Supplemental table1. Primer pairs used for amplification and Sanger sequencing of detected region.

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| Primer | Sequence | Expected length |
| F0 | 5' CTTCGTGGACATCTTCGTATTG 3' | 1593 |
| R0 | 5' TGGTAGCTCTTCGGTAGTAGCC 3' |
| F1 | 5' CTTCGCAGCGTGTAGCAGGTGA 3' | 1369 |
| R1 | 5' AGCGGTGAACCAAGACGCAGTA 3' |