

Covid-19 was uploaded to NCBI in 2018 by the Institute of Military Medicine Nanjing on January 5th, 2018 that 100% matches today's current Covid-19

This virus originated in a lab, created as a bioweapon by China. Here is your smoking gun that China has been developing viral weapons of mass destruction and is using this weapon to conduct biological warfare on humanity. Whether intentionally released or accidental, China is guilty of biological weapons research and all infected countries must hold China accountable for its crimes against humanity. Evidence of it matching 100% Covid-19 can be found when you run the blast and results are included in this PDF.

Go to this URL and look it up yourself on the Viral Genome database and then on the right hand side click Run Blast and you will find many 100% matches to the current 2019 Covid-19 virus.

<https://www.ncbi.nlm.nih.gov/protein/AVP78033.1>

envelope protein [Bat SARS-like coronavirus]

GenBank: AVP78033.1

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LOCUS	AVP78033	75 aa	linear	VRL 05-FEB-2020
DEFINITION	envelope protein [Bat SARS-like coronavirus].			
ACCESSION	AVP78033			
VERSION	AVP78033.1			
DBSOURCE	accession MG772933.1			
KEYWORDS	.			
SOURCE	Bat SARS-like coronavirus			
ORGANISM	Bat SARS-like coronavirus Viruses; Riboviria; Nidovirales; Coronidovirineae; Coronaviridae; Orthocoronavirinae; Betacoronavirus; Sarbecovirus.			
REFERENCE	1 (residues 1 to 75)			
AUTHORS	Hu,D., Zhu,C., Ai,L., He,T., Wang,Y., Ye,F., Yang,L., Ding,C., Zhu,X., Lv,R., Zhu,J., Hassan,B., Feng,Y., Tan,W. and Wang,C.			
TITLE	Genomic characterization and infectivity of a novel SARS-like coronavirus in Chinese bats			
JOURNAL	Emerg Microbes Infect 7 (1), 154 (2018)			
PUBMED	30209269			
REMARK	Publication Status: Online-Only			
REFERENCE	2 (residues 1 to 75)			
AUTHORS	Hu,D.			
TITLE	Direct Submission			
JOURNAL	Submitted (05-JAN-2018) Institute of Military Medicine Nanjing Command, Nanjing, Institute of Military Medicine Nanjing Command, Nanjing, NO. 293 East Zhongshan Road, Nanjing, JiangSu 210002, China			
FEATURES	Location/Qualifiers			

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Job Title	gb AVP78033.1
RID	6MPR43WP016 <i>Search expires on 03-13 23:53 pm</i> Download All ▾
Program	BLASTP ? Citation ▾
Database	nr See details ▾
Query ID	AVP78033.1
Description	envelope protein [Bat SARS-like coronavirus]
Molecule type	amino acid
Query Length	75
Other reports	Distance tree of results Multiple alignment MSA viewer ?

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	Description	Max Score	Total Score	Query Cover	E value	Per. Ident	Accession
<input checked="" type="checkbox"/>	envelope protein [Severe acute respiratory syndrome coronavirus 2]	144	144	100%	2e-43	100.00%	YP_009724392.1
<input checked="" type="checkbox"/>	envelope protein [Severe acute respiratory syndrome coronavirus 2]	142	142	100%	1e-42	98.67%	QHZ00381.1
<input checked="" type="checkbox"/>	E protein [Severe acute respiratory syndrome-related coronavirus]	137	137	100%	1e-40	94.74%	APO40581.1
<input checked="" type="checkbox"/>	envelope protein [Bat SARS-like coronavirus RsSHC014]	134	134	100%	1e-39	94.74%	AGZ48809.1
<input checked="" type="checkbox"/>	envelope protein, partial [Severe acute respiratory syndrome coronavirus 2]	134	134	93%	1e-39	100.00%	QID88859.1
<input checked="" type="checkbox"/>	small envelope protein [Coronavirus BtRI-BetaCoV/SC2018]	134	134	100%	2e-39	94.74%	QDF43816.1
<input checked="" type="checkbox"/>	protein E [Severe acute respiratory syndrome-related coronavirus]	134	134	100%	3e-39	94.74%	NP_828854.1
<input checked="" type="checkbox"/>	small membrane protein [Bat SARS coronavirus HKU3-7]	133	133	100%	5e-39	93.42%	ADE34757.1
<input checked="" type="checkbox"/>	small envelope protein [Bat SARS-like coronavirus]	133	133	100%	6e-39	93.42%	ATO98160.1
<input checked="" type="checkbox"/>	envelope protein E [SARS coronavirus GD01]	132	132	100%	7e-39	93.42%	AAP51230.1
<input checked="" type="checkbox"/>	small envelope protein [BtRs-BetaCoV/HuB2013]	132	132	100%	8e-39	93.42%	AIA62312.1
<input checked="" type="checkbox"/>	envelope protein E [SARS coronavirus GD01]	132	132	100%	9e-39	93.42%	AAP51230.1

