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Correction: Identification of a novel *ANK1* gene variant c.1504-9G>A and its mechanism of intron retention in hereditary spherocytosis

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KEYWORDS

ANK1, hereditary spherocytosis, intron retention, whole-exome sequencing, minigene splicing assay

A Correction on
Identification of a novel ANK1 gene variant c.1504-9G>A and its mechanism of intron retention in hereditary spherocytosis

by Xiong T, Xu Z, Wan Q, Chen F, Ye Y, Wang H and Wu C (2024). Front. Genet. 15:1390924. doi: 10.3389/fgene.2024.1390924

In the originally published article, the *ANK1* gene variant was incorrectly reported using transcript NM_020475.2 in the **Section of 2.2.3** Pathogenic variant confirmation. The correct reference transcript should be NM_001142446.

A correction has been made to the section Materials and methods, **subsection 2.2.3** Pathogenic variant confirmation:

"The reference for the *ANK1* DNA sequence was NCBI transcript version NM_001142446".

The original article has been updated.

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