Virulence Factors and Genotyping of Vibrio parahaemolyticus

Eun-Gyoung Lim, Young-Hee Kim, Ji-Yung Mun, Yang-Hyo Oh and Yung-Bu Kim

Department of Microbiology, College of Medicine, Pusan National University, Pusan, Korea

Six strains of Vibrio parahemolyticus isolated from diarrheal patients and 12 strains from seawater were serotyped and analyzed for biochemical characteristics, antibiotic sensitivity and detection of toxR, gyrB, tdhl, and trh2 genes. Arbitrarily-primed polymerase chain reaction was performed on the 6 strains from patients and the results were as followed. The V. parahemolyticus isolated from patients belonged to 5 different serotypes: O4:K8, O4:KUT, O6:K18, O10:K71 and O3:K6, but those isolates from sea water belonged to 5 different serotypes: O1:KUT, O2:KUT, O3:K45, O4:K37 and OUT:KUT. All strains explained have different serotypes depending on the different source. Both toxR and gvrB genes were detected from all strains isolated. As for control the 2 strains of serotype O3:K6 and 6 strains isolated from patients, serotype O3:K6 were resistant to oxacillin, peniciffin, and vancomycin. All strains were sensitive to chloramphenicol and tetracycline yet the antibiogram type showed 5

groups from I to V. The *trh*1 was detected both from serotype O4:KUT and O6:K18 from patients and the *trh*2 was also detected from one strain from each O10:K71 and O1:KUT isolated from patients and seawater respectively. The *tdh* gene was only detected from two strains of O3:K6 from patients of 1998. The *tdh*, *trh*1 and *trh*2 were not detected from 7 strains out of 12 strains isolated from seawater whereas the production titer of TDH isolated from patients showed from 2048 times to 4096 times.

Four strains of the serotype O3:K6 isolated from Korea, India and Japan as well as 3 strains from Korean patients were tested by AP-PCR to classify serotypes. As for its result the amplicon showed the same in the 4 strains of the serotype O3:K6 whereas the four strains of different serotype from patients are so difference as to explain no inter-relations at all. The result explains that the serotype O3:K6 is the same genes regardless from where it is isolated.