

Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review

Hassane Zouhal, Juan Del Coso, Ayyappan Jayavel, Claire Tourny, Guillaume Ravé, Nidhal Jebabli, Cain C T Clark, Benjamin Barthélémy, Anthony C Hackney, Abderraouf Ben Abderrahman

▶ To cite this version:

Hassane Zouhal, Juan Del Coso, Ayyappan Jayavel, Claire Tourny, Guillaume Ravé, et al.. Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review. Journal of Sport and Health Science, 2023, 12 (3), pp.359-368. 10.1016/j.jshs.2021.07.003 . hal-03334082

HAL Id: hal-03334082 https://hal.science/hal-03334082v1

Submitted on 25 Oct 2021

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial - NoDerivatives 4.0 International License





Available online at www.sciencedirect.com

ScienceDirect



Journal of Sport and Health Science 00 (2021) 1-10

Review

Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review

Hassane Zouhal^a,*, Juan Del Coso^b, Ayyappan Jayavel^c, Claire Tourny^d, Guillaume Ravé^e, Nidhal Jebabli^f, Cain C.T. Clark^g, Benjamin Barthélémy^e, Anthony C. Hackney^h, Abderraouf Ben Abderrahman^f

^a M2S (Laboratoire Mouvement, Sport, Santé)-EA 1274, Department of Sport Sciences, University of Rennes, Rennes F-35000, France ^b Rey Juan Carlos University, Centre for Sport Studies, Madrid 28032, Spain

^c SRM College of Physiotherapy, SRM Institute of Science and Technology, SRM Nagar, Kattankulathur 603203, India ^d Department of Sport Sciences, University of Rouen, Mont Saint Aignan, CETAPS EA 3832, F-76821, France

^e Toulouse Football Club, Toulouse 31000, France

^f Higher Institute of Sport and Physical Education, Ksar-Said, University of Manouba, Tunis 2010, Tunisia

^g Centre for Intelligent Healthcare, Coventry University, Coventry CV1 5FB, UK

^h Department of Exercise & Sport Science, Department of Nutrition, University of North Carolina, Chapel Hill, NC 27514, USA

Received 26 February 2021; revised 13 May 2021; accepted 18 June 2021

2095-2546/© 2021 Published by Elsevier B.V. on behalf of Shanghai University of Sport. This is an open access article under the CC BY-NC-ND license (http://creativecommons.org/licenses/by-nc-nd/4.0/)

Abstract

Background: The aim of this study was to review, systematically, evidence concerning the link between the *ACTN3* R577X polymorphism and the rates and severity of non-contact injuries and exercise-induced muscle damage in athletes and individuals enrolled in exercise training programs.

Methods: A computerized literature search was performed in the electronic databases PubMed, Web of Science, and SPORTDiscus, from inception until November 2020. All included studies compared the epidemiological characteristics of non-contact injury between the different geno-types of the *ACTN3* R577X polymorphism.

Results: Our search identified 492 records. After the screening of titles, abstracts, and full texts, 13 studies examining the association between the *ACTN3* genotypes and the rate and severity of non-contact injury were included in the analysis. These studies were performed in 6 different countries (Spain, Japan, Brazil, China, Republic of Korea, and Italy) and involved a total participant pool of 1093 participants. Of the studies, 2 studies involved only women, 5 studies involved only men, and 6 studies involved both men and women. All the studies included were classified as high-quality studies (≥ 6 points in the Physiotherapy Evidence Database (PEDro) scale score). Overall, evidence suggests there is an association between the *ACTN3* R577X genotype and non-contact injury in 12 investigations. Six studies observed a significant association between *ACTN3* R577X polymorphism and exercise induced muscle damage: two with non-contact ankle injury, three with non-contact muscle injury, and one with overall non-contact injury. *Conclusion*: The present findings support the premise that possessing the *ACTN3* XX genotype may predispose athletes to a higher probability of

Conclusion: The present indings support the premise that possessing the *ACIN3* XX genotype may predispose athletes to a higher probability of some non-contact injuries, such as muscle injury, ankle sprains, and higher levels of exercise-induced muscle damage.

Keywords: α-actinin-3 deficiency; Athletic performance; Exercise-related injury; Muscle injury; Single nucleotide polymorphism

1. Introduction

The protein α -actinin-3 is a structural component of the Z-disc encoded by the *ACTN3* gene. The main role of α -actinin-3 is anchoring actin thin filaments to assist in maintaining the myofibrillar array, which contributes to regulating muscle length and tension during contraction.¹ Interestingly,

E-mail address: hassane.zouhal@univ-rennes2.fr (H. Zouhal).

 α -actinin-3 is expressed only in fast muscle fibers (all fast-glycolytic type 2X fibers and 50% of fast oxidative type 2A fibers),² suggesting a specific function for powerful and fast muscle contractions. A common genetic variant in the *ACTN3* (rs1815739, also known as the R577X polymorphism) leads to the replacement of an arginine (R) with a premature stop codon (X) at amino acid 577. Consequently, individuals with the *ACTN3* XX genotype are α -actinin-3 protein deficient due to the lack of protein expression. In contrast, homozygous individuals for the R allele (RR genotype) or heterozygote individuals (RX

Peer review under responsibility of Shanghai University of Sport.

^{*} Corresponding author.

https://doi.org/10.1016/j.jshs.2021.07.003

ARTICLE IN PRESS

genotype) express α -actinin-3, although recently it has been found that the capacity to express α -actinin-3 is dose-dependent for RR *vs.* RX individuals.³ The deficiency of α -actinin-3 does not entail any disease or clinical condition but has been demonstrated to produce some potentially negative phenotypes in humans, such as lower muscle strength,^{4,5} reduced muscle volume,^{6,7} impaired capacity to tolerate the strain produced by explosive muscular actions,⁸ and/or an association with decreased bone mineral density.⁹

In athletes, several investigations have confirmed that α -actinin-3 deficiency due to the ACTN3 XX genotype may negatively influence sprint and power performance.¹⁰ Lower performance in sport disciplines requiring near-to-maximal production of strength or power has been associated with decreased capacity of the ability of muscle fibers to produce powerful contractions in the absence of α -actinin-3.⁷ This notion is supported by several investigations that found a higher occurrence of the ACTN3 RR variant in elite sprint/power athletes when compared with nonathletes.¹ In contrast, a higher-than-expected frequency of the ACTN3 XX genotype has been found in some groups of elite endurance athletes,^{9,11} although the overrepresentation of the XX genotype has not been replicated in other cohorts of elite athletes.^{12,13} Collectively, the information obtained from studies investigating the frequency of the different ACTN3 genotypes in cohorts of elite athletes points toward a negative influence of the XX genotype on elite sprint/power-based exercise (hence, a positive influence of the R allele on this type of exercise) with little or no effect of the XX genotype on endurance-based exercise.

