

Cloning and protein expression of *Actinobacillus actinomycetemcomitans* cytolethal distending toxin subunit CdtA

Sun-Young Ko¹, Dong-Keun Jeong¹, So-Hyun Ryu¹,
and Hyung-Seop Kim^{1,2}

1. Department of Periodontology, College of Dentistry, Chonbuk National University
2. Reseach Institute of Oral Bio-Science, Chonbuk National University

I. INTRODUCTION

Periodontitis is a persistent bacterial infection, which causes chronic inflammation that leads to the destruction of the tooth supporting tissues. A wide variety of bacterial species are implicated in the pathogenesis of the disease. One of these is the Gram-negative facultative anaerobe *Actinobacillus actinomycetemcomitans*, a species highly associated with localized aggressive periodontitis. This form of the disease occurring in adolescents and young adults is characterized by rapid and severe loss of supporting alveolar bone and connective tissue attachment¹. It remains unclear that the general pathogenic mechanisms and immune involvement in periodontitis, in particular the bacterial viru-

lence factors responsible for the proposed immunosuppression in this disease. *A. actinomycetemcomitans* occasionally responsible for non-oral infections including endocarditis, bacteremia, pericarditis, septicemia, pneumonia, infectious arthritis, osteomyelitis, synovitis, skin infections, urinary tract infections, and abscesses². It has been estimated that approximately 0.6% of cases of infective endocarditis are caused by *A. actinomycetemcomitans*³. Studies suggest that bacterial virulence factors act to impair host mechanisms and play significant roles in infectious diseases associated with *A. actinomycetemcomitans*⁴⁻⁶. These virulence factors included leukotoxin, lipopolysaccharide(LPS), and cytotoxin-cytolethal distending toxin(CDT).

CDT was first documented as a toxin oc-

* Correspondence: Hyung-Seop Kim, Department of Periodontology, College of Dentistry, Chonbuk National University, 634-18, gumam-dong Dukjin-gu, Chonju, Korea, Tel: +82-63-250-2115, Fax: +82-63-250-2259 (E-mail: cbuperio@chonbuk.ac.kr)

curing in *Escherichia coli* strains by Johnson and Lior in 1987^{15,26,27}. CDT has been identified and cloned from *A. actinomycetemcomitans*⁷⁻⁹, and various other pathogenic bacteria¹⁰⁻¹⁸. The term CDT reflected the unique properties of this protein causing progressive cell distension and cytotoxicity to cultured cells such as CHO, HeLa, and Hep-2.

Another unique phenotype of the CDT-intoxicated cells is cell cycle arrest at the transition from G2 to M phase. *A. actinomycetemcomitans* is the only oral bacterium known to produce a cytolethal distending toxin (CDT), which causes growth arrest in mammalian cells¹⁹⁻²².

CDT is prevalent in *A. actinomycetemcomitans* strains. Ahmed et al. (2001) found that 43 of 50 strains from periodontitis patients contained all three *cdt* genes and expressed CDT activity. In another study, PCR of subgingival plaque samples revealed that 13 of 106 diseased sites in 146 patients with aggressive and chronic periodontitis contained *A. actinomycetemcomitans* expressing all three *cdt* genes²⁴. Fabris et al. (2002) reported that 39 of 40 *A. actinomycetemcomitans* isolates from a mix of healthy and periodontal diseased subjects expressed activity that caused the distension of CHO cells.

The genes of *cdt* have been isolated as *cdtA*, *-B*, and *-C*, tandemly located in the *cdt* locus. Most recent studies indicate that all three components, *cdtA*, *-B*, and *-C*, are required for full activity²⁸⁻³². Pull-down as-

say revealed that the active CDT holotoxin is a heterotrimer, consisting of CdtB as the enzymatically active subunit, and CdtA and CdtC which mediate the delivery of CdtB into host cells²⁸⁻³¹.

