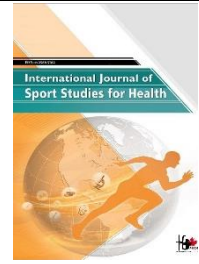


## International Journal of Sport Studies for Health

Journal Homepage



# Advancing Human Performance: Genomic Innovations and Ethical Imperatives in Sports Science

Danial. Kahrizi<sup>1\*</sup> <sup>1</sup> Biotechnology Department, Tarbiat Modares University, Tehran Iran

\* Corresponding author email address: dkahrizi@modares.ac.ir

## Article Info

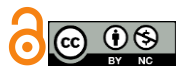
## Article type:

Letter to the Editor

## How to cite this article:

Kahrizi, D. (2025). Advancing Human Performance: Genomic Innovations and Ethical Imperatives in Sports Science . *International Journal of Sport Studies for Health*, 8(4), 1-3.

<http://dx.doi.org/10.61838/kman.intjssh.8.4.5>



© 2025 the authors. Published by KMAN Publication Inc. (KMANPUB), Ontario, Canada. This is an open access article under the terms of the Creative Commons Attribution-NonCommercial 4.0 International (CC BY-NC 4.0) License.

## A B S T R A C T

The rapid convergence of genomic technologies and sports science is redefining the boundaries of human athletic potential. As a researcher at the intersection of molecular biology and sports medicine, I applaud your journal's commitment to this transformative field and wish to expand upon critical developments in gene expression regulation, CRISPR-Cas9 applications, and next-generation sequencing (NGS) that demand scholarly attention and ethical scrutiny.

**Keywords:** *Genomics, Gene Expression, Next-Generation Sequencing (NGS), Polygenic Risk Scores (PRS), Athletic Performance*

## 1. Gene Expression Plasticity: Beyond Hereditary Determinism

Athletic prowess is no longer viewed through the narrow lens of static genetic inheritance. Cutting-edge research reveals that polymorphisms in ACTN3 (rs1815739) and PPARA (rs4253778) modulate fast-twitch muscle fiber composition and lipid metabolism, respectively (1). Epigenetic regulation-particularly DNA methylation at HIF1A promoters under hypoxic conditions-demonstrates

how environmental stimuli dynamically shape performance phenotypes (2). A 2024 study further identified exercise-induced histone acetylation at the PGC-1 $\alpha$  locus, enhancing mitochondrial biogenesis in endurance athletes (3). These findings challenge the dichotomy of "nature versus nurture," emphasizing a continuum of gene-environment interplay.

## 2. CRISPR-Cas9: Therapeutic Promise vs. Enhancement Ethics

Article history:

Received 22 March 2025

Revised 18 April 2025

Accepted 23 April 2025

Published online 01 October 2025

Therapeutic applications of CRISPR, such as MSTN knockout for sarcopenia treatment or COL5A1 repair to mitigate tendinopathy risk, offer groundbreaking solutions for injury rehabilitation (4). However, the potential misuse of germline editing to engineer "designer athletes" with enhanced erythropoietin (EPO) expression or myostatin deficiency raises profound ethical concerns (5). The World Anti-Doping Agency's (WADA) 2025 provisional ban on gene-editing technologies in competitive sports underscores the urgency of global regulatory frameworks (6). Science Translational Medicine editorial advocates for an international consortium to monitor CRISPR applications, balancing innovation with competitive integrity (7).

### 3. NGS and Polygenic Risk Scores: Personalization and Pitfalls

NGS-driven polygenic risk scores (PRS) now enable stratification of athletes by injury susceptibility (e.g., COL1A1 variants and ACL rupture risk) and metabolic efficiency (8). Elite teams increasingly integrate whole-genome sequencing with AI-driven analytics to optimize training loads and nutritional interventions (9). For instance, a 2024 trial in The Lancet Digital Health demonstrated a 23% reduction in hamstring injuries among soccer players using PRS-guided regimens. Yet, disparities in access to genomic technologies risk exacerbating inequalities between resource-rich and developing nations. The absence of standardized protocols for data anonymization further complicates ethical implementation (10).