In addition to performance, the absence of α -actinin-3 has been linked to a higher probability of injury¹⁴ in several sport and exercise scenarios. For example, soccer (football) players with the *ACTN3* XX genotype suffered a higher incidence of non-contact muscle injuries when compared to players with the RR or the RX genotype.^{15,16} A higher probability of muscle-type injuries was also found in XX compared to RR runners.¹⁷ On the other hand, at least 1 investigation has reported that athletes with the RR or the RX genotype had an increased risk of muscle injury when practicing various sport activities.¹⁸ Additionally, there is an increase in passive hamstring stiffness in R-allele individuals compared to XX counterparts,¹⁹ although the link between muscle stiffness and increased muscle injury risk has not yet been established.^{14,20} The most consistent specific findings posit a link between the XX genotype and ankle injuries^{21–23} and higher levels of exercise-induced muscle damage.^{24–26} These associations suggest a negative impact of α -actinin-3 deficiency on muscle and ligament capacity to endure the forces generated during exercise and point toward an increased susceptibility to contraction-induced damage.^{8,9} Despite the extant literature indicating an influence of the *ACTN3* genotype on exercise-related non-contact injuries, there is no previous investigation, to our knowledge, that has systematically reviewed the influence of the *ACTN3* R577X polymorphism on non-contact injuries. Hence, the aim of this investigation was to review evidence systematically concerning the link between the *ACTN3* R577X polymorphism and the rates and severity of non-contact injuries and exercise-induced muscle damage in athletes and individuals enrolled in exercise-training programs.

2. Methods

2.1. Eligibility criteria

This systematic review was conducted in accordance with the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement.27 Only studies that examined the link between the ACTN3 R577X polymorphism and the rate or severity of non-contact injuries in individuals who were at any level/sex/age and currently participating in a sport or exercise-training program were included. The following inclusion criteria were established for studies of this topic (Table 1). The studies (1) were available in peer-reviewed journals; (2) involved participants aged ≥ 14 years; (3) included trained athletes (frequency of training ≥ 5 times/week) and individuals enrolled in exercise training programs (i.e., nonathletes); (4) used validated methods for the characterization of injury epidemiology (e.g., injury severity, occurrence and incidence); (5) obtained epidemiological data about non-contact injuries developed during sport competitions and exercise practice; and (6) full text available in English. Studies were excluded if they (1) did not contain an experiment study design with the original data, such as books, systematic or narrative reviews, case studies, or opinion pieces; (2) did not meet the minimum requirements to classify the sample as exercisers (frequency of training <3times/week); or (3) were not written in English.

Table 1

Inclusion criteria according to the Population, Intervention, Comparison, Outcomes and Study (PICOS) approach.

Criteria	Inclusion	Exclusion
1	Original article published in peer-reviewed journals	Books, reviews, case studies, opinion pieces, non-peer-reviewed journals
2	Involved professional, amateur, or recreational athletes or individuals practicing exercise in regular programs	Untrained individuals
3	Participants >14 years old, described as young and/or senior athletes	Participants <14 years old, described as child athletes
4	Analyzed the effects of the <i>ACTN3</i> R577X polymorphism on the rates or severity of non-contact injuries	A gene polymorphism other than ACTN3 R577X analyzed
5	Used quantification of injuries (e.g., injury severity, occurrence, and incidence)	Rates or severity of non-contact injuries not reported
6	Full text available in English	Full text not accessible in English

Please cite this article as: Hassane Zouhal et al., Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review, Journal of Sport and Health Science (2021), https://doi.org/10.1016/j.jshs.2021.07.003

ACTN3 genotype and sport injury

2.2. Literature search strategy

Literature searches were conducted in 3 electronic databases including PubMed, Web of Science, and SPORTDiscus, from inception until November 2020. No year restriction was used for the search strategy, in an effort to obtain all studies on the topic. The following key terms (and synonyms searched for by the MeSH database) were included and combined using the operators "AND", "OR", "NOT": ((Alpha-actinin-3 OR ACTN3) OR ("ACTN3 gene" OR "ACTN3 R577X polymorphism" OR "ACTN3 R577X" OR "ACTN3 R577R" OR "ACTN3 577XX genotype*" OR "ACTN3 577RR genotype*" OR "ACTN3 577RX genotype*") AND ("injuries" OR "injury" OR "muscle injury*" OR "non-contact injury" OR "strain" OR "soreness*" OR "sprain*" OR "contusion" OR "fracture" OR "dislocation" OR "concussion") AND (correlation study OR association OR relationship)). The references used in the identified studies were searched as well, in order to identify additional relevant research papers. The search for published studies was independently performed by 2 authors (AJ and ABA), and disagreements were resolved through discussion.

2.3. Study selection

The screening and study selection were performed by 2 authors based on the above-mentioned inclusion and exclusion criteria. If the title and abstract of the article showed any potential relevance in terms of assessing the influence of the *ACTN3* genotype on the rate and severity of non-contact injuries, the full text was examined. A third author (HZ) was consulted if the 2 investigators (AJ and ABA) responsible for the study selection were not able to reach an agreement on the inclusion of an article. Fig. 1 depicts the details of the study-selection methodology.