A study of Lee et al. (2003) with *C. jejuni* showed both CdtA and CdtC, but not CdtB, were able to bind individually to the surface of HeLa cells using a common receptor³³. Mao and DiRienzo (2002) demonstrated that only CdtA was able to bind to the surface of CHO cells. Following internalization conversely, no cytotoxic effects were observed. On the contrary, the introduction of CdtB or CdtC into the cells did cause cell distention and eventually cell death. This study urges one to conclude only CdtA is responsible for the attachment to the cell and the CdtC subunit assists CdtB in the cytolethal activity of the holotoxin³⁴.

The precise role of the CdtA protein in the mechanism of action of the holotoxin is unknown at the present time. The ultimate goal of this study is to develop an edible vaccine against *A. actinomycetemcomitans*, causative agent for LAP. As an initial step, I cloned *cdtA* gene from *A. actinomycetemcomitans* and expressed CdtA protein.

II. MATERIALS AND METHODS

1. Materials

All chemicals and plastic wares were purchased from Sigma (St. Louis, MO) and Falcon Labware (Frankline Lakes, NJ),

respectively. Oligonucleotides were ordered from COSMO. T vector and Tag polymerase was supplied from Promega(USA). Restriction enzymes were obtained from BioLabs (New England).

2. Bacterial culture

A. actinomycetemcomitans Y4 was grown in Luria-Bertani(LB) broth medium for 24 h at 37°C in an atmosphere of 5% CO₂ in air and 200µl of overnight culture was transferred into 5ml of a fresh medium, leaving for further incubation.

3. Preparation of genomic DNA

About 1 ml of an overnight culture of *A. actinomycetemcomitans* Y4 was collected and added to a 1.5 ml microcentrifuge tube. The tube was centrifuged at 13,000~16,000 ×g for 2 minutes to pellet the cells. Remove the supernatant. After addition 600 µl of Nucleus Lysis Solution, gently pipeted until the cell are resuspended. The mixture was further incubated at 80°C for 5 minutes to lyse the cells; then cooled to room temperature. After addition 3 µl of RNase Solution to the cell lysate, the sample was again incubated at 37°C for 15~60 minutes

then cooling to room temperature. Finally added 200 µl of Protein Precipitation Solution was added with vigorous vortexing. The vortexed sample was incubated on ice for 5 minutes, then centrifuged at 13,000~16,000 ×g for 3 minutes.

4. PCR amplification of *cdtA* gene

The genomic DNA was used as templates for amplifying *cdtA* gene in 50µl of PCR reaction containing Taq polymerase, dNTP, and oligonucleotides, primers, 10×PCR buffer. The primers specific for *cdtA* gene were designed and synthesized as described in Table 1. The genomic DNA was denatured by 5 min incubation at 94°C and then target genes were amplified for 30 cycles of PCR reaction (1 min 94°C, 1 min 63°C, 1 min 72°C). The PCR product was then treated for 10 min at 72°C and stored at 4°C.

5. Cloning of *cdtA* gene

PCR products were ligated into pGM-T vector following manufacture's protocol and the ligation mixtures were transformed into competent *E. coli* Top10 cells. The resulting antibiotics resistant bacteria were selected from agar plates and further screened by

Table 1. List of oligonucleotides used to amplify *cdtA* genes. The underlined sequences are the locations of restriction enzyme sites.

	Sequence
<i>cdtA</i> (F)	ATG <u>CGG</u> <u>ATC</u> <u>CAT</u> GAG TGA CTA TTC TCA G
<i>cdtA</i> (R)	ATG <u>CCT</u> <u>GCA</u> <u>GTT</u> AAT TAA CCG CTG TTG C

checking the plasmid from each bacterial colony. First, plasmids were compared in size with that of control vector in agarose gels to reveal the presence of any insert DNA. Second, digestions of plasmids with restriction enzymes were performed to further prove the presence of target genes. The plasmids containing expected insert size were then sequenced to confirm the integrity of target gene. The DNA sequencing was performed by Xenotech (Korea).

6. Sequence analysis of *cdtA* gene from *A. actinomycetemcomitans* (Y4)

The DNA sequence of cloned genes was analyzed by comparing with the previously known sequences. For *cdtA* sequence analysis, the known sequence from *A. actinomycetemcomitans* (Y4)(GenBank GI9955924) was used as standard. The internet site (www.ualberta.ca) were used to generate sequence alignment between two genes.

