

# Transmission of foot-and-mouth disease SAT2 viruses at the wildlife-livestock interface of two major transfrontier conservation areas

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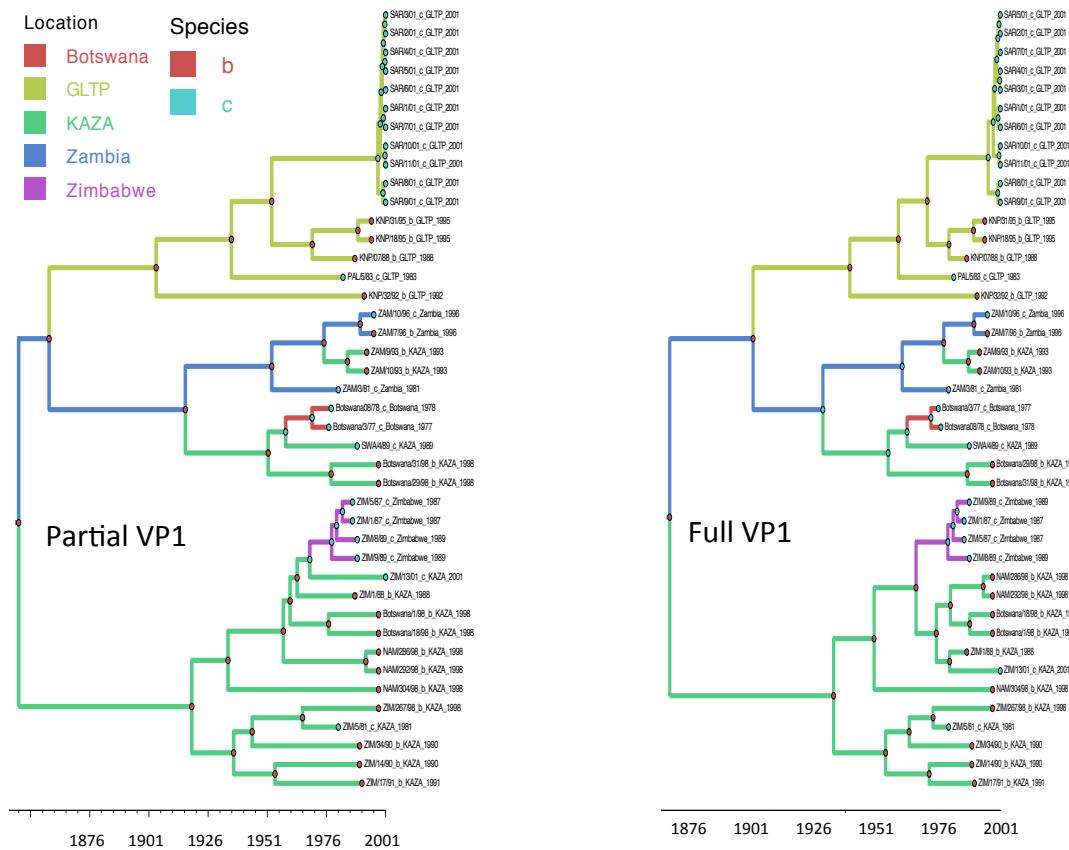


Image S1. The left panel depicts the MCC tree of FMDV SAT2 phylogeny reconstructed using a subset of virus with available full VP1 sequencing. MCC tree on the left panel shows the MCC tree reconstructed using only partial VP1 sequences of the same viruses.