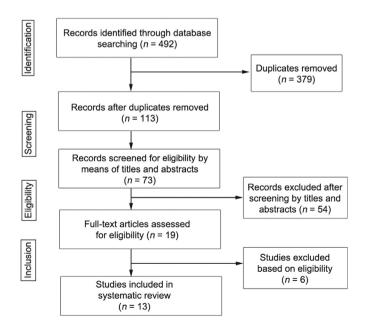


Fig. 1. Selection of research articles (n=13) included in this systematic review using the recommendations in the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) statement.²⁶

2.3.1. Data extraction

Once the inclusion/exclusion criteria were applied, data extraction was conducted to collect information about participants, intervention, comparisons, outcomes, and study design (PICOS) following PRISMA methodology. The following relevant data from each study were extracted: study details (author, year of publication, country, duration of follow-up), study population (sample size, age, participants' sex), sport or main exercise activity performed by the participants, the method used to determine the genotyping of ACTN3 R577X (rs1815739), injury definition, workload (external and internal load parameters), and measures of association (i.e., relative risk (RR) or odds ratio (OR). Where possible, these associations were extracted directly from the original article. For articles in which this information was not present, associations were calculated using raw data, if available. Load parameters were classified as external or internal based on the International Olympic Committee consensus statement on load in sport and risk of injury.²⁸

2.3.2. Quality assessment

The methodological quality of the included studies was assessed using the Physiotherapy Evidence Database (PEDro) scale.²⁹ A cut-off point of 6 points in the PEDro scale was used to discriminate high-quality studies, and only studies with scores above this threshold were considered for the systematic review.²⁶ Two authors (AJ and ABA) independently assigned a score to each study using the PEDro scale, and disagreements were resolved by consulting a third researcher (HZ).

3. Results

3.1. Study selection and description of the included studies

Our initial search identified 492 records (Fig. 1), and after the screening of titles, abstracts, and full texts, respectively, 13 studies were included in our final analysis. Accordingly, the characteristics of the participants, the follow-up periods, the injury definitions, and the methods used to determine the genotyping of *ACTN3* R577X (rs1815739) polymorphism are shown in Table 2. These studies were performed in 6 different countries (Spain, Japan, Brazil, China, Republic of Korea, and Italy), involving a total participant pool of 1093 participants, where 20 represents the lowest number and 257 the highest number of participants within the studies. Two studies involved only women, 5 studies involved only men, and 6 studies involved both men and women. All the studies included were classified as high-quality studies (≥ 6 in the PEDro scale score) (Table 3).

3.2. Evaluation of the effects of ACTN3 R577X genotype on non-contact injuries

The results of best evidence synthesis of the effects of *ACTN3* R577X genotype on non-contact injuries are presented in Table 4. For 12 studies,^{12–15,17,18,20–22,27,29,30} evidence existed of an association between the *ACTN3* R577X genotype and non-contact injury, and for the 1 remaining study, no effect was observed.¹⁹ From the total, 6 studies observed a significant

Please cite this article as: Hassane Zouhal et al., Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review, Journal of Sport and Health Science (2021), https://doi.org/10.1016/j.jshs.2021.07.003

Table 2

Characteristics of the studies that examined the association between the ACTN3 R577X polymorphism and the rates and severity of non-contact injuries in highly trained athletes and in individuals enrolled in exercise training programs.

Study	Country	Population/sample size/sex/age (years, mean \pm SD or range)/sport and level of practice	Follow-up	Injury data collection	Genetic testing
Pimenta et al. (2012) ³³	Brazil	37 male professional soccer players Age: 24.8 \pm 1.7	Before and after eccentric exercise	Exercise-induced muscle damage (eccen- tric exercise) Post-exercise serum CK and α -actin immediately after (post), 2- and 4-h post- eccentric exercise	Extraction of genomic DNA from the peripheral blood samples. A DNA fragment carrying the exo 16 from the <i>ACTN3</i> gene was amplified from the genomic DNA, and the following initiators were used: 5'-CTGTTGCCTGTGGTAAGTGGG-3'; reverse, 5'-TGGTCACAGTATGCAGGAGGG-3 correlated to the adjacent intronic sequences.
Iwao-Koizumi et al. (2014) ¹⁸	Japan	99 female sports students Average age: 19.7 Sports: football, softball, basketball, and badmin- ton Level of training: future professional athletes	_	A questionnaire was developed and filled out by each participant. The questionnaire asked whether the participant had suffered severe injuries during sports activity in the past.	DNA was collected from a few drops of saliva on water-soluble paper (Mishima Dissolve Paper, 60 MDP; Nippon Paper Papylia Co., Ltd., Tokyo, Japan), Genotyping using TaqMan assay (Taq- Man [®] SNP Genotyping Assays; Life Technologies Carlsbad, CA, USA).
Kim et al. (2014) ²¹	Republic of Korea	97 elite ballet dancers and 203 normal female adults Age: 23.1 ± 1.3	-	Structured injury questionnaire and testing for injury risks on the joints	MGB TaqMan [®] SNP Genotyping assay method was used to analyze <i>ACTN3</i> R577X polymorphism (rs1815739) from the extracted DNA.
Shang et al. (2015) ²²	China	142 males from a Chinese army infantry division Age: 21.0 \pm 0.2	Past 1 year	Non-contact ankle sprains were diagnosed by an experienced clinician according to the standard validated criteria.	DNA was extracted from blood cells by standard procedures by the kit manufacturer (Promega, Madison, WI, USA). Genotyping was done using a TaqMan allele discrimination assay that used the 5 nuclease activity of Taq polymerase.
Belli et al. (2017) ²⁶	Brazil	20 men and women athletes Age: 40.5 ± 1.2	Before and after an official 37.1 km adventure race (22.1 km mountain biking, 10.9 km trek- king, 4.1 km water trekking, 30 m rope course, and orienteering)	Post-exercise muscle damage	Genomic DNA was isolated from the buffy coat of centrifuged whole blood, using the QIAamp DNA Blood Mini Kit (Qiagen, Valencia, CA, USA) according to the manufacturer's instructions. Gen- otyping of <i>ACTN3</i> R577X (rs1815739) polymor- phism was conducted using a TaqMan SNP Genotyping Assay (Applied Biosystems, Foster City, CA, USA).
Del Coso et al. (2017) ²⁴	Spain	23 experienced men and women triathletes Age: 36.4 ± 5.2	Before and after the race	The changes in serum CK (CK–MM iso- form) were measured in the blood sam- ples, and muscle pain was measured with a visual analogue scale $(0-10 \text{ cm})$.	DNA was isolated from the whole blood obtained before the race (QIAamp [®] DNA Blood Mini Kit, QIAGEN, The Netherlands) according to the man- ufacturer's protocol. Genotyping was performed using a TaqMan [®] SNP genotyping assay (Life Technologies TM).
Del Coso et al. (2017) ²⁵	Spain	71 healthy, experienced (3 years) marathon runners, men and women Age: 42.7 \pm 8.9	Before and after the race	Post-exercise muscle damage; post-exer- cise serum CK	Genomic DNA was isolated from the whole blood obtained before the race (QIAamp [®] DNA Blood Mini Kit) according to the manufacturer's protocol. Genotyping was performed using a TaqMan [®] SNF genotyping assay (Life Technologies TM).

genotyping assay (Life TechnologiesTM).

