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Correction

Correction: Yenmis et al. The Distribution of Sport Performance Gene Variations Through COVID-19 Disease Severity. *Diagnostics* 2025, *15*, 701

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Error in Figure

In the original publication [1], there was a mistake in Figure 1 as published. The corresponding image file for Figure 1c had been inadvertently saved under the label "a" on the storage device, leading to the incorrect image being incorporated into the manuscript during figure preparation. The corrected Figure 1 and figure legend appear below. The authors state that the scientific conclusions are unaffected. This correction was approved by the Academic Editor. The original publication has also been updated.

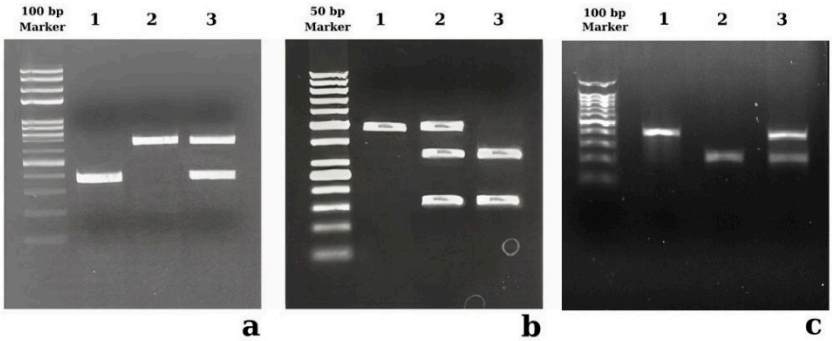


Figure 1. The restriction enzyme patterns of ACE rs4646994 (**a**), PPRGC1A rs8192678 (**b**), and ACTN3 rs1815739 (**c**) polymorphisms. (**a**) The first lane is a 100 bp size marker, lane 1 is del/del (homozygous), lane 2 is ins/ins (homozygous), and lane 3 is ins/del (heterozygous). (**b**) The first lane is a 50 bp size marker, lane 1 is TT (homozygous), lane 2 is TC (heterozygous), and lane 3 is CC (homozygous). (**c**) The first lane is a 100 bp size marker, lane 1 is CC (homozygous), lane 2 is TT (homozygous), and lane 3 is CT (heterozygous).

Check for updates

Received: 26 May 2026
Accepted: 12 June 2026
Published: 15 July 2026

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Reference

1. Yenmis, G.; Kallenci, I.; Dokur, M.; Koc, S.; Yalinkilic, S.B.; Atak, E.; Demirbilek, M.; Arkan, H. The Distribution of Sport Performance Gene Variations Through COVID-19 Disease Severity. *Diagnostics* **2025**, *15*, 701. [CrossRef] [PubMed]

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