Mem. Fac. Fish., Kagoshima Univ. Vol. 27, No. 1, pp. 73~78 (1978)

Microflora in the Digestive Tract of Marine Fish-II

Similarity between the Isolates and Vibrio Standard Strains

> Taizo SAKATA*, Jun OKABAYASHI* and Daiichi KAKIMOTO*

Abstract

Vibrio species predominated in the intestines of marine fishes (26 strains) and Vibrio standard strains (15 strains) were examined using 120 bacteriological tests. The results were subjected to calculate the similarity values. There was a considerable number of differential characters between both groups. Phenotypic data obtained in this experiment indicated that the similarity among intestinal Vibrio species from different fishes was relatively higher than that between fish Vibrio and Vibrio standard strains.

It is well known that the indigenous microflora of marine fishes is made up of Vibrio species exclusively. Vibrio species are also commonly isolated from the coastal environment. In the Bergey's Manual¹⁾ (8th ed.) are listed only 5 species for the genus Vibrio, nemely, V. cholerae, V. parahaemolyticus, V. anguillarum, V. fischeri and V. costicola. The objective of this study is to characterize the intestinal Vibrios from marine fishes and to determine the taxonomic relationship with Vibrio standard strains (type strains).

Materials and Methods

Bacterial strains. Bacterial strains used in this study are listed in Table 1. The fish strains were isolated from yellowtail (*Seriola*) cultured in Kinko Bay (1974, 1975), skipjack (*Katsuwonus*) and tuna (*Thunus*) caught in the southwest Pacific Ocean (1975, 1976). The representative strains were selected among the predominant groups in the intestines as being based on the main characteristics and employed in the subsequent experiments. *Vibrio* standard strains were supplied by Dr. TAKAGI, Hokkaido University.

Media used and cultural conditions. The compositions of basal media and cultural conditions were described previously²). All test cultures were inoculated from one day old culture in broth medium by use of a sterile needle.

Morphological and Physiological tests. The regular tests in bacteriology were

^{*} Laboratory of Microbiology, Faculty of Fisheries, Kagoshima University.

Fish sample	Yea	ar	Bacterial count	Isolated	Se	elected
Yellowtail	2.0 kg	1974	5.9×10 ⁶ /g	50	Y I *	3 strains
Yellowtail	0.4	1975	1.0×10 ⁷	54	ΥI	4
			1.0×107	47	ΥS	4
			1.6×10 ⁶	38	ΥF	3
Skipjack	4.3	1975	1.9×10 ⁶	22	ΚI	2
Skipjack	2.3	1975	8.4×10 ⁶	31	ΚI	4
Tuna	12.0	1976	2. 0×10 ⁶	28	ΤI	6
Vibrio standaro	d strains					
Vibrio fischeri						2
Vibrio anguillarum						3
Vibrio alginolyticus						3
	Vibr	io paraha	emolyticus			7
Total strains studied					41	

m 11 4

* Y I, intestine of yellowtail; YS, stomach of yellowtail

YF, food of yellowtail; KI, intestine of skipjack

T I, intestine of tuna.

carried out according to the standard methods described by HARRIGAN et al.³⁾.

Biochemical tests. Hydrolysis of macromolecules was tested using the methods described previously²⁾. Zone of hydrolysis was detected after incubation for 3-7 days by flooding with mercuric chloride solution, Lugol solution and alcohol for gelatin, starch and alginate hydrolysis, respectively. Fermentation of carbohydrates was examined in Durham fermentation tubes and medium containing 1.0 % peptone (Daigo Eiyo), half strength ASW, 1.0% carbohydrate and indicator (BTB). The ability to decarboxylate lysine, ornithine, arginine and glutamic acid was examined by using the method of MOELLER⁴). The ability to utilize nitrogen sources was determined by using the medium containing (g/l) glucose (10), KH₂PO₄ (1), MgSO₄•7H₂O (0.5) KCl (0.2), NaCl (15) and nitrogen source (1), pH 7.5. The sensitivity to various antibiotics was tested by a disc method using tri-discs (Eiken).

Results and Discussion

The predominant Vibrio species (26 strains) isolated from 3 kinds of marine fishes were compared with Vibrio standard strains including V. parahaemolyticus (7 strains), V. alginolyticus (3), V. fischeri (2) and V. anguillarum (3). The similarity values among them were calculated from the results obtained with 120 bacteriological tests. They were identical with respect to acid production from glucose, 0/129 sensitivity and chitin hydrolysis. These tests, however, are characteristics of the genus Vibrio.