(continued on next page)

ARTICLE IN PRESS

4

H. Zouhal et al.

Study	Country	Population/sample size/sex/age (years, mean \pm SD or range)/sport and level of practice	Follow-up	Injury data collection	Genetic testing
Massidda et al. (2019) ¹⁶	Italy	257 male professional soccer players Age: 19.4 ± 5.2	1–5 years follow-up	Physical complaint occurring during prac- tice that prevented a player from partici- pating in training or match play for at least 1 day after the day of the onset. Ultrasound and magnetic resonance imag- ing scans were used to morphologically classify the injuries.	DNA was extracted from a buccal swab and a com- mercially available kit (QIAGEN, Hilden, Ger- many) was used. Concentration of extracted DNA was determined by fluorometric method (through Qubit by Invitro- gen, Thermo Fisher Scientific, Waltham, MA, USA).
Miyamoto et al (2018) ⁴²	Japan	76 male sports university students Age: 21.2 ± 2.8 Sport: multiple sports Level of training: regular training/recreationally active	-	Past hamstring-muscle strain injury	DNA was extracted from saliva and collected with a DNA self-collection kit (Oragene, DNA Genotek, Ontario, Canada) according to the manufacturer's protocol. DNA was quantified using a spectrophotometer (Eppendorf Bio Photometer Plus, Eppendorf, Tokyo, Japan).
Del Coso et al. 2017) ³¹	Spain	67 healthy and experienced marathon runners (male, all Spanish Caucasians) Low CK, $n = 36$; age = 43.0 ± 8.1 High CK, $n = 31$; age = 41.4 ± 9.4	Immediately after the race	Post-exercise muscle damage	A DNA fragment carrying the exon 16 from the ACTN3 gene was amplified from the genomic DNA and the following initiators were used: 5- CTGTTGCCTGTGGTAAGTGGG-3; reverse, 5- TGGTCACAGTATGCAGGAGGG-3, correlated to the adjacent intronic sequences.
Clos et al. (2019) ¹⁵	Spain	43 male professional soccer players Age: 27.5 \pm 1.2	7 years follow-up	Injury rate, injury severity, and injury recovery times were established. Injury severity was established according to the days a player needed to be absent from training and competition: mild, 1–15 days; moderate, 16–30 days; severe, more than 30 days.	Genomic DNA isolation was performed using a QIAmp DNA Blood Minikit (Qiagen). DNA quantity was measured with a Nano-Drop ND-1000 Spectrophotometer (Thermo Fisher Sci- entific Inc., Waltham, MA, USA).
Del Coso et al. (2020) ³⁴	Spain	22 experienced men and women triathletes Age: 35.4 ± 4.3 Low CK, $n = 10$ High CK, $n = 12$	Before and after the race	Post-exercise muscle damage	A DNA fragment carrying the exon 16 from the <i>ACTN3</i> gene was amplified from the genomic DNA and the following initiators were used: 5'-CTGTTGCCTGTGGTAAGTGGG-3'; reverse, 5'-TGGTCACAGTATGCAGGAGGG-3', correlated to the adjacent intronic sequences.
Moreno et al. (2020) ¹⁷	Spain	139 (XX: 32, RX: 67, RR: 40) Healthy marathon runners, men and women Age: 41.3 ± 10.2	1 year preceding to marathon	Physical complaints/visible damage to any part of lower limb assessed by qualified medical/healthcare practitioner	DNA was isolated using an organic-based DNA extraction method adapted to Amicon1 Ultra 0.5- mL columns, including a final concentration step to 50 μL.

Abbreviations: CK = creatine-kinase; CK-MM = CK skeletal muscle.

6

Physiotherapy Evidence Database (PEDro) score of some long-term studies

Table 3

						PEDro items						
Study	Eligibility criteria Random- Concealed- allocation allocation	Random- allocation	Random- Concealed- allocation allocation	Group homogeneity Blinded Blinded therapists Blinded Drop Intention to Between-group Point estimates subjects assessor out treat analyses comparison and variability 5%	Blinded subjects	Blinded therapists	Blinded assessor	Drop out 5%	Intention to Between-gro treat analyses comparison	Between-group comparison	Point estimates and variability	Score
Pimenta et al. $(2012)^{33}$ Iwao-Koizumi et al. $(2014)^{18}$ Kim et al. $(2014)^{21}$ Shang et al. $(2015)^{25}$ Belli et al. $(2017)^{26}$ Del Coso et al. $(2017)^{25}$ Del Coso et al. $(2017)^{25}$ Del Coso et al. $(2017)^{31}$ Massidda et al. $(2018)^{42}$ Miyamoto et al. $(2018)^{42}$ Clos et al. $(2019)^{15}$ Miyamoto et al. $(2019)^{15}$ Clos et al. $(2019)^{15}$	•••••	000000000000000000000000000000000000000	•••••	•••••	•••••	000000000000000000000000000000000000000	•••••	000000000000000000000000000000000000000	000000000000000000000000000000000000000	•••••	•••••	~~~~~

association between the *ACTN3* R577X polymorphism and exercise-induced muscle damage,^{24–26,31–33} 2 studies observed association between *ACTN3* R577X polymorphism and noncontact ankle injury,^{21,22} and 1 study observed association with non-contact sports injury (i.e., muscle strain and ligament rupture).¹⁸ Additionally, 2 studies^{17,34} observed significant associations between *ACTN3* R577X polymorphism and non-contact lower-limb injury, and 1 study with non-contact musculoskeletal injury.¹⁵ Only 1 study reported no significant association between *ACTN3* R577X polymorphism and passive stiffness or hamstring strain injury (Table 4).¹⁹

4. Discussion

Characterizing the influence of the ACTN3 R577X genotype on exercise-related sport injuries is pertinent because the ACTN3 XX genotype, which produces α -actinin-3 deficiency, has been repeatedly associated with a reduction in sports performance in sprint and power-based disciplines.^{35–37} The main outcomes of this investigation indicate that most studies of the effect of ACTN3 R577X polymorphism on injury epidemiology indicate an association of the XX genotype with a deleterious effect on injury, expressed either by increased incidence or severity, as compared to RX and RR genotypes. However, current evidence suggests that the higher likelihood of exercise-related injury in XX athletes is specific to muscle injury, ligament injury, and an exacerbation of muscle damage. It can be concluded that possessing the ACTN3 XX genotype may predispose athletes to a higher probability of muscle and ligament injury during exercise. The evidence linking ACTN3 genotype and injury epidemiology is still emerging, so it is premature to recommend the determination of athletes' ACTN3 genotype for the prediction of injury risk.