### 4. Toward a Multidisciplinary Framework for Responsible Innovation

The integration of multi-omics (metabolomics, proteomics) with wearable biometrics promises unparalleled insights into performance optimization. However, this demands collaboration beyond academia: sports federations, bioethicists, and policymakers must jointly address challenges such as informed consent in genetic testing and the delineation of therapeutic vs. enhancement thresholds. Initiatives like the IOC's 2024 Global Summit on Genomic Equity highlight the need for inclusive policies that democratize access while safeguarding privacy (11).

### 5. Conclusion

As CRISPR and NGS technologies advance, the sports science community stands at a crossroads. Will we embrace these tools to foster equitable health advancements, or

permit a new era of "genetic elitism"? I urge your journal to spearhead this discourse through dedicated special issues and expert panels, ensuring that genomic innovation aligns with the core values of athleticism: fairness, inclusivity, and respect for human dignity.

### Authors' Contributions

Not applicable.

### Declaration

In order to correct and improve the academic writing of our paper, we have used the language model ChatGPT.

### Transparency Statement

Not applicable.

### Acknowledgments

None.

### Declaration of Interest

The author reports no conflict of interest.

### Funding

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

### Ethical Considerations

Not applicable.

### References

1. Bulgay C, Cepicka L, Dalip M, Yıldırım S, Ceylan HI, Yılmaz Ö, et al. The relationships between ACTN3 rs1815739 and PPARA-α rs4253778 gene polymorphisms and athletic performance characteristics in professional soccer players. BMC sports science, medicine & rehabilitation. 2023;15(1):121. [PMID: 37749582] [PMCID: PMC10518950] [DOI]
2. Earhart ML, Blanchard TS, Strowbridge N, Sheena R, McMaster C, Staples B, et al. Heatwave resilience of juvenile white sturgeon is associated with epigenetic and transcriptional alterations. Sci Rep. 2023;13(1):15451. [PMID: 37723229] [PMCID: PMC10507091] [DOI]
3. Li J, Zhang S, Li C, Zhang X, Shan Y, Zhang Z, et al. Endurance exercise-induced histone methylation modification involved in skeletal muscle fiber type transition and mitochondrial biogenesis. Sci Rep. 2024;14(1):21154. [PMID: 39256490] [PMCID: PMC11387812] [DOI]
4. Chae SY, Jeong E, Kang S, Yim Y, Kim JS, Min DH. Rationally designed nanoparticle delivery of Cas9

ribonucleoprotein for effective gene editing. Journal of controlled release : official journal of the Controlled Release Society. 2022;345:108-19. [PMID: 35247491] [DOI]

5. Bojarczuk A. Ethical Aspects of Human Genome Research in Sports-A Narrative Review. Genes (Basel). 2024;15(9). [PMID: 39336807] [PMCID: PMC11430849] [DOI]

6. Ren X, Shi Y, Xiao B, Su X, Shi H, He G, et al. Gene Doping Detection From the Perspective of 3D Genome. Drug testing and analysis. 2025. [PMID: 39757126] [DOI]

7. Xue Y, Shang L. Governance of Heritable Human Gene Editing World-Wide and Beyond. Int J Environ Res Public Health. 2022;19(11). [PMID: 35682323] [PMCID: PMC9180052] [DOI]

8. Chikowore T, Läll K, Micklesfield LK, Lombard Z, Goedecke JH, Fatumo S, et al. Variability of polygenic prediction for body mass index in Africa. Genome medicine. 2024;16(1):74. [PMID: 38816834] [PMCID: PMC11140909] [DOI]

9. Suwinski P, Ong C, Ling MHT, Poh YM, Khan AM, Ong HS. Advancing Personalized Medicine Through the Application of Whole Exome Sequencing and Big Data Analytics. Front Genet. 2019;10:49. [PMID: 30809243] [PMCID: PMC6379253] [DOI]

10. Badr J, Motulsky A, Denis JL. Digital health technologies and inequalities: A scoping review of potential impacts and policy recommendations. Health policy (Amsterdam, Netherlands). 2024;146:105122. [PMID: 38986333] [DOI]

11. Magee P, Ienca M, Farahany N. Beyond neural data: Cognitive biometrics and mental privacy. Neuron. 2024;112(18):3017-28. [PMID: 39326392] [DOI]