

Supplementary figure S6. Identification of the inverted CAAT box (shown with green), an upstream promoter element (blue), the canonical TATA box (orange), an initiator element (yellow), and the 2 downstream activating elements (grey and pink) in the putative major late promoter region of porcine adenovirus serotype 5 (species *Porcine mastadenovirus C*) strains HNF-70, GES7 and Z11. The virus serotype (isolate)/GenBank accession number are shown for PAdV-5 reference strain HNF-70, whilst the virus name/host/country/year have been mentioned for strains GES7 and Z11. A '*' denotes an identical nucleotide (nt) residue. The nt positions correspond to those of the complete genome sequence of strain HNF-70.

PAdV-5_(HNF-70)/AF289262 GES7/Pig/DOM/2020 Z11/Pig/DOM/2021	<div style="display: flex; justify-content: space-between;"> 5055 5114 </div> CGTCATCGTCTCCACATCTAGAAATGTGATTGGTTTATATTGTATGTCACGTGATCAG CGTCATCGTCTCCACATCTAGAAATGTGATTGGTTTATATTGTATGTCACGTGATCAG CGTCATCGTCTCCACATCTAGAAATGTGATTGGTTTATATTGTATGTCACGTGATCAG *****
PAdV-5_(HNF-70)/AF289262 GES7/Pig/DOM/2020 Z11/Pig/DOM/2021	<div style="display: flex; justify-content: space-between;"> 5115 5174 </div> ATTTGCCATATAAAGGCGTGGCTTCTTCGGCGGCCTCATTGTCCTCGTTGGAGACGTCGG ATTTGCCATATAAAGGCGTGGCTTCTTCGGCGGCCTCATTGTCCTCGTTGGAGACGTCGG ATTTGCCATATAAAGGCGTGGCTTCTTCGGCGGCCTCATTGTCCTCGTTGGAGACGTCGG *****
PAdV-5_(HNF-70)/AF289262 GES7/Pig/DOM/2020 Z11/Pig/DOM/2021	<div style="display: flex; justify-content: space-between;"> 5175 5234 </div> CGACAGGTGGGTAGGCTACGACAAATCTGGCATGATGTCCGCGCTGAAATTGTCTGTCT CGACAGGTGGGTAGGCTACGACAAATCTGGCATGATGTCCGCGCTGAAATTGTCTGTCT CGACAGGTGGGTAGGCTACGACAAATCTGGCATGATGTCCGCGCTGAAATTGTCTGTCT *****
PAdV-5_(HNF-70)/AF289262 GES7/Pig/DOM/2020 Z11/Pig/DOM/2021	<div style="display: flex; justify-content: space-between;"> 5235 5294 </div> CTATGAAGGATGAAGATTTGACAGAGTACTTGCCCCGGAGATGTTCTTTGAGCTCTCTT CTATGAAGGATGAAGATTTGACAGAGTACTTGCCCCGGAGATGTTCTTTGAGCTCTCTT CTATGAAGGATGAAGATTTGACAGAGTACTTGCCCCGGAGATGTTCTTTGAGCTCTCTT *****