4.1. Overall rate of injury

Note: circle : No; dark dot : Yes.

In the studies investigating all types of non-contact exerciserelated injuries in athletes, the evidence indicates a higher injury incidence in RR individuals (defined as those possessing the R allele) when compared to XX individuals.^{17,18} Iwao-Koizumi et al.¹⁸ found that in female athletes training to become professional athletes, those with the R allele had a 2.52-fold higher likelihood of suffering a severe injury than did those with the X allele.¹⁸ Similarly, Moreno et al.¹⁷ found that among amateur runners preparing for a marathon, those with the RR genotype had a 1.79-fold higher likelihood of suffering a running-related injury when compared to their XX counterparts, whereas the likelihood of injury increased to 1.93 when comparing RR and RX athletes. Injury incidence in these marathon runners tended to be higher in RR runners; overall injury incidence was not different in phenotypes (2.78 injuries per 1000 h of running) and in RX runners (1.65 and 1.94 injuries/1000 h of running, respectively), although this difference was not statistically significant. The higher overall injury incidence in RR athletes, despite the ability to fully express α -actinin-3 in their fast-type muscle fibers,³ may be related to other phenotypes found in RR athletes. RR athletes habitually present higher levels of muscle strength and power than their XX counterparts, 38,39 which may lead to

ARTICLE IN PRESS

ACTN3 genotype and sport injury

Table 4

Results of best evidence synthesis of association between ACTN3 R577X genotype and non-contact injuries.

Studies	Main study outcomes	Best evidence	synthesis
		Presence of association	Level of evidence
Pimenta et al. $(2012)^{33}$	ACTN3 R577X polymorphism and muscle damage	Yes	High
	After eccentric training, XX athletes presented higher levels for CK		
	(4-h post) and α -actin (post and 2-h post) compared to RR and RX.		
	XX athletes were more susceptible to muscle damage after eccentric		
L K : : (1 (2014) ¹⁸	exercise compared to RR and RX athletes.	37	TT' 1
Iwao-Koizumi et al. (2014) ¹⁸	ACTN3 R577X polymorphism and non-contact sports injury	Yes	High
	RR genotype and R allele frequency was higher than XX frequency in		
Kim et al. (2014) ²¹	athletes who experienced muscle injury. <i>ACTN3</i> R577X polymorphism and injury risk	Yes	Uich
Killi et al. (2014)	ACTN3 polymorphism was associated with ballerinas' performance	1 05	High
	capacity.		
	Injury frequency in the ankle in XX ballerinas was 4.65 times higher		
	than in RR and RX counterparts.		
Shang et al. (2015) ²²	ACTN3 R577X polymorphism and non-contact ankle sprain	Yes	High
	The frequency of the XX genotype was significantly higher among the		6
	acute ankle sprains group than the control group (52.1% vs. 41.1%).		
Belli et al. (2017) ²⁶	ACTN3 R577X polymorphism and muscle damage	Yes	High
	Athletes with the XX genotype presented a higher magnitude of muscle		-
	damage than RX and RR after an adventure race.		
Del Coso et al. $(2017)^{24}$	ACTN3 R577X polymorphism and exercise induced muscle damage	Yes	High
	X-allele triathletes presented greater signs of exercise-induced muscle		
	damage during a half-ironman race than RR homozygotes. XX and RX		
	(X-allele group) presented higher self-reported values of lower limb		
25	muscle pain than RR.		
Del Coso et al. $(2017)^{25}$	ACTN3 R577X polymorphism and exercise induced muscle damage	Yes	High
	In comparison to RR homozygotes, X-allele carriers for the R577X		
	polymorphism of the ACTN3 gene presented higher values for typical		
	markers of exercise-induced muscle damage during a competitive		
Massidda et al. (2019) ¹⁶	marathon. <i>ACTN</i> 3 R577X polymorphism and non-contact lower limb injury	Yes	High
Wassidda et al. (2019)	The ACTN3 R577X polymorphism and non-contact lower limb lightly	1 05	riigii
	severity of muscle injuries in professional football players; players		
	with the ACTN3 577XX genotype have higher odds of having muscle		
	injuries than their RR counterparts.		
Miyamoto et al. (2018) ⁴²	ACTN3 R577X polymorphism and passive stiffness and hamstring	No	High
,	strain injury		6
	NS difference between genotypes for the frequency of past hamstring		
	strain injury ($p = 0.587$)		
Del Coso et al. $(2017)^{31}$	ACTN3 R577X polymorphism and exercise induced muscle damage	Yes	High
	Marathoners with a lower CK response after a marathon had a more		
	favorable polygenic profile than marathoners with high serum CK		
15	concentrations.		
Clos et al. $(2019)^{15}$	ACTN3 R577X polymorphism and non-contact musculoskeletal injury	Yes	High
	NS difference between genotypes for non-contact musculoskeletal		
	soft-tissue injury rate		
$D_{1}(C_{1}) = (1 (2020))^{34}$	XX are the ones with the higher injury rate.	37	TT' 1
Del Coso et al. $(2020)^{34}$	ACTN3 R577X polymorphism and exercise induced muscle damage	Yes	High
	Total genotype score was higher in low-CK responders than in high- CK responders. A favorable polycopic grafile can contribute to reduc		
	CK responders. A favorable polygenic profile can contribute to reduc- ing the level of muscle damage developed during endurance exercise.		
Moreno et al. (2020) ¹⁷	<i>ACTN3</i> R577X polymorphism and non-contact lower limb injury	Yes	High
wordto et al. (2020)	XX runners had a higher frequency of sudden injuries ($p = 0.024$) and	105	mgn
	the OR for muscle-type injury was 2.0 (95%CI: $0.51-7.79$) times		
	higher than RR.		
	nfidence interval; CK = creatine-kinase; NS = not significant; OR = odds ratio.		

sport-specific actions with a greater degree of speed/power.³⁹ In fact, this is one of the assumptions that may explain the higher frequency of RR athletes participating in strength- and power-based sports.⁸ RR athletes may have a lower range of motion and muscle flexibility because lower-ankle dorsiflexion⁷ and

lower-trunk flexibility^{40,41} have been found in RR *vs.* RX–XX genotypes. A greater volume or magnitude of sport-specific actions performed at higher intensity, coupled with reduced joint mobility, may explain the higher injury rates of RR athletes compared to RX–XX athletes. Nevertheless, the studies of

this topic are scant, and the evidence is still not sufficient to link clearly the *ACTN*3RR genotype with a higher injury rate. Future investigations are needed to affirm whether RR individuals are more prone to overall sport-related injuries, particularly in high-intensity sports.