The comparison between Vibrio isolates and standard strains based on main

Character		Fish isolates	V. standards	
Colony form	Rough	10	0	
•	Smooth	14	12	
	Swarm	2	3	
Growth at pH	4.5	14	0	
-	5.5	26	14	
	9.5	26	15	
Growih at 42 C		10	0	
Oxidase		22	15	
Catalase		17	13	
Indole		2	9	
Nitrate reductio	on	23	15	
V. P.		16	3	
M. R.		24	15	
Hugh & Leifson	F+G*	12	0	
0	F	14	15	
0/129 sensitivity	y	25	15	
Urease		6	0	
Total strains		26	15	

Table 2. Physiological Chacteristics

F+G, acid and gas production, F, acid production.

Carbohydrates	Isolates		V. stand.	
	F + G*	F*	F + G*	F
Glucose	14	11	0	15
Xvlose	1	3	0	0
Arabinose	0	0	0	0
Rhamnose	0	1	0	0
Fructose	15	9	0	15
Galactose	16	6	0	8
Mannose	15	11	0	15
Cellobiose	4	10	0	12
Sucrose	2	5	0	11
Maltose	11	7	0	8
Lactose	0	1	0	1
Trehalose	11	3	0	15
Raffinose	0	0	0	0
Dextrin	9	6	0	12
Starch	8	6	0	12
Glycogen	19	0	2	10
Inulin	4	18	0	8
Adonitol	0	0	0	0
Mannitol	1	3	0	15
Sorbitol	0	0	0	3
Inositol	0	0	0	0
Salicin	0	0	0	0
Total strains	26		15	;

Table 3. Fermentation of Carbohydrates

F+G, acid and gas production F, acid production from carbohydrates.

Compounds	Isolates	V. standards	
KNO3	1	2	
NH ₄ Cl	13	12	
$\rm NH_4 NO_3$	13	12	
Casamino acid	25	13	
Phe	2	7	
Tyr	6	9	
Trp	2	9	
Asp	7	10	
Asn	20	12	
Lys	9	13	
Met	4	10	
Thr	12	12	
Ilu	2	9	
Glu	9	10	
Gln	3	10	
Pro	2	8	
Arg	15	12	
Gly	9	9	
Ser	11	12	
Cys	14	10	
Ala	5	10	
Leu	6	11	
Val	11	12	
His	14	10	
Total strains	26	15	

Table 4. Nitrogen Source Utilization

Table 5. Hydrolysis of Macromolecules

Macromolecules	Fish isolates	Vibrio standards
Casein	9	13
Gelatin	11	15
Starch	1	13
Glycogen	1	13
Tributyrin	26	15
Tween80	12	14
Chitin	23	13
Cellulose	0	0
alginate	0	0
Total strains	26	15

characteristics is shown in Table 2, 3, 4, and 5. There was a considerable number of differential characters between fish *Vibrio* and *Vibrio* standard strains including colony form on agar plate, growth at pH 4.5, growth at 42 C, gas production from carbohydrates, ability to utilize amino acids, ability to hydrolyze casein and starch and so on. *Vibrio* species predominated in fish intestine had the tendency to be resistant to low pH (pH 4.5) and high temperature (42 C). Moreover, most of them were unable to hydrolyze casein, starch and glycogen and unable to utilize some kinds of amino acids as sole nitrogen source



in contrast to *Vibrio* standard strains. Similarity diagram shown in Fig. 1 indicates that there is relatively higher similarity among the predominant *Vibrio* strains in the intestine of marine fishes than that between *Vibrio* species isolated from fishes and *Vibrio* standard strains.

SIMIDU et al.⁵ analyzed 114 strains of *Vibrio* and allied genera isolated from sea water, normal marine fish and diseased fish and divided them into two large, distinct groups (phenons). The *Vibrio* species isolated from the marine fish intestines in this study showed a considerable similarity with phenon 1 proposed by SIMIZU et al.⁵ in spite of diversity in casein, gelatin and starch hydrolysis.

Acknowledgement

The authors are grateful to Dr. K. TAKAGI, Hokkaido University, who kindly provided *Vibrio* standard strains.

A part of this study was supported by a Reseach Grant of the Ministry of Education in Japan (1975, no 07169).

References

- 1) SHEWAN, J. M. and M. VERON (1974): Bergey's Manual of Determinative Bacteriology. 8th edition. 340-345. The Williams and Wilkins Co., Baltimore.
- 2) SAKATA, T., M. NAKAJI and D. KAKIMOTO (1978): Microflora in the digestive tract of marine fish—I. *This Mem.*, in press.
- 3) HARRIGAN, W. F., and E. M. MARGAREL (1966): Laboratory methods in microbiology. Acad., Press, London, New York.
- 4) MOELLER, V. (1955): Acta Pathol. Microbiol. Scand., 36, 158.
- 5) SIMIDU, U. and E. KANEKO (1973): A numerical taxonomy of Vibrio and Aeromonas from normal and diseased marine fish. *Bull. Jap. Soc. Sci. Fish.*, 39, 689-703.