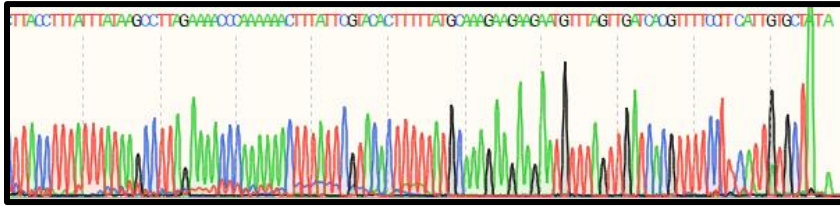
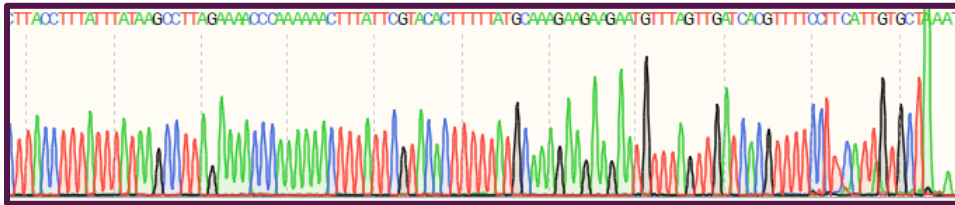


(A) Merlot WT 294 bp fragment



(B) Merlot WB 294 bp fragment



(C) Merlot WB 3901 bp fragment

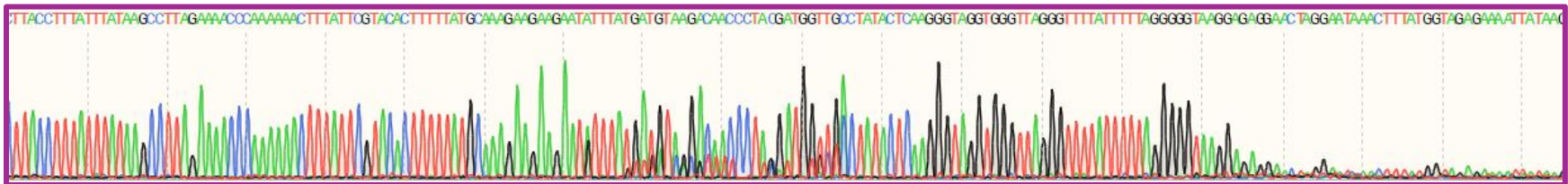


Figure S8. Sanger sequencing data of purified fragments from the region around the insertion within GSVIVG01008260001 for the 294 bp fragments amplified in (A) Merlot WT, (B) Merlot WB, as well as the (C) 3901 bp fragment in Merlot WB. Sanger sequencing data generated using the reverse VvSCD1 primer are shown. The sequences shown all start at identical locations within WT sequence and end at the end of the sequence generated through Sanger sequencing. The purple arrow shows approximately where the insertion sequence begins in the Merlot WB 3901 bp fragment.