the evaluation of athletes for genetically determined cardiac conditions that may predispose to malignant arrhythmias, heart failure, and sudden cardiac death. High-level exercise can lead to electrical and structural cardiac remodelling which mimics inherited cardiac conditions (ICCs). Differentiation between 'athlete's heart' and pathology can be challenging and often requires the whole armamentarium of available investigations. Genetic studies over the last 30 years have identified many of the genetic variants that underpin ICCs and technological advances have transformed genetic testing to a more readily available and affordable clinical tool which may aid diagnosis, management, and prognosis. The role of genetic testing in the evaluation and management of athletes with suspected cardiac conditions is often unclear beyond the context of specialist cardio-genetics centres. This document is aimed at physicians, nurses, and allied health professionals involved in the athlete's care. With the expanding role and availability of genetic testing in mind, this document was created to address the needs of the broader sports cardiology community, most of whom work outside specialized cardio-genetics centres, when faced with the evaluation and management of athletes with suspected ICC. The first part of the document provides an overview of basic terminology and principles and offers guidance on the appropriate use of genetic testing in the assessment of such athletes. It outlines key considerations when contemplating genetic testing, highlighting the potential benefits and pitfalls, and offers a roadmap to genetic testing. The second part of the document presents common clinical scenarios in Sports Cardiology practice, outlining the diagnostic, prognostic, and therapeutic implications of genetic testing, including impact on exercise recommendations. The scope of this document does not extend to a comprehensive description of the genetic basis, investigation, or management of ICCs.

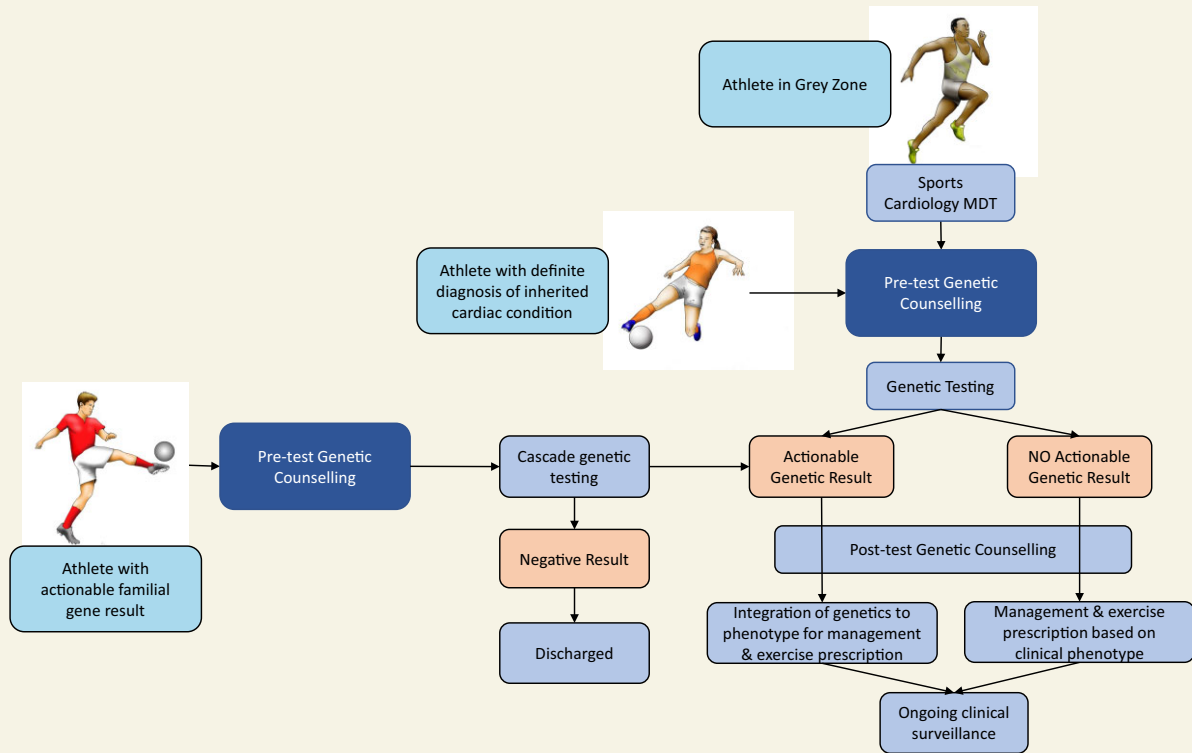
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A scientific statement from the Sports Cardiology and Exercise Section of the European Association of Preventive Cardiology, the European Heart Rhythm Association, the ESC Working group of myocardial and pericardial diseases, the ESC Council on Cardiovascular Genomics, the European Society of Human Genetics, and the Association for European Paediatric and Congenital Cardiology. © The Author(s) 2022. Published by Oxford University Press on behalf of the European Society of Cardiology.

