

DECLARATION

I declare that this dissertation, which I hereby submit for the degree of  
**Development of novel seminested polymerase chain reaction  
assays for detecting toxigenic *Vibrio cholerae* and *Shigella*  
spp. in water**

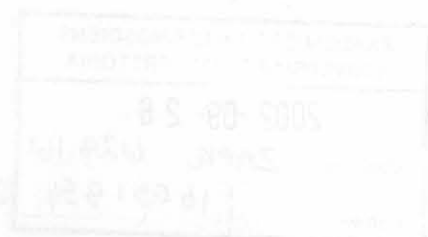
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by

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Submitted in partial fulfilment of the requirements for the degree of  
**Magister Scientiae**  
in the Faculty of Natural and Agricultural Sciences  
University of Pretoria  
Pretoria  
December 2001



ACKNOWLEDGEMENTS

**DECLARATION**

I declare that the dissertation, which I hereby submit for the degree Magister Scientiae at the University of Pretoria, is my own work and has not previously been submitted by me for a degree at another university.

Dr. S. N. Venter for his guidance and for convincing me that this study was possible.

My son, who never complained.

\_\_\_\_\_  
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Thanks to my father for his help and enduring patience.

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Dr. S.N. Venter for his guidance and for convincing me that this study was possible.

My son, who never complained.

Kevin Murray for his help and enduring patience.

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Rapid and sensitive polymerase chain reaction (PCR)-based assays were developed for the detection of toxigenic *Vibrio cholerae* and virulent *Shigella* spp. in environmental water samples. In contrast to conventional nested and semi-nested PCR assays, the newly developed pit-strip seminested PCR assays are performed in a total of 2 amplification cycles. Oligonucleotide primers were designed to specifically amplify sequences within the cholera toxin gene (*ctxA/ctxB*), *ctxA/ctxB* and the toxin-converting phage (*ctxΦ*) of virulent *Shigella* spp. By using agarose gel electrophoresis, detection of the PCR-amplified products, a detection limit of  $1.3 \times 10^3$  cfu of *V. cholerae* and  $1.8 \times 10^3$  cfu of *Shigella flexneri* were obtained from crude cell lysates. However, by coupling the PCR assays with an enrichment culture procedure, the sensitivity of the

## SUMMARY

### Development of novel seminested polymerase chain reaction assays for detecting toxigenic *Vibrio cholerae* and *Shigella* spp. in water

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Rapid and sensitive polymerase chain reaction (PCR)-based assays were developed for the detection of toxigenic *Vibrio cholerae* and virulent *Shigella* spp. in environmental water samples. In contrast to conventional nested and seminested PCR assays, the newly developed pit-stop seminested PCR assays are performed in a total of 30 amplification cycles. Oligonucleotide primers were designed to specifically amplify sequences within the cholera toxin gene (*ctxA*) of *V. cholerae* and the invasion plasmid antigen gene (*ipaH*) of virulent *Shigella* spp. By using agarose gel electrophoresis for detection of the PCR-amplified products, a detection limit of  $1.3 \times 10^3$  cfu of *V. cholerae* and  $1.6 \times 10^3$  cfu of *Shigella flexneri* were obtained from crude cell lysates. However, by coupling the PCR assays with an enrichment culture procedure, the sensitivity of the

assays was improved and as few as 4 cfu and 1.6 cfu of *V. cholerae* and *S. flexneri* organisms, respectively, could be detected in pure culture. Analysis of seeded environmental and drinking water samples yielded detection limits of as few as one *V. cholerae* organism/ml and less than 14 cfu/ml *S. flexneri* in some samples.

The basic PCR detection protocol was modified to incorporate a membrane filtration step in order to concentrate large volumes of water samples. This allowed for the detection of low numbers of viable *V. cholerae* and *Shigella* spp. in naturally contaminated waters. By making use of seeded environmental, and drinking water samples, detection limits of 1 cfu/100 ml of *V. cholerae* and 8 cfu/100 ml of *Shigella* cells were obtained. The detection protocol was successfully applied to the analysis of naturally contaminated environmental water samples. The results obtained in this study indicate that the newly developed pit-stop seminested PCR assays, combined with an enrichment step, fulfills the requirements of simplicity and sensitivity for use in an environmental laboratory and may represent significant tools in monitoring water sources for the presence of *V. cholerae* and *Shigella* spp.

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