BMC Medical Genomics

CORRECTION

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Correction to: Compound heterozygous variants including a novel copy number variation in a child with atypical ataxia-telangiectasia: a case report

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Following publication of the original article [1], the authors reported an error in Fig. 2.

Figure 2 was replaced as the authors identified an error in part C that was introduced during the peer review process. This error may have affected the interpretation of the results, and therefore, the figure was replaced with the earlier correct version. The original article [1] has been corrected.

(See figure on next page.)

Figure 2 **a** Chromatograms of *ATM* sequence in the proband (top), the patient's father (middle), and the patient's mother (bottom) showing an SNV of c.742C > T (p.Arg248Ter) from the father. **b** Analysis of the next-generation sequencing data using VisCap. **c** SNP array analysis of the chromosome from the proband (top), the patient's mother (middle), and the patient's father (bottom) showing a novel CNV by the deletion of exons 24–40 from the mother. *SNV* single-nucleotide variation, *SNP* single-nucleotide polymorphism, *CNV* copy number variation

The original article can be found online at https://doi.org/10.1186/s12920-021-01053-3.

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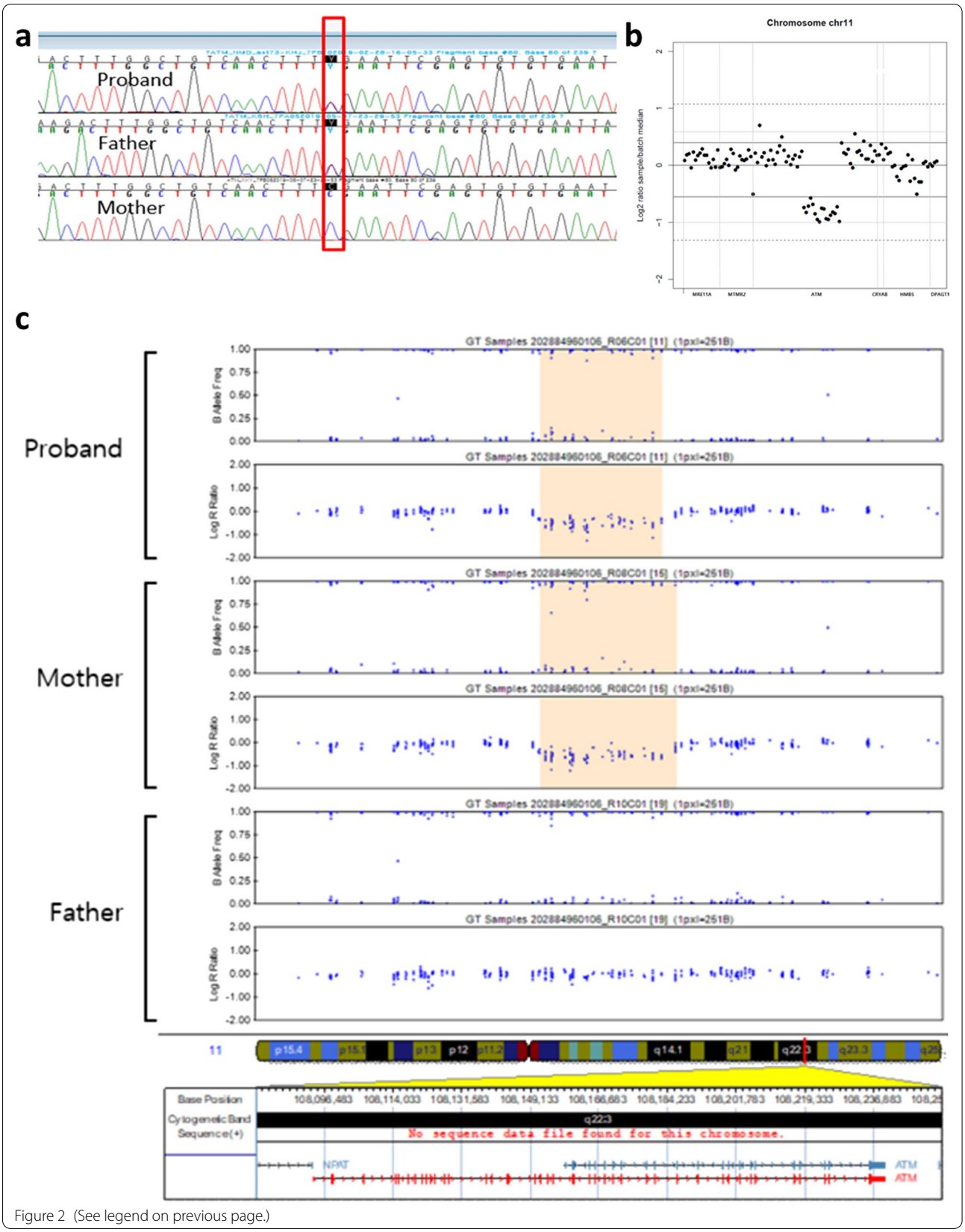

Figure 2 (See legend on previous page.)

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