

Veterinary Research Communications

Canine circovirus and Canine adenovirus type 1 and 2 in dogs with parvoviral enteritis

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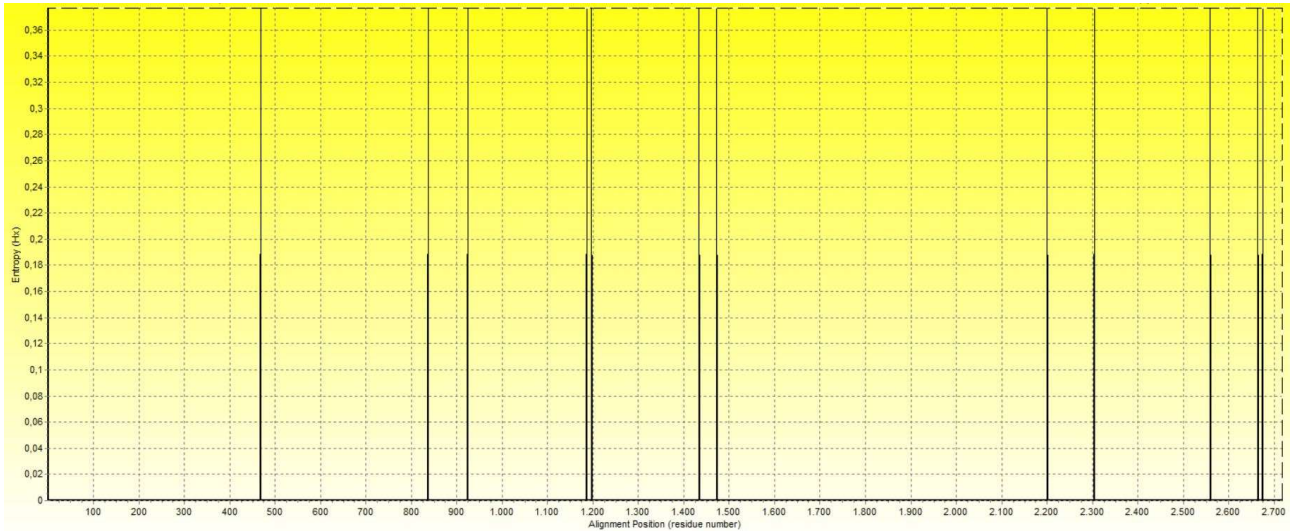
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Online Resource 5a Graphical representation of the hexon gene nucleotide sequence entropy obtained with the entropy ($H(x)$) plot function implemented in BioEdit. The plot shows the variability of the different nucleotide residues in the hexon gene alignment. The analysis was performed on the CAdV-2 nucleotide sequences generated in this study and on CAdV-2 reference strains which had available both hexon and fiber genes sequences (**Online Resource 2**)



Online Resource 5b Graphical representation of the fiber gene nucleotide sequence entropy obtained with the entropy ($H(x)$) plot function implemented in BioEdit. The plot shows the variability of the different nucleotide residues in the fiber gene alignment. The analysis was performed on the CA Δ V-2 nucleotide sequences generated in this study and on CA Δ V-2 reference strains which had available both hexon and fiber genes sequences (**Online Resource 2**)

