

RESEARCH ARTICLE

Applying Personal Genetic Data to Injury Risk Assessment in Athletes

Gabrielle T. Goodlin¹, Andrew K. Roos², Thomas R. Roos², Claire Hawkins³, Sydney Beach³, Stephen Baur¹, Stuart K. Kim^{1*}

1 Departments of Developmental Biology and Genetics, Stanford University Medical Center, Stanford, CA, 94305, United States of America, **2** Division of Epidemiology, Department of Health Research and Policy, Stanford University School of Medicine, Stanford, CA, 94305, United States of America, **3** Department of Human Biology, Stanford University, Stanford, CA, 94305, United States of America

* stuartkm@stanford.edu



CrossMark
click for updates

OPEN ACCESS

Citation: Goodlin GT, Roos AK, Roos TR, Hawkins C, Beach S, Baur S, et al. (2015) Applying Personal Genetic Data to Injury Risk Assessment in Athletes. PLoS ONE 10(4): e0122676. doi:10.1371/journal.pone.0122676

Academic Editor: Alejandro Lucia, Universidad Europea de Madrid, SPAIN

Received: May 2, 2014

Accepted: February 24, 2015

Published: April 28, 2015

Copyright: © 2015 Goodlin et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data is available upon request from Stanford University School of Medicine Department of Genetics; please contact Dr. Stuart Kim: Beckman Center B300 279 Campus Drive Stanford, CA 94309 t:(650) 725-7671 f:(650) 725-7739 stuartkm@stanford.edu

Funding: This work was supported by a Spectrum grant from NIH/Stanford University (#S13-064). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Abstract

Recent studies have identified genetic markers associated with risk for certain sports-related injuries and performance-related conditions, with the hope that these markers could be used by individual athletes to personalize their training and diet regimens. We found that we could greatly expand the knowledge base of sports genetic information by using published data originally found in health and disease studies. For example, the results from large genome-wide association studies for low bone mineral density in elderly women can be re-purposed for low bone mineral density in young endurance athletes. In total, we found 124 single-nucleotide polymorphisms associated with: anterior cruciate ligament tear, Achilles tendon injury, low bone mineral density and stress fracture, osteoarthritis, vitamin/mineral deficiencies, and sickle cell trait. Of these single nucleotide polymorphisms, 91% have not previously been used in sports genetics.

We conducted a pilot program on fourteen triathletes using this expanded knowledge base of genetic variants associated with sports injury. These athletes were genotyped and educated about how their individual genetic make-up affected their personal risk profile during an hour-long personal consultation. Overall, participants were favorable of the program, found it informative, and most acted upon their genetic results.

This pilot program shows that recent genetic research provides valuable information to help reduce sports injuries and to optimize nutrition. There are many genetic studies for health and disease that can be mined to provide useful information to athletes about their individual risk for relevant injuries.

Introduction

The regular demands of training and competition during a season make professional, collegiate, and recreational athletes alike highly susceptible to injury. From 1988 through 2003, the incidence rate of injuries in NCAA athletes was 15.47 per 1,000 athlete exposures[1]. Recreational distance running causes high numbers of injuries, with incidence rates estimated

Competing Interests: The authors have declared that no competing interests exist.

between 30% and 75% per year[[2,3](#)]. Similarly, up to 74.8% of age-group triathletes that participate in Ironman-distance races will be injured at least once each training season[[4,5](#)]. Avoiding injuries and staying healthy is key to a team or individual athlete's success[[6](#)].

Recent genetic research has revealed specific markers associated with increased risk for sports injuries and performance-related conditions[[7,8,9](#)]. Use of this genetic information may aid in the development of tailored injury prevention program for athletes, which could provide a new edge for successful competition[[10](#)]. For example, 100 genetic loci involved in sports performance were analyzed for a professional soccer team in the English Premier League, and the English Institute of Sport expressed interest in providing genetic testing to Britain's Olympic athletes[[11](#)]. As of 2012, at least thirteen direct-to-consumer genomic companies offer genetic testing to athletes and pilot studies of the genomic profiling of soccer players is underway [[10,12,13](#)].

In this paper, we examine the possibility of applying genetic knowledge about health and disease to predicting risk of sports injury in athletes. We performed a literature search on genetic studies in three categories of health that had not been previously used in the context of sports genetics: stress fracture, vitamin and mineral deficiencies, and osteoarthritis. We found a large number of DNA polymorphisms (113) in these categories, along with 11 polymorphisms that had been previously associated with athletic injuries (anterior cruciate ligament rupture, Achilles tendon injury, and sickle cell trait). We chose to focus on injuries whose risk factors could be addressed by modifications in training or diet, rather than non-actionable injuries, such as concussions. Through our literature search, we identified 113 SNPs that have previously not been discussed in relation to sports injuries and performance. A new sports genetic database was constructed, and used to help customize injury prevention and training for fourteen athletes. Fourteen elite triathletes from the Stanford University Triathlon Team were genotyped and given information that could lead to individualized changes in training routines and injury prevention programs.

Materials and Methods

Genome-wide association studies (GWAS) from the NHGRI GWAS Catalog and candidate gene studies from PubMed (MEDLINE) were selected for their relevance to athletic health: bone mineral density, osteoarthritis, calcium levels, vitamin E, vitamin D, vitamin B12, folate pathway vitamin levels, homocysteine, and phytosterol levels. Single nucleotide polymorphisms (SNPs) from genome-wide studies that reached genome-wide significance were accepted, along with SNPs from candidate gene studies. The studies were rated according to the criteria established for the assessment of cumulative evidence on genetic associations and effect size estimates were based on odds ratios reported for published SNPs associated with each category (OR 1.0–1.3 = small effects, 1.3–2.0 = medium effects, and >2.0 = large effects)[[14](#)].

We wanted to test our sports genetics database on a small set of elite athletes and recruited nineteen athletes on the Stanford University Triathlon Team based on their commitment to training and athletic ability ([Table 1](#)). Athletic ability was defined as those who qualified to compete at the 2012 USAT Collegiate National Championship or equivalent. Athletic injuries were defined according to the NCAA standard of missing at least one day of practice or competition due to a diagnosed injury[[1](#)]. At Stanford, triathlon is a club sport that is scored by individual performances, which minimizes ethical complications arising from genetic discrimination affecting athletic scholarships or starting lineups. This set of elite athletes is too small to perform a rigorous trial to show the effectiveness of the sports genetics program in reducing injury incidence. Our goal was to simply observe their responses to the new genetic knowledge. Of the nineteen recruited, fourteen consented to participate in this pilot program

Table 1. Summary of Athlete Cohort.

