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Publisher Correction: The chloroplast protein HCF164 is predicted to be associated with *Coffea* S_H9 resistance factor against *Hemileia vastatrix*

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The original version of this Article contained an error in Figure 3, panels B and C, where the results were incorrectly displayed.

The original Article has been corrected.

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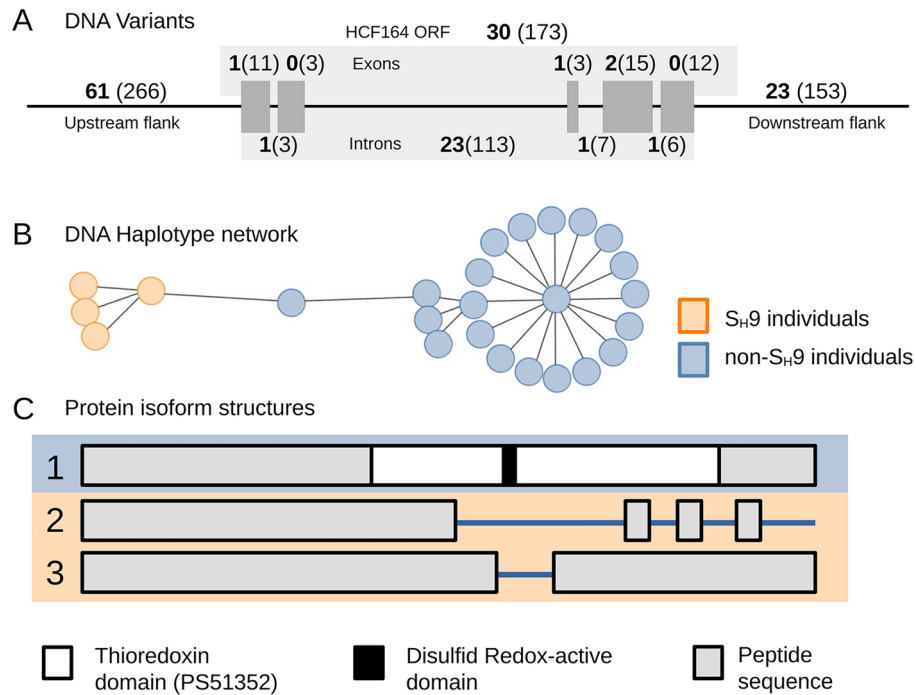



Figure 3. Analysis of the HCF164 sequence in chromosome 7c (*C. canephora*-derived sug-genome): **(A)** Depiction of the variants identified in the genomic region encoding the HCF164 protein in chromosome 7c following the reference genome annotation (GCA_003713225). The diagram shows the ORF (composed of five exons represented as grey rectangles) and 2 kbp upstream and downstream flanking regions. The numbers in brackets represent the number of variants identified in the 25 studied individuals, whereas the numbers in bold represent variants potentially associated with the SH9 factor (that is, variants exclusively found in SH9 individuals). **(B)** Haplotype network of the genomic region encoding the HCF164 protein (including 2 kbp flanking regions) obtained for the 25 studied individuals. **(C)** Schematic view of the alignment of the three HCF164 protein isoforms predicted in the 25 studied individuals. The thioredoxin domain and the redox-active disulphide center are highlighted in white and black, respectively. Orange represents SH9 individuals and blue represents non-S_{H9} individuals.

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