Molecular Detection of Zonula Occludens Toxin (zot) Genes in Vibrio Cholerae O1 using PCR

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Abstract
Background: Zot (Zonula occludens toxin) is one of the secretion toxins of Vibrio cholerae in small intestine that binds to certain receptors in the epithelial cells and causes a change in the structure of tight junction. The purpose of this research is rapid detection of zot enterotoxin gene using PCR.

Materials and Methods: The genomic DNA was extracted by DNA isolation kit and gene amplification was carried out by the zot gene-specific primers. Then, PCR products were investigated by electrophoresis on 1.2% agarose gel stained by ethidium bromide. Also, the specificity of primers was measured using bacterial samples other than V. cholerae, such as enterotoxigenic Escherichia coli (ETEC), Salmonella typhi and Aeromonas hydrophila. The sensitivity of the PCR reaction was also evaluated using serial dilutions of V. cholerae O1 concentration (cfu/ml).

Results: The data showed that the designed primers specificity for zot gene was successful and the sensitivity of this method was determined about 142 cfu/ml.

Conclusion: In conclusion, this molecular detection can be used as a simple diagnostic kit in clinical laboratories for identification of V. cholerae.

Keywords: Vibrio cholera; zot; PCR; diagnosis

Introduction
Vibrio cholerae, the causative agent of cholera, is an important agent for causing severe diarrheal disease in many parts of the world. According to severe watery diarrhea and rapid dehydration, this disease may lead to death of the patients (1, 2). Cholera is a seasonal disease and occurs especially in the warm seasons of the year in areas where poor health is present (3). Therefore, correct, reliable and rapid detection of cholera toxin (CT)-producing V. cholerae is very important. Today, many identification methods are carried out for detection of V. cholerae such as conventional culture, biochemical, immunological and molecular-based assays (4-7). Although isolation and identification of V. cholerae by culture of a stool specimen is the gold standard for the laboratory detection of cholera, this method is time consuming and laborious, requiring more than three days (8). On the other hand, this method and some commercially available kits cannot distinguish between CT and the heat-labile enterotoxin (LT) of Escherichia coli (9). Also, due to nutrient starvation and physical stress, this bacterium may enter a viable but non-culturable (VBNC) state. Therefore, this procedure may explain the failure of traditional culture methods for isolation and detection of this infectious agent from contaminated water and food samples implicated in foodborne outbreaks (10, 11). In this field, molecular diagnosis can be a key strategy and is used for successful detection of this infectious agent. Zonula occludens toxin (zot), is a main toxin from V. cholerae that regulates tight junction permeability, reversibly. Since the location of the zot coding gene is upstream of cholera toxin, zot can play a synergistic role in the development of symptoms (12, 13). In this work, we used PCR for identification of zot enterotoxin gene in order to develop a rapid and correct detection kit.

Materials and methods

DNA Extraction
V. cholerae (O1 serogroup), enterotoxigenic Escherichia coli (ETEC) and Aeromonas hydrophila were obtained from Bou-Ali Hospital in Tehran. All Strains were grown for 24 h in LB broth, aerobically. Then, the bacterial
sedimentation was done by centrifugation at 9000 g for 5 minutes and finally, genomic DNA was purified using a DNA extraction kit (DNP TM Kit, SinaClon BioScience) according to manufacturer’s protocol.

**Primer design**
In this study, zot gene sequence was examined in GenBank and primers were designed according to this sequence. For this, one pair of primers was designed for specific amplification of this gene (Forward: 5′-CGGCTTTGTGTCCAAGATGG-3′ and reverse: 5′-CGCTGCAAAGGTATCGAACA-3′). Accordingly, a specific band of 246 bp was amplified.

**PCR Reaction**
PCR was performed with forward and reverse primers, the final volume for each reaction was 5 μl, which included 60 ng (1 μl) of DNA template, 2.5μM (0.25 μl) each of the forward and reverse primers, 2.5 μl master mix of Taq DNA polymerase (AmpliQon PCR Kit, Denmark) and 1 μl of distilled water. The program of thermocycler for reaction was prepared: initial denaturation at 94 °C for 5 minutes and 30 cycles including denaturation (95 °C, 1 min), annealing (55 °C, 1 min), extension (72 °C, 1 min) and final extension at 72 °C for 5 minutes. After completion of the reaction, PCR products were examined using 1.2% agarose gel electrophoresis.