7. Expression of recombinant CdtA protein in bacterial expression system

In order to express cloned genes in *E. coli* system, gene fragments of *cdtA* were subcloned to pRSET expression vector. BamHI and PstI were used to transfer *cdtA* gene. The ligation mixtures were transformed in TOP10 host cell. The ampicillin resistant bacteria were selected from LB agar and transformed BL21(DE3) cell.

Transformed BL21(DE3) bacterial colonies were inoculated and grown overnight at 37°C in LB agar containing 100µg/ml of ampicillin. The cells will be further grown for 4h in the presence of 1mM IPTG to induce expression of the fusion protein.

8. Isolation of recombinant CdtA protein

The cells were obtained by centrifugation at 6000rpm for 10minutes. The cells were suspended in phosphate buffer saline (PBS). The suspension was sonicated for three 30seconds bursts on ice. The sonicated samples were centrifuged at 13,000rpm for 10 minutes. The supernatant fluid was then passed through a 0.45 micron filter to obtain the soluble fractions. Cell pellets were washed with PBS containing 0.01% Triton X-100 and resuspended in PBS to obtain the inclusion body.

9. Analysis of CdtA protein (SDS-PAGE, Western blot)

The factions containing recombinant protein (6×His-tagged protein) were identified by SDS-PAGE. Bacteria collected from 1ml of IPTG-induced culture were mixed with 100 µl of gel loading buffer and heated in a boiling water bath for 5min. Samples were applied to a 10~20% polyacrylamide gel (25 µg of protein per lane). After electrophoresis, protein bands were transferred to a nitrocellulose membrane. The membrane was

blocked with 3% BSA in TBS buffer (20mM Tris-HCl, pH 7.6, 0.8% NaCl) for 1h at room temperature with shaking. The blocked membrane was added Ni-NTA tagged with AP (alkaline phosphate) and then washed in TBS buffer. The membrane was added BCIP/NBT to induce developing reaction and incubated in the dark. The membrane was rinsed with dH₂O to stop the developing reaction.

III. RESULTS

1. PCR product of *cdtA* gene

The genomic DNA from *A. actinomycetemcomitans* Y4 was isolated and used as templates for amplifying *cdtA* genes. The primer specific for *cdtA* gene was separately designed based on the published sequences. PCR was performed yielding 0.6 kb *cdtA* (Figure 1).

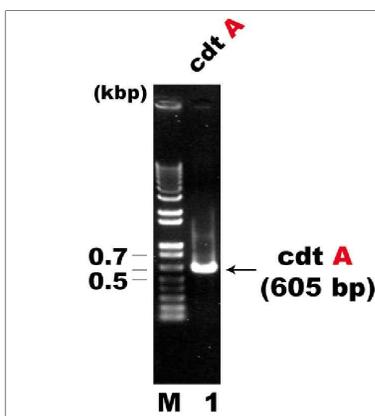


Figure 1. PCR amplification of *A. actinomycetemcomitans* Y4 genes encoding *cdtA*. The genes for *cdtA* were amplified by PCR using the genomic DNA linearized by restriction enzyme digestion.

2. Cloning into T-vector of *cdtA*

PCR products were ligated into T vector, yielding T vector cloned with *cdtA*. The cloned T vectors were transformed into an *E. coli* strain, Top 10. Plasmids were prepared from antibiotics-resistant colonies and separated in agarose gel (Figure 2). In T-*cdtA* agarose gel, the 4th lane band is lower than the others because it does not include insert DNA.

