

Detection of Recombinant Hare Myxoma Virus in Wild Rabbits (*Oryctolagus cuniculus algirus*)

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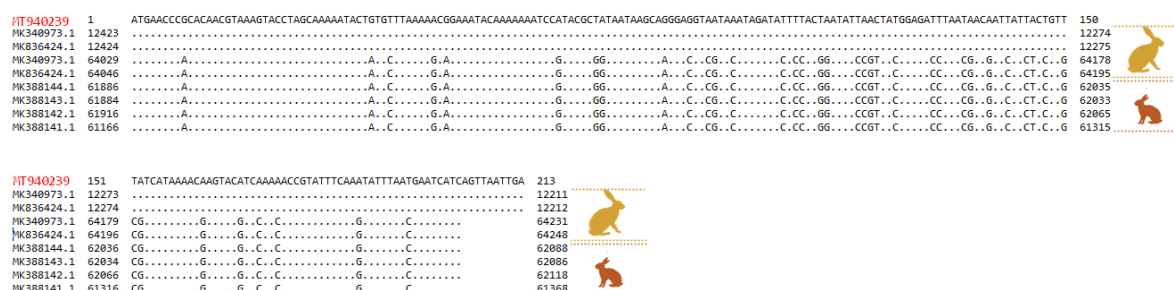


Figure S1. Nucleotide alignment between the M066L from sequence MT940239 with the homologous sequences of M066L from other hare recombinant strains (MK836424 and MK340973) (first to third sequences) and with the M066R from hare and classic Myxoma virus (MYXV) strains from rabbit (fourth to ninth sequences). The hare yellow icon delimits the sequences from strains isolated from hares, and the rabbit orange icon delimits the rabbit strains. The nucleotide position relative to the query (MT940239) indicates the insertion location (12423nt or 12424nt), and the location of M066R sequence (61184nt to 64046nt) in the complete genome.

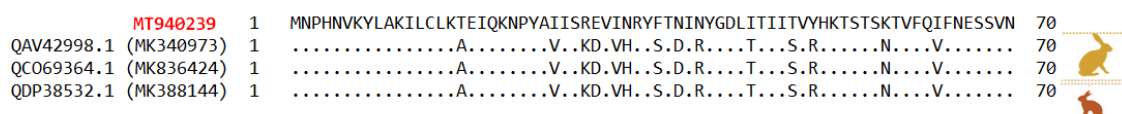


Figure S2. Alignment of the truncated M066L protein sequence from a natural recombinant MYXV (deduced from sequence MT940239) with the homologous regions of M066R proteins deduced from hare recombinant MYXV strains (MK836424 and MK340973) and classic MYXV strain (MK388144).