To summarize: When accounting for all types of injury, RR athletes may be more prone to sport-specific injury because the presence of α -actinin-3 in their muscles likely entails the production of more forceful and powerful contractions.

4.2. Muscle injury and exercise-induced muscle damage

4.2.1. Muscle injury

When investigating only non-contact muscle-type injuries, the evidence indicates that XX athletes are more prone to this specific type of injury. For example, Massidda et al.¹⁶ found that, among male professional football players, the odds were 2.66 times higher for those with the XX genotype to suffer a muscle injury than for their RR counterparts, with no difference between RR and RX football players. Additionally, the XX players had a 2.13-fold higher probability of having a severe injury compared with RR players, despite similar training exposure among genotypes. Similar results were shown by Clos et al.¹⁵ in a sample of professional football players; those with the XX genotype presented a 1.84-fold greater likelihood of suffering a muscle-type injury, although there was no difference in severity.¹⁵ In a sample of amateur marathoners, Moreno et al.¹⁷ showed that XX runners presented a 2.0-fold greater likelihood of muscle-type injuries when compared to RR runners. This was evident only for muscle-type injuries because the probability of the overall injury (i.e., including all types of injuries) was higher in RR vs. XX individuals. On the other hand, Miyamoto et al.⁴² did not find any effect of the ACTN3 genotype on the frequency of hamstring strain injuries in a sample of young male athletes. However, this investigation was based only on a specific body location, and that could have reduced the likelihood of finding differences among the genotypes. Collectively, these investigations indicate that α -actinin-3 deficiency, due to homozygosity in the ACTN3 R577X polymorphism, may entail a higher probability of non-contact muscle injuries. Del Coso et al.^{7,10} proposed that this could be explained by the lower capacity of α -actinin-3-deficient muscle fibers to endure physical stress. Interestingly, α -actinin-3 deficiency is habitually compensated for by a higher amount of α -actinin-2.⁴³ The current data concerning differing muscleinjury incidence among genotypes indirectly suggests that the role of α -actinin-3 is different from the role of α -actinin-2, which may explain why these muscle sarcomeric isoforms of α -actinin have remained through evolution; however, further work to elucidate the veracity of this claim is warranted.

4.2.2. Exercise-induced muscle damage

Additional evidence suggesting that α -actinin-3 deficiency, due to the *ACTN3* XX genotype, results in muscle fibers with less capacity to endure stress during exercise was found in those investigations that assessed muscle damage. When compared to RR counterparts, carriers of the *ACTN3* X allele presented higher values of indirect markers of muscle damage, such as (1) reduced jump height;^{24,25} (2) postexercise or pre- to postexercise changes in serum creatine kinase or myoglobin concentrations;^{33,39} and (3) self-reported muscle-pain values.²⁵ The results are similar when comparing XX athletes to those with the R allele,²⁶ suggesting that the R allele may have a dose-response effect, playing a protective role to prevent exercise-induced muscle damage (i.e., RR > RX > XX). Higher markers of exercise-induced muscle damage in XX individuals have also been found in some non-athletic populations,⁴⁴ although this is not always the case.⁴⁵ These outcomes indicate that α -actinin-3 has a positive role in modulating the capacity of muscle to endure potentially damaging contractions during exercise, ³⁷ whereas α -actinin-3 deficiency due to homozygosity in the ACTN3 R577X polymorphism may entail a higher probability of muscle injury/muscle damage during exercise. Of note, ACTN3 is one of several genes involved in the response to exercise-induced muscle damage; the capacity of skeletal muscle to endure the physical stress imposed by exercise may be associated with possessing favorable genotypes in several genes.^{24,32}

To summarize: When accounting for muscle-specific injury and exercise-induced muscle damage, α -actinin-3 deficiency resulting from homozygosity in the null-X allele of the *ACTN3* gene was associated with a higher incidence of injury as well as with injuries of higher severity.

4.3. Ligament injury

In the current review, 2 independent investigations determined that XX individuals are more prone to ligament injury, specifically in the ankle.^{21,22} According to the odds ratio analysis performed by Kim et al.,²¹ elite ballerinas with the XX genotype had a 4.7-fold higher risk of ankle injury than the RR- and RX-genotyped ballerinas, although this greater risk of injury was not present in other body locations, such as the knee, pelvis, or back. In an investigation done by Shang et al.,²² the frequency of the XX genotype in lance corporals with ankle sprain was higher than that in a control group without this type of injury. Interestingly, this investigation revealed that the XX genotype is associated not only with the likelihood of injury but also with the severity; the presence of the X allele resulted in greater gravity of an injury (e.g., ankle sprains from Grade I to Grade III). As previously suggested,³⁶ the higher likelihood of ligament injury in α -actinin-3-deficient athletes might be related to a dysfunction in the capacity of the muscle to hold the joint during sport-specific actions rather than to an effect of the XX genotype on the characteristics of the ligament. This is because, to date, there is no evidence suggesting that ACTN3 R577X polymorphism has any impact on ligament tissue. XX athletes may be more prone to ligament injury during exercise, although this evidence exists for the ankle joint only.

To summarize: Further replication investigations are needed to confirm this finding and to expand the perspectives to other joints.