3. Sequence analysis of cloned genes

After checking the presence of any insert in the ligated plasmid, we confirmed the integrity of cloned *cdtA* sequence by nucleotide sequencing. For *cdtA* sequence analysis, the corresponding sequence from *A. actinomycetemcomitans* Y4 was compared with that of the previously-cloned gene. The rep-

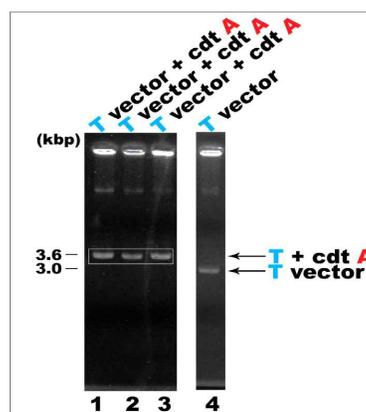


Figure 2. Screening of plasmids cloned with PCR-amplified *cdtA* genes. The genes encoding *cdtA* were cloned into T vector and bacterial transformants were screened for the presence of insert DNA.

```

Aa-Y4 ATGAGTGACTATTCTCAGCCTGAACTCTCAATCTGATTTAGCACCTAAATCTTCAACAACACAAATCCAAACCCCAACCCCT 80
r-cdtA ATGAGTGACTATTCTCAGCCTGAACTCTCAATCTGATTTAGCACCTAAATCTTCAACAACACAAATCCAAACCCCAACCCCT 80

Aa-Y4 ATTATCAAAGCATCTTCAATGCCATTGAATTTGCTCTCTTCATCCAAGAATGGACAGGTATCGCCGCTTGAACCATCAA 160
r-cdtA ATTATCAAAGCATCTTCAATGCCATTGAATTTGCTCTCTTCATCCAAGAATGGACAGGTATCGCCGCTTGAACCATCAA 160

Aa-Y4 ACTTTATGACTTTGATGGGACAAAATGGGGCACTGTTGACTGCTCTGGGCGCTAGCAAACGCAATGGTTATGGGCTTAT 240
r-cdtA ACTTTATGACTTTGATGGGACAAAATGGGGCACTGTTGACTGCTCTGGGCGCTAGCAAACGCAATGGTTATGGGCTTAT 240

Aa-Y4 CCCAATATATATTCGCAGGACTTTGGAAATATTCGTAATGGGAAGATAGAACCCTGGTAAACACCGTGAAATATTTTCGTTT 320
r-cdtA CCCAATATATATTCGCAGGACTTTGGAAATATTCGTAATGGGAAGATAGAACCCTGGTAAACACCGTGAAATATTTTCGTTT 320

Aa-Y4 TGTTAATCAATCTTTAGGTACATGTATTGAAGCTTACGGTAATGGTTTAAATTCATGATACTTGTAGTCTGGACAAATTAG 400
r-cdtA TGTTAATCAATCTTTAGGTACATGTATTGAAGCTTACGGTAATGGTTTAAATTCATGATACTTGTAGTCTGGACAAATTAG 400

Aa-Y4 CACAAGAGTTTGAAGTATTACCTACTGATAGTGGTTCGGTTGTCATTAAAAGTGTCTCACAGGACCTTGTGTCACCTTAT 480
r-cdtA CACAAGAGTTTGAAGTATTACCTACTGATAGTGGTTCGGTTGTCATTAAAAGTGTCTCACAGGACCTTGTGTCACCTTAT 480

Aa-Y4 AATCCTGTAAGTCCAACATATTTATTCACAGTTACATTTCAACTTGTGATGGCGCAACAGAACCAATTACGTGATCAAAC 560
r-cdtA AATCCTGTAAGTCCAACATATTTATTCACAGTTACATTTCAACTTGTGATGGCGCAACAGAACCAATTACGTGATCAAAC 560

Aa-Y4 ATGGTATCTCGCTCCTCCTGTATTAGAAGCAACAGCGGTTAATTAA 606
r-cdtA ATGGTATCTCGCTCCTCCTGTATTAGAAGCAACAGCGGTTAATTAA 606

```

Figure 3. Sequence alignment between the cloned *cdtA* gene and the corresponding region of the reported *cdtA* gene. The sequence of the cloned *cdtA* gene was compared to that of known *cdtA* gene.

