

Supporting information

Molecular Detection of Reticuloendotheliosis virus 5' Long Terminal Repeat Integration in the Genome of Avipoxvirus Field Strains from Different Avian Species in Egypt

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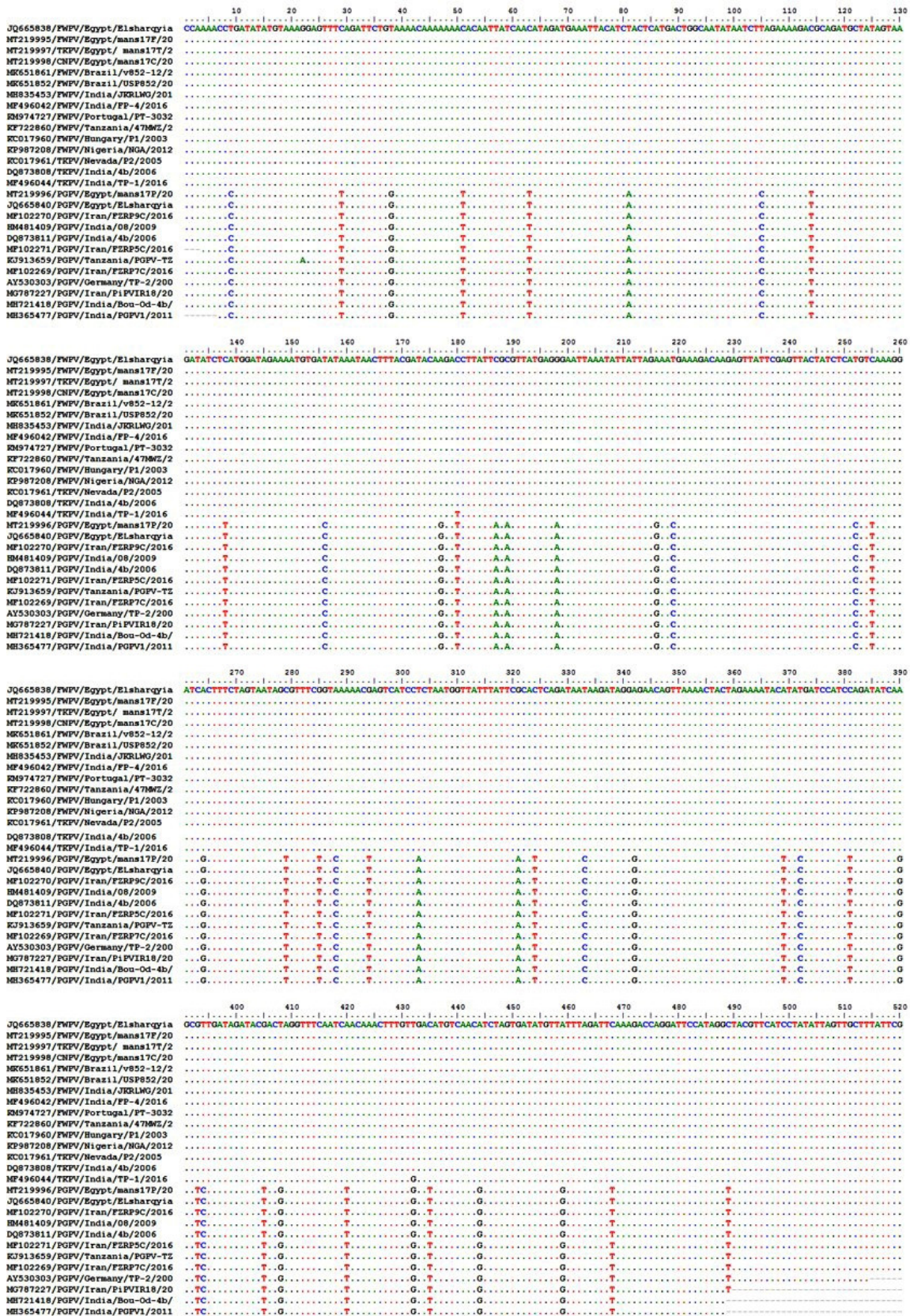
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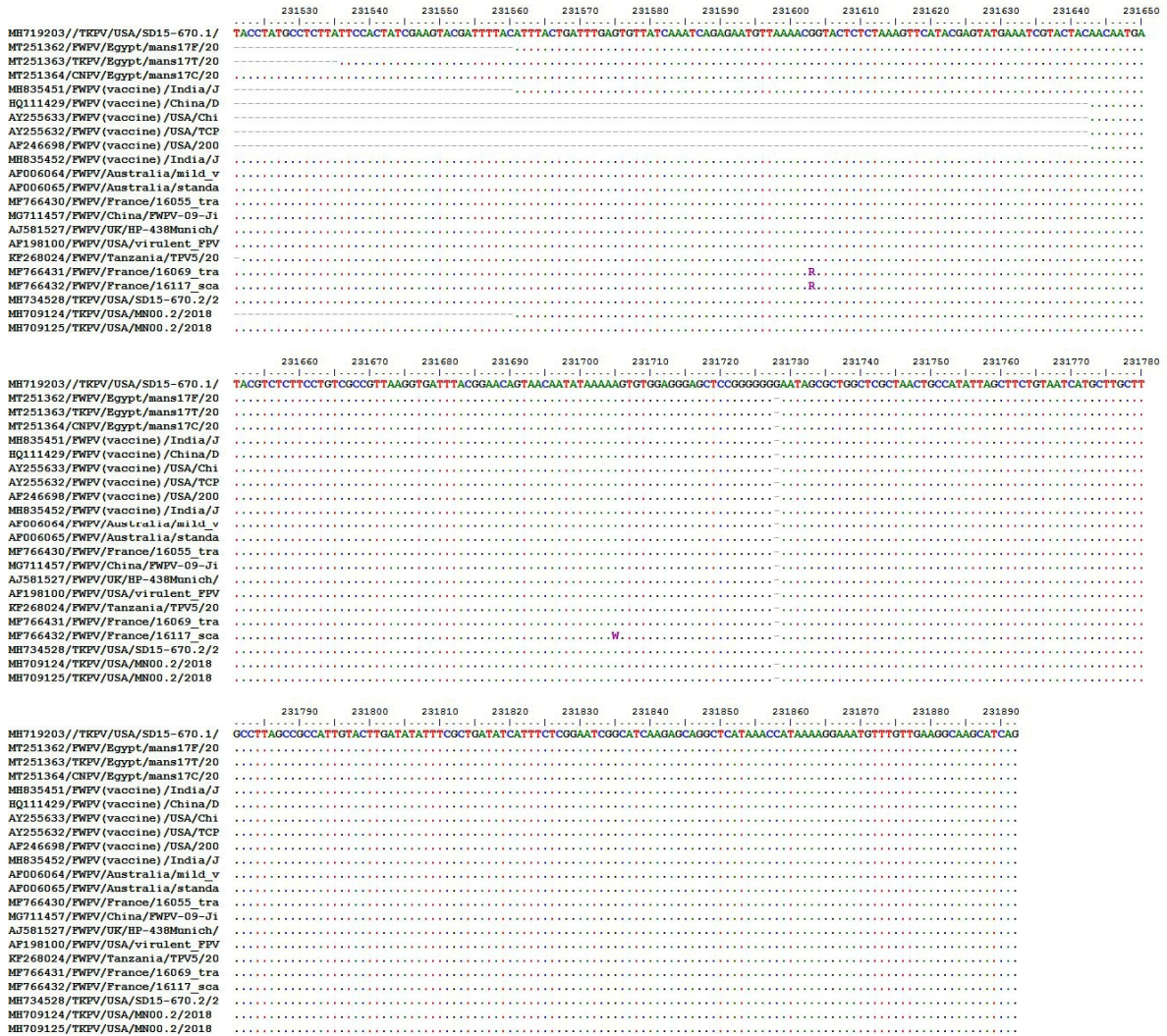
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Supplementary Figure S1. The BioEdit multiple alignments of amplified *P4b* gene nucleotide sequences of our APVs isolates compared to other reference APVs sequences retrieved from GenBank. PGPV strains have 44 nucleotide substitutions in comparison with FWPV and TKPV strains.



Supplementary Figure S2. The BioEdit multiple alignments of amplified REV-5'LTR nucleotide sequences detected in our APVs isolates in comparison to other reference REV-5'LTR sequences available in GenBank. All aligned sequences are identical to each other without any nucleotide substitution.