5. Conclusion

The current systematic review indicates that most studies of the effect of the ACTN3 gene relative to injury epidemiology

Please cite this article as: Hassane Zouhal et al., Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review, Journal of Sport and Health Science (2021), https://doi.org/10.1016/j.jshs.2021.07.003

ACTN3 genotype and sport injury

in athletes have found an association between the ACTN3 genotype and injury risk and severity. Specifically, possessing the ACTN3 XX genotype may predispose athletes to a higher probability of some non-contact injuries, such as muscle injury, ankle sprains, and higher levels of exercise-induced muscular damage. Notwithstanding this interesting finding, we acknowledge that this conclusion is based on a limited number of studies (n = 13). Evidence is still emerging and, thus far, has been obtained from studies with very heterogeneous samples of athletes, with different methodologies used to quantify injury incidence and epidemiology and, in most cases, with low or small samples sizes, all of which can complicate interpretations of data.⁴⁶ Despite the preliminary evidence for the notion that the XX genotype might be associated with a higher risk of injury issues, the common view is that there is no current clinical application for genetic testing in the area of exercise prescription and injury prevention.^{47,48} The evidence herein suggests that this may not be the case. Future research into this topic is warranted in order to produce more convincing evidence-based approaches to either support or refute ACTN3 genotyping for injury prevention.

Authors' contributions

HZ and JDC were involved in the conceptualization of the study, data analysis, and the writing of the manuscript; AJ, CT, GR, NJ, CCTC, BB, and ABA were involved in the data assessment, data analysis, and the writing of the manuscript; ACH was involved in the writing of the manuscript. All authors have read and approved the final version of the manuscript, and agree with the order of presentation of the authors.

Competing interests

The authors declare that they have no competing interests.

References

- Houweling PJ, Papadimitriou ID, Seto JT, et al. Is evolutionary loss our gain? The role of ACTN3 p.Arg577Ter (R577X) genotype in athletic performance, ageing, and disease. Hum Mutat 2018;39:1774–87.
- North KN, Beggs AH. Deficiency of a skeletal muscle isoform of alphaactinin (alpha-actinin-3) in merosin-positive congenital muscular dystrophy. *Neuromuscul Disord* 1996;6:229–35.
- Hogarth MW, Garton FC, Houweling PJ, et al. Analysis of the ACTN3 heterozygous genotype suggests that α-actinin-3 controls sarcomeric composition and muscle function in a dose-dependent fashion. Hum Mol Genet 2016;25:866–77.
- Erskine RM, Williams AG, Jones DA, Stewart CE, Degens H. The individual and combined influence of ACE and ACTN3 genotypes on muscle phenotypes before and after strength training. Scand J Med Sci Sports 2014;24:642–8.
- Walsh S, Liu D, Metter EJ, Ferrucci L, Roth SM. ACTN3 genotype is associated with muscle phenotypes in women across the adult age span. J Appl Physiol 2008;105:1486–91.
- Zempo H, Tanabe K, Murakami H, Iemitsu M, Maeda S, Kuno S. ACTN3 polymorphism affects thigh muscle area. Int J Sports Med 2010;31:138–42.
- Del Coso J, Moreno V, Gutiérrez-Hellín J, et al. ACTN3 R577X genotype and exercise phenotypes in recreational marathon runners. Gene (Basel) 2019;10:413. doi:10.3390/genes10060413.

- Seto JT, Lek M, Quinlan KG, et al. Deficiency of α-actinin-3 is associated with increased susceptibility to contraction-induced damage and skeletal muscle remodeling. *Hum Mol Genet* 2011;20:2914–27.
- Yang N, Schindeler A, McDonald MM, et al. α-Actinin-3 deficiency is associated with reduced bone mass in human and mouse. *Bone* 2011;49:790–8.
- Del Coso J, Hiam D, Houweling P, Pérez LM, Eynon N, Lucía A. More than a "speed gene": ACTN3 R577X genotype, trainability, muscle damage, and the risk for injuries. Eur J Appl Physiol 2019;119:49–60.
- Ma F, Yang Y, Li X, et al. The association of sport performance with ACE and ACTN3 genetic polymorphisms: a systematic review and meta-analysis. *PLoS One* 2013;8:e54685. doi:10.1371/journal.pone.0054685.
- Ahmetov II, Egorova ES, Gabdrakhmanova LJ, Fedotovskaya ON. Genes and athletic performance: An update. *Med Sport Sci* 2016;61:41–54.
- Kikuchi N, Miyamoto-Mikami E, Murakami H, et al. ACTN3 R577X genotype and athletic performance in a large cohort of Japanese athletes. *Eur J Sport Sci* 2016;16:694–701.
- Watsford ML, Murphy AJ, McLachlan KA, et al. A prospective study of the relationship between lower body stiffness and hamstring injury in professional Australian rules footballers. *Am J Sports Med* 2010;38:2058–64.
- Clos E, Pruna R, Lundblad M, Artells R, Esquirol Caussa J. ACTN3 single nucleotide polymorphism is associated with non-contact musculoskeletal soft-tissue injury incidence in elite professional football players. *Knee Surg Sport Traumatol Arthrosc* 2019;27:4055–61.
- Massidda M, Voisin S, Culigioni C, et al. ACTN3 R577X polymorphism is associated with the incidence and severity of injuries in professional football players. *Clin J Sport Med* 2019;29:57–61.
- Moreno V, Areces F, Ruiz-Vicente D, Ordovás JM, Del Coso J. Influence of the ACTN3 R577X genotype on the injury epidemiology of marathon runners. *PLoS One* 2020;15: e0227548. doi:10.1371/journal.pone.0227548.
- Iwao-Koizumi K, Ota T, Hayashida M, et al. The ACTN3 gene is a potential biomarker for the risk of non-contact sports injury in female athletes. *J Mol Biomark Diagn* 2014;**S6**:002. doi:10.4172/2155-9929.S6-002.
- Miyamoto K, Aoki W, Ohtani Y, et al. Peptide barcoding for establishment of new types of genotype–phenotype linkages. *PLoS One* 2019;14: e0215993. doi:10.1371/journal.pone.0215993.
- 20. Kay AD, Richmond D, Talbot C, Mina M, Baross AW, Blazevich AJ. Stretching of active muscle elicits chronic changes in multiple strain risk factors. *Med Sci Sports Exerc* 2016;48:1388–96.
- Kim JH, Jung ES, Kim CH, Youn H, Kim HR. Genetic associations of body composition, flexibility and injury risk with ACE, ACTN3 and COL5A1 polymorphisms in Korean ballerinas. *J Exerc Nutrition Biochem* 2014;18:205–14.
- Shang X, Li Z, Cao X, et al. The association between the ACTN3 R577X polymorphism and noncontact acute ankle sprains. *J Sports Sci* 2015;33:1775–9.
- Qi B, Liu JQ, Liu GL. Genetic association between ACTN3 polymorphism and risk of non-acute ankle sprain. *Genet Mol Res.* 2016. doi:10.4238/gmr15048962.
- Del Coso J, Salinero JJ, Lara B, et al. ACTN3 X-allele carriers had greater levels of muscle damage during a half—ironman. *Eur J Appl Physiol* 2017;117:151–8.
- Del Coso J, Valero M, Salinero JJ, et al. ACTN3 genotype influences exercise-induced muscle damage during a marathon competition. *Eur J Appl Physiol* 2017;117:409–16.
- Belli T, Crisp AH, Verlengia R. Greater muscle damage in athletes with ACTN3 R577X (RS1815739) gene polymorphism after an ultra-endurance race: A pilot study. *Biol Sport* 2017;34:105–10.
- Moher D, Shamseer L, Clarke M, et al. Preferred reporting items for systematic review and meta-analysis protocols (PRISMA-P) 2015 statement. Syst Rev 2015;4:1. doi:10.1186/2046-4053-4-1.
- Soligard T, Schwellnus M, Alonso JM, et al. How much is too much? (Part 1) International Olympic Committee consensus statement on load in sport and risk of injury. *Br J Sports Med* 2016;50:1030–41.
- Maher CG, Sherrington C, Herbert RD, Moseley AM, Elkins M. Reliability of the PEDro scale for rating quality of randomized controlled trials. *Phys Ther* 2003;83:713–21.

