

Genomes	NCBI Nucleotide Accession Number https://www.ncbi.nlm.nih.gov/nucleotide
<i>Human adenovirus type 2</i>	J01917.1
<i>Human adenovirus type 5</i>	AC_000008.1
<i>Human adenovirus type 12</i>	X73487
<i>Autographa californica Nuclear Polyhedrosis Virus</i>	NC_001623.1
<i>Hepatitis B Virus</i>	NC_003977.2
<i>HIV-1 HXB2</i>	K03455.1
<i>Escherichia virus lambda</i>	NC_001416.1
<i>Escherichia virus Mu</i>	NC_000929.1
<i>Escherichia virus T4</i>	NC_000866.4
<i>Acidianus rod-shaped virus 1</i>	NC_009965.1
<i>Nitrosopumilus spindle-shaped virus isolate NSV1</i>	NC_048199.1
<i>Sulfolobus turreted icosahedral virus</i>	NC_005892
<i>Alternaria brassicicola endornavirus isolate 1</i>	NC_026136.1
<i>Carrot mottle mimic umbravirus (CMoMV) dsRNA1</i>	NC_001726.1
<i>Fig badnavirus 1</i>	NC_017830.1
<i>Candidatus Carsonella ruddii</i> strain BT chromosome	CP024798.1
<i>Escherichia coli</i> K-12	NC_000913.3
<i>Homo sapiens</i> mitochondrion	NC_012920.1
SARS-CoV-2	NC_045512.2
SARS-CoV-2 <i>Spike</i> region original: 21,563-25,384nt	NC_045512.2
SARS-CoV-2 <i>Spike</i> region delta: 21,561-25,376nt*	MZ437368.1

Table S1 - NCBI Nucleotide Accession Numbers of nucleotide sequences analyzed in this article.