

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

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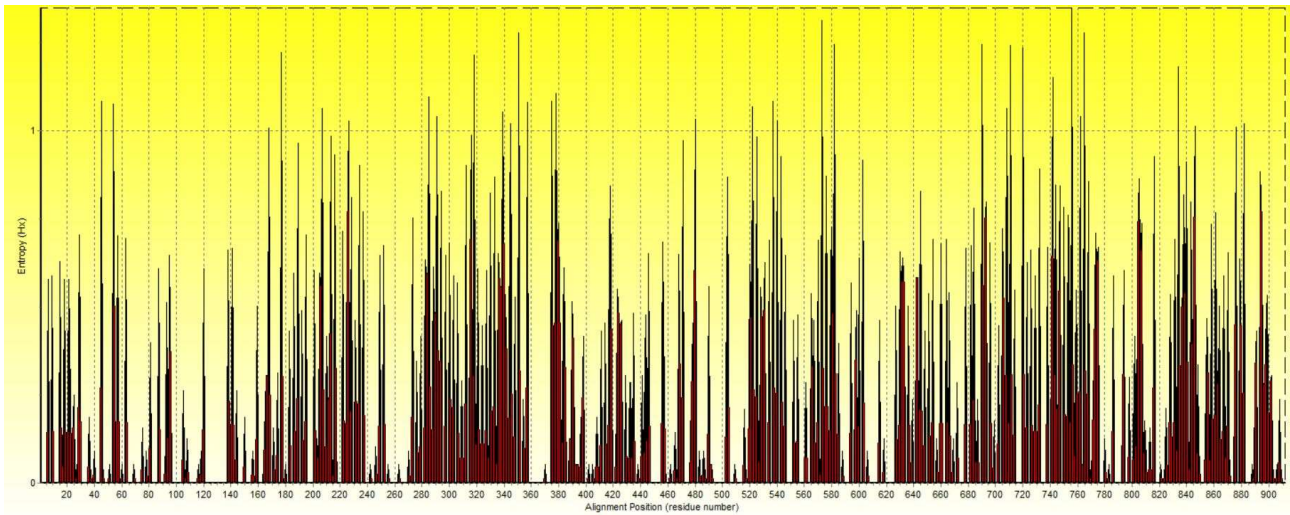
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Online Resource 3a Graphical representation of the replicase (Rep) gene nucleotide sequence entropy obtained with the entropy ($H(x)$) plot function implemented in BioEdit. The plot shows the variability of the different nucleotide residues in the Rep gene alignment. The analysis was performed on the CanineCV nucleotide sequences generated in this study and on CanineCV reference strains which had available both Rep and capsid (Cap) genes sequences



Online Resource 3b Graphical representation of the capsid (Cap) 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 Cap gene alignment. The analysis was performed on the CanineCV nucleotide sequences generated in this study and on CanineCV reference strains which had available both replicase (Rep) and Cap genes sequences