	N (%)	Age (yrs)	Varsity (%) *	Training (hrs/week) **	Injury 2012 (%) ***	Injury 2013 (%) ***
All	14 (100%)	24.8 [20–38]	85.7%	13.1 [5–20]	71.4%	33.3%
Male	9 (64.3%)	25.6 [20–38]	88.9%	-	66.6%	22.2%
Female	5 (35.7%)	23.6 [20–29]	80.0%	-	80.0%	40.0%

Note: Numbers listed are the mean from all athletes in the cohort and numbers in brackets are the min-max range for those characteristics.

*: Varsity status is defined as whether or not the athlete participated on an NCAA Div 1, II, or III Varsity level team (swimming, cross-country, track-and-field, crew) for at least 1 year prior to being on the Stanford Triathlon Team.

**: Self-reported average training hours per week during the collegiate season from Sept 1st, 2012 to May 1st, 2013. Training was periodized, includes off days plus rest weeks, includes structured team plus individual workouts, and is not limited by discipline.

***: Injury status is defined as any athlete who was limited in their training, could not participate in team workouts, or was unable to race for at least one day due to a diagnosed injury from Sept 1st to Aug 31st of each season.

doi:10.1371/journal.pone.0122676.t001

and athletes were genotyped through 23andMe, a direct-to-consumer company. The raw data file was used to identify the genotype at rsIDs that were found to be associated with sports injury risk based on our data mining of published literature. We conducted an hour-long personal consultation for each athlete, during which we explained how their personal genetic data affected their risk in each of the six sports injury categories. If an athlete was at risk in any category, we provided information about interventions to reduce that risk.

Three and twelve months after their consultation meeting, athletes were sent surveys using REDCap (Research Electronic Data Capture) software[15] to assess genetic information sharing, psychological effects, behavioral changes in training and/or diet, and to track injuries. Ten and thirteen out of fourteen participants filled out the three-month and twelve-month surveys, respectively ([S1 File](#)).

Ethics Statement

The pilot program protocol was approved by the Research Compliance Office at Stanford University (IRB Protocol 24576). Written informed consent was obtained from all athletes to participate in the study, as well as verbal consent to use their deidentified information for study purposes.

Results

The literature review identified a total of 124 SNPs in the following categories: bone mineral density (BMD), osteoarthritis, vitamin and mineral deficiencies, ACL rupture, Achilles tendon injury, and sickle cell trait (SCT) ([Fig 1](#)).

We analyzed the raw genotype files for SNPs associated with the six types of injuries identified in the extensive literature review ([Table 2](#)). These injury categories were selected for their relevance to athletes ([Table 3](#)), and because they are actionable; that is, athletes can modify their training and/or diet to reduce their overall risk. After the raw genotype files were analyzed, athletes went through an hour-long confidential consultation session. If an athlete was found to be at increased risk for a specific injury category, they were given information on preventative measures that could be taken for that specific injury. [Fig 2](#) shows an example of the profiles given to the athletes.

The first injury category, rupture of the ACL, is one of the most severe injuries sustained in active populations[16]. Candidate gene studies performed on athletes have found SNPs in the *COL1A1*, *COL5A1*, *COL12A1*, and *MMP12* genes that are associated with ACL tears

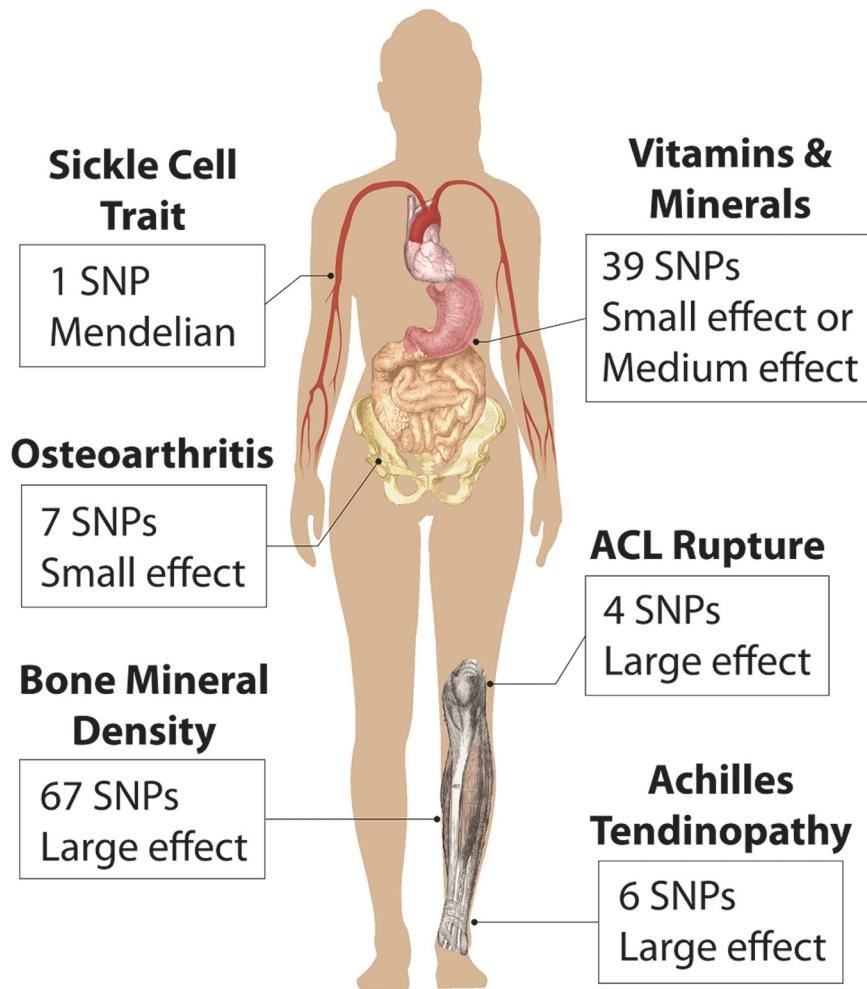


Fig 1. Areas of Interest for Genetic Markers. Six sports related categories were tested in athletes that relate to different injuries or attributes in different locations of the human body. For each category we list the number of associated single nucleotide polymorphisms (SNPs) that we reported on as well as the overall effect size, based on odds ratios or β -coefficients, for having a genetic risk in that category.

doi:10.1371/journal.pone.0122676.g001

[16,17,18,19,20,21]. Although these studies were performed with small sample sizes, the SNPs associated with ACL rupture have high allelic odds ratios between 2.4 and 50. One Stanford tri-athlete had a genetic score that indicated protection against ACL tears, while several athletes were found to be at risk. As this injury is rare in endurance athletes, participants did not express concern about their genetic predisposition for ACL tears during their consultations. Dynamic warm-up programs that target proper landing and cutting form, core strength, and hamstring strength have been shown to be effective at reducing ACL injuries[22].

