

**Exploring the Pathogenic Potential of *Vibrio vulnificus* Isolated from Seafood Harvested Along the
Mangaluru Coast, India**

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Table S1: Biochemical and physiological characteristics of *V. vulnificus*

Biochemical trait	Result (n=21)
Indole test	+
Salt tolerance 0%	-
Salt tolerance 3%	+
Salt tolerance 8%	-
Salt tolerance 11%	-
Citrate test	+
Arginine decarboxylase	-
Lysine decarboxylase	+
Ornithine decarboxylase	Variable
Mannitol fermentation	+
Lactose fermentation	Variable
Oxidase test	+
Catalase test	-
O/F	+/+
Lecithinase activity	-
Starch hydrolysis	+
Caseinase activity	+
Hemolysis (human erythrocyte)	+
Gelatinase activity	+
Siderophore	+

Table S2: Genotypic attributes of *Vibrio vulnificus* analysed in this study

Genotypic attributes	Result (n=21)
<i>viuB</i> (vulnibactin siderophore)	+
<i>vcgC</i> (Virulence correlated gene, Clinical type)	+
<i>vcgE</i> (Virulence correlated gene, Environmental type)	-
HP1 (CPS Operon allele 1)	+
HP2 (CPS Operon allele 2)	-
<i>vwhA</i> (<i>Vibrio vulnificus</i> hemolysin A)	+
<i>rtxA1</i> (Repeats in toxin)	+

Table S3: Details of isolation of *V. vulnificus* from seafood

Type of sample	Total no of sample	No of samples positive for <i>V. vulnificus</i>
Bivalves (clams, oyster & mussel)	44	19
Crustaceans (crab, shrimp)	16	2
Fin fishes (sardine, mackerel, sole fish)	6	-
Marine sediment	4	-
Total	70	21

Table S4: Details of sampling location and coordinates

Sampling station	Coordinates
Mulki	13°5'18.6936" N 74°46'57.396" E
Mangalore old port	12°51'37.2024" N 74°49'54.9336" E
Sasihithlu	13°3'58.1724" N 74°46'47.5176" E
Ullala	12°49'53.4972" N 74°50'14.0208" E
Manjeshwara	12°42'21.1896" N 74°53'16.4832" E
Bengre	12°52'58.5192" N 74°49'5.0808" E

Table S5: Details of primers used for species specific confirmation, genotyping and virulence gene detection

Primer code	Primer sequence (5'-3')	Product size (bp)	Annealing temperature (°C)	Reference
<i>gyrB</i>	(F): GTCCGAAGTGGAATCCTTCA (R): TGGTTCTTACGGTTACGGCC	285	63	Kumar et al. 2006
<i>vvhA</i>	(F): GACTATCGCATCAACAACCG (R): AGGTAGCGAGTATTACTGCC	704	55	Lee et al. 1998
<i>rtxA1</i>	(F): TGAAGGTGTCGTGGTTACA (R): GTCAGGTCCTTCTCGACAGC	510	60	Acharya et al. 2013
<i>viuB</i>	(F): GGTTGGGCACTAAAGGCAGAT (R): TCGCTTTCTCCGGGGCGG	316	60	Panicker et al. 2004
<i>vcgC</i>	(F): AGCTGCCGATAGCGATCT (R): CGCTTAGGATGATCGGTG	278	55	Rosche et al., 2005
<i>vcgE</i>	(F): CTCAATTGACAATGATCT (R): CGCTTAGGATGATCGGTG	278	55	
HP1	(F): TTTGGGTTTGAAAGGCTTG (R): GTGCCTTTGCGAATTTTGAT	342	50	Han et al. 2009
HP2	(F): TTCCATCAAACATCGCAGAA (R): CTTTTGTCCGGCTTCTATGC	152	50	

Table S6: Details of primers used for real time PCR

Primer code	5'-3'	Product size (bp)	Reference
<i>rtxA1</i>	CCGCATGGTCAAGTAAGGTT TGTCAGGCAATGAGAAGCAC	175	This study
<i>vvhA</i>	CTATCGTGCACGCTTTGGTA ACCGTTTTGTCACCGTTCTC	213	This study
<i>vvpE</i>	AGCATGGAGCAAAGAAGCAT CCTTGCAATGCTCTGTCGTA	183	This study
<i>Flp</i>	TGACAAAAGCGATTGAGTGG CAGACATAGCAACGGCAATAATC	114	This study
<i>ompU</i>	ACACCGTATAGGCCGTCTTG CTGGTGTTAACGCTGCTGAA	166	This study
<i>pilA</i>	GAGTTAATGATAGTGGTGGCG CAGTGTTATGGTACCAAGAGC	225	This study