GenPept ▾

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envelope protein [Bat SARS-like coronavirus]

GenBank: AVP78033.1

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LOCUS AVP78033 75 aa linear VRL 05-FEB-2020

DEFINITION envelope protein [Bat SARS-like coronavirus].

ACCESSION AVP78033

VERSION AVP78033.1

DBSOURCE accession [MG772933.1](#)

KEYWORDS .

SOURCE Bat SARS-like coronavirus

ORGANISM [Bat SARS-like coronavirus](#)

Viruses; Riboviria; Nidovirales; Coronaviridae; Orthocoronavirinae; Betacoronavirus; Sarbecovirus.

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AUTHORS Hu,D., Zhu,C., Ai,L., He,T., Wang,Y., Ye,F., Yang,L., Ding,C., Zhu,X., Lv,R., Zhu,J., Hassan,B., Feng,Y., Tan,W. and Wang,C.

TITLE Genomic characterization and infectivity of a novel SARS-like coronavirus in Chinese bats

JOURNAL Emerg Microbes Infect 7 (1), 154 (2018)

PUBMED [30209269](#)

REMARK Publication Status: Online-Only

REFERENCE 2 (residues 1 to 75)

AUTHORS Hu,D.

TITLE Direct Submission

JOURNAL Submitted (05-JAN-2018) Institute of Military Medicine Nanjing Command, Nanjing, Institute of Military Medicine Nanjing Command, Nanjing, NO. 293 East Zhongshan Road, Nanjing, JangSu 210002, China

FEATURES Location/Qualifiers

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 /name="E"
 CDS 1..75
 /coded_by="MG772933.1:26150..26377"

ORIGIN

1 mysfvseetg tlivnsvllf lafvvflvlt lailtalrlc ayccnivnvs lvkpsfyvys
 61 rvknlnssrv pdllv

//

Analyze this sequence

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RID

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Query ID

AVP78033.1

Description

envelope protein [Bat SARS-like coronavirus]

Molecule type

amino acid

Query Length

75

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Max Total Query E Per.

