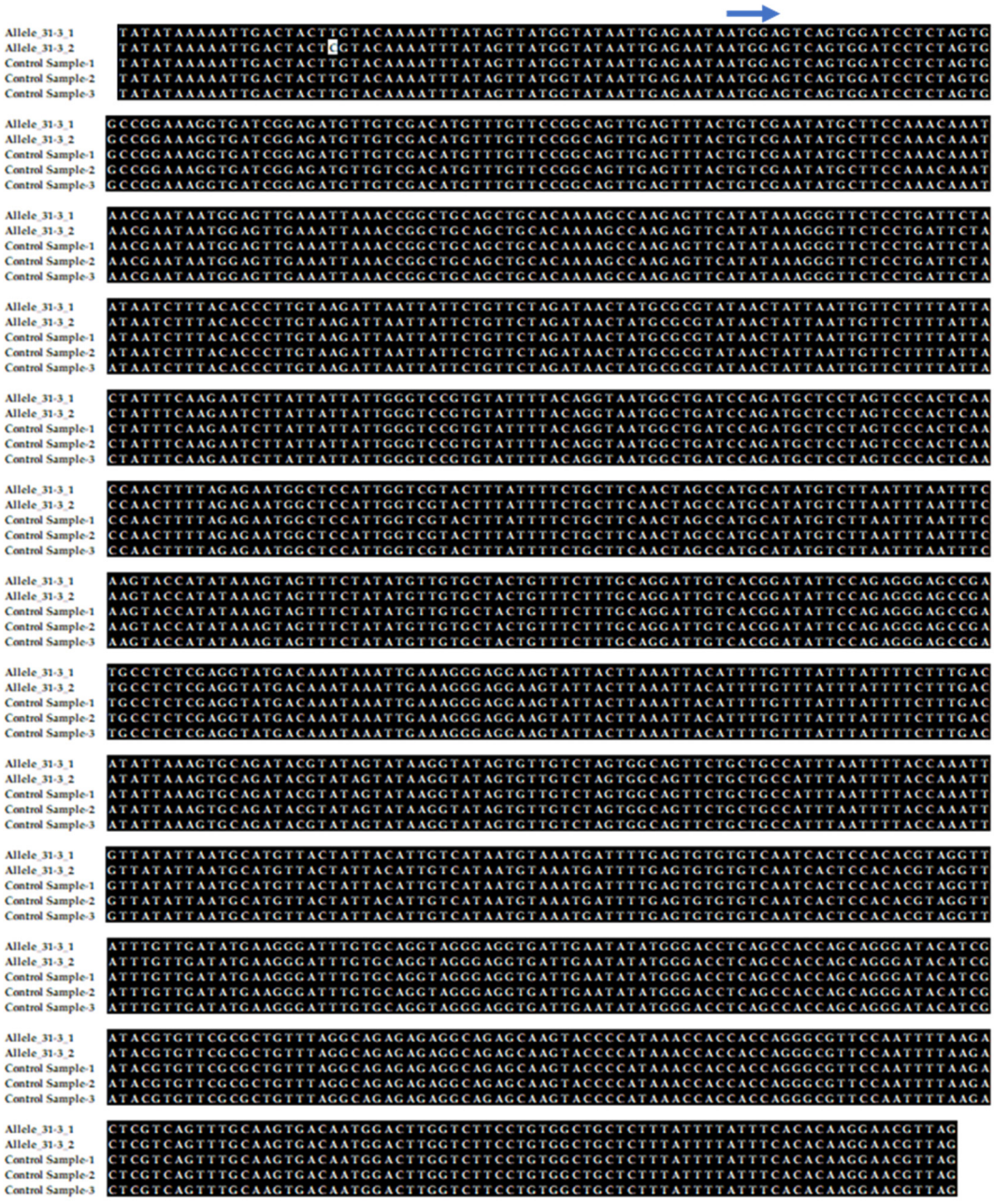


**Figure S2**. Mutation identified in the upstream region of *SmMFT*-2 gene of mutant sample, 31-3. The mutated base is displayed in white background. Black background indicates regions with identical bases. The blue arrow indicates the region of start codon (ATG).