

Correction

Correction: Jaramillo-Valverde *et al.* Exploratory Study on Genetic Variants Related to Hydatidosis Susceptibility and Albendazole Pharmacogenetics in the Cusco Region in Peru. 2025; 17(3): 40566

Luis Jaramillo-Valverde^{1,*}, Marlon Yuri Garcia-Paitan¹, Dolly Landeo¹, Saul J. Santivañez¹, Ramon Cacacabelos^{1,2}

¹School of Medicine, Universidad Continental, 12001 Huancayo, Peru

²EuroEspes Biomedical Research Center, Institute of Medical Science and Genomic Medicine, 15165 Bergondo, Corunna, Spain

*Correspondence: ljaramillo@continental.edu.pe (Luis Jaramillo-Valverde)

Academic Editor: Gustavo Caetano-Anollés

Submitted: 14 January 2026 Accepted: 10 May 2026 Published: 17 June 2026

The authors would like to correct several errors in the *Exploratory Study on Genetic Variants Related to Hydatidosis Susceptibility and Albendazole Pharmacogenetics in the Cusco Region in Peru* (Frontiers in Bioscience-Scholar, 2025; 17(3): 40566), as errors were introduced in the annotation of single-nucleotide polymorphism (SNP) reference identifiers, SNP positions, substituted amino acid annotations and SNP genotype annotation.

The authors would like to request the following corrections to this paper [1]:

(1) SNP reference identifiers:

rs2779249 → rs2069705 (4 occurrences throughout the full text, including once each in Table 1 and Table 2);

rs11568821 → rs3761548 (5 occurrences throughout the full text, including once each in Table 1 and Table 2);

rs2266780 → rs3745274 (6 occurrences throughout the full text, including once each in Table 1, Table 2, and Table 3); In addition, the sentence “Polymorphisms in CYP2B6, such as CYP2B66 (rs3745274) and CYP2B69 (rs28399499), affect albendazole metabolism” should be corrected to: “Polymorphisms in CYP2B6 (rs3745274) and CYP2B69 (rs28399499) affect albendazole metabolism.”

rs2266782 → rs2279343 (6 occurrences throughout the full text, including once each in Table 1, Table 2, and Table 3);

rs8330 → rs2070673 (6 occurrences throughout the full text, including once each in Table 1, Table 2, and Table 3).

(2) SNP positions:

+221 → +154 (once in each of Table 1 and Table 2);

−298 → 20230 (once in each of Table 1 and Table 2).

(3) Substituted amino acid annotation:

MII → MIT (2 occurrences throughout the full text, including once in Table 1).

(4) SNP genotype annotation:

In Table 2, for *VDR* rs7975232 (intron 8), the homozygous genotype was incorrectly labeled as “TT: 0.150 (0.043–0.403)” and should be corrected to “CC: 0.150 (0.043–0.403)”.

The authors declare that these corrections do not change the results or conclusions of this paper. We sincerely apologize for having these errors in the article and apologize for any inconvenience caused.

The authors apologize for these errors. This correction has been approved by the Editor-in-Chief of the journal.

References

- [1] Jaramillo-Valverde L, Garcia-Paitan MY, Landeo D, Santivañez SJ, Cacacabelos R. Exploratory Study on Genetic Variants Related to Hydatidosis Susceptibility and Albendazole Pharmacogenetics in the Cusco Region in Peru. *Frontiers in Bioscience Scholar*. 2025; 17: 40566. <https://doi.org/10.31083/FBS40566>