	Description	Score	Score	Cover	value	Ident	Accession
■	envelope protein [Severe acute respiratory syndrome coronavirus 2]	144	144	100%	2e-43	100.00%	YP_009724392.1
■	envelope protein [Severe acute respiratory syndrome coronavirus 2]	142	142	100%	1e-42	98.67%	QHZ00381.1
■	E protein [Severe acute respiratory syndrome-related coronavirus]	137	137	100%	1e-40	94.74%	APO40581.1
■	envelope protein [Bat SARS-like coronavirus RsSHC014]	134	134	100%	1e-39	94.74%	AGZ48809.1
■	envelope protein, partial [Severe acute respiratory syndrome coronavirus 2]	134	134	93%	1e-39	100.00%	QID88859.1
■	small envelope protein [Coronavirus BtRI-BetaCoV/SC2018]	134	134	100%	2e-39	94.74%	QDF43816.1
■	protein E [Severe acute respiratory syndrome-related coronavirus]	134	134	100%	3e-39	94.74%	NP_828854.1
■	small membrane protein [Bat SARS coronavirus HKU3-7]	133	133	100%	5e-39	93.42%	ADE34757.1
■	small envelope protein [Bat SARS-like coronavirus]	133	133	100%	6e-39	93.42%	ATO98160.1
■	envelope protein E [SARS coronavirus GD01]	132	132	100%	7e-39	93.42%	AAP51230.1
■	small envelope protein [BtRs-BetaCoV/HuB2013]	132	132	100%	8e-39	93.42%	AIA62312.1
■	envelope protein E [SARS coronavirus BJ182-12]	132	132	100%	9e-39	93.42%	ACB69908.1
■	envelope protein [SARS coronavirus ExoN1]	132	132	100%	1e-38	93.42%	ACZ72023.1
■	envelope protein [Bat coronavirus BM48-31/BGR/2008]	132	132	100%	1e-38	92.11%	YP_003858586.1
■	RecName: Full=Envelope small membrane protein; Short=E protein; Short=sM	132	132	100%	1e-38	93.42%	Q0Q473.1
■	envelope protein [SARS coronavirus HKU-39849]	132	132	100%	1e-38	93.42%	ADC35514.1
■	small membrane protein [Bat SARS coronavirus HKU3-8]	132	132	100%	1e-38	92.11%	ADE34768.1
■	envelope protein [Bat SARS-like coronavirus YNLF_34C]	132	132	100%	1e-38	93.42%	AKZ19089.1
■	envelope protein [Rhinolophus affinis coronavirus]	132	132	100%	1e-38	93.42%	AHX37560.1
■	envelope protein [SARS coronavirus ExoN1]	132	132	100%	2e-38	93.42%	ACZ72198.1
■	envelope protein [SARS coronavirus Sino3-11]	132	132	100%	2e-38	93.42%	AAR23255.1
■	envelope protein [Bat SARS CoV Rf1/2004]	131	131	100%	2e-38	93.42%	ABD75313.1
■	small envelope protein [BtRf-BetaCoV/JL2012]	131	131	100%	2e-38	92.11%	AIA62280.1
■	envelope protein [Bat coronavirus Rp/Shaanxi2011]	131	131	100%	2e-38	93.42%	AGC74167.1
■	envelope protein [SARS coronavirus MA15 ExoN1]	131	131	100%	2e-38	93.42%	ACZ72243.1
■	small envelope protein [BtRf-BetaCoV/SX2013]	131	131	100%	3e-38	92.11%	AIA62302.1
■	small envelope protein [SARS coronavirus TJ01]	130	130	100%	5e-38	92.11%	ABM92862.1
■	envelope protein [SARS coronavirus ExoN1]	130	130	100%	7e-38	93.42%	ACZ72257.1
■	small envelope protein [Bat SARS-like coronavirus]	130	130	100%	9e-38	93.42%	ATO98184.1
■	small envelope E protein [SARS coronavirus TW-GD1]	128	128	100%	5e-37	93.42%	AAS44718.1
■	envelope protein [Bat coronavirus]	128	128	100%	6e-37	90.79%	ANA96030.1
■	envelope protein [SARS coronavirus Sino1-11]	124	124	100%	1e-35	92.11%	AAR23247.1
■	small envelope protein [SARS coronavirus BJ01]	124	124	100%	2e-35	89.47%	AAS48454.1
■	envelope protein [SARS coronavirus ExoN1]	108	108	98%	4e-29	93.33%	ACZ72168.1
■	Chain A, Envelope small membrane protein [Severe acute respiratory syndrom	103	103	82%	3e-27	88.71%	5X29_A
■	Chain A, Envelope small membrane protein [Severe acute respiratory syndrom	100	100	77%	3e-26	91.38%	2MM4_A
■	putative envelope protein [Zaria bat coronavirus]	86.7	86.7	100%	1e-20	57.14%	ADY17913.1
■	envelope protein [Bat Hp-betacoronavirus/Zhejiang2013]	84.0	84.0	94%	2e-19	54.93%	YP_009072442.1
■	RecName: Full=Envelope small membrane protein; Short=E protein; Short=sM	56.6	56.6	94%	1e-08	42.67%	Q0Q4E8.1