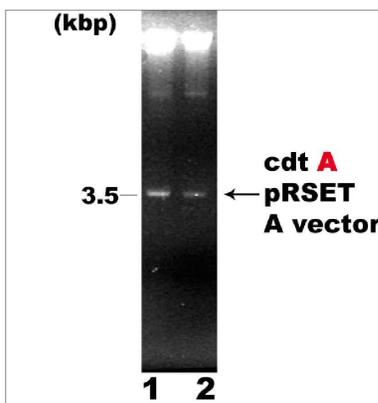


Figure 4. Screening of plasmids of *cdtA* cloned in pRSET vector. The pRSET-*cdtA* plasmid was transformed in BL21(DE3) bacterial strains, and their transformants were screened for the presence of recombinant expression vector.

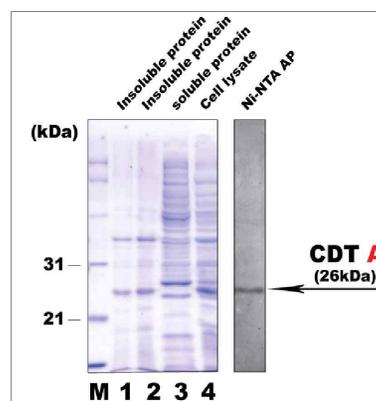


Figure 5. *CdtA* insoluble protein was confirmed and when testing with AP conjugated Ni-NTA, *CdtA* protein of 26kDa was observed.

representative sequence alignment was shown in Figure 3. There was no miss match at all.

4. Subcloning into expression vector

The *cdtA* gene was subcloned into pRSET vector according to the procedures suggested by the manufacturer(Qiagen). And the pRSET-*cdtA* transformed into BL21(DE3) host cells. The *cdtA* plasmids were isolated and purified for the induction of recombinant protein. Plasmids with expected *cdtA* gene were prepared from antibiotics-resistant colonies and separated in agarose gel(Figure 4).

5. Expression of recombinant CdtA

After IPTG induction of pRSET-cdt A Expression vector, SDS-PAGE & Western blot proceeded. As a result, CdtA insoluble protein was confirmed and when testing with AP conjugated Ni-NTA, CdtA protein of 26kDa was observed(Figure 5).

IV. DISCUSSION

Actinobacillusactinomycetemcomitans has been implicated not only in the pathogenesis of serious periodontal disease¹⁾, but also in a variety of human systemic diseases such as endocarditis, meningitis and osteomyelitis²⁾. Some cytotoxic factors produced by *A. actinomycetemcomitans* have been studied at the molecular level, and CDT from this pathogen has been identified and cloned^{7,8,9)}.

The CDT action was first described in a

culture of eukaryotic cells challenged with supernatants from *Campylobacter* spp. and enteropathogenic *E. coli*^{15,26,27)}. Both species caused growth arrest of the intoxicated cells, along with slow distension, over a period of 2 to 5 days. It is now known that CDT is produced by a number of Gram-negative pathogenic bacteria, including *Haemophilus ducreyi*, *Shigella dysenteriae*, *Helicobacter hepaticus*, as well as *A. actinomycetemcomitans*¹⁰⁻¹⁸⁾. The growth inhibition and morphological alterations caused by the toxin have been demonstrated in fibroblastic^{18,26,27)}, epithelial or endothelial cells^{18,35,36,37,38)}, as well as T- and B-lymphocytes^{19,39,40)}.

The periodontium is composed of a number of distinct cell types including epithelial cells, fibroblasts, cementoblasts and osteoblasts. From a functional standpoint, the periodontium can be broadly broken down into two tissue compartments referred to as gingiva and the attachment apparatus. Gingiva, which is composed of epithelial cells and the adjacent connective tissue, is attached to a tooth via hemidesmosomes formed by cells of the junctional epithelium. This provides a physical barrier that protects the underlying attachment apparatus from the potentially harmful host as well as microbial and/or environmental factors found within the oral cavity proper. The connective tissues of the periodontium are the gingival connective tissue and the periodontal ligament (PDL) tissue. Gingival fibroblasts (GF) comprise the major cell population of the gingival connective tissue and are respon-