The second injury category, Achilles tendon injury, includes degenerative and painful conditions that affect athletes in a wide range of sports, including up to 18.5% of runners [23,24,25]. Candidate gene studies have found variants within the *MMP3*, *COL5A1*, *COL1A1*, *CASP-8*, and *GDF5* genes to be associated with Achilles tendon pathologies[26,27,28,29,30,31]. Interestingly, the polymorphism rs1800012 is located within a regulatory region of the promoter of *COL1A1*, which encodes type I collagen, the major protein component of tendons and ligaments[32,33]. The rare T allele increases expression of the gene, whereas the G allele lowers expression and is associated with increased risk for Achilles tendon rupture (OR = 11)[18].

Table 2. Summary of Findings from the Genetic Literature Review.

Injury/Trait	Number of SNPs	Effect Size *	Level of Evidence **	Study Type	References
ACL Rupture	4	Large	Weak	Candidate Gene	[16–21]
Achilles Tendon Injuries	6	Large	Weak	Candidate Gene	[26–31]
Bone Mineral Density	67	Large	Strong	GWAS	[41–44]
Osteoarthritis	7	Small	Strong	GWAS	[45, 48–49]
Vitamin/Mineral Deficiencies	39	-	-	-	-
Iron Biomarkers	8	Small	Moderate	GWAS	[51, 52]
Vitamin E	3	Small	Moderate	GWAS	[53, 54]
Vitamin D	6	Medium	Moderate	GWAS	[55, 56]
Calcium	2	Small	Strong	GWAS	[57, 58]
Magnesium	4	Small	Moderate	GWAS	[59]
B Vitamins	7	Medium	Moderate	GWAS	[60, 61]
Homocysteine	8	Small	Moderate	GWAS	[62, 63]
Phytosterols	3	Small	Moderate	GWAS	[64]
Sickle Cell Trait	1	Mendelian	Strong	Candidate Gene	[67, 68]

*: Effect size estimates are based on odds ratios reported for published SNPs associated with each category. Odds ratios of 1.0–1.3 are small effects, 1.3–2.0 are medium effects, and greater than 2.0 are large effects. Varsity status is defined as whether or not the athlete participated on an NCAA Div I, II, or III Varsity level team (swimming, cross-country, track-and-field, crew) for at least 1 year prior to being on the Stanford Triathlon Team.

**: Level of evidence based on criteria for assessment of cumulative evidence of genetic associations from Ioannidis et al 2008 [14].

doi:10.1371/journal.pone.0122676.t002

Standard practices to prevent Achilles tendon injuries include stretching to increase flexibility, plus eccentric strengthening of the gastrocnemius and soleus muscles[34,35,36]. Three triathletes were identified with a genetic risk for Achilles tendon injuries, two of which had a history of lower leg muscle issues (gastrocnemius and flexor digitorum longus strains). To reduce the risk of an Achilles tendon injury, these athletes began gastrocnemius/soleus eccentric strengthening, calf and Achilles stretches, and shifted to running on softer surfaces.

The third injury category, stress fracture, is a common overuse injury in repetitive sports, and affects up to 20% of female athletes and military recruits[37]. Bone mineral density is a major determinant in a bone's ability to withstand loading and low BMD is an intrinsic risk factor for stress fractures in athletes[37,38]. It has been shown that genetics determine 50–85% of the variance in BMD[39]. Active, young females reporting a family history of osteoporosis or low BMD were almost twice as likely to develop a stress fracture[40]. Estrada *et al.*

Table 3. Relevance of Selected Injuries and Conditions to Athletes.

Injury/Trait	Relevance to Athletes
Anterior Cruciate Ligament Rupture	A severe sports-related injury sustained in multiple landing and cutting/pivoting sports.
Achilles Tendinopathy	Common tendinopathies among athletes from repeated overloading of the tendon.
Low Bone Mineral Density	A major risk factor for stress fractures and stress reactions among athletes.
Osteoarthritis	The onset of osteoarthritis is related to joint injuries, which are common among athletes.
Vitamin/Mineral Deficiencies	Minor mineral deficiencies and chronic low vitamin levels can impair athletic performance.
Sickle Cell Trait	SCT can cause potentially fatal complications for athletes when training under extreme and stressful conditions.

doi:10.1371/journal.pone.0122676.t003

Genetic Profile

Trait/Condition	Summary
Bone Mineral Density	Average risk for low bone mineral density
Osteoarthritis	Average risk for osteoarthritis
Achilles Tendinopathy	Increased risk for Achilles tendinopathy
ACL Rupture	Average risk for ACL rupture
Calcium Levels	Increased risk for low calcium levels
Vitamin D	Increased risk for vitamin D deficiency
Vitamin E	Decreased risk for vitamin E deficiency
B Vitamins	Slightly increased risk for B12, B9 and B6 vitamin deficiencies
Magnesium Levels	Average for magnesium levels
Homocysteine Levels	Decreased risk for elevated homocysteine levels
Phytosterol Levels	Average for phytosterol levels
Iron Status	Increased risk for low iron
Sickle Cell Trait	No sickle cell trait

Fig 2. Example Summary of an Athlete's Genetic Profile. Each athlete was given information related to the categories tested. The summary page gives four color-coded risk levels for each trait: decreased risk (green), average (black), slightly increased risk (yellow), or increased risk (red). Further information, including background information, injury mechanism, genetic basis, and prevention strategies was accessible by clicking on the category in the side menu.

doi:10.1371/journal.pone.0122676.g002

performed a large meta-analysis on lumbar spine and femoral neck bone mineral density, including 17 GWA studies involving a total of 32,961 individuals. They identified 64 SNPs associated with bone mineral density at genome-wide significance. An algorithm was developed summing the individual contributions from 63 of these SNPs into one combined genetic score. These scores were divided into five different risk categories based on the population distribution[41]. Individuals in the highest risk category have 1.56 increased odds for osteoporosis, and 1.60 increased odds for fracture. Conversely, individuals in the lowest category were protected against osteoporosis and fracture and had decreased odds of 0.38 and 0.54, respectively[41]. Three additional studies found three SNPs that are also significantly associated with bone mineral density or fracture that were not included in the 2012 Estrada *et al.* paper[42,43,44].

We reported the combined genetic score developed by Estrada *et al.*, as well as the individual SNP results, for a total of 66 SNPs. Two participants had genetic scores on the extreme ends of the risk groups established by Estrada *et al.* The triathlete with the worst genetic risk score had a history of three previous stress fractures during his/her career as a collegiate runner. A DEXA scan, which directly measures BMD, indicated osteopenia for this athlete. The triathlete with the most protective score had no history of stress fracture, and only one stress reaction so far as a cross-country runner. Triathletes can easily redistribute training volume to cycling or swimming if they are predisposed to running injuries, such as stress fractures.

The fourth category, osteoarthritis, is a major cause of pain and disability in the elderly and is the most common form of arthritis worldwide[45]. Sports and high-impact activities lead to cartilage lesions, which eventually lead to early-onset osteoarthritis[46,47]. There are seven SNPs associated with osteoarthritis at genome-wide significance[45,48,49]. These SNPs, however, only have allelic odds ratios between 1.1 and 1.2, which indicates a small effect on osteoarthritic risk. Given the strong link between joint injury and subsequent development of osteoarthritis, a focus on preventing joint injuries in young athletes with high risk could prevent or delay the onset of disease. Two Stanford triathletes were heterozygous at all seven loci, which have an additive allele effect [45,48,49]. This information could affect an athlete's decision regarding: 1) surgical treatment for a joint injury, 2) training plan (intensity or duration) when they are older, and 3) the age at which they retire from competition.

The fifth category includes SNPs associated with vitamin and mineral deficiencies. Exercise results in biochemical adaptations in muscle that place metabolic pathways under stress and increase the need for micronutrients. Routine training increases the loss of micronutrients, thus requiring greater intake of micronutrients for building, maintaining, and repairing lean body mass[50]. Moreover, proper levels of calcium and vitamin D are important for maintaining bone health, while iron, zinc, and the vitamin B complex are important for hematological function.

We reported on 39 SNPs associated with levels of: iron status biomarkers, circulating vitamin E, circulating vitamin D, serum calcium, magnesium, vitamin B12, vitamin B6, vitamin B9, phytosterol, and plasma homocysteine[51,52,53,54,55,56,57,58,59,60,61,62,63,64]. Meta-analyses on iron biomarkers included over 6,600 individuals, and the associated SNPs explain 0.7%-2.2% of the variance in serum iron levels[51,52]. Wang *et al.* performed a GWAS on vitamin D involving 33,996 individuals and found SNPs that explain 1–4% of the variance in serum levels[56]. O'Seaghdha *et al.* performed a GWAS on 20,611 individuals and found SNPs that account for 0.54% of the variance in serum calcium concentrations[57]. Genetic predispositions for vitamin or mineral deficiencies can prompt athletes to consume adequate amounts of micronutrients. Research has shown that many athletes have suboptimal magnesium levels, which can compromise athletic performance[65]. One of the athletes we tested had a genetic risk for magnesium deficiency. As an endurance athlete who needs to replenish micronutrients during activity, this athlete switched to an electrolyte drink that had higher levels of potassium and magnesium.

The last category, sickle cell trait, is a Mendelian condition caused by the inheritance of one normal allele of the hemoglobin gene (hemoglobin A), and one abnormal allele (hemoglobin S, rs334)[66,67,68]. In individuals heterozygous for the hemoglobin β chain mutant allele, red blood cells can sickle when the athlete is subjected to strong stressors, such as high altitude, extreme heat, dehydration, and exhaustion. This sickling can lead to gross hematuria, splenic infarction, and exertional rhabdomyolysis, which can be fatal[69]. No triathlete was positive for sickle cell trait. The NCAA currently recommends that athletes with sickle cell trait engage in slow and gradual conditioning regimens, rest and recover adequately between repetitions, stop activity immediately upon experiencing symptoms, stay optimally hydrated, maintain proper asthma treatment, refrain from exercise during acute illness, use supplemental oxygen as needed, and seek prompt medical care when experiencing unusual distress[70].

Twelve of fourteen athletes were recommended training interventions. The 3-month and 12-month questionnaires showed that eight of those twelve athletes acted upon their genetic results. Two of the remaining four who had not started an intervention indicated that they plan on taking action in the future. Moreover, seven believe that the implemented changes have had, or will have, a positive impact on their athletic careers. Importantly, after one year, four athletes were still actively following their interventions. In the year preceding this pilot study,

ten of the fourteen total athletes (71%) sustained an injury ([Table 1](#)). However, during the one-year follow up period, only four of the twelve athletes (33%) for whom we had full data sustained an injury ([Table 1](#)).

Discussion

There is a rich compendium of genetic studies that can be used to study sports injury risk in elite athletes. By including genetic studies of health and disease, we have greatly expanded the database of DNA variations that can be applied to sports genetics. We performed a literature search and identified 124 SNPs that have been associated with six different risk categories: ACL rupture, Achilles tendon injury, low bone mineral density, osteoarthritis, vitamin and mineral deficiencies, and sickle cell trait. To our knowledge, 113 of these SNPs have not previously been used in sports genetics[[7,8,10,71](#)].

We looked at large genetic studies investigating health traits that could be of interest to athletes. Most published gene association studies investigate disease in the general population rather than athletes. For example, studies on bone mineral density and vitamin/mineral deficiencies have not been previously used in sports genetics. Estrada *et al.* studied low bone mineral density, osteoporosis, and fractures in older women. We propose that these findings may also be relevant for athletes. Training creates a physical stress that, together with genetic predispositions, may lead to an injury that normally only occurs when older. Genetic information linked to vitamin and mineral deficiencies can be useful to help optimize micronutrient intake. Testing for sickle cell trait is already mandatory for all Division I and Division II NCAA athletes. We included the rs334 variant in our report because the DNA test is more accurate and cost-effective than the hemoglobin solubility tests recommended by the NCAA[[72](#)].

We began by recruiting elite collegiate triathletes for several reasons. First, triathletes are at increased risk for developing a wide range of injuries due to the three different disciplines of the sport. Second, Stanford University triathletes are club athletes in an individually scored sport, and hence ethical considerations regarding student-athlete scholarships and starting line-ups were not a concern. Athletes were individually motivated to participate (or not) in the genetic testing program, and independently chose whether or not to share their genetic results with third parties. Finally, these athletes are highly dedicated to athletic performance and could benefit from a new competitive edge. Although the genetic associations described here only contribute a small amount to overall risk for injury, they may still be important to elite athletes looking for every edge they can get. Over the 2011 and 2012 ITU World Triathlon Series race seasons, the average time difference between 1st and 10th for the Elite Men was 1.5 minutes, or about 1.4% of the total race time for each event[[73](#)]. Optimization of training and diet from individual genetic information could make a difference for some of these athletes.

Nevertheless, the small effect from genetic testing for athletic injury may mean that it is currently premature for use in the general athletic population[[74](#)]. Injury etiologies are complex combinations of extrinsic and intrinsic factors. In the future, better understanding of the multi-factorial etiologies of sports-related injuries will help guide the use of genetic information in athletic performance and in the clinical management of 'high-risk' individuals[[75,76](#)].

This pilot program was limited to only fourteen athletes, which is too small to perform a rigorous trial to show the effectiveness of the sports genetics program in reducing injury incidence and to draw meaningful conclusions about the effectiveness of the genetic testing. Our goal was to simply observe the athletes' responses to the new genetic knowledge. Nevertheless, there was positive feedback about the benefits of genetic testing. Fourteen of the nineteen athletes that were contacted decided to participate. Of those that replied to the 3- and 12-month follow up questionnaires, all responded positively to the program and thirteen indicated that they found

the genetic results both useful and informative. Moreover, all but one athlete reported sharing their genetic information with their coaches, trainers, physicians and/or teammates, highlighting that genetic information can be used to benefit athletes through multiple channels. Eight out of twelve athletes that were recommended training changes actually modified their training based on their genetic data. Besides prompting athletes to include new modifications in their training or diet, genetic knowledge may also increase compliance with currently prescribed prehabilitation strategies[77]. Genetic testing for sports injury risk could expand to include other types of elite athletes. Long-distance runners and rowers could benefit from this program as these athletes are also at risk for stress fractures, one of the major categories covered by our genetic testing. Knee ligament sprains and Achilles tendon injuries are common to many sports, such as baseball, basketball, soccer, and football. Athletes in these sports could benefit from genetic information regarding ACL rupture and Achilles tendon injury.

Conclusion

There is a rich source of new genetic knowledge that can be applied to athletic injury risk. Many health risks, such as low bone mineral density in the elderly, may also be pertinent to athletes. Athletes push themselves to the limit, which increases their chance for injury, and small effects in physiology can have meaningful consequences in competition. Through our literature search, we identified 113 SNPs that have previously not been discussed in relation to sports injuries and performance. A new sports genetic database was constructed, and used to help customize training and injury prevention for fourteen triathletes. In the future, additional categories of injury can be included in the sports genetics database, and personalized genetic consultation can be expanded to other types of athletes.

Supporting Information

S1 File. Three- and Twelve-month Follow-up Survey Questions. Text copy of survey questions distributed to athletes three and twelve months after initial results consultation.
(PDF)

Acknowledgments

We would like to thank members of the Kim lab for helpful discussion, athletes and coaches of the Stanford University Triathlon Team for participating in our program, Gordon Matheson for valuable advice and guidance, John Ioannidis for scientific guidance, and the Stanford Center for Clinical Informatics (Stanford CTSA UL1RR025744 from NIH/NCRR).

Author Contributions

Conceived and designed the experiments: GTG SKK AKR TRR. Performed the experiments: GTG SKK AKR TRR CH S. Beach S. Baur. Analyzed the data: GTG SKK AKR TRR. Contributed reagents/materials/analysis tools: GTG SKK AKR TRR CH S. Beach S. Baur. Wrote the paper: GTG SKK AKR TRR CH S. Beach S. Baur.

References

1. Hootman JM, Dick R, Agel J. Epidemiology of collegiate injuries for 15 sports: summary and recommendations for injury prevention initiatives. *J Athl Train.* 2007; 42: 311–319. PMID: [17710181](#)
2. van Mechelen W. Running injuries. A review of the epidemiological literature. *Sports Med.* 1992; 14: 320–335. PMID: [1439399](#)

3. van Gent RN, Siem D, van Middelkoop M, van Os AG, Bierma-Zeinstra SM, Koes BW. Incidence and determinants of lower extremity running injuries in long distance runners: a systematic review. *Br J Sports Med.* 2007; 41: 469–480; discussion 480. PMID: [17473005](#)
4. O'Toole ML, Hiller WD, Smith RA, Sisk TD. Overuse injuries in ultraendurance triathletes. *Am J Sports Med.* 1989; 17: 514–518. PMID: [2789479](#)
5. Egermann M, Brocail D, Lill CA, Schmitt H. Analysis of injuries in long-distance triathletes. *Int J Sports Med.* 2003; 24: 271–276. PMID: [12784169](#)
6. Eirale C, Tol JL, Farooq A, Smiley F, Chalabi H. Low injury rate strongly correlates with team success in Qatari professional football. *Br J Sports Med.* 2013; 47: 807–808. doi: [10.1136/bjsports-2012-091040](#) PMID: [22904292](#)
7. Wang G, Padmanabhan S, Wolfarth B, Fuku N, Lucia A, Ahmetov II, et al. Genomics of elite sporting performance: what little we know and necessary advances. *Adv Genet.* 2013; 84: 123–149. doi: [10.1016/B978-0-12-407703-4.00004-9](#) PMID: [24262098](#)
8. Maffulli N, Margiotti K, Longo UG, Loppini M, Fazio VM, Denaro V. The genetics of sports injuries and athletic performance. *Muscles Ligaments Tendons J.* 2013; 3: 173–189. PMID: [24367777](#)
9. Cupeiro R, Benito PJ, Maffulli N, Calderon FJ, Gonzalez-Lamuno D. MCT1 genetic polymorphism influence in high intensity circuit training: a pilot study. *J Sci Med Sport.* 2010; 13: 526–530. doi: [10.1016/j.jsmams.2009.07.004](#) PMID: [19850519](#)
10. Kambouris M, Ntalouka F, Ziogas G, Maffulli N. Predictive genomics DNA profiling for athletic performance. *Recent Pat DNA Gene Seq.* 2012; 6: 229–239. PMID: [22827597](#)
11. Wagner JK. Sidelining Gina: The Impact of Personal Genomics and Collective Bargaining in Professional Sports. *VA Sports & Ent LJ.* 2013; 12: 1–19.
12. Wagner JK, Royal CD. Field of Genes: An Investigation of Sports-Related Genetic Testing. *J Pers Med.* 2012; 2: 119–137. doi: [10.3390/jpm2030119](#) PMID: [25562204](#)
13. Kambouris M, Del Buono A, Maffulli N. Genomics DNA profiling in elite professional soccer players: a pilot study. *Trans Med @ UniSa.* 2014; 9: 18–22. PMID: [24809029](#)
14. Ioannidis JP, Boffetta P, Little J, O'Brien TR, Uitterlinden AG, Vineis P, et al. Assessment of cumulative evidence on genetic associations: interim guidelines. *Int J Epidemiol.* 2008; 37: 120–132. PMID: [17898028](#)
15. Harris PA, Taylor R, Thielke R, Payne J, Gonzalez N, Conde JG. Research electronic data capture (REDCap)—a metadata-driven methodology and workflow process for providing translational research informatics support. *J Biomed Inform.* 2009; 42: 377–381. doi: [10.1016/j.jbi.2008.08.010](#) PMID: [18929686](#)
16. Posthumus M, September AV, Keegan M, O'Cuinneagain D, Van der Merwe W, Schwellnus MP, et al. Genetic risk factors for anterior cruciate ligament ruptures: COL1A1 gene variant. *Br J Sports Med.* 2009; 43: 352–356. doi: [10.1136/bjsm.2008.056150](#) PMID: [19193663](#)
17. Khoschnau S, Melhus H, Jacobson A, Rahme H, Bengtsson H, Ribom E, et al. Type I collagen alpha1 Sp1 polymorphism and the risk of cruciate ligament ruptures or shoulder dislocations. *Am J Sports Med.* 2008; 36: 2432–2436. doi: [10.1177/0363546508320805](#) PMID: [18669982](#)
18. Collins M, Posthumus M, Schwellnus MP. The COL1A1 gene and acute soft tissue ruptures. *Br J Sports Med.* 2010; 44: 1063–1064. doi: [10.1136/bjsm.2008.056184](#) PMID: [19193665](#)
19. Posthumus M, September AV, O'Cuinneagain D, van der Merwe W, Schwellnus MP, Collins M. The COL5A1 gene is associated with increased risk of anterior cruciate ligament ruptures in female participants. *Am J Sports Med.* 2009; 37: 2234–2240. doi: [10.1177/0363546509338266](#) PMID: [19654427](#)
20. Posthumus M, September AV, O'Cuinneagain D, van der Merwe W, Schwellnus MP, Collins M, et al. The association between the COL12A1 gene and anterior cruciate ligament ruptures. *Br J Sports Med.* 2010; 44: 1160–1165. doi: [10.1136/bjsm.2009.060756](#) PMID: [19443461](#)
21. Posthumus M, Collins M, van der Merwe L, O'Cuinneagain D, van der Merwe W, Ribbans WJ, et al. Matrix metalloproteinase genes on chromosome 11q22 and the risk of anterior cruciate ligament (ACL) rupture. *Scand J Med Sci Sports.* 2012; 22: 523–533. doi: [10.1111/j.1600-0838.2010.01270.x](#) PMID: [21410539](#)
22. Daneshjoo A, Mokhtar AH, Rahnama N, Yusof A. The effects of injury preventive warm-up programs on knee strength ratio in young male professional soccer players. *PLoS One.* 2012; 7: e50979. doi: [10.1371/journal.pone.0050979](#) PMID: [23226553](#)
23. Puddu G, Ippolito E, Postacchini F. A classification of Achilles tendon disease. *Br J Sports Med.* 1976; 4: 145–150.
24. Kader D, Saxena A, Movin T, Maffulli N. Achilles tendinopathy: some aspects of basic science and clinical management. *Br J Sports Med.* 2002; 36: 239–249. PMID: [12145112](#)

25. Lopes AD, Hespanhol Junior LC, Yeung SS, Costa LO. What are the main running-related musculo-skeletal injuries? A Systematic Review. *Sports Med.* 2012; 42: 891–905. doi: [10.2165/11631170-00000000-00000](https://doi.org/10.2165/11631170-00000000-00000) PMID: [22827721](#)
26. Raleigh SM, van der Merwe L, Ribbans WJ, Smith RK, Schwellnus MP, Collins M. Variants within the MMP3 gene are associated with Achilles tendinopathy: possible interaction with the COL5A1 gene. *Br J Sports Med.* 2009; 43: 514–520. doi: [10.1136/bjsm.2008.053892](https://doi.org/10.1136/bjsm.2008.053892) PMID: [19042922](#)
27. September AV, Cook J, Handley CJ, van der Merwe L, Schwellnus MP, Collins M. Variants within the COL5A1 gene are associated with Achilles tendinopathy in two populations. *Br J Sports Med.* 2009; 43: 357–365. doi: [10.1136/bjsm.2008.048793](https://doi.org/10.1136/bjsm.2008.048793) PMID: [18443036](#)
28. Mokone GG, Schwellnus MP, Noakes TD, Collins M. The COL5A1 gene and Achilles tendon pathology. *Scand J Med Sci Sports.* 2006; 16: 19–26. PMID: [16430677](#)
29. Mokone GG, Gajjar M, September AV, Schwellnus MP, Greenberg J, Noakes TD, et al. The guanine-thymine dinucleotide repeat polymorphism within the tenascin-C gene is associated with achilles tendon injuries. *Am J Sports Med.* 2005; 33: 1016–1021. PMID: [15983124](#)
30. Posthumus M, Collins M, Cook J, Handley CJ, Ribbans WJ, Smith RK, et al. Components of the transforming growth factor-beta family and the pathogenesis of human Achilles tendon pathology—a genetic association study. *Rheumatology.* 2010; 49: 2090–2097. doi: [10.1093/rheumatology/keq072](https://doi.org/10.1093/rheumatology/keq072) PMID: [20360039](#)
31. Nell EM, van der Merwe L, Cook J, Handley CJ, Collins M, September AV. The apoptosis pathway and the genetic predisposition to Achilles tendinopathy. *J Orthop Res.* 2012; 30: 1719–1724. doi: [10.1002/jor.22144](https://doi.org/10.1002/jor.22144) PMID: [22588838](#)
32. O'Brien M. Structure and metabolism of tendons. *Scand J Med Sci Sports.* 1997; 7: 55–61. PMID: [9211604](#)
33. Fujii K, Yamagishi T, Nagafuchi T, Tsuji M, Kuboki Y. Biochemical properties of collagen from ligaments and periaricular tendons of the human knee. *Knee Surg Sports Traumatol Arthrosc.* 1994; 2: 229–233. PMID: [8536046](#)
34. Mafi N, Lorentzon R, Alfredson H. Superior short-term results with eccentric calf muscle training compared to concentric training in a randomized prospective multicenter study on patients with chronic Achilles tendinosis. *Knee Surg Sports Traumatol Arthrosc.* 2001; 9: 42–47. PMID: [11269583](#)
35. Alfredson H, Pietila T, Jonsson P, Lorentzon R. Heavy-load eccentric calf muscle training for the treatment of chronic Achilles tendinosis. *Am J Sports Med.* 1998; 26: 360–366. PMID: [9617396](#)
36. Maffulli N, Sharma P, Luscombe KL. Achilles tendinopathy: aetiology and management. *J R Soc Med.* 2004; 97: 472–476. PMID: [15459257](#)
37. Bennell K, Matheson G, Meeuwisse W, Brukner P. Risk factors for stress fractures. *Sports Med.* 1999; 28: 91–122. PMID: [10492029](#)
38. Lewiecki EM, Baim S, Bilezikian JP, Eastell R, LeBoff MS, Miller PD. 2008 Santa Fe Bone Symposium: update on osteoporosis. *J Clin Densitom.* 2009; 12: 135–157. doi: [10.1016/j.jocd.2009.02.002](https://doi.org/10.1016/j.jocd.2009.02.002) PMID: [19426925](#)
39. Ralston SH, Uitterlinden AG. Genetics of osteoporosis. *Endocr Rev.* 2010; 31: 629–662. doi: [10.1210/er.2009-0044](https://doi.org/10.1210/er.2009-0044) PMID: [20431112](#)
40. Field AE, Gordon CM, Pierce LM, Ramappa A, Kocher MS. Prospective study of physical activity and risk of developing a stress fracture among preadolescent and adolescent girls. *Arch Pediatr Adolesc Med.* 2011; 165: 723–728. doi: [10.1001/archpediatrics.2011.34](https://doi.org/10.1001/archpediatrics.2011.34) PMID: [21464375](#)
41. Estrada K, Styrkarsdottir U, Evangelou E, Hsu YH, Duncan EL, Ntzani EE, et al. Genome-wide meta-analysis identifies 56 bone mineral density loci and reveals 14 loci associated with risk of fracture. *Nat Genet.* 2012; 44: 491–501. doi: [10.1038/ng.2249](https://doi.org/10.1038/ng.2249) PMID: [22504420](#)
42. Zhang LS, Hu HG, Liu YJ, Li J, Yu P, Zhang F, et al. A follow-up association study of two genetic variants for bone mineral density variation in Caucasians. *Osteoporos Int.* 2012; 23: 1867–1875. doi: [10.1007/s00198-011-1863-z](https://doi.org/10.1007/s00198-011-1863-z) PMID: [22159821](#)
43. Paternoster L, Lorentzon M, Vandenput L, Karlsson MK, Ljunggren O, Kindmark A, et al. Genome-wide association meta-analysis of cortical bone mineral density unravels allelic heterogeneity at the RANKL locus and potential pleiotropic effects on bone. *PLoS Genet.* 2010; 6: e1001217. doi: [10.1371/journal.pgen.1001217](https://doi.org/10.1371/journal.pgen.1001217) PMID: [21124946](#)
44. Jin H, Evangelou E, Ioannidis JP, Ralston SH. Polymorphisms in the 5' flank of COL1A1 gene and osteoporosis: meta-analysis of published studies. *Osteoporos Int.* 2011; 22: 911–921.
45. Evangelou E, Valdes AM, Kerkhof HJ, Styrkarsdottir U, Zhu Y, Meulenbelt I, et al. Meta-analysis of genome-wide association studies confirms a susceptibility locus for knee osteoarthritis on chromosome 7q22. *Ann Rheum Dis.* 2011; 70: 349–355. doi: [10.1136/ard.2010.132787](https://doi.org/10.1136/ard.2010.132787) PMID: [21068099](#)

46. Bennell K, Hunter DJ, Vicenzino B. Long-term effects of sport: preventing and managing OA in the athlete. *Nat Rev Rheumatol.* 2012; 8: 747–752. doi: [10.1038/nrrheum.2012.119](https://doi.org/10.1038/nrrheum.2012.119) PMID: [22847471](#)
47. Caine DJ, Golightly YM. Osteoarthritis as an outcome of paediatric sport: an epidemiological perspective. *Br J Sports Med.* 2011; 45: 298–303. doi: [10.1136/bjsm.2010.081984](https://doi.org/10.1136/bjsm.2010.081984) PMID: [21330645](#)
48. Identification of new susceptibility loci for osteoarthritis (arcOGEN): a genome-wide association study. *Lancet.* 2012; 380: 815–823. doi: [10.1016/S0140-6736\(12\)60681-3](https://doi.org/10.1016/S0140-6736(12)60681-3) PMID: [22763110](#)
49. Valdes AM, Evangelou E, Kerkhof HJ, Tamm A, Doherty SA, Kisand K, et al. The GDF5 rs143383 polymorphism is associated with osteoarthritis of the knee with genome-wide statistical significance. *Ann Rheum Dis.* 2011; 70: 873–875. doi: [10.1136/ard.2010.134155](https://doi.org/10.1136/ard.2010.134155) PMID: [20870806](#)
50. Rodriguez NR, Di Marco NM, Langley S. American College of Sports Medicine position stand. Nutrition and athletic performance. *Med Sci Sports Exerc.* 2009; 41: 709–731. doi: [10.1249/MSS.0b013e31890eb86](https://doi.org/10.1249/MSS.0b013e31890eb86) PMID: [19225360](#)
51. Oexle K, Ried JS, Hicks AA, Tanaka T, Hayward C, Bruegel M, et al. Novel association to the proprotein convertase PCSK7 gene locus revealed by analysing soluble transferrin receptor (sTfR) levels. *Hum Mol Genet.* 2011; 20: 1042–1047. doi: [10.1093/hmg/ddq538](https://doi.org/10.1093/hmg/ddq538) PMID: [21149283](#)
52. Kullo IJ, Ding K, Jouni H, Smith CY, Chute CG. A genome-wide association study of red blood cell traits using the electronic medical record. *PLoS One.* 2010; 5: e13011. doi: [10.1371/journal.pone.0013011](https://doi.org/10.1371/journal.pone.0013011) PMID: [20927387](#)
53. Major JM, Yu K, Wheeler W, Zhang H, Cornelis MC, Wright ME, et al. Genome-wide association study identifies common variants associated with circulating vitamin E levels. *Hum Mol Genet.* 2011; 20: 3876–3883. doi: [10.1093/hmg/ddr296](https://doi.org/10.1093/hmg/ddr296) PMID: [21729881](#)
54. Johansen CT, Wang J, Lanktree MB, Cao H, McIntyre AD, Ban MR, et al. Excess of rare variants in genes identified by genome-wide association study of hypertriglyceridemia. *Nat Genet.* 2010; 42: 684–687. doi: [10.1038/ng.628](https://doi.org/10.1038/ng.628) PMID: [20657596](#)
55. Ahn J, Yu K, Stolzenberg-Solomon R, Simon KC, McCullough ML, Gallicchio L, et al. Genome-wide association study of circulating vitamin D levels. *Hum Mol Genet.* 2010; 19: 2739–2745. doi: [10.1093/hmg/ddq155](https://doi.org/10.1093/hmg/ddq155) PMID: [20418485](#)
56. Wang TJ, Zhang F, Richards JB, Kestenbaum B, van Meurs JB, Berry D, et al. Common genetic determinants of vitamin D insufficiency: a genome-wide association study. *Lancet.* 2010; 376: 180–188. doi: [10.1016/S0140-6736\(10\)60588-0](https://doi.org/10.1016/S0140-6736(10)60588-0) PMID: [20541252](#)
57. O'Seaghhdha CM, Yang Q, Glazer NL, Leak TS, Dehghan A, Smith AV, et al. Common variants in the calcium-sensing receptor gene are associated with total serum calcium levels. *Hum Mol Genet.* 2010; 19: 4296–4303. doi: [10.1093/hmg/ddq342](https://doi.org/10.1093/hmg/ddq342) PMID: [20705733](#)
58. Kapur K, Johnson T, Beckmann ND, Sehmi J, Tanaka T, Katalik Z, et al. Genome-wide meta-analysis for serum calcium identifies significantly associated SNPs near the calcium-sensing receptor (CARS) gene. *PLoS Genet.* 2010; 6: e1001035. doi: [10.1371/journal.pgen.1001035](https://doi.org/10.1371/journal.pgen.1001035) PMID: [20661308](#)
59. Meyer TE, Verwoert GC, Hwang SJ, Glazer NL, Smith AV, van Rooij FJ, et al. Genome-wide association studies of serum magnesium, potassium, and sodium concentrations identify six loci influencing serum magnesium levels. *PLoS Genet.* 2010; 6: e1001045. doi: [10.1371/journal.pgen.1001045](https://doi.org/10.1371/journal.pgen.1001045) PMID: [20700443](#)
60. Lin X, Lu D, Gao Y, Tao S, Yang X, Feng J, et al. Genome-wide association study identifies novel loci associated with serum level of vitamin B12 in Chinese men. *Hum Mol Genet.* 2012; 21: 2610–2617. doi: [10.1093/hmg/ddz062](https://doi.org/10.1093/hmg/ddz062) PMID: [22367966](#)
61. Tanaka T, Scheet P, Giusti B, Bandinelli S, Piras MG, Usala G, et al. (2009) Genome-wide association study of vitamin B6, vitamin B12, folate, and homocysteine blood concentrations. *Am J Hum Genet.* 2009; 84: 477–482. doi: [10.1016/j.ajhg.2009.02.011](https://doi.org/10.1016/j.ajhg.2009.02.011) PMID: [19303062](#)
62. Pare G, Chasman DI, Parker AN, Zee RR, Malarstig A, Seedorf U, et al. Novel associations of CPS1, MUT, NOX4, and DPEP1 with plasma homocysteine in a healthy population: a genome-wide evaluation of 13 974 participants in the Women's Genome Health Study. *Circ Cardiovasc Genet.* 2009; 2: 142–150. doi: [10.1161/CIRCGENETICS.108.829804](https://doi.org/10.1161/CIRCGENETICS.108.829804) PMID: [20031578](#)
63. Lange LA, Croteau-Chonka DC, Maruelle AF, Qin L, Gaulton KJ, Kuzawa CW, et al. Genome-wide association study of homocysteine levels in Filipinos provides evidence for CPS1 in women and a stronger MTHFR effect in young adults. *Hum Mol Genet.* 2010; 19: 2050–2058. doi: [10.1093/hmg/ddq062](https://doi.org/10.1093/hmg/ddq062) PMID: [20154341](#)
64. Teupser D, Baber R, Ceglarek U, Scholz M, Illig T, Gieger C, et al. Genetic regulation of serum phytosterol levels and risk of coronary artery disease. *Circ Cardiovasc Genet.* 2010; 3: 331–339. doi: [10.1161/CIRCGENETICS.109.907873](https://doi.org/10.1161/CIRCGENETICS.109.907873) PMID: [20529992](#)
65. Bohl CH, Volpe SL. Magnesium and exercise. *Crit Rev Food Sci Nutr.* 2002; 42: 533–563. PMID: [12487419](#)

66. Ghansah A, Rockett KA, Clark TG, Wilson MD, Koram KA, Oduro AR, et al. Haplotype analyses of haemoglobin C and haemoglobin S and the dynamics of the evolutionary response to malaria in Kassena-Nankana District of Ghana. *PLoS One*. 2012; 7: e34565. doi: [10.1371/journal.pone.0034565](https://doi.org/10.1371/journal.pone.0034565) PMID: [22506028](#)
67. Ingram VM. Abnormal human haemoglobins. III. The chemical difference between normal and sickle cell haemoglobins. *Biochim Biophys Acta*. 1959; 36: 402–411. PMID: [13852872](#)
68. Ashley-Koch A, Yang Q, Olney RS. Sickle hemoglobin (HbS) allele and sickle cell disease: a HuGE review. *Am J Epidemiol*. 2000; 151: 839–845. PMID: [10791557](#)
69. Eichner ER. Sports medicine pearls and pitfalls—sickle cell trait and athletes: three clinical concerns. *Curr Sports Med Rep*. 2007; 6: 134–135. PMID: [19202657](#)
70. 2012–2013 NCAA Sports Medicine Handbook. In: Klossner D, editor. 23 ed. Indianapolis, Indiana: NCAA. 2012. PMID: [2012](#)
71. Pruna R, Artells R, Ribas J, Montoro B, Cos F, Muñoz C, et al. Single nucleotide polymorphisms associated with non-contact soft tissue injuries in elite professional soccer players: influence on degree of injury and recovery time. *BMC Musculoskelet Disord*. 2013; 14: 221. doi: [10.1186/1471-2474-14-221](https://doi.org/10.1186/1471-2474-14-221) PMID: [23890452](#)
72. Ryan K, Bain BJ, Worthington D, James J, Plews D, Mason A, et al. Significant haemoglobinopathies: guidelines for screening and diagnosis. *Br J Haematol*. 2010; 149: 35–49. doi: [10.1111/j.1365-2141.2009.08054.x](https://doi.org/10.1111/j.1365-2141.2009.08054.x) PMID: [20067565](#)
73. World Triathlon Series Results. North Vancouver, BC: International Triathlon Union. 2013.
74. Schwellnus MP. Genetic biomarkers and exercise-related injuries: current clinical applications? *Br J Sports Med*. 2013; 47: 530–532. doi: [10.1136/bjsports-2013-092515](https://doi.org/10.1136/bjsports-2013-092515) PMID: [23637115](#)
75. Collins M. Genetic risk factors for soft-tissue injuries 101: a practical summary to help clinicians understand the role of genetics and 'personalised medicine'. *Br J Sports Med*. 2010; 44: 915–917. doi: [10.1136/bjsm.2009.058040](https://doi.org/10.1136/bjsm.2009.058040) PMID: [19553227](#)
76. Pitsiladis Y, Wang G, Wolfarth B, Scott R, Fuku N, Mikami E, et al. Genomics of elite sporting performance: what little we know and necessary advances. *Br J Sports Med*. 2013; 47: 550–555. doi: [10.1136/bjsports-2013-092400](https://doi.org/10.1136/bjsports-2013-092400) PMID: [23632745](#)
77. Schneider KI, Schmidtke J. Patient compliance based on genetic medicine: a literature review. *J Community Genet*. 2014; 5: 31–48. doi: [10.1007/s12687-013-0160-2](https://doi.org/10.1007/s12687-013-0160-2) PMID: [23934761](#)