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## Graphical Abstract



Flow chart demonstrating pathways for athletes to proceed with cardiac genetic testing depending whether they have a known diagnosis of inherited cardiac condition (ICC), they are an athlete in the grey zone, or they have a known actionable (pathogenic or likely pathogenic) genetic result in their family. Multidisciplinary should consist of individuals with expertise in sports cardiology, cardiac genetics and the diagnosis, and management of ICCs and genetic counsellors.

**Keywords** Genetic testing • Athletes • Sudden cardiac death • Inherited cardiac conditions • Channelopathies • Cardiomyopathies

## Introduction

### Background

Sports Cardiology practice commonly involves the evaluation of athletes for genetically determined cardiac conditions that may predispose to malignant arrhythmias, heart failure, and sudden cardiac death (SCD). High-level exercise can lead to electrical and structural cardiac remodelling which mimics inherited cardiac conditions (ICCs), often referred as the 'grey zone'. Differentiation between 'athlete's heart' and pathology can be challenging and often requires the whole armamentarium of available investigations.<sup>1–5</sup> Genetic studies over the last 30 years have identified many of the genetic variants that underpin ICCs and technological advances have transformed genetic testing to a more readily available and affordable clinical tool which may aid diagnosis, management, and prognosis. The role of genetic testing in the evaluation and management for athletes with suspected cardiac conditions is often unclear beyond the context of specialist cardio-genetics centres.

### Aim of the document and target audience

This document is aimed at physicians, nurses, and allied health professionals involved in the athlete's care. With the expanding role and availability of genetic testing in mind, this document was created to address the needs of the broader sports cardiology community, most of whom work outside specialized cardio-genetics centres, when faced with the evaluation and management of athletes with suspected ICC. The first part of the document provides an overview of basic terminology and principles and offers guidance on the appropriate use of genetic testing in the assessment of such athletes. It outlines key considerations when contemplating genetic testing, highlighting the potential benefits and pitfalls, and offers a roadmap to genetic testing summarized in the Graphical abstract. The second part of the document presents common clinical scenarios in Sports Cardiology practice, outlining the diagnostic, prognostic, and therapeutic implications of genetic testing, including subsequent impact on exercise recommendations. The scope of this document does not extend to a comprehensive description of the genetic basis, investigation, or management of ICCs.

## Cardiac genetic testing

### Genetic testing in the inherited cardiac conditions

Most ICCs are inherited in an autosomal dominant manner, meaning that there is a 50% risk of transmission to offspring. Depending on the disease in question, genetic testing may identify a disease-causing (pathogenic or likely pathogenic) genetic variants in up to 70% of patients.<sup>6</sup> The common ICCs, their prevalence and the approximate yield of genetic testing are shown in *Table 1*. The likelihood of finding a causal genetic variant is highest in patients with known familial disease and lowest in older patients and individuals with atypical clinical features.

Genetic counselling is essential prior to any individual undergoing genetic testing and should be performed by trained health care professionals, ideally working within a multidisciplinary team,<sup>6,10</sup> in specialized centres. This is particularly important in complex cases. The aim is to help patients understand and manage the psychological, social, professional, ethical, and legal implications of a genetic test, as well as gather information for other family members, including cardiac and non-cardiac symptoms and autopsy reports. When a causative genetic variant is identified in the index case, often referred as 'proband', relatives can be genetically tested for the same variant, a process known as 'predictive or cascade testing'. If the variant is present, a baseline clinical evaluation is necessary to look for the clinical phenotype and manage according to established protocols. Asymptomatic carriers with no clinical phenotype, referred as 'genotype positive-phenotype negative', can be offered clinical surveillance and reproductive advice.<sup>11</sup> If the variant is absent, relatives can be discharged but should be advised to seek reassessment if new symptoms develop or if new clinically relevant data emerge in the family.<sup>11</sup> A more cautious approach may be considered in situations where the genetic basis of the disease is more complex. A cautious approach is also necessary when considering genetic testing in children when issues of consent, the long-term implications of a positive genetic test, and age-related disease penetrance need to be considered.

In families where there has been a sudden cardiac death, comprehensive macroscopic and histopathological evaluation of the heart, ideally by a cardiac pathologist, is required for an accurate assessment as to the potential cause of death.<sup>12,13</sup> Genetic analysis using DNA from post-mortem tissue or blood samples, also referred as 'molecular autopsy', can be invaluable and assist in the risk assessment of surviving relatives.<sup>12–15</sup> A diagnosis of an ICC or the absence of an identifiable cause, also referred as sudden arrhythmic death syndrome (SADS) will guide genetic and familial evaluation. Similar to clinical testing, molecular autopsy results must always be interpreted in tandem with the results of the post-mortem examination and in accordance with consensus criteria for assigning pathogenicity to genetic variants.<sup>12,13,16,17</sup>

### Nomenclature, techniques, and result interpretation

Cardiac genetic testing has been transformed with the advent of next-generation sequencing (NGS) techniques. These sequencing technologies provide high-throughput of millions of DNA fragments

thereby allowing rapid analysis of large sections of DNA as well as providing more complete coverage of larger genes, such as the ryanodine receptor (*RYR2*) and titin (*TTN*) which have been historically challenging to comprehensively sequence.<sup>18</sup>

In many diagnostic laboratories, genetic testing is performed with NGS-panels which include sequencing genes known to be associated with the disease diagnosed or suspected. More comprehensive approaches include: whole-exome sequencing (WES), which encompasses all coding regions of the genome and whole-genome sequencing (WGS), which involves sequencing of all coding and non-coding regions of the DNA. Whole-genome sequencing provides a comprehensive dataset at the expense of significant challenges in terms of volume and interpretation of data and financial cost.<sup>19,20</sup> It is important to acknowledge that when performing WGS or WES, only variants in genes relevant to the phenotype should be reported.

Genetic testing results are not binary but represent a spectrum of pathogenicity. Variants are grouped in five classes according to guidelines of the American College of Medical Genetics and the Association of Molecular Pathologists (ACMG/AMP).<sup>21</sup> When there is sufficient evidence to consider a variant as the cause of disease, it is classified as pathogenic (Class 5) or likely pathogenic (Class 4). When the evidence indicates that the variant is unlikely to cause disease, it is classified as benign (Class 1) or likely benign (Class 2). When there is insufficient or conflicting evidence, the variant is classified as a variant of uncertain significance (VUS) (Class 3).<sup>21</sup> The classification of a variant is based on a combination of variables: (i) frequency of the variant in large population databases such as the genome aggregation database (gnomAD),<sup>22</sup> (ii) presence of the variant in individuals with the respective disease in genetic testing result databases (e.g. Clinvar), (iii) predicted effect of the variant using computational (in-silico) models (e.g. SIFT, Polyphen-2, mutation taster), (iv) functional data assessing the effect of the variant, and (v) segregation analysis of the variant in families.

Careful variant interpretation and classification is essential to avoid diagnostic errors and should be performed in experienced centres. Only pathogenic and likely pathogenic variants should be viewed as causal variants and used for cascade genetic testing of relatives. If a variant is classified as a VUS, then the variant cannot be used to support a suspected diagnosis or for cascade genetic testing of relatives. For the purpose of sports cardiology practice, a VUS should be ignored, similar to benign or likely benign variants. Some variants may be of future clinical relevance but this needs to be explored in the context of specialist cardio-genetics clinics.<sup>23</sup> It is worth noting that a negative genetic test does not exclude a diagnosis in the presence of a clear or highly suspicious phenotype.

### Genetic testing in Sports Cardiology

Identification, evaluation, and management of ICCs are an integral part of Sports Cardiology practice, as ICCs are implicated as a leading cause of SCD in young (<35 years) athletes.<sup>24</sup> Hypertrophic cardiomyopathy (HCM) accounts for the majority of deaths in a long-standing national registry in the USA,<sup>25</sup> while arrhythmogenic cardiomyopathy (ACM) has been reported as the main cause of death in the North East of Italy.<sup>26</sup> Studies in the UK and Australia indicate that SADS, characterized by a negative or normal post-mortem, is more common than previously thought, suggesting that inherited

arrhythmia syndromes such as long QT syndrome (LQTS), catecholaminergic polymorphic ventricular tachycardia (CPVT), and Brugada syndrome (BrS) account for a significant proportion of SCDs in young and athletic individuals.<sup>27,28</sup> In an attempt to minimize such tragedies, preventive efforts include targeted evaluation of high-risk individuals, as well as wider screening of low-risk individuals.<sup>24</sup> Athletes with cardiac symptoms, a family history of an ICC or SCD at a young age and those with electrocardiogram (ECG) or structural traits suggestive of cardiac disease should be referred to specialist centres for comprehensive assessment.

Similar to the general population, genetic testing in sports cardiology has primarily been used in athletes with a clear phenotype of an ICC in order to facilitate familial cascade testing, or for predictive testing in athletes with a family history of an ICC and a known causative variant.<sup>24</sup> Recently, a broader scope was proposed, where, genetic testing is used for diagnostic and prognostic purposes, including guiding medical therapy and exercise prescription.<sup>7</sup> The utility of genetic testing for diagnostic purposes is particularly relevant in sports cardiology, as the effects of exercise on electrical and structural cardiac remodelling often pose significant challenges and differentiation between 'athlete's heart' and ICC requires the use of several diagnostic tools (Figure 1). Athletes may exhibit increased left ventricular (LV) wall thickness, dilatation of the left ventricle and right ventricle (LV/RV) with associated impairment in function or prominent LV trabeculations<sup>24</sup> (Figure 1). Compared with the general population, athletes are also more likely to demonstrate longer QT intervals and a higher prevalence of T-wave inversion, partial, or complete right bundle branch block and early repolarization patterns.<sup>24,29,30</sup>

Similar to the core principles of genetic testing in any setting, genetic testing for diagnostic purposes should only be used in athletes after careful consideration of a number of factors in the context of a multidisciplinary team (MDT) and after appropriate pre-test counselling.<sup>6,10</sup> The physician should assess the pre-test probability of a positive test based on the suspected condition and strength of the phenotype and consider the potential prognostic and therapeutic implications. For some conditions, such as LQTS, the utility of genetic testing in the athlete can be broad as the identification of different genetic subtypes (LQT 1–3) can inform the risk of arrhythmic events, identify potential triggers to be avoided and help to target medical therapies and plan exercise advice.<sup>6,31</sup> The utility of genetic testing for ICCs commonly encountered in clinical practice is outlined in Table 1.

Genetic testing is incorporated into contemporary exercise guidelines which aim to promote safe participation in exercise through tailored exercise prescription for elite and recreational athletes with ICC.<sup>24</sup> For most conditions, risk stratification and exercise prescription are dictated by the clinical phenotype, allowing for a liberal approach in athletes with a positive genotype but no or very mild phenotype.<sup>24</sup> A notable exception is ACM, where individuals carrying a casual variant, even in the absence of phenotypic evidence of disease, are restricted from the competitive sport as they have a greater chance of progression to overt disease, heart failure, and life-threatening arrhythmias with high-level athletic activity.<sup>24,31</sup>

## Pathway for genetic testing in athletes

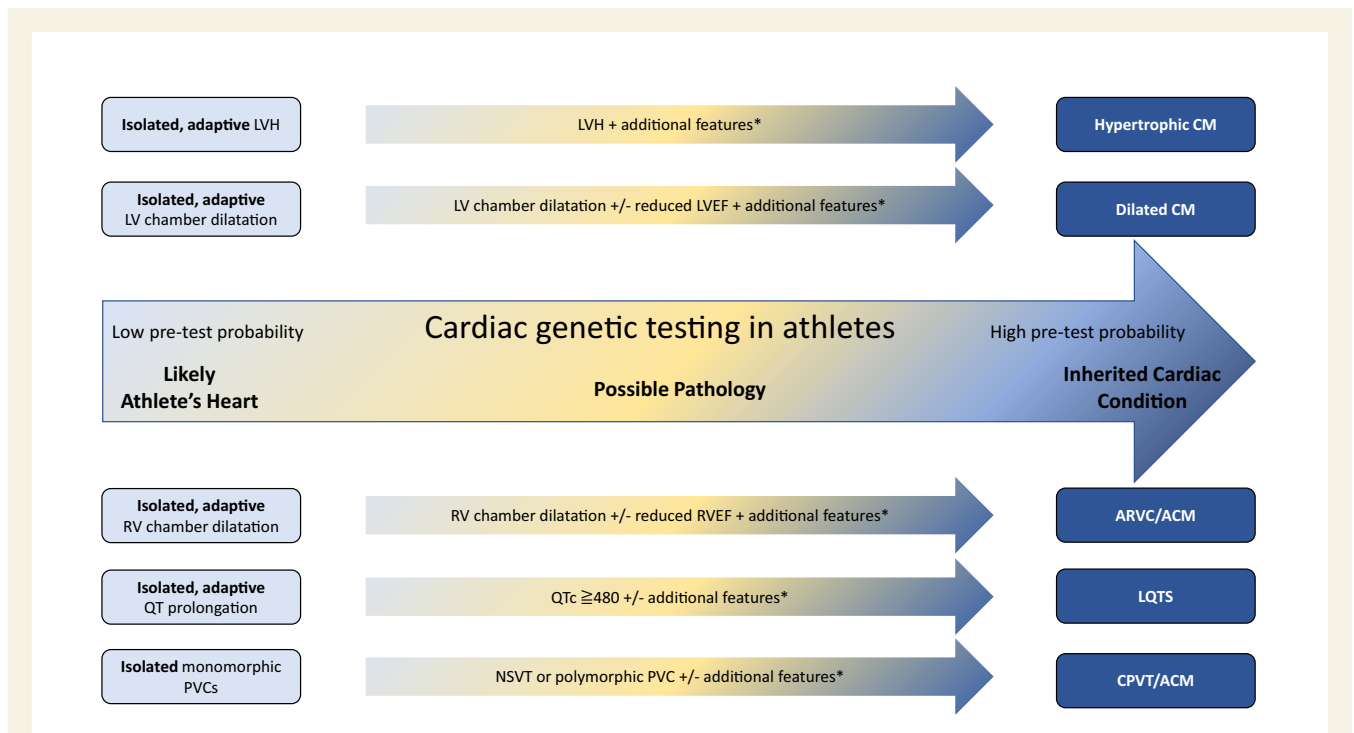
Genetic testing of athletes requires an MDT approach in specialized centres. Individuals with expertise in sports cardiology, cardiac

genetic testing and the diagnosis and management of ICCs should be involved. Key components are outlined in the graphical abstract and include the following:

- (1) *Comprehensive clinical evaluation* to ascertain whether findings indicate physiological or pathological changes in the athlete. Sports physicians, cardiologists, cardiac physiologists, genetic nurses, and clinical geneticists all require expertise pertinent to their roles in the diagnosis of ICCs.<sup>8</sup> In the absence of a clear or highly suggestive phenotype or family history of an ICC there will be a low pre-test probability for identifying a monogenic cause.
- (2) *Pre-test genetic counselling* provides information and psychological support to the athlete and should be performed by appropriately trained health professionals.<sup>24,25</sup> Athletes are supported to make an informed decision, having explored all possible outcomes and respective implications of genetic testing for them and their family. Issues relating to competitive sport participation, insurance, and broader financial and ethical issues should be explored.
- (3) *Expertise in cardiac genetic testing* is important in determining that the correct genes are tested for the right phenotype, as well as accurate interpretation of findings. This requires testing by approved clinical laboratories in collaboration with the MDT. Confining sequencing to high confidence genes<sup>32–35</sup> and careful interpretation of variants minimizes the risk of misclassification of genetic variants and inappropriate feedback to the individual.<sup>36–38</sup> With ever-changing knowledge of genetic variants, reclassification is always a possibility. This can include upgrading a VUS to a likely pathogenic or pathogenic variant or downgrading a variant from likely pathogenic or pathogenic to VUS or even likely benign/benign. The laboratories and MDT should, therefore, have mechanisms in place to ensure variants are periodically reviewed.
- (4) *Post-test genetic counselling* provides the results and reviews potential implications with the athlete. Psychological support may be necessary when an athlete has dealt with significant uncertainty or stress or in cases where there has been exclusion from competitive sports.<sup>39–42</sup> Furthermore, there can be symptoms of anxiety and posttraumatic stress in athletes with a family history of premature SCD, a diagnosis of an ICC or where an implantable cardioverter defibrillator is recommended.<sup>43–46</sup>
- (5) *Clinical follow-up and family evaluation* will be necessary for athletes with a definite or possible diagnosis of an ICC.<sup>16</sup>

## Ethical and legal aspects of genetic testing in athletes

There are specific ethical and legal aspects of genetic testing that need to be considered in competitive and especially professional athletes prior to undertaking the test. The decision about whether to pursue genetic testing should be made following pre-test genetic counselling. Genetic counselling should include all the key stakeholders with whom the athlete wishes to share the decision-making process (family, club doctor, club representative, sporting association) but at the same time, it is important to ensure that decisions are made by the athlete without undue duress. Relevant parties should be fully informed of the possible results and the potential implications prior to undertaking the test. Post-test counselling is critical given the potential psychosocial, financial, and



**Figure 1** Diagrammatic representation of the utility of genetic testing in Sports Cardiology for some of the common clinical scenarios including the 'grey zone'. The system is based on the pre-test probability of a positive yield across the spectrum with on the left adaptive changes due to exercise where genetic testing should not be used and on the right an established diagnosis of an inherited cardiac condition when genetic testing should be performed. A multidisciplinary team discussion is paramount when genetic testing is considered in the 'grey zone' depicted by the arrows. \*Additional features for individual phenotypes as reported in the main text.

**Table 1** Prevalence, estimated genetic yield, and utility of genetic testing in athletic individuals for the most commonly encountered cardiomyopathies and channelopathies (modified from Wilde et al.<sup>6</sup> and Gray et al.<sup>7</sup>)

| Disease under investigation in the athlete | Estimated prevalence <sup>6,8,9</sup> | Estimated genetic testing yield (%) | Diagnostic utility in the athlete | Prognostic utility in the athlete | Therapeutic implications (medical/device) | Impact of exercise on disease expression <sup>a</sup> | Impact on exercise recommendations <sup>b</sup> |
|--|---------------------------------------|-------------------------------------|-----------------------------------|-----------------------------------|---|---|---|
| LQTS                                       | 1 in 2000                             | 70                                  | +++                               | +++                               | +++                                       | +   | ++ <sup>c</sup>                                 |
| CPVT                                       | 1 in 10 000                           | 60                                  | +++                               | +                                 | +   | ++  | ++  |
| BrS  | 1 in 2000                             | 10                                  | +                                 | +                                 | +   | +/-   | -   |
| HCM  | 1 in 500                              | 50                                  | +++                               | ++                                | ++  | +/-   | -   |
| ACM  | 1 in 10 000                           | 50                                  | +++                               | ++                                | ++  | ++  | +++   |
| DCM  | 1 in 500                              | 20                                  | ++                                | +++                               | ++  | +   | +   |

Table scale (-, +, ++, +++) indicates likely utility of genetic testing with '-' indicating unlikely to be helpful and '+++' indicating highly likely to be helpful. LQTS, long QT syndrome; CPVT, catecholaminergic polymorphic ventricular tachycardia; BrS, Brugada syndrome; HCM, hypertrophic cardiomyopathy; ACM, arrhythmogenic cardiomyopathy; DCM, dilated cardiomyopathy.

<sup>a</sup>Degree to which exercise influences how the disease phenotype manifests in the athlete.

<sup>b</sup>How a genetic testing result can influence exercise prescription. The scoring represents a consensus view from the authors based on evidence and existing recommendations.

<sup>c</sup>Specific genotype of LQTS can influence exercise recommendations (e.g. swimming with LQT1).

mental health implications, particularly if the athlete is excluded from play.<sup>23</sup>

The implications relating to insurance for the athlete and their club are largely defined by the country in which the athlete resides. As a general rule, insurance companies require the athlete to disclose any family history but do not require them to disclose the results of genetic testing. In

athletes with a family history of ICC proceeding with predictive testing can be beneficial as the absence of the familial variant effectively excludes the condition and thereby may improve their insurance assessment. It is also worthwhile noting that for some conditions (such as LQTS) genetic testing can also be beneficial by facilitating genotype-directed management and prognosis and ultimately even return-to-play.<sup>23,47-49</sup>

## Special considerations in children and adolescent athletes

### General considerations

In the paediatric athlete genetic testing for ICC should adhere to the same principles as the adult athlete,<sup>50</sup> but requires some specific considerations. Children may not fully understand the future implications of genetic testing. Depending on the child's age, genetic counselling in expert paediatric centres may be necessary. Specialist centres will be able to facilitate a more comprehensive approach to genetic counselling, including the assistance of a child mental health specialist if required. Indeed, the psychological impact of a positive genetic test result may be significant for the child, especially if this leads to sports exclusion even in the absence of a phenotype.

In children with a highly suggestive or established ICC phenotype, diagnostic genetic testing may confirm the diagnosis, facilitate familial cascade screening and in some cases aid with risk stratification.<sup>51</sup> In phenotype negative children with a familial diagnosis of inherited channelopathies, such as LQTS and CPVT, cascade genetic testing may aid with risk stratification and the introduction of effective prophylactic therapies, given the risk of sudden death in the early stages of life. Genetic testing may influence sports participation even in the absence of established phenotype in both LQTS and CPVT where a positive genetic result is sufficient for diagnosis.<sup>24,31,52</sup> On the contrary, the timing of cascade genetic testing in phenotype negative children with a familial diagnosis of inherited cardiomyopathy requires a more individualized approach, depending on the impact of genetic testing on patient management and risk stratification. This cautious approach takes into consideration the significant psychological and social implications of a genetic diagnosis and needs to be balanced with the benefits of a positive genotype, which may lead to more comprehensive clinical evaluation, more frequent monitoring for evidence of disease expression, and offer a better understanding of the individual risk related to sport participation and therefore steer decisions regarding the pursue of a sporting career. There is no strong evidence to date demonstrating that, similar to adults, high-level exercise expedites disease expression in paediatric ACM. However, the possible impact of strenuous exercise, as well as the possibility of sports disqualification if an ACM disease-causing variant were identified, should be taken into consideration when discussing ACM genetic testing in a young athlete.<sup>24,31,53</sup> In addition, certain genotypes such as filamin-C (*FLNC*), desmoplakin (*DSP*), Transmembrane Protein 43 (*TMEM43*), or lamin A/C (*LMNA*) are highly arrhythmogenic and the arrhythmia can pre-date the cardiomyopathy phenotype so early genetic diagnosis may be important.<sup>10</sup>

### Athletes with congenital heart disease

Congenital heart disease (CHD) can occur as a chromosomal syndrome (12%), a heritable syndrome with Mendelian pattern (8%), or a sporadic mutation (80%) with variable penetrance and heritability. A genetic origin has been established for a small proportion (< 5%) of CHDs.<sup>54</sup> Consequently, genetic testing in athletes with CHD should be contemplated in cases with: (i) a strong family history of CHD, (ii) where the syndromic disease is suspected, or (iii) in specific phenotypes with increased risk for cardiac events and known genetic association [e.g. atrial septal defects with arrhythmias (*NKX2.5*)].<sup>55</sup>

## Common clinical scenarios

### Left ventricular hypertrophy and T-wave inversion

Left ventricular hypertrophy (LVH) is a well-documented trait of athletic adaptation. In most cases, the hypertrophy does not exceed 15 mm and is associated with LV cavity dilatation (eccentric remodelling).<sup>24,56</sup> It is more prevalent in black compared with white athletes, males compared with females and individuals who participate in high endurance sports.<sup>56,57</sup> Further evaluation for HCM is necessary, in the absence of hypertension or use of performance-enhancing substances, when the degree of LVH exceeds 15 mm, it is out of proportion to the athlete's demographics and athletic activity or is associated with additional features suggestive of cardiomyopathy.<sup>24,53</sup> Such features include cardiac symptoms, family history of SCD in a first-degree relative under the age of 40 years, ECG abnormalities, asymmetric hypertrophy, and eventually late-gadolinium enhancement suggesting cardiomyopathy on cardiac magnetic resonance (CMR).

In HCM, genetic testing is indicated for athletes with an unequivocal diagnosis and can be used for predictive testing in athletes with a known familial genetic result.<sup>10,53</sup> Occasionally, after an informed discussion with the athlete, genetic testing may be useful in athletes with a highly suggestive but not diagnostic HCM phenotype. The role of genetic testing in risk stratification for HCM is limited and does not usually influence decisions relating to sport participation in individuals with no or mild LVH and no conventional risk factors for SCD.<sup>53</sup>

T-wave inversion on the ECG of an athlete may be a normal variant or may be indicative of underlying heart disease.<sup>29,58–60</sup> For individuals judged to have pathological T-wave inversion, comprehensive clinical evaluation, including CMR imaging, may identify a cardiomyopathy in up to 41% of athletes.<sup>10</sup> In contrast, the diagnostic yield of genetic testing in this setting, is limited to 10% and in most cases is associated with a clinical phenotype.<sup>60</sup> Therefore, the use of routine genetic testing in asymptomatic athletes with T-wave inversion is not advisable in the absence of a family history or other features suggestive of an ICC.

### Ventricular cavity dilatation

Dilation of the left ventricle is a common feature of the athlete's heart, but when it is associated with mildly reduced systolic function and/or non-ischemic scar on CMR, it falls into the grey zone between physiological remodelling and dilated cardiomyopathy (DCM).<sup>53</sup> Comprehensive clinical evaluation in most cases suffices for the differential diagnosis of these two entities, however, genetic testing can aid in prognostic information and management decisions for certain genotypes.<sup>10</sup>

Genetic testing should be contemplated in any athlete who has been diagnosed with a familial DCM, especially when clinical features point to specific genotypes: conduction abnormalities (*LMNA*<sup>61</sup> and *SCN5A*<sup>62</sup>), disproportionate arrhythmic burden and/or multiple cases of sudden cardiac arrest, or SCD within the family (*LMNA*,<sup>59</sup> *SCN5A*,<sup>62</sup> *FLNC*,<sup>63</sup> *DSP*,<sup>64</sup> *BAG3*, *PLN*, *TTN*,<sup>65</sup> and *RBM20*<sup>66</sup>). Although the impact of exercise on arrhythmic DCM forms is not established, recent data from small cohorts suggest that

high-intensity exercise may be contributing to worse outcomes in *LMNA* patients<sup>67</sup> and may exacerbate skeletal muscle dysfunction in *FLNC* carriers.<sup>68</sup> An athlete with mild left ventricular dilatation, no or mild left ventricular dysfunction, prominent non-ischemic myocardial fibrosis, and significant burden of ventricular arrhythmia is most likely affected by a left-dominant ACM and a molecular genotyping for ACM-causing gene mutation, either desmosomal or non-desmosomal is needed to achieve a definite diagnosis, guide management, and prognosis.<sup>24,31,53,69</sup> Finally, cascade genetic testing should be offered in apparently healthy athletes with a familial diagnosis of DCM when an actionable variant has been identified in the family.<sup>70</sup>

## Prolonged QT interval

Athletic individuals tend to exhibit longer QT intervals compared with sedentary counterparts. Consequently, corrected QT (QTc) intervals of up to 470 ms for male and 480 ms for female athletes are accepted as normal. Although a QTc of  $\geq 500$  ms is commonly associated with congenital LQTS, the diagnosis may be challenging in those with lower QTc values in the absence of symptoms or documented family history. Moreover, recently a study suggested the existence of an acquired, exercise-induced QT prolongation phenotype, which is typical of LQTS, but reverts back to normal after a period of de-training.<sup>30</sup> In the cases so far described no arrhythmic events were recorded, however, more data are needed to fully understand the arrhythmic risk in such individuals.<sup>30</sup> A correct diagnosis in athletes is crucial as arrhythmia-related triggers often include increased adrenergic activity, such as exercise or emotional stress,<sup>71</sup> and effective treatments are available.<sup>72</sup>

Genetic testing is an invaluable tool both for diagnosing LQTS as well as informing risk and guiding medical therapy and exercise prescription. The genetic yield is  $>70\%$  in individuals with congenital LQTS<sup>73</sup> and can identify individuals with incomplete penetrance, who display a normal QTc,<sup>71,73–74</sup> but still have a 10% risk of experiencing a cardiac event by the age 40 years.<sup>75</sup> Genetic testing should be offered to all athletes with a familial diagnosis of LQTS and those athletes with a QTc of  $\geq 500$  ms. Genetic testing should also be considered in athletes with a QTc  $\geq 480$  ms regardless of personal and family history. If negative, in the absence of symptoms and/or family history, re-evaluation of the athlete after a period of de-training of at least 3 months and clinical screening of first-degree family members in a specialized cardio-genetic centre, may help distinguish between physiological adaptation and LQTS.<sup>30</sup> Genetic testing in athletes with a QTc  $>440$  ms but  $<480$  ms is justified only in the context of additional features suggestive of a LQTS diagnosis, including symptoms, congenital deafness, family history of unexplained SCD, T-wave notching, documented polymorphic arrhythmias, paradoxical prolongation of the QT interval during exercise, or T-wave alternans.

## Ventricular arrhythmias

Premature ventricular beats in an athlete are fairly common and usually benign. Ventricular arrhythmias (VAs) in young athletes can be classified as common or uncommon, according to their morphology (i.e. site of origin), complexity, response to exercise, and associated clinical findings.<sup>76–78</sup> Genetic testing should only be considered following comprehensive clinical phenotyping,<sup>23</sup> when an underlying

ICC is suspected. In particular, exercise-induced polymorphic or bi-directional VAs should raise suspicion of CPVT.<sup>8</sup> Other ICCs that can present with exercise-induced VAs include ACM,<sup>64,78,79</sup> DCM,<sup>80</sup> and more rarely HCM.<sup>81</sup> The presence of isolated VAs with benign features (i.e. most often left bundle branch block, inferior axis) should not prompt genetic testing.<sup>31</sup>

## Aortic dilatation

Aortic aneurysm is present in up to 1% of the Western population. Thoracic aortic aneurysm and dissection (TAAD) occurs more frequently in young adults compared with the abdominal aneurysms of the elderly. Approximately 20% of TAAD patients have a positive family history of presumed genetic aetiology, with  $>30$  genes currently identified. Cases are typically divided into syndromic and non-syndromic forms.<sup>82–84</sup> The diagnostic yield of genetic testing is estimated to be 20–30% with higher yield in patients with family history.<sup>85,86</sup> In the case of bicuspid aortic valve (BAV) related aortopathy, the diagnostic yield of genetic testing is  $\sim 5\%$ .<sup>86</sup>

Athletes tend to exhibit slightly larger aortic dimensions compared with sedentary individuals. A dilated aortic root, however, beyond conventional normal limits is not a characteristic of the athlete's heart and athletes do not typically demonstrate progressive aortic dilatation.<sup>87,88</sup> Therefore, an athlete presenting with aortic dilatation should undergo counselling and genetic testing according to the general population criteria<sup>83,84,86</sup>: (i) Aortic root diameter  $> 40$  mm in the absence of clear aetiology; (ii) associated features of connective tissue diseases, (iii) TAA with positive family history of aortic dissection ( $<60$  year) or SCD  $<45$  years, (iv) BAV with TAA. When a genetic mutation causing TAA is identified in the family predictive testing is advised in 1st degree relatives, from the age of 10 years in non-syndromic forms and earlier in syndromic presentations. In athletes with a rapid progression of aortic dilatation ( $>3$  mm/year), genetic testing can be considered. In athletes with BAV with normal diameters of the ascending aorta, genetic testing is of little additive value, as diagnostic yield is low.

## Conclusions

Genetic testing in athletes may aid diagnosis, inform arrhythmic risk and prognosis, guide management, including informing athlete's exercise prescription and ultimately facilitate safe 'return to play' to recreational or competitive sport. As genetic testing becomes more widely available there is an increasing expectation that it is considered as part of a comprehensive cardiac assessment in athletes when appropriate. It is, therefore, imperative that health care professionals involved in the athlete's care have an understanding of the indications, as well as the strengths and limitations of genetic testing. An MDT through liaison with an experienced cardio-genetics centre will ensure that both the physician and the athlete are supported. Appropriate pre- and post-test counselling will ensure that the athlete and all the stakeholders understand the potential implications of genetic testing in terms of ethical, legal, and financial repercussions.

## Authors' contributions

S.C., B.G., and M.P. contributed to the conception and design of the work. All authors contributed to drafting and critically revising the manuscript. All authors gave final approval and agree to be accountable for all aspects of the work ensuring integrity and accuracy.

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