**Results**

**PCR Detection**
The extracted DNA from *V. cholerae* culture was evaluated using electrophoresis on 1.2% agarose gel (Figure 1A). The measured DNA value with NanoDrop instrument was 62 ng / μl. To carry out the PCR reaction, extracted DNA was used as template. PCR reaction was done. After amplification and analysis with 1.2% agarose gel, a clear band was observed at position 246 bp, which confirmed presence of zot gene in the *V. cholerae* sample (Figure 1B).

**Specificity and sensitivity of primers in PCR detection**
To confirm the specificity of primers, DNA templates were selected from samples of *V. cholerae*, ETEC (enterotoxigenic *Escherichia coli*), *Salmonella typhi* and *Aeromonas hydrophila*.

**Specificity and sensitivity of primers**
To evaluate the specificity of the designed primers, genomic DNA samples were selected as DNA templates from *V. cholerae* O1 (Inaba Serotyping), ETEC (enterotoxigenic *Escherichia coli*), *Salmonella typhi* and *Aeromonas hydrophila* were selected as DNA templates. Moreover, to determine the sensitivity of the test, a sample of *V. cholerae* was diluted sequentially from 1.42×10⁷ cfu/ml to 1.42×10⁻⁴ cfu/ml, and PCR was performed with each of these dilutions. Finally, PCR products were examined with 1.2% agarose gel electrophoresis.
The electrophoresis on the agarose gel showed a clear band for *V. cholerae* at 246 bp, confirming the specificity of primers designed for zot gene in *V. cholerae*. In samples other than Vibrio, which were used as a template, they were not amplified and were observed as primer dimers (Figure 2). To confirm the PCR sensitivity, serial dilutions were prepared from 1.42×10^3 cfu/ml to 1.42×10^4 cfu/ml of *V. cholerae* genome and considered as a template for the reaction. After analysis of PCR products, it was found that the sensitivity of this assay is about 142 cfu/ml (Figure 3).

**Discussion**

In this work, we developed a simple diagnostic molecular assay for detection of zot gene from *V. cholera*. For this, the specificity of primers was studied using bacterial samples other than *V. cholerae* including enterotoxigenic *E. coli*, *Salmonella typhi* and *Aeromonas hydrophila*, which data showed that the specificity of primers for *V. cholerae* is successful. In addition, the sensitivity of the PCR technique was investigated by providing different dilutions of the *V. cholerae* genome, and the limit of detection was determined about 142 cfu/ml. Conventional microbiological methods for detection of *V. cholerae* are time consuming and expensive. Molecular assays such as PCR are very specific compared to conventional methods and suitable alternative to traditional methods (14, 15). The rapid diagnosis of cholera and its timely treatment is very important (16). Cholera is one of the infectious and acute diseases of the gastrointestinal tract that is associated with reduced water and electrolyte imbalances in the patient's body (17). Toxigenic strains of *V. cholerae* have played an important role in the development of cholera epidemics (18). zot is the second toxin of *V. cholerae*, which increases the permeability of bowel intestinal mucosa by altering the zonula occludens joints (19, 20). The change caused by zot enterotoxin is reversible and limited to the small intestine (21). In the present work, by designing specific primers for the zot gene, a PCR technique was used to amplify this gene. In 2011, Chua et al., used a triplex PCR method to identify toxic and non-toxigenic *V. cholerae*. The diagnostic limit of this method was reported 2×10^3 cfu (3). In 2012, Mehrabadi et al., used the multiplex PCR method to identify toxigenic *V. cholerae*, which had a diagnostic limit of 10-100 cfu (15). In 2014, Barzamini et al., used a PCR method to detect ctxA gene and their diagnostic limit was estimated about 40 cfu /ml (14). In 2015, Zeinoddini et al., with new method, triplex dot blotting assay, detected *V. cholerae* in sensitivity of 10 cfu (6).

**Conclusion**

In conclusion, according to this molecular detection we could develop a simple diagnostic kit for clinical laboratories in order to identify *V. cholerae*.

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**Author Contributions**

The authors have similar contributions.

**Conflict of Interest**

The authors declare no conflict of interest.

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