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CORRECTION

Correction: Exome sequencing reveals *IFT172* variants in patients with non-syndromic cholestatic liver disease

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There are errors in the S3 Table. In the S3B Table, the predicted translation of the heterozygous ABCB11 variant c.3589C>G found in the patient F39CE53 should shave been p.(Leu1197Val). In S3C Table, the variant found in the patient M11RO526 should be c.1798A>T and in patient F4RO528 should be c. 2545T>C. Please view the correct S3 Table below.

Supporting information

S3 Table. Variants detected by initial Sanger sequencing in 60 excluded patients.
(DOCX)

Reference

1. Neřoldová M, Ciara E, Slatinská J, Fraňková S, Lišková P, Kotalová R, et al. (2023) Exome sequencing reveals *IFT172* variants in patients with non-syndromic cholestatic liver disease. PLOS ONE 18(7): e0288907. https://doi.org/10.1371/journal.pone.0288907

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