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CORRIGENDA

In vitro antisense therapeutics for a deep intronic mutation causing Neurofibromatosis type 2

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Post publication the authors realized that they had made some errors in their article for which they would like to apologise.
g.74409 T>A the mutation at the genomic level should be termed: g.74408 T>A, and the sequence of the scheme depicted in Figure 1b, contains a couple of single nucleotide changes. A revised figure is shown below.

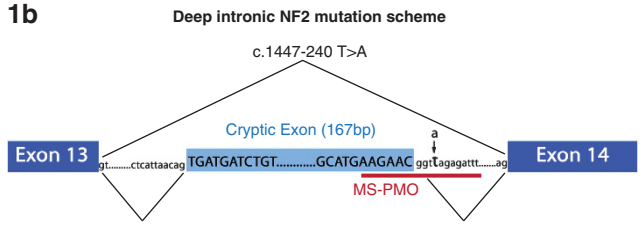

Figure 1 Deep intronic *NF2* mutation description. (**a**) Analysis of patient fibroblasts showed a proportion of *NF2* transcripts containing the inclusion of a cryptic exon (NF2 CEI) compared with the WT NF2 mRNA (NF2 WT) (upper panel). Forward sequence of cryptic exon inclusion is shown (bottom panel). (**b**) Schematic representation of the identified *NF2* deep intronic mutation and MS-PMO location. Constitutive and cryptic exons are represented by dark and light gray boxes, respectively. The boundaries of the cryptic inserted exon are shown in uppercase; flanking intronic sequences are shown in lowercase. MS-PMO sequence is underlined. Mutated nucleotide is shown in bold and the nucleotide change is indicated by an arrow.

Return of whole-genome sequencing results in paediatric research: a statement of the P^{3}G international paediatrics platform

Bartha Maria Knoppers, Denise Avard, Karine Sénécal and Ma'n H Zawati along with the P^{3}G International Paediatrics Platform members

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