ARTICLE IN PRESS

- Moher D, Liberati A, Tetzlaff J, Altman DG, Group PRISMA. Preferred reporting items for systematic reviews and meta-analyses: The PRISMA statement. *Ann Intern Med* 2009;151:264–9.
- Del Coso J, Valero M, Salinero JJ, Lara B, Gallo-Salazar C, Areces F. Optimum polygenic profile to resist exertional rhabdomyolysis during a marathon. *PloS One* 2017;12: e0172965. doi:10.1371/journal.pone.0172965.2.
- 32. Semenova EA, Miyamoto-Mikami E, Akimov EB, et al. The association of HFE gene H63D polymorphism with endurance athlete status and aerobic capacity: Novel findings and a meta-analysis. *Eur J Appl Physiol* 2020;120:665–73.
- Pimenta EM, Coelho DB, Cruz IR, et al. The ACTN3 genotype in soccer players in response to acute eccentric training. *Eur J Appl Physiol* 2012;112:1495–503.
- **34.** Del Coso J, Salinero JJ, Lara B, et al. Polygenic profile and exerciseinduced muscle damage by a competitive half-ironman. *J Strength Cond Res* 2020;**34**:1400–8.
- **35.** Yang N, MacArthur DG, Gulbin JP, et al. ACTN3 genotype is associated with human elite athletic performance. *Am J Hum Genet* 2003;**73**:627–31.
- 36. Papadimitriou ID, Lucia A, Pitsiladis YP, et al. ACTN3 R577X and ACE I/D gene variants influence performance in elite sprinters: A multi-cohort study. *BMC Genomics* 2016;17:285. doi:10.1186/s12864-016-2462-3.
- Pickering C, Kiely J. ACTN3: More than just a gene for speed. Front physiol 2017;8: 1080. doi:10.3389/fphys.2017.01080.
- Kikuchi N, Nakazato K, Min SK, Ueda D, Igawa S. The ACTN3 R577X polymorphism is associated with muscle power in male Japanese athletes. *J Strength Cond Res* 2014;28:1783–9.
- **39.** Pimenta EM, Coelho DB, Veneroso CE, et al. Effect of ACTN3 gene on strength and endurance in soccer players. *J Strength Cond Res* 2013;**27**:3286–92.

- Zempo H, Fuku N, Murakami H, Miyachi M. The relationship between alpha-actinin-3 gene R577X polymorphism and muscle flexibility. *Juntendo Med J* 2016;62(Suppl.1):118. doi:10.14789/jmj.62.s118.
- Kikuchi N, Zempo H, Fuku N, et al. Association between ACTN3 R577X polymorphism and trunk flexibility in 2 different cohorts. *Int J Sports Med* 2017;38:402–6.
- 42. Miyamoto N, Miyamoto-Mikami E, Hirata K, Kimura N, Fuku N. Association analysis of the ACTN 3 R577X polymorphism with passive muscle stiffness and muscle strain injury. *Scand J Med Sci Sports* 2018;28:1209–14.
- Seto JT, Quinlan KG, Lek M, et al. ACTN3 genotype influences muscle performance through the regulation of calcineurin signaling. *J Clin Invest* 2013;123:4255–63.
- 44. Vincent B, Windelinckx A, Nielens H, et al. Protective role of alpha-actinin-3 in the response to an acute eccentric exercise bout. J Appl Physiol (1985) 2010;109:564–73.
- Clarkson PM, Hoffman EP, Zambraski E, et al. ACTN3 and MLCK genotype associations with exertional muscle damage. J Appl Physiol (1985) 2005;99:564–9.
- Baltazar-Martins G, Gutiérrez-Hellín J, Aguilar-Navarro M, et al. Effect of ACTN3 genotype on sports performance, exercise-induced muscle damage, and injury epidemiology. *Sports (Basel)* 2020;8:99. doi:10.3390/ sports8070099.
- Pickering C, Kiely J, Grgic J, Lucia A, Del Coso J. Can genetic testing identify talent for sport? *Genes (Basel)* 2019;10:972. doi:10.3390/ genes10120972.
- Vlahovich N, Hughes DC, Griffiths LR, et al. Genetic testing for exercise prescription and injury prevention: AIS-Athlome consortium-FIMS joint statement. *BMC Genomics* 2017;18:818. doi:10.1186/ s12864-017-4185-5.

Please cite this article as: Hassane Zouhal et al., Association between ACTN3 R577X genotype and risk of non-contact injury in trained athletes: A systematic review, Journal of Sport and Health Science (2021), https://doi.org/10.1016/j.jshs.2021.07.003

10