E protein [Tylonycteris bat coronavirus HKU4]	56.2	56.2	94%	2e-08	42.67%	AWH65882.1
small membrane protein [Tylonycteris bat coronavirus HKU4]	56.2	56.2	94%	2e-08	42.67%	YP_001039958.1
envelope protein [Bat coronavirus]	56.2	56.2	85%	2e-08	46.88%	ANA96044.1
E protein [Tylonycteris bat coronavirus HKU4]	54.7	54.7	94%	7e-08	42.67%	AWH65904.1
E [Betacoronavirus Erinaceus/VMC/DEU/2012] [Betacoronavirus Erinaceus/VM]	53.9	53.9	97%	1e-07	44.87%	YP_009513016.1
envelope protein [Hypsugo bat coronavirus HKU25]	51.2	51.2	94%	1e-06	40.00%	ASL68947.1
E [Hedgehog coronavirus 1] [Hedgehog coronavirus 1]	51.2	51.2	97%	2e-06	43.59%	QCC20719.1
envelope protein [Hypsugo bat coronavirus HKU25]	50.8	50.8	78%	2e-06	47.46%	ASL68958.1
small envelope protein [BtVs-BetaCoV/SC2013]	50.8	50.8	94%	2e-06	38.67%	AHY61342.1
small envelope protein [Bat coronavirus]	50.4	50.4	94%	3e-06	38.67%	BBJ36013.1
small envelope protein [Bat coronavirus]	50.4	50.4	94%	3e-06	38.67%	YP_009361862.1
envelope protein [Erinaceus hedgehog coronavirus HKU31]	50.1	50.1	80%	5e-06	45.00%	QGA70697.1
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envelope protein [Middle East respiratory syndrome-related coronavirus]	49.3	49.3	94%	8e-06	37.33%	AUM60019.1
envelope protein [Middle East respiratory syndrome-related coronavirus]	48.5	48.5	94%	2e-05	37.33%	ATQ39391.1
E protein [Coronavirus Neoromicia/PML-PHE1/RSA/2011]	47.4	47.4	94%	5e-05	36.00%	AIG13101.1
E protein [Middle East respiratory syndrome-related coronavirus]	46.6	46.6	94%	1e-04	36.00%	AVV62531.1
envelope protein [Middle East respiratory syndrome-related coronavirus]	45.8	45.8	94%	3e-04	36.00%	ALA49390.1
E protein [Middle East respiratory syndrome-related coronavirus]	45.1	45.1	94%	4e-04	36.00%	AGV08472.1
E protein [Middle East respiratory syndrome-related coronavirus]	45.1	45.1	94%	5e-04	36.00%	AHZ65623.1
hypothetical protein [Escherichia coli]	45.1	45.1	94%	5e-04	36.00%	WP_071995508.1
envelope protein [Middle East respiratory syndrome-related coronavirus]	45.1	45.1	94%	5e-04	36.00%	ASU90334.1
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envelope protein [Ferret enteric coronavirus]	44.3	44.3	90%	8e-04	31.88%	ASR18940.1
envelope protein [Middle East respiratory syndrome-related coronavirus]	44.3	44.3	94%	9e-04	36.00%	ASU90554.1
E protein [Middle East respiratory syndrome-related coronavirus]	43.1	43.1	94%	0.003	34.67%	ATG84684.1
E protein [Middle East respiratory syndrome-related coronavirus]	42.7	42.7	94%	0.003	34.67%	AGV08540.1
small membrane protein [Pipistrellus bat coronavirus HKU5]	42.7	42.7	80%	0.004	40.00%	YP_001039967.1
envelope protein [Pipistrellus abramus bat coronavirus HKU5-related]	42.7	42.7	80%	0.005	40.00%	QHA24692.1
envelope glycoprotein [Ferret enteric coronavirus]	41.2	41.2	86%	0.015	33.33%	ADD49360.1
envelope protein [Ferret systemic coronavirus]	39.3	39.3	86%	0.075	33.33%	ASR18948.1
envelope protein [Ferret coronavirus]	38.9	38.9	86%	0.100	31.82%	YP_009256199.1
envelope protein [Ferret coronavirus]	37.7	37.7	86%	0.35	31.82%	BAV31351.1
envelope protein [Ferret coronavirus]	37.0	37.0	90%	0.56	31.88%	BAS25712.1
envelope protein [Mink coronavirus strain WD1127]	37.0	37.0	86%	0.58	28.79%	YP_009019184.1
envelope protein [Alphacoronavirus Mink/China/1/2016]	36.6	36.6	86%	0.81	28.79%	AVY53337.1

 hypothetical protein EGR_05687 [Echinococcus granulosus]	36.6	36.6	73%	2.1	34.55%	XP_024350632.1
 envelope protein [Mink coronavirus strain WD1133]	35.4	35.4	86%	2.6	28.79%	ADI80524.1

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