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Corrigendum: Genome-wide association study identifying variants related to performance and injury in high-performance athletes

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KEYWORDS

genome-wide association, genetics, DNA, lower limb musculoskeletal injury

A Corrigendum on Genome-wide association study identifying variants related to performance and injury in high-performance athletes

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In the published article, there was an error in [Figure 1](#) as published. While the Figure legend is correct, the image is incorrect. The corrected [Figure 1](#) and its caption appear below.

In the published article, there was an error in [Figure 2](#) as published. While the Figure legend is correct, the image is incorrect. The corrected [Figure 2](#) and its appear below.

The authors apologize for these errors and state that this does not change the scientific conclusions of the article in any way.

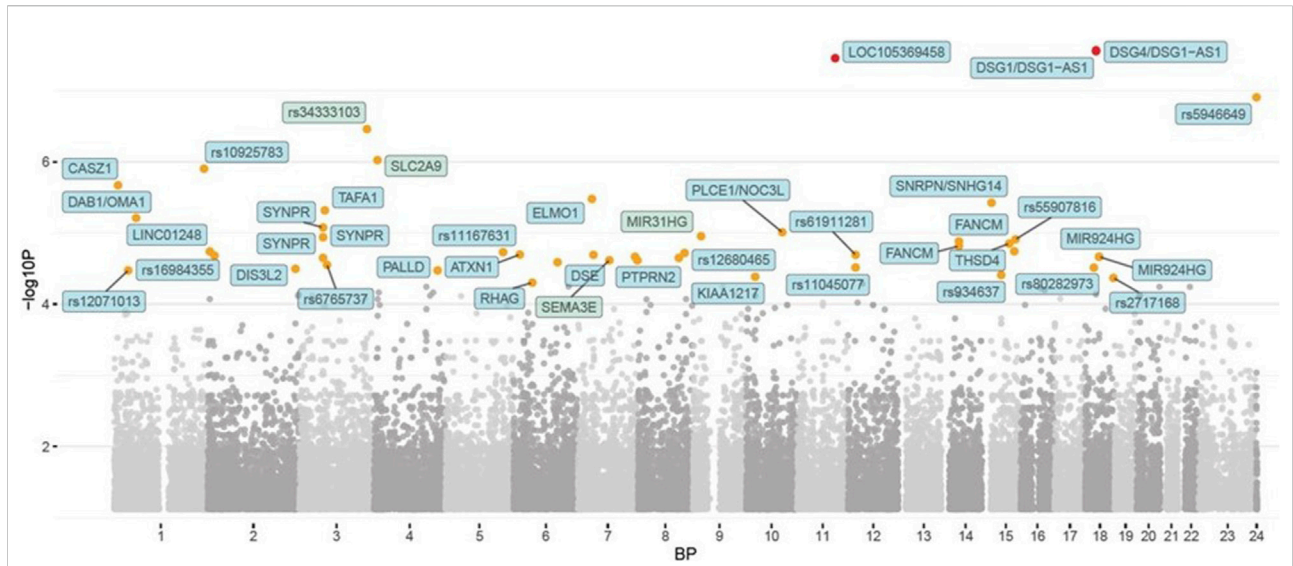


FIGURE 1
 Manhattan plot representing the p values of the genome-wide association in reaching the podium (medalist) or not. The orange dots represent $p < 10^{-5}$ while the red dots represent $p < 10^{-8}$ (i.e., strong genome-wide significance).

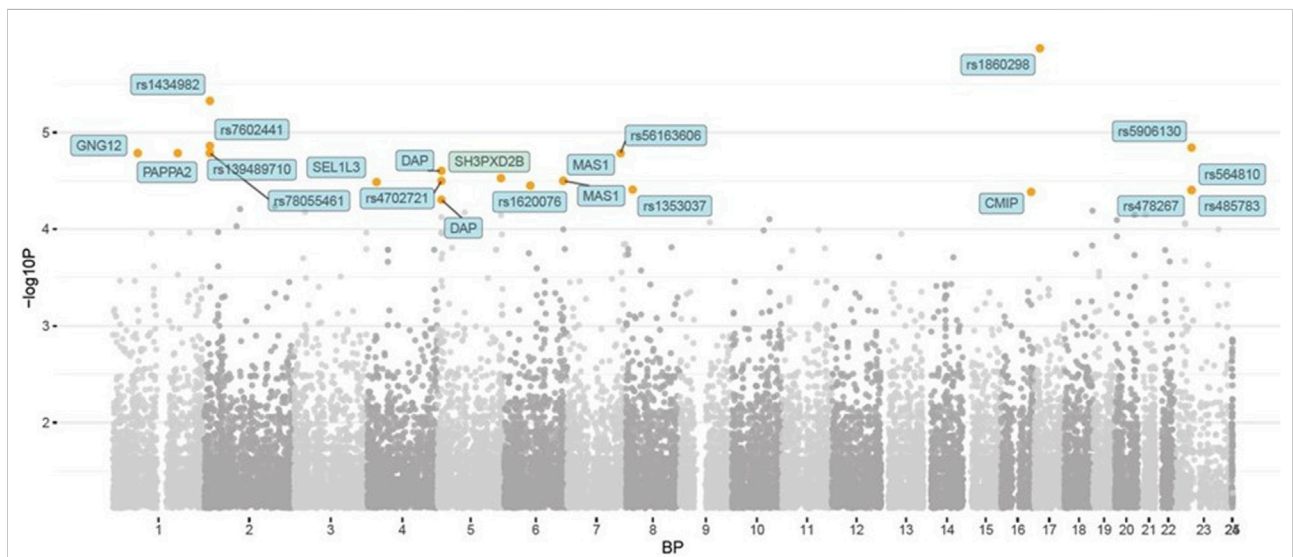


FIGURE 2
 Manhattan plot representing the p values of the genome-wide association in being injured or not. The orange dots represent $p < 10^{-5}$ while the red dots (N/A) represent $p < 10^{-8}$ (i.e., strong genome-wide significance).