Corrigendum

Biallelic inactivation of \textit{REV7} is associated with Fanconi anemia

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The nomenclature for the \textit{REV7} mutation was incorrectly noted in two sentences in the second paragraph of Results and Discussion and in Figure 1H. The correct sentences and figure part are below.

Whole exome sequencing (WES) on genomic DNA from the EGF123 proband identified a homozygous \textit{REV7} variant, c.254T>A.

The c.254T>A \textit{REV7} is a variant based on a survey of publicly accessible variant databases.

In addition, the fourth sentence of the Abstract was incorrect. The correct sentence is below.

Patient-derived cells demonstrated an extended FA phenotype, which included increased chromosome breaks and G\textsubscript{2}/M accumulation upon exposure to DNA crosslinking agents, \(\gamma\text{H2AX}\) and 53BP1 foci accumulation, and enhanced p53/p21 activation relative to cells derived from healthy subjects.

The authors regret the errors.