

Your search is limited to records that include: Porcine epidemic diarrhea virus (taxid:28295)

Job Title	R2 ...
RID	BR89R9A1016 Search expires on 06-30 08:47 am
Program	BLASTN
Database	ref_viruses_rep_genomes
Query ID	lcl Query_4363
Description	R2 ...
Molecule type	dna
Query Length	31

Descriptions

Descriptions

Description	Scientific Name	Max Score	Total Score	Query Cover	E value	Per. Ident	Acc. Len	Accession
Porcine epidemic diarrhea virus, complete genome	Porcine epidemic diarrhea virus	58.4	58.4	100%	1e-12	100.00%	28033	NC_003436.1

Graphic Summary

Distribution of the top 1 Blast Hits on 1 subject sequences

Query					
1	6	12	18	24	30

Alignments

Alignment view

Pairwise

☐ CDS feature

Restore defaults

Porcine epidemic diarrhea virus, complete genome
Sequence ID: **NC_003436.1** Length: 28033 Number of Matches: 1
Range 1: 26188 to 26218

Score	Expect	Identities	Gaps	Strand	Frame
58.4 bits(31)	1e-12()	31/31(100%)	0/31(0%)	Plus/Minus	
Query 1	GACAATTGTTGTAGTGGCCTTGGCGAC				
Sbjct 26218	GACAATTGTTGTAGTGGCCTTGGCGA				

Taxonomy

Reports

Lineage

Organism	Blast Name	Score	Number of Hits	Description
Porcine epidemic diarrhea virus	viruses	58.4	1	Porcine epidemic diarrhea virus hits

Organism

Description	Score	E value	Accession
Porcine epidemic diarrhea virus [viruses]			
Porcine epidemic diarrhea virus, complete genome	58.4	1e-12	NC_003436

Taxonomy

Taxonomy	Number of hits	Number of Organisms	Description
Porcine epidemic diarrhea virus	1	1	Porcine epidemic diarrhea virus hits