



CORRECTION



Correction: Identification of a novel Wilson disease gene mutation frequent in Upper Austria: a genetic and clinical study

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The original version of this article unfortunately contained a mistake. In the article, we described a “new” ATP7B variant (p.R816S) as a common mutation in Upper Austria,

mistakenly considering the last 3 nucleotides of exon 9 to code for Arginine. In fact, the third nucleotide is the first of the adjacent intron. Reanalysis of the sample by NGS showed that this variant was in fact 2447+1G>T. This variant is listed as likely pathogenic in ClinVar (<https://www.ncbi.nlm.nih.gov/clinvar/>).