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Rahma Mkaouar, Zied Riahi, Jihene Marrakchi, Nessrine Mezzi, Lilia Romdhane, et al.. Corrigendum: Current phenotypic and genetic spectrum of syndromic deafness in Tunisia: paving the way for precision auditory health. *Frontiers in Genetics*, 2024, 15, pp.1437233. 10.3389/fgene.2024.1437233 . pasteur-04691609

HAL Id: pasteur-04691609

<https://pasteur.hal.science/pasteur-04691609v1>

Submitted on 9 Sep 2024

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EDITED AND REVIEWED BY
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RECEIVED 23 May 2024
ACCEPTED 27 May 2024
PUBLISHED 25 June 2024

CITATION
Mkaouar R, Riahi Z, Marrakchi J, Mezzi N, Romdhane L, Boujemaa M, Dallali H, Sayeb M, Lahbib S, Jaouadi H, Boudabbous H, Zekri L, Chargui M, Messaoud O, Elyounsi M, Kraoua I, Zaouak A, Turki I, Mokni M, Boucher S, Petit C, Giraudet F, Mbarek C, Besbes G, Halayem S, Zainine R, Turki H, Tounsi A, Bonnet CRYSTEL, Mrad R, Abdelhak S, Trabelsi M and Charfeddine C (2024), Corrigendum: Current phenotypic and genetic spectrum of syndromic deafness in Tunisia: paving the way for precision auditory health.
Front. Genet. 15:1437233.
doi: 10.3389/fgene.2024.1437233

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Corrigendum: Current phenotypic and genetic spectrum of syndromic deafness in Tunisia: paving the way for precision auditory health

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KEYWORDS

syndromic deafness (SD), spectrum of SDs, next-generation sequencing (NGS), early detection, under-diagnosis, Tunisia

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In the published article, there was an error in the legend for Figure 4 as published. The gene names corresponding to color degradation for the Waardenburg syndrome pie chart are incorrect.

The corrected legend appears below: The SOX10 gene should be at the top (darkest blue), and the EDNRB gene should be at the bottom (lightest blue). Consequently, the number of patients should be 8 for the SOX10 gene and 2 for the EDNRB gene.

In addition, there was an error in the caption of Figure 4 in the published version of the manuscript: “FIGURE 4: Genetic and mutational spectrum of syndromic deafness in the Tunisian population. Deafness syndromes with known molecular etiologies are represented by pie charts. Each syndromic deafness is indicated with a different color. Color degradation has been adopted to specify

the gene(s) associated with each syndrome. The number of variants identified in each gene is provided.”

The corrected Figure 4 appears below:

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

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