

>*Vibrio vulnificus* isolate 9 16S ribosomal RNA gene, partial sequence

Sequence ID: AY245189.1 Length: 1250

Range 1: 385 to 1083, Score:1242 bits (672), Expect:0.0,

Identities:691/700(99%), Gaps:2/700(0%), Strand: Plus/Plus

Query 2	AGCACCGGCTAACTCCGTGCCAGCAGCCGCGTAATACGGAGGGTGCGAGCGTTAATC-G	60
Sbjct 385	AGCACCGGCTAACTCCGTGCCAGCAGCCGCGTAATACGGAGGGTGCGAGCGTTAATCGG	444
Query 61	GAATTACTGGGCGTAAAGCGCATGCAGGTGGTTTGTTAAGTCAGATGTGAAAGCCCCGGGG	120
Sbjct 445	GAATTACTGGGCGTAAAGCGCATGCAGGTGGTTTGTTAAGTCAGATGTGAAAGCCCCGGGG	504
Query 121	CTCAACCTCGGAATTGCATTTGAAACTGGCAGACTAGAGTACTGTAGAGGGGGGTAGAAT	180
Sbjct 505	CTCAACCTCGGAATAGCATTTGAAACTGGCAGACTAGAGTACTGTAGAGGGGGGTAGAAT	564
Query 181	TTCAGGTGTAGCGGTGAAATGCGTAGAGATCTGAAGGAATACCGGTGGCGAAGGCGGCCCC	240
Sbjct 565	TTCAGGTGTAGCGGTGAAATGCGTAGAGATCTGAAGGAATACCGGTGGCGAAGGCGGCCCC	624
Query 241	CCTGGACAGATACTGACACTCAGATGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCC	300
Sbjct 625	CCTGGACAGATACTGACACTCAGATGCGAAAGCGTGGGGAGCAAACAGGATTAGATACCC	684
Query 301	TGGTAGTCCACGCCGTAAACGATGTCTACTTGGAGGTTGTGGCCTTGAGCCGTGGCTTTC	360
Sbjct 685	TGGTAGTCCACGCCGTAAACGATGTCTACTTGGAGGTTGTGGCCTTGAGCCGTGGCTTTC	744
Query 361	GGAGCTAACGCGTTAAGTAGACCGCCTGGGGAGTACGGTCGCAAGATTAAAACTCAAATG	420
Sbjct 745	GGAGCTAACGCGTTAAGTAGACCGCCTGGGGAGTACGGTCGCAAGATTAAAACTCAAATG	804
Query 421	AATTGACGGGGGGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACGCGAAGA	480
Sbjct 805	AATTGACGGGGGGCCCGCACAAGCGGTGGAGCATGTGGTTTAATTCGATGCAACGCGAAGA	864
Query 481	ACCTTACCTACTCTTGACATCCAGAGAACTTCCAGAGATGGATTGGTGCCTTCGGGAAC	540
Sbjct 865	ACCTTACCTACTCTTGACATCCAGAGAACTTCCAGAGATGGATTGGTGCCTTCGGGAAC	924
Query 541	TCTGAGACAGGTGCTGCATGGCTGTCGTCAGCTCGTGTTGTGAAATGTTGGGTAAAGTCC	600
Sbjct 925	TCTGAGACAGGTGCTGCATGGCTGTCGTCAGCTCGTGTTGTGAAATGTTGGGTAAAGTCC	984
Query 601	CGCAACGAGCGCAACCCTTATCCTTGTTTGCCAGCGAGTAATGTCGGGAACCTCAGGGAG	660
Sbjct 985	CGCAACGAGCGCAACCCTTATCCTTGTTTGCCAGCACTTCGGGT - GGGAACCTCAGGGAG	1043
Query 661	ACTGCCGGTGATAAACCGGAGGAAGGTGGGGACGACGTCA	700
Sbjct 1044	ACTGCCGGTGATAAACCGGAGGAAGGTGGGGACGACGTCA	1083

Figure Supplementary S1. The 16S rDNA of partial nucleotide sequence comparison using BLAST in NCBI Genebank database