sible for the production of the extracellular matrix of the tissue in health and disease⁴¹). The PDL is a physically small but very important connective tissue, since it links the alveolar bone to the tooth surface, providing the appropriate tooth support in response to mechanical loading⁴²). It consists of a variety of fibroblast cell populations (most commonly designated as PDL cells) with different functional characteristics^{43,44}). Both the gingival connective tissue and PDL are likely to be targeted by *A. actinomycetemcomitans*⁴⁵,⁴⁶). Belibasakis et al. (2002) demonstrated that CDT of *A. actinomycetemcomitans* is involved in the non-lethal inhibition of proliferation of human periodontal ligament fibroblasts and human gingival fibroblasts. Belibasakis et al. (2004) suggested that inhibition was associated with cell-cycle arrest at both the G0/G1 and G2/M phases of growth⁴⁷). However, Kanno et al. (2005) suggested that it is extremely difficult to evaluate these results because: 1) these investigators did not attempt to identify either cdt genes or Cdt polypeptides in their test strain *A. actinomycetemcomitans* HK 1519; 2) their bacterial extract was a crude preparation contaminated with significant levels of LPS (6.5 µg/ml); 3) the concentration of extract used in their assays was not reported; and 4) effects on cell cycle and evaluation of DNA damage, standard measures of CDT activity, were not determined. Kanno et al.(2005) showed that Human periodontal ligament fibroblasts (HPLF) are resistant to the cytotoxic effects of the *A. actino-*

mycetemcomitans CDT. Belibasakis et al. (2005) demonstrated that *A. actinomycetemcomitans* induces IL-6 production in GF by a mechanism largely independent of its Cdt and *A. actinomycetemcomitans*-induced RANKL expression in GF occurs independently of IL-1, IL-6, TNF-α or PGE2.

The crystal structure of the holotoxin from *Haemophilus ducreyi* reveals that CDT consists of an enzyme of the DNase-I family, bound to two ricin-like lectin domains. The lectin subunits form a deeply grooved, highly aromatic surface. The holotoxin possesses a steric block of the CdtB active site by means of a non-globular extension of the CdtC subunit⁵⁰). *A. actinomycetemcomitans* CdtA has a signal peptidase II recognition site that is normally found on lipoproteins, suggesting that this protein may be anchored to the bacterial outer membrane⁷). CdtA labelled with either a fluorophore³⁴) or with biotin³³) has been found to localize to selected areas of the plasma membrane of mammalian cells.

V. CONCLUSION

It is not clear what role CdtA plays in the final activity of CDT, and further work is required to fully identify the role of CdtA and its mechanism of host cell binding. The vaccine development to target *A. actinomycetemcomitans* *cdtA* and the early immunization during the time span of the window of infectivity will have a synergistic effect on suppressing internalization of *A. actinomycetemcomitans* CDT. For this pur-

pose, I cloned *cdtA* from *A. actinomycetemcomitans* (Y4), and subcloned *cdtA*, expressed and purified to homogeneity the CdtA protein as 6×His tag fusion protein. Next, I will try to investigate the biological activities of CdtA in bacterial pathogenesis, and also to produce the CdtA antibody for vaccine development.

VI. REFERENCES

1. Slots J, Ting M. *Actinobacillus actinomycetemcomitans* and *Porphyromonas gingivalis* in human periodontal disease: occurrence and treatment. *Periodontol* 2000 1999;20:82–121.
2. Van Winkelhoff AJ, Slots J. *Actinobacillus actinomycetemcomitans* and *Porphyromonas gingivalis* in nonoral infections. *Periodontol*. 2000 1999;20:122-35
3. Das M, Badley AD, Cockerill FR, Steckelberg JM, Wilson WR. Infective endocarditis caused by HACEK microorganisms. *Annu. Rev. Med.* 1997;48:25-33
4. Shenker BJ. Immunologic dysfunction in the pathogenesis of periodontal diseases. *J Clin Periodontol* 1987;14:489-498.
5. Wilson M, Henderson B. Virulence factors of *Actinobacillus actinomycetemcomitans* relevant to the pathogenesis of inflammatory periodontal diseases. *FEMS Microbiol Rev* 1995;17:365-379.
6. Henderson B, Ward JM, Nair SP, Wilson M. Molecular pathogenicity of the oral opportunistic pathogen *Actinobacillus actinomycetemcomitans*. *Annu Rev Microbiol* 2003; 57:29-55.
7. Mayer MP, Bueno LC, Hansen EJ, DiRienzo JM. Identification of a cytolethal distending toxin gene locus and features of a virulence-associated region in *Actinobacillus actinomycetemcomitans*. *Infect Immun* 1999; 67:1227-1237.
8. Shenker BJ, McKay T, Datar S et al. *Actinobacillus actinomycetemcomitans* immunosuppressive protein is a member of the family of cytolethal distending toxins capable of causing a G2 arrest in human T cells. *J Immunol* 1999;162:4773-4780.
9. Sugai M, Kawamoto T, Peres SY et al. The cell cycle-specific growth-inhibitory factor produced by *Actinobacillus actinomycetemcomitans* is a cytolethal distending toxin. *Infect Immun* 1998;66:5008-5019.
10. Cope LD, Lumbly S, Latimer JL et al. A diffusible cytotoxin of *Haemophilus ducreyi*. *Proc Natl Acad Sci U S A* 1997; 94:4056-4061.
11. Okuda J, Fukumoto M, Takeda Y, Nishibuchi M. Examination of diarrheagenicity of cytolethal distending toxin: suckling mouse response to the products of the *cdtABC* genes of *Shigella dysenteriae*. *Infect Immun* 1997;65:428-433.
12. Pickett CL, Pesci EC, Cottle DL et al. Prevalence of cytolethal distending toxin production in *Campylobacter jejuni* and relatedness of *Campylobacter sp.* *cdtB* gene. *Infect Immun* 1996;64:2070-2078.
13. Scott DA, Kaper JB. Cloning and sequencing of the genes encoding *Escherichia coli* cytolethal distending toxin. *Infect Immun* 1994;62:244-251.
14. Young VB, Knox KA, Schauer DB. Cytolethal distending toxin sequence and activity in the enterohepatic pathogen *Helicobacter hepaticus*. *Infect Immun* 2000; 68:184-191.
15. Johnson WM, Lior H. Response of Chinese ovary hamster cells to a cytolethal distending toxin (CDT) of *Escherichia coli* and possible misinterpretation as heat-la-

- bile (LT) enterotoxin. FEMS Microbiol. Lett. 1987a;43:19-23.
16. Lara-Tejero M, Galan JE. Cytolethal distending toxin: limited damage as a strategy to modulate cellular functions. Trends Microbiol 2002;10:147-52.
 17. Dreyfus L, Burns D, Barbieri J et al. Cytolethal Distending Toxin. In Bacterial Protein Toxins. Washington, DC, USA: ASM Press; 2003:257-70.
 18. Cortes-Bratti X, Frisan T, Thelestam M. The cytolethal distending toxins induce DNA damage and cell cycle arrest. Toxicon 2001;39:1729-36.
 19. Shenker BJ, McKay T, Datar S et al. *Actinobacillus actinomycetemcomitans* immunosuppressive protein is a member of the family of cytolethal distending toxins capable of causing a G2 arrest in human T cells. J Immunol 1999;162:4773-80.
 20. Sugai M, Kawamoto T, Peres SY et al. The cell cycle-specific growth-inhibitory factor produced by *Actinobacillus actinomycetemcomitans* is a cytolethal distending toxin. Infect Immun 1998;66:5008-19.
 21. Mayer MP, Bueno LC, Hansen EJ, DiRienzo JM. Identification of a cytolethal distending toxin gene locus and features of a virulence-associated region in *Actinobacillus actinomycetemcomitans*. Infect Immun 1999; 67:1227-37.
 22. Yamano R, Ohara M, Nishikubo S et al. Prevalence of cytolethal distending toxin production in periodontopathogenic bacteria. J Clin Microbiol 2003;41:1391-8
 23. Ahmed HJ, Svensson LA, Cope LD et al. Prevalence of cdtABC genes encoding cytolethal distending toxin among *Haemophilus ducreyi* and *Actinobacillus actinomycetemcomitans* strains. J Med Microbiol 2001;50:860-864.
 24. Tan KS, Song KP, Ong G. Cytolethal distending toxin of *Actinobacillus actinomycetemcomitans*. Occurrence and association with periodontal disease. J Periodont Res 2002;37:268-272.
 25. Fabris AS, DiRienzo JM, Wikstrom M, Mayer MP. Detection of cytolethal distending toxin activity and cdt genes in *Actinobacillus actinomycetemcomitans* isolates from geographically diverse populations. Oral Microbiol Immunol 2002;17:231-238.
 26. Belibasakis G, Johansson A, Wang Y et al. Inhibited proliferation of human periodontal ligament cells and gingival fibroblasts by *Actinobacillus actinomycetemcomitans*: involvement of the cytolethal distending toxin. Eur J Oral Sci 2002;110:336-73.
 27. Hassane DC, Lee RB, Pickett CL. *Campylobacter jejuni* cytolethal distending toxin promotes DNA repair responses in normal human cells. Infect Immun 2003; 71:541-5.
 28. Lara-Tejero M, Galan JE. CdtA, CdtB, and CdtC form a tripartite complex that is required for cytolethal distending toxin activity. Infect. Immun. 2001;69:4358-4365
 29. Saiki K, Konishi K, Gomi T et al. Reconstitution and purification of cytolethal distending toxin of *Actinobacillus actinomycetemcomitans*. Microbiol. Immunol. 2001; 45:497-506
 30. Wising C, Svensson LA, Ahmed HJ et al. Toxicity and immunogenicity of purified *Haemophilus ducreyi* cytolethal distending toxin in a rabbit model. Microb. Pathog. 2002;33:49-62
 31. Shenker BJ, Besack D, McKay T et al. *Actinobacillus actinomycetemcomitans* cytolethal distending toxin (Cdt): evidence that the holotoxin is composed of three subunits: CdtA, CdtB, and CdtC. J. Immunol. 2004;172:410-417
 32. Avenaoud P, Castroviejo M, Claret S et al.

- Expression and activity of the cytolethal distending toxin of *Helicobacter hepaticus*. Biochem. Biophys. Res. Commun. 2004; 318:739-745
33. Lee RB, Hassane DC, Cottle DL, Pickett CL. Interactions of *Campylobacter jejuni* cytolethal distending toxin subunits CdtA and CdtC with HeLa cells. Infect. Immun. 2003;71:4883-4890.
 34. Mao X, DiRienzo JM. Functional studies of the recombinant subunits of a cytolethal distending holotoxin. Cell. Microbiol. 2002; 4:245-255.
 35. Svensson LA, Tarkowski A, Thelestam M, Lagergard T. The impact of *Haemophilus ducreyi* cytolethal distending toxin on cells involved in immune response. Microb Pathog 2001;30:157-66.
 36. DiRienzo JM, Song M, Wan LS, Ellen RP. Kinetics of KB and HEp-2 cell responses to an invasive, cytolethal distending toxin-producing strain of *Actinobacillus actinomycetemcomitans*. Oral Microbiol Immunol 2002;17:245-51.
 37. Hickey TE, McVeigh AL, Scott DA et al. *Campylobacter jejuni* cytolethal distending toxin mediates release of interleukin-8 from intestinal epithelial cells. Infect Immun 2000;68:6535-41.
 38. Cortes-Bratti X, Chaves-Olarte E, Lagergard T, Thelestam M. The cytolethal distending toxin from the chancroid bacterium *Haemophilus ducreyi* induces cell-cycle arrest in the G2 phase. J Clin Invest 1999;103:107-15
 39. Sato T, Koseki T, Yamato K et al. p53-independent expression of p21(CIP1/WAF1) in plasmacytic cells during G(2) cell cycle arrest induced by *Actinobacillus actinomycetemcomitans* cytolethal distending toxin. Infect Immun 2002;70:528-34.
 40. Nalbant A, Chen C, Wang Y, Zadeh HH. Induction of T-cell apoptosis by *Actinobacillus actinomycetemcomitans* mutants with deletion of ltxA and cