Rationale and design of the PROspective ATHletic Heart (Pro@Heart) study: long-term assessment of the determinants of cardiac remodelling and its clinical consequences in endurance athletes

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ABSTRACT
Background Exercise-induced cardiac remodelling (EICR) results from the structural, functional and electrical adaptations to exercise. Despite similar sports participation, EICR varies and some athletes develop phenotypic features that overlap with cardiomyopathies. Training load and genotype may explain some of the variation; however, exercise ‘dose’ has lacked rigorous quantification. Few have investigated the association between EICR and genotype.

Objectives (1) To identify the impact of training load and genotype on the variance of EICR in elite endurance athletes and (2) determine how EICR and its determinants are associated with physical performance, health benefits and cardiac pathology.

Methods The Pro@Heart study is a multicentre prospective cohort trial. Three hundred elite endurance athletes aged 14–23 years will have comprehensive cardiovascular phenotyping using echocardiography, cardiac MRI, 12-lead ECG, exercise-ECG and 24-hour-Holter monitoring. Genotype will be determined using a custom cardiomyopathy gene panel and high-density single-nucleotide polymorphism arrays. Follow-up will include online tracking of training load. Cardiac phenotyping will be repeated at 2, 5, 10 and 20 years.

Results The primary endpoint of the Pro@Heart study is the association of EICR with both training load and genotype. The latter will include rare variants in cardiomyopathy-associated genes and polygenic risk scores for cardiovascular traits. Secondary endpoints are the incidence of atrial and ventricular arrhythmias, physical performance and health benefits and their association with training load and genotype.

Conclusion The Pro@Heart study is the first long-term cohort study to assess the impact of training load and genotype on EICR.

Trial registration number NCT05164328; ACTRN12618007162688.

Key messages
What is already known on this topic
► There is a large unexplained variability in the extent of exercise-induced cardiac remodelling among highly trained endurance athletes.
► Endurance athletes may present with structural, functional and electric features that overlap with cardiomyopathies.
► Endurance athletes have an increased risk of developing atrial fibrillation.

What this study adds
► Insights on the impact of training load and genotype on the variability of exercise-induced cardiac remodelling and its prognostic impact.
► A new gold standard for training load quantification in the field of sports cardiology.
► Knowledge on the phenotypic features that predispose endurance athletes to a higher physical performance and more health benefits; but also to cardiac disease and arrhythmias such as atrial fibrillation.

How this study might affect research, practice or policy
► Stimulate other long-term prospective collaborative trials in the field of sports cardiology that combine comprehensive cardiovascular phenotyping, training load quantification and genotyping.
► Provide a better understanding of the normal spectrum and variability of the athlete’s heart for clinicians.
► Aid clinical practice in identifying athletes at risk of cardiac disease and arrhythmias.

INTRODUCTION
‘Exercise-induced cardiac remodelling’ (EICR) comprehends the structural, functional and electrical adaptations of the heart in response to the metabolic and mechanical demands of exercise.1 Although cross-sectional and some short-term prospective studies have enhanced our understanding...
of EICR, numerous questions concerning the spectrum, determinants and the prognosis of EICR remain unanswered.

The determinants of EICR
Training load
Athletes participating in highly dynamic and static sports (eg, cycling and rowing), accomplishing more training hours per week and performing at the highest level have the largest hearts. However, not all athletes engaging in the same sport and competing at a similar level remodel equally. In a group of 174 non-elite male runners, for example, 16% had concentric left ventricular (LV) remodelling, 11% had non-dilated concentric LV hypertrophy (LVH), 4% had non-dilated eccentric LVH, 9% had dilated eccentric LVH and the remaining 60% had normal LV geometry. A 2004 study by Abergel et al showed a wide scatter of LV internal diameter at end-diastole, maximal wall thickness and LV end-systolic volume indices, indicating that training intensification of training induced significant cardiac biventricular volumes and mass. In seasoned athletes, endurance training in sedentary individuals increased longitudinal trials with supervised exercise programmes, external and internal loads. External loads are objective and record a wealth of data including time, distance, speed, power output and heart rate during exercise. With an electronic training diary linked to the athlete’s heart monitoring and sports GPS device, external and internal training loads from every single training session can be exported and expressed across multiple time domains (eg, per week, month, years).

Genotype
Where training load fails to explain the variability in cardiac structure and function between athletes, genetics may provide insights.

Genetic make-up plays an essential role in cardiac structure and function. The most notable impact of genetics on cardiac morphology is in cardiomyopathies such as hypertrophic cardiomyopathy (HCM), dilated cardiomyopathy (DCM) and ARVC. These are typically caused by single rare variants in genes that encode cardiomyocyte components such as sarcomere and desmosomal proteins. However, the genetic heterogeneity, pleiotropy, variable penetrance and the variable impact of environmental factors result in an important phenotypic variability.

Exercise-induced ARVC serves as a prime example. Our research group has contributed to the recognition of exercise-induced ARVC, characterised by a combination of potentially life-threatening ventricular arrhythmias originating from a dilated and mildly dysfunctional right ventricle (RV) at rest or during exercise. Genetic testing in these athletes revealed a lower than expected prevalence of desmosomal gene mutations, thereby implying a significant contribution of intense exercise to the resulting phenotype. Furthermore, in patients labelled with ‘gene-elusive’ ARVC a higher prevalence of athletes and more intense exercise history was reported. The terms ‘exercise-induced’ and ‘gene-elusive’ ARVC may be considered synonyms, with the former Australian/Belgian terminology emphasising the common feature of high doses of intense endurance exercise and the latter Baltimore terminology emphasising the observation that genetic mutations are seldom identified. As not all endurance athletes develop exercise-induced/gene-elusive ARVC, a solely exercise-induced ARVC is disputed. It remains to be investigated which factors, including genotype, predispose certain athletes towards a phenotypic expression of ARVC when exposed to exercise. The gene-exercise relation is already evidenced by the exacerbated disease progression in genotype positive ARVC due to the mechanical stress of exercise. Interactions between mechanical stress and genetic susceptibility have also been reported for peripartum cardiomyopathy. Although assumed an acquired disorder, recent data have shown an increased prevalence of rare cardiomyopathy gene variants, including truncating TTN variants (TTNv), in women with peripartum cardiomyopathy compared with a reference population. Further, studies in a zebrafish TTNv model have shown reduced tolerance of mechanical stress in titin-deficient hearts. Aside from rare gene
variants, the potential role of more common variants should be emphasised. As an example, common polymorphisms associated with cardiac structure and function have been linked to incident DCM in the general population and may influence the phenotype among carriers of TTN truncating variants.

Genetics also play an essential role in the onset of arrhythmias. There is growing evidence on the role of a genetic predisposition in the onset of AF. Research in cases of early-onset AF and families with AF has identified numerous variants in genes coding for ion channels, transcription factors and myocardial structure, which could be a primary cause of AF, regardless of exercise. Genome-wide associations studies in large populations with AF and controls have identified >100 chromosomal loci associated with AF in which common variants are present. As the prevalence of AF in athletes exceeds that of the general population, the interaction between genetics and environmental factors (eg, exercise) is thought to play a crucial role in phenotypic expression. Of particular interest are the genes encoding for mechanical stress-sensitive cardiac ion channels in which stimuli such as hypertension or endurance exercise alter channel activity. The KCNQ1 gene for example encodes for the cardiac I_Ks channel and a missense mutation was detected in a family with hypertension, atrial dilation and atrial fibrillation. Other mechanical stress-sensitive ion channels include two-pore domain K+ channels (eg, TREK1, TREK2, TRAAK), transient receptor potential channels (eg, TRPM4, TRPC3, TRPC6) and the Piezo1 channel.

Sex
Besides training load and genetics, cross-sectional studies suggest sex-differences in the type and extent of EICR. Absolute measures of LV and RV dimensions and mass are smaller in females, but when corrected for body surface some echocardiographic studies have reported larger indexed end-diastolic diameters in female athletes than in males. A recent study using cardiac MR (CMR) showed lower indexed LV and RV volumes in adolescent and adult female athletes with a higher tendency towards eccentric remodelling as evaluated by volume to mass ratios. Functionally, while no sex-differences were noted for LVEF and RVEF in adolescent athletes, RVEF was lower in adult males compared females. The latter could suggest that RV function remains stable from adolescence to adulthood in females but decreases in males. Using deformation imaging, LV and RV global longitudinal strain was higher at rest and increased more during exercise in female athletes compared with males, thus further supporting a potential higher susceptibility towards RV dysfunction in male athletes. Whether the sex-differences in EICR contribute to the reduced arrhythmogenic risk in female athletes remains unclear.

Ethnicity
As ethnic diversity in sports has grown over the years so has the relevance and understanding of the role of ethnic background on EICR. While ventricular dilation is common in white athletes with 14% having an LV diameter of >60 mm on transthoracic echocardiography, significant hypertrophy such as an LV wall thickness of >12 mm is uncommon. In black athletes the degree of ventricular dilation is similar to white athletes whereas an in increased in LVWT >12 mm is more frequent and seen in up to 12% of black athletes. Black athletes generally present with a higher systolic blood pressure at rest and during exercise which could explain the higher prevalence of LVH. Additionally LV hypertrabeculation is more frequent in black athletes than in white athletes. These differences in structural remodelling between white and black athletes is also illustrated on the ECG with anterior T-wave inversions being more common in black athletes and considered a normal ethnic variant. With regard to East Asian athletes a slight distinction can be made between Chinese and Japanese athletes. The former have shown similar cardiac dimensions as compared with white athletes whereas the latter have greater LV end diastolic dimensions. In Arab athletes, in comparison to white and black athletes, smaller cardiac dimensions have been measure with a similar degree of LVH as white athletes. Finally in Pacific Islanders a higher LV mass was seen when compared with white athletes. Whether differences in cardiac remodelling between ethnic population is based on differences in genotype, training load or other factors remains to be explored.

The prognosis of EICR
Despite the many health benefits of endurance exercise, athletes are not granted immunity from cardiovascular disease.

Atrial fibrillation
Atrial fibrillation is the most common arrhythmia and has a U-shaped dose–response relationship with endurance exercise. On the one hand, low to moderate intensity exercise has been associated with a decreased risk of AF. On the other hand, a higher incidence of AF was seen when endurance exercise was performed more frequently (ie, >4/week) and longer (ie, >5 hours/week) or when a lifetime exercise history of >2000 hours was met. A 2009 meta-analysis calculated that endurance athletes are 5.3 times more likely to develop AF than controls. Part of the propensity for AF in high-level endurance athletes is thought to arise from the structural and functional atrial adaptations. A recent study by Trivedi et al investigated the differences in atrial structural and functional properties between healthy non-athletes, non-athletes with paroxysmal AF, healthy endurance athletes and athletes with paroxysmal AF. Diastolic dysfunction, LA dilation and reduced LA strain characterised non-athletes with AF whereas athletes, regardless of AF, had a normal diastolic function, increased LA volumes and
decreased LA strain. However, athletes with AF had lower LA emptying fraction and LA expansion index, which are suggestive of atrial cardiomyopathy.57 This study highlights the difficulties in distinguishing the healthy athlete’s dilated atrium for one that is predisposed to AF.

Ventricular arrhythmias and sudden cardiac death
Sudden cardiac death (SCD) is an uncommon event in athletes and occurs primarily through ventricular arrhythmias. The incidence of SCD ranges between 0.4 and 8.4 per 100 000 athlete-years.58 ARVC and other inherited cardiomyopathies such as HCM and DCM can be highly arrhythmogenic and are among the most common causes of SCD in young athletes.59–60 A particular challenge in sports cardiology is the overlapping phenotype of EICR and these cardiomyopathies. Although research has granted clinicians strategies to distinguish disease from physiological adaptation a better understanding of the extent and mechanisms of EICR is required.23–61–65

Another putative risk factor associated with SCD in athletes is myocardial fibrosis.60 Reports on myocardial scarring in athletes are heterogeneous regarding prevalence, location, extent, aetiology and clinical significance. Coronary artery disease (CAD)-patterns and non-CAD subepicardial/midmyocardial fibrosis have been associated with potentially life-threatening arrhythmias such as non-sustained ventricular tachycardia but also ventricular tachycardia leading to sudden death or an appropriate shock by an implanted defibrillator.66–67 In contrast, hinge point fibrosis associated with cumulative exercise exposure, such as years of training and the lifetime number of races and race distances is considered a benign consequence of RV overload.16–18

The majority of SCD in athletes are classified as sudden unexplained deaths in which the role of plain EICR as potential triggers and/or substrates of arrhythmias is unknown.68–70

Physical performance and health benefits
Despite the increased risk of arrhythmias in some highly trained individuals, there is no denying the numerous health benefits associated with endurance exercise in the general population. Endurance exercise is associated with a lower prevalence and better control of cardiovascular risk factors such as arterial hypertension, dyslipidaemia, obesity and diabetes mellitus.68–71 Increased longevity is the strongest health outcome associated with exercise. Regular physical exercise at low to moderate intensity, even in small doses, reduces all-cause and cardiovascular mortality.72–73 These findings can be extended to higher levels of exercise with Olympic athletes demonstrating an increased life expectancy of up to 6.5 years compared with the general population.74–78 Research in Tour de France participants from 1947 until 2012 has similarly reported a 41% lower mortality rate in cyclists for both malignancies and cardiovascular causes.79 However, the cited research is limited by its retrospective design and focus on elite athletes who on average were over 20 years old thereby being potentially affected by the ‘survival of the fittest’ selection bias.

The time required to cover a given distance defines physical performance of an endurance athlete. Physiologically, the latter can be brought back to the velocity and power generated by the athlete, with VO₂max being one of the major determinants. Elite endurance athletes have the highest VO₂max, with values of more than 70 and up to 85 mL/kg/min, equalling 150%–200% of the VO₂max seen in healthy active young individuals.80 Alongside capillary density and mitochondrial density, the high levels of VO₂max have also been attributed to EICR such as increased cardiac size and stroke volume.81 For the elite athlete, the ultimate measure of physical performance is the level of competitiveness, the number of finals reached and Olympic Games participated in, or the amount of victories, trophies and medals obtained. It is unknown whether and/or which features of EICR, including pathology such as AF and myocardial fibrosis, affect VO₂max and physical performance. How much do alterations in training regimen explain differences in physical performance? To which extent does genotype play a role in predetermining gold medallists and Olympic champions? Finally, looking at both genotype and phenotype the question raises whether you can outtrain unfavourable genetics?

In conclusion, much of the variability, determinants and the outcome of EICR has yet to be determined. Prospective long-term studies that investigate EICR in large cohorts of athletes are needed to improve knowledge in these areas of sports cardiology. The Pro@Heart study is a multicentre prospective cohort trial designed to assess and follow-up all aspects of EICR in young competitive endurance athletes. The interplay between training load and genotype will be investigated and insights on the long-term outcome of EICR will be gained.

THE PRO@HEART STUDY
Trial design
The Pro@Heart (Prospective Athlete’s Heart) study is an international multicentre prospective cohort trial with collaborators in Australia (Baker Heart and Diabetes Institute Melbourne, St Vincent’s Hospital Melbourne, University of Adelaide, Royal Adelaide Hospital, Royal Melbourne Hospital and Victor Chang Cardiac Research Institute) and Belgium (University Hospitals Leuven, University Hospital Antwerp and Jessa Hospital Hasselt). (ClinicalTrials.gov Registry Identifier NCT05164328—Australia New Zealand Clinical Trials Registry Identifier ACTRN12618000716268).

Objectives and hypothesis
The primary objective of the Pro@Heart study is to investigate the impact of training load (ie, accurately evaluated in term of type, frequency, duration and intensity) and genotype on the variability of structural, functional and electrical EICR in young competitive endurance athletes. The second objective is to determine how EICR,
training load and genotype are associated with physical performance, health benefits and cardiac pathology (e.g., exercise-related cardiomyopathies and arrhythmias) during follow-up over several decades. The hypotheses are that: (1) Genetic factors contribute to variability in EICR beyond training load and, (2) Athletes with extreme EICR have a higher risk of arrhythmias. The study hypotheses are illustrated in Figure 1.

Study population and eligibility criteria
Athletes aged 14–23 years are recruited from elite endurance sports programmes and organisations including Cycling Vlaanderen, Triathlon Vlaanderen, Belgian Cycling, Cycling Australia, Rowing Australia and Athletics Australia and from sport performance centres including Nottebohm Antwerpen, Bakala Academy Leuven and Adlon Hasselt. Volunteers have also been recruited by means of ‘word of mouth’ and social media campaigns within elite endurance sporting circles. Non-athletes consist of age and gender matched university and college students recruited from Australian and Belgian universities in addition to social media campaigns. Subjects will be included based on the eligibility criteria listed in Box 1. Informed consent will be obtained from the participants or their legal guardian.

A young age at inclusion will ensure a baseline evaluation at the beginning of the athlete’s career, at a time point of relative low cumulative training load. Moreover, it will allow close follow-up of the evolution of EICR over years, in correlation with training load, genotype, cardiovascular disease and athletic performance. The age range of 14–23 years was considered the best compromise between enrolling athletes with relatively low cumulative training load and sufficient commitment to high-level endurance training, minimising the risk of dropout. To ensure a high level of athleticism, individuals must be competing at a national or international level for at least 2 years in a sport with a high dynamic and moderate-high static training components performed for sustained periods. Hence triathlon, cycling, distance running (≥1500 m), rowing, swimming of 400 m or more and cross-country skiing (which includes biathlon and Nordic combined) were sports eligible for inclusion. The distances for running and swimming were selected based on the characteristic parabolic pacing profile of endurance exercise (i.e., a fast-paced start, a slower and stable-paced middle section and finally a sprint). As cardiac volumes are similar between sedentary individuals and subjects exercising <3 hours per week, a cut-off of <3 hours per week of physical activity defined non-athletes.

Study procedures
Baseline evaluation includes medical history, review of medication and supplements, physical examination with blood pressure measurement on both arms and blood samples for biochemistry and DNA isolation. Genetic variants will be identified using a custom cardiomyopathy gene panel and the Axiom Precision Medicine Diversity Array (Thermo Fisher Scientific, Waltham, Massachusetts, USA). Rare variants will be assessed using the American College of Medical Genetics and Genomics guidelines.
College of Medical Genetics and Genomics pathogenicity criteria. Polygenic risk scores for various traits will be derived from suites of common variants. A resting 12-lead ECG will be recorded and interpreted in accordance with international recommendations.

Two-dimensional and three-dimensional transthoracic echocardiograms (TTE) and CMR with the administration of gadolinium as a contrast agent including native and postcontrast T1 mapping and will be performed at rest to assess ventricular systolic and diastolic function, atrial function, extracellular volume and myocardial tissue characterisation (ie, myocardial fibrosis). Cardiopulmonary exercise testing for maximal oxygen consumption measurement (VO2max) including 12-lead ECG recording will evaluate athletic performance and exercise-related repolarisation abnormalities and arrhythmias. Dual-energy x-ray absorptiometry will assess body composition. Twenty-four-hour Holter monitoring will determine heart rate limits and the prevalence of arrhythmias. In addition to resting cardiac imaging, all recruits will also undergo exercise imaging to evaluate contractile reserve, using either two-dimensional TTE or CMR.

Follow-up will include annual telephone communication enquiring about clinical events as well as continuous monitoring of training load using an electronic training diary. The duration, distance, altitude, speed, power output and heart rate during training sessions recorded by sports GPS trackers will be exported to a big data platform. Measured external and internal training loads will be combined into a single parameter quantifying training load as a composite of duration and intensity, such as the Bannister training impulse (TRIMP) and derivatives. The permission to track all sports activities using a central coach profile will be asked to the athlete or their legal guardian. All exported raw data will be pseudonymised in accordance with current general data protection regulation guidelines.

The study flow chart is presented in figure 2. A detailed overview of study procedures is provided in online supplemental appendix.

Endpoints
The primary endpoint of the Pro@Heart study is the association of EICR (eg, ventricular hypertrophy, dilation, reduced function and/or myocardial fibrosis) with both training load, quantified as a combination of duration and intensity, and genotype such as the prevalence of rare and common variants in cardiomyopathy-associated genes and polygenic risk scores. Secondary endpoints are the (1) long-term incidence of arrhythmias such as AF and atrial flutter as well as ventricular ectopic beats, non-sustained and sustained ventricular tachycardia, (2) the physical performance (eg, VO2 max and race results) and (3) the health benefits (eg, longevity, cardiovascular risk factors) and their association with the different phenotypes of EICR, training load and genotype.

STATISTICAL CONSIDERATIONS
Sample size and power calculations
The study is powered for primary and secondary objectives.

Regarding the primary objectives, using an average indexed LV end-diastolic volume of 113±19mL/m² from preliminary data and considering that 43% of the variance in LVEDV can be explained by all single-nucleotide polymorphisms (SNP) on a genotyping array, a population of 220 athletes would render a statistical power over 90%.

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end-diastolic volume of 125±22mL/m² and considering the same variance in RVEDV to be explained by all SNPs on a genotyping array, a population of 220 athletes would give a statistical power of >90%.

For the secondary objective of AF, based on the 2009 meta-analysis by Abdulla and Nielsen reporting a prevalence of AF of 23% in 655 athletes and 12.5% in 895 non-athletes aged 51±9 years, a sample size of 300 endurance athletes and 150 non-athletic controls would provide 80% power and 5% probability of a type I error in detecting a difference in AF prevalence between athletes and non-athletes during long-term follow-up.

Statistical analysis
Groups will be defined based on the types of EICR being investigated. Differences between baseline and follow-up will be compared using either a related-samples Wilcoxon signed rank test or a paired-samples t-test. Independent samples will be compared using either a Mann-Whitney U test or an independent-samples t-test as appropriate. Categorical data will be compared using a Related-Samples McNemar test. In case of three or more groups being defined repeated-measures ANOVA or a Friedman test will be used to analyse the continuous outcomes. Cochrane Q test will be used for dichotomous outcomes. The risk of type I errors due to multiple comparison will be assessed and corrections using the Bonferroni method or Benjamini-Hochberg procedure will be considered. Multiple linear, non-linear and logistic regression analysis will be used to determine the association between cardiac remodelling (including adverse remodelling), the presence of potential pathological variants in cardiomyopathy associated genes and training load. Differences between gender and ethnicities will be addressed either by subgroup analyses comparing athletes with different gender and ethnicity or by incorporating gender and ethnicity as an independent variable in multivariate analyses. Cumulative event rates will be calculated and presented using Kaplan-Meier time-to-event curves.

TRIAL STATUS
The Pro@Heart trial has been approved by the ethical committees of the University Hospitals Leuven, University Hospitals Antwerp, Jessa Ziekenhuis Hasselt and Baker Heart and Diabetes Institute. The first subject gave informed consent and took part in the trial on 23 June 2015. To date, over 220 athletes have been included, more than 70 athletes have had their 2-year follow-up appointment and the 5-year follow-up visits are being scheduled. Inclusions are ongoing. Trial results will be communicated through publications. Published articles will be available to the community on www.proatheteartbe.

CONCLUSION
Cross-sectional and short-term longitudinal trials have offered a better understanding of EICR and exposed the large variability in clinical phenotypes, including cardiovascular pathology but also the health and performance benefits. However, few studies have investigated the potential determinants and the long-term prognosis of EICR in a prospective manner. While training load may explain some of this variability, to date, the quantification of training load has primarily been reliant on subjective recall. The role of genotype in cardiomyopathies and certain cardiac arrhythmias is clear but its importance in EICR and arrhythmias in athletes remain to be investigated.

The Pro@Heart study is a multicentre prospective cohort study that will combine comprehensive phenotyping with long-term follow-up to provide new insights on the spectrum and variability of EICR as well as how training load and genetics determine cardiac structure, functional and electrical properties, including arrhythmias in endurance athletes.

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