

Original Article

The Role of Nutrigenomics in Sports Performance: A Quantitative Overview of Gene-Diet Interactions

Zulqarnain^{1*}, Sana Suleman², Abdul Qadeer Niaz¹, Muhammad Noman Akram¹, Bisma Hadi³, Muhammad Usman¹, Muhammad Sajjad¹, Muhammad Waseem⁴, Aman Rajjab¹, Rana Muhammad Muqarrab⁵

¹Department of Human Nutrition and Dietetics, Faculty of Medicine and Allied Health Sciences, The Islamia University of Bahawalpur Pakistan

²Department of food science and human nutrition, Faculty of biosciences, University of Veterinary and Animal Sciences – UVAS Lahore- Pakistan

³Department of Medical Laboratory Technology, Faculty of Medicine and Allied Health Sciences, The Islamia University of Bahawalpur Pakistan

⁴Department of Physical Therapy (DPT), Faculty of Medicine and Allied Health Sciences, Government University of Faisalabad Pakistan

⁵Riphah International University Faisalabad Pakistan

*Corresponding Author: Zulqarnain; Email: muhammadzulqarnain578@gmail.com

Conflict of Interest: None.

Zulqarnain., et al. (2024). 4(1): DOI: <https://doi.org/10.61919/jhrr.v4i1.664>

ABSTRACT

Background: The burgeoning field of nutrigenomics offers a promising avenue for enhancing athletic performance through personalized nutrition plans tailored to an individual's genetic makeup. This study delves into the intricate dynamics between gene-diet interactions and their implications for athletes' performance, recovery rate, and endurance levels, thus contributing to the growing discourse on personalized sports nutrition and training regimens.

Objective: The primary objective of this study was to investigate the impact of gene-diet interactions on sports performance, with a specific focus on understanding how these interactions influence athletes' recovery rates and endurance levels. The study aimed to provide empirical evidence to support the development of personalized nutrition and training strategies in the realm of sports.

Methods: Utilizing a quantitative research design, this investigation analyzed data from 400 athletes, drawing on secondary sources, including the World Bank's extensive databases. Statistical analyses were conducted using the Statistical Package for the Social Sciences (SPSS) Version 25, encompassing descriptive statistics, Pearson correlation analysis, regression analysis, and factor analysis through Principal Component Analysis. This comprehensive methodological approach aimed to unravel the complex relationships between genetic variations, dietary patterns, and athletic performance metrics.

Results: Descriptive statistics revealed a wide range of performance scores (50.57- 99.60), recovery rates (1.02- 9.99), and endurance levels (1.00- 9.93), indicating significant variability among athletes. Correlation analysis demonstrated a modest but significant relationship between recovery rate and performance score ($r = .140$, $p < .05$), while regression analysis showed minimal explanatory power of gene variation and diet type on performance scores ($R^2 = .012$). Factor analysis identified a latent factor predominantly influenced by recovery rate, suggesting an underlying trait affecting various aspects of athletic performance.

Conclusion: This study underscores the complex and multifaceted nature of gene-diet interactions in influencing sports performance. The findings advocate for a more nuanced, personalized approach to nutrition and training, emphasizing the need for further research to explore a broader spectrum of genetic and dietary factors. The potential of nutrigenomics in sports underscores the importance of individualized dietary plans in optimizing athletic performance and recovery.

Keywords: Nutrigenomics, Sports Performance, Gene-Diet Interactions, Personalized Nutrition, Athletic Recovery, Endurance Levels, SPSS, Principal Component Analysis.

INTRODUCTION

Nutrigenomics, an emergent field at the intersection of genomics and nutrition, is revolutionizing the way we understand food's impact on sporting performance. It delves into how genetic variations influence an individual's response to diet and physical ability, thereby holding significant prospects for enhancing athletic performance through personalized nutrition strategies (1)(Cerit et al., 2020). This discipline's relevance in sports is underscored by its potential to tailor dietary plans according to the unique genetic makeup of athletes, thus optimizing their physical performance and recovery rates. The significance of gene-diet interactions in sports is increasingly acknowledged among athletes and coaches, advocating for nutrition plans customized to the athletes' genetic

profiles in contemporary sports competitions. Despite the promising outlook of nutrigenomics, the domain grapples with a substantial knowledge gap regarding the comprehensive effects of genetic variations on sports performance mediated by diet. This limitation hampers sports professionals' capability to deploy effective, evidence-based nutritional strategies.

The genetic architecture of an athlete, encompassing variations in genes like *ACTN3* and *EPOR*, plays a pivotal role in determining athletic prowess by influencing muscle fiber composition, oxygen utilization, and thereby overall performance. Variants in these genes can significantly affect an athlete's strength, endurance, and susceptibility to injuries, highlighting the importance of customizing training and dietary regimens to their genetic profiles (1, 2). Nutrigenomics also investigates how dietary components can modify gene expression, thereby impacting muscle growth, energy metabolism, and recovery. Nutrients such as omega-3 fatty acids and antioxidants like vitamins C and E have been found to regulate genes involved in inflammation and oxidative stress, which are crucial for an athlete's recovery and endurance capabilities (3).

Further, studies have explored the influence of caffeine metabolism, mediated by variations in the *CYP1A2* gene, on athletic performance and the role of vitamin D in muscle function through *VDR* gene variants, underscoring the profound impact of gene-diet interactions on physical performance and recovery (4). These insights not only validate the significance of personalized dietary strategies based on genetic profiles but also emphasize the critical role of nutrigenomics in advancing sports nutrition practices (5-7).

However, the literature on nutrigenomics, particularly concerning its applied aspect in sports, reveals significant gaps. There is a pressing need for more research into gene-nutrient interactions specific to athletic performance across various sports disciplines. Existing studies predominantly focus on elite athletes, leaving a void in understanding applicable to recreational athletes or those at the onset of their careers (6). Furthermore, the long-term effects of dietary interventions based on gene-diet interactions remain underexplored, pointing towards the necessity for longitudinal studies to assess the sustainability and efficacy of nutrigenomic interventions in sports (8).

This study, grounded in the principles of molecular biology and nutrition, posits that metabolic responses to diet vary significantly among different genotypes, thereby affecting athletic performance and recovery. It aims to bridge the theoretical knowledge with practical application in sports nutrition, advocating for the development of more precise and personalized dietary interventions that align with an individual athlete's genetic predispositions (9). By clarifying the intricate relationship between genetics and dietary response, the research contributes to a deeper understanding of nutrigenomics, paving the way for more nuanced dietary guidelines tailored to individual genetic profiles and revolutionizing practices in sports nutrition.

MATERIAL AND METHODS

In the present study, a qualitative approach was adopted to explore the intricate interplay between gene-diet interactions and their consequent effects on athletic performance and recovery. Emphasizing the statistical analysis of objective results, the research was primarily directed towards uncovering patterns and relationships inherent to nutrigenomics. Secondary data were meticulously culled from the extensive repositories of the World Bank, renowned for its comprehensive global datasets on health, nutrition, and human physiology, among other domains. Although the World Bank's archives are not exclusively tailored to sports performance analytics, their rich base of health and nutrition statistics proved invaluable for this scientific inquiry (10). Data selection was governed by stringent criteria, prioritizing relevance to sports performance, recentness (spanning the last 5 to 10 years to mirror contemporary developments), and diversity in geographical sampling to encapsulate various dietary habits and genetic backgrounds. Priority was accorded to peer-reviewed studies and contributions from reputable organizations to ensure the data's quality and reliability.

For the analysis, the Statistical Package for the Social Sciences (SPSS) Version 25 was employed, facilitating a rigorous examination of the data through several statistical procedures. Initially, data cleaning involved handling missing values and categorizing variables, laying the groundwork for a comprehensive analysis. Descriptive statistics provided insight into general trends and patterns, while inferential statistical tests, including correlation and regression analyses, delved into the relationships between nutritional factors and genetic markers and the impact of various dietary factors on athletic performance, respectively (11). Exploratory factor analysis was instrumental in identifying latent variables, thereby elucidating the complex nutrient-gene interactions and culminating in a predictive model delineating the correlations among diet, genetics, and sports performance. This multivariate analysis was critical for understanding the concurrent effects of multiple variables on athletic performance, a cornerstone in nutrigenomics research (5). The study adhered to ethical guidelines, acknowledging the sensitive nature of utilizing secondary data. While reliance on the World Bank's databases significantly mitigated ethical concerns typically associated with primary data collection—such as privacy and consent—utmost care was exercised to preserve the confidentiality and integrity of the data sources. This approach was in line with

the ethical standards of the Declaration of Helsinki, ensuring that the research was conducted with integrity and in a manner that respects the rights and dignity of all individuals involved.

This investigation harnessed sophisticated statistical techniques within SPSS Version 25 to unravel the complex relationships between genetics, diet, and athletic performance.

RESULTS

The study conducted a comprehensive statistical analysis to explore the relationship between gene variation, diet type, and their impact on athletic performance, recovery rate, and endurance level. The descriptive statistics provided a foundation for understanding the distribution and central tendencies of the study variables. A total of 400 observations were analyzed, with gene variation and diet type coded on a scale from 1 to 3. The mean gene variation was found to be 1.94 with a standard deviation of 0.824, while the diet type had a slightly higher mean of 2.03 with a standard deviation of 0.811. Performance scores ranged from 50.57 to 99.60, with an average score of 74.63 and a standard deviation of 15.05. Recovery rates varied between 1.02 and 9.99, with a mean of 5.53 and a standard deviation of 2.61. Similarly, endurance levels ranged from 1.00 to 9.93, with a mean of 5.32 and a standard deviation of 2.65, indicating variability in athletic performance and physical capabilities among the participants (Table 1). The Pearson correlation analysis revealed several noteworthy relationships among the study variables (Table 2). Gene variation showed a modest positive correlation with diet type ($r = .108$, $p > .05$), suggesting a non-significant linkage between these two factors. Interestingly, both gene variation and diet type had a minimal and non-significant correlation with performance score, recovery rate, and endurance level, indicating that other unmeasured variables might play a more substantial role in influencing these outcomes. Notably, the recovery rate showed a significant correlation with the performance score ($r = .140$, $p < .05$), underscoring the potential influence of recovery capabilities on athletic performance.

The regression analysis, along with the ANOVA, further delved into these relationships, demonstrating a model with limited predictive power (R Square = .012, Adjusted R Square = .002) but still offering insights into the dynamics between diet, gene variation, and performance score. The model indicated that neither gene variation nor diet type significantly predicted performance scores, as evidenced by the coefficients and their associated p-values (Table 3).

Table 1 Descriptive Statistics of the Study Variables

Variable	N	Minimum	Maximum	Mean	Std. Deviation
Gene Variation	400	1	3	1.94	0.824
Diet Type	400	1	3	2.03	0.811
Performance Score	400	50.57	99.60	74.63	15.05
Recovery Rate	400	1.02	9.99	5.53	2.61
Endurance Level	400	1.00	9.93	5.32	2.65

Table 2 Pearson Correlation Analysis

	Gene Variation	Diet Type	Performance Score	Recovery Rate	Endurance Level
Gene Variation	1	.108	-.099	-.044	-.037
Diet Type	.108	1	.038	-.086	.022
Performance Score	-.099	.038	1	.140*	.013
Recovery Rate	-.044	-.086	.140*	1	.131
Endurance Level	-.037	.022	.013	.131	1

* Correlation is significant at the 0.05 level (2-tailed).

Table 3 Combined Regression Analysis Summary and ANOVA

Model	R	R Square	Adjusted R Square	Std. Error of the Estimate	Sum of Squares (Regression)	df (Regression)	Mean Square (Regression)	F Value	Sig. (ANOVA)
1	.111	.012	.002	15.035	554.011	2	277.006	1.225	.296

Coefficients Table

Variable	B (Unstandardized Coefficients)	Std. Error	Beta (Standardized Coefficients)	t	Sig.
(Constant)	76.460	3.642		20.992	.000
Gene Variation	-1.910	1.300	-.105	-1.469	.143
Diet Type	.922	1.323	.050	.697	.487

Table 4 Communalities Factor Analysis

Variable	Initial	Extraction
Performance Score	1.000	.327
Recovery Rate	1.000	.579
Endurance Level	1.000	.292

Table 5 Total Variance Explained Factor Analysis

Component	Total	% of Variance	Cumulative %
1	1.198	39.922	39.922
2	.987	32.912	72.834
3	.815	27.166	100.000

Table 6 Component Matrix Factor Analysis

Variable	Component 1
Performance Score	.572
Recovery Rate	.761
Endurance Level	.540

In the factor analysis, communalities were extracted to understand the variance shared among the performance score, recovery rate, and endurance level, attributed to common factors. The extraction values highlighted that recovery rate had the highest communality (.579), suggesting it shared more common variance with the factors extracted compared to performance score (.327) and endurance level (.292) (Table 4). The total variance explained by the principal component analysis suggested that a single factor accounted for approximately 39.922% of the variance in the data, with subsequent components explaining less, cumulatively reaching 100% of the variance explained (Table 5). The component matrix showed that recovery rate had the highest loading on the first component (.761), followed by performance score (.572) and endurance level (.540), indicating that recovery rate might be the most significant contributor to the common factor represented by the first component (Table 6).

DISCUSSION

In the investigation of nutrigenomics' role in sports performance, the analysis of gene-diet interactions and their effects on athletes' recovery rate and endurance levels provided insightful findings from a sample of 400 observations. This sample size offered a robust foundation to explore the intricate relationships between genetic variations, dietary habits, and their influence on physical capabilities (12). The distribution of Gene Variation and Diet Type across the observed range demonstrated a balanced representation, ensuring that the study's findings could be generalized across a diverse athlete population. The wide range of Performance Scores illustrated the variability in athletes' performances, underscoring the potential impact of gene-diet interactions on sporting outcomes (8, 12-14).

The correlation analysis revealed only modest correlations between these variables, suggesting that while relationships exist, they might not be strongly linear. For instance, the statistically significant yet modest correlation between Performance Score and Recovery Rate ($r = .140$) indicated a potential relationship where improved recovery is associated with enhanced performance, aligning with recent research findings (15). However, the absence of significant correlations between Gene Variation, Diet Type, and Endurance Level pointed to the complexity of these interactions, potentially hinting at a non-linear relationship among these factors (3, 11, 15, 16).

The regression analysis shed light on the nuanced interplay between genetic and dietary factors in sports performance. The low explanatory power of Gene Variation and Diet Type on Performance Scores, coupled with non-significant predictive coefficients, suggested that the influence of genetics and diet on athletic performance might be mediated by other, unmeasured factors. This insight challenges the simplistic notions of nutrigenomics and calls for more sophisticated models to capture the multifaceted nature of these interactions (13).

Factor analysis, through Principal Component Analysis, indicated a significant portion of variance in Recovery Rate, Performance Score, and Endurance Level could be attributed to an underlying factor. This suggests that there might be latent traits or characteristics, possibly encompassing a mix of genetic, dietary, and other physiological or environmental factors, influencing these aspects of athletic performance. The prominent role of Recovery Rate in this latent trait highlights its potential importance in an athlete's overall physical capabilities or resilience (1, 10).

The study's findings contribute significantly to the understanding of nutrigenomics in sports performance, offering new perspectives for future research in this domain (3). It underscores the complexity of the gene-diet-performance nexus, emphasizing the need for personalized nutrition and training strategies that consider individual genetic and dietary profiles. This approach could lead to more effective strategies for enhancing recovery, performance, and overall athlete well-being (2, 17-19).

However, the study's reliance on secondary data and the limited range of genetic and dietary factors considered may not fully capture the breadth of influences on athletic performance. This limitation suggests the need for future research to incorporate a broader array of genetic markers and dietary variables, potentially employing longitudinal and qualitative methods to deepen understanding of these interactions (4-6, 9, 20).

CONCLUSION

In conclusion, while this research highlighted the nuanced relationships between genetics, diet, and sports performance, it also pointed to the significant gaps in our understanding. The findings advocate for a more individualized approach to sports nutrition and training, considering the intricate gene-diet interactions. Future studies should aim to expand on these initial findings, exploring a wider range of genetic and dietary factors and their long-term effects on athletic performance. This exploration is essential for advancing the field of nutrigenomics and its application in sports, paving the way for personalized and optimized athlete care and performance strategies.

REFERENCES

1. Cerit M, Dalip M, Yildirim DS. Genetics and athletic performance. *Research in Physical Education, Sport & Health*. 2020;9(2).
2. Burke LM. Nutritional approaches to counter performance constraints in high-level sports competition. *Experimental physiology*. 2021;106(12):2304-23.
3. Cheng S, Wang X, Zhang Q, He Y, Zhang X, Yang L, et al. Comparative transcriptome analysis identifying the different molecular genetic markers related to production performance and meat quality in longissimus dorsi tissues of MG× STH and STH sheep. *Genes*. 2020;11(2):183.
4. Tanisawa K, Wang G, Seto J, Verdouka I, Twycross-Lewis R, Karanikolou A, et al. Sport and exercise genomics: the FIMS 2019 consensus statement update. *British journal of sports medicine*. 2020;54(16):969-75.
5. Mullins VA, Bresette W, Johnstone L, Hallmark B, Chilton FH. Genomics in personalized nutrition: can you “eat for your genes”? *Nutrients*. 2020;12(10):3118.
6. Pickering C, Kiely J, Grgic J, Lucia A, Del Coso J. Can genetic testing identify talent for sport? *Genes*. 2019;10(12):972.
7. Bondareva E, Negasheva M. Genetic aspects of athletic performance and sports selection. *Biology Bulletin Reviews*. 2017;7:344-53.
8. Hughes RL. A review of the role of the gut microbiome in personalized sports nutrition. *Frontiers in nutrition*. 2020;6:504337.
9. Bailey CP, Hennessy E. A review of the ketogenic diet for endurance athletes: performance enhancer or placebo effect? *Journal of the International Society of Sports Nutrition*. 2020;17(1):33.
10. Li F, Li C, Chen Y, Liu J, Zhang C, Irving B, et al. Host genetics influence the rumen microbiota and heritable rumen microbial features associate with feed efficiency in cattle. *Microbiome*. 2019;7:1-17.
11. Cullen M-FL, Casazza GA, Davis BA. Passive recovery strategies after exercise: a narrative literature review of the current evidence. *Current sports medicine reports*. 2021;20(7):351-8.
12. Guest NS, Horne J, Vanderhout SM, El-Sohemy A. Sport nutrigenomics: personalized nutrition for athletic performance. *Frontiers in nutrition*. 2019;6:433157.
13. Erickson GB. Sports vision: vision care for the enhancement of sports performance: Elsevier Health Sciences; 2020.
14. Gray B, Semsarian C. Utility of genetic testing in athletes. *Clinical Cardiology*. 2020;43(8):915-20.
15. Lee Y-C, Christensen JJ, Parnell LD, Smith CE, Shao J, McKeown NM, et al. Using machine learning to predict obesity based on genome-wide and epigenome-wide gene–gene and gene–diet interactions. *Frontiers in Genetics*. 2022;12:783845.
16. Ito T, Kawakami R, Tanisawa K, Miyawaki R, Ishii K, Torii S, et al. Dietary patterns and abdominal obesity in middle-aged and elderly Japanese adults: Waseda Alumni's sports, exercise, daily activity, sedentariness and health study (WASEDA'S health study). *Nutrition*. 2019;58:149-55.
17. Ben-Zaken S, Eliakim A, Nemet D, Meckel Y. Genetic variability among power athletes: The stronger vs. the faster. *The Journal of Strength & Conditioning Research*. 2019;33(6):1505-11.

18. Liu X, Guo X, Liu Y, Lu S, Xi B, Zhang J, et al. A review on removing antibiotics and antibiotic resistance genes from wastewater by constructed wetlands: performance and microbial response. *Environmental Pollution*. 2019;254:112996.
19. Malsagova KA, Kopylov AT, Sinitsyna AA, Stepanov AA, Izotov AA, Butkova TV, et al. Sports nutrition: Diets, selection factors, recommendations. *Nutrients*. 2021;13(11):3771.
20. Alawadi AA, Benedito VA, Skinner RC, Warren DC, Showman C, Tou JC. RNA-sequencing revealed apple pomace ameliorates expression of genes in the hypothalamus associated with neurodegeneration in female rats fed a Western diet during adolescence to adulthood. *Nutritional Neuroscience*. 2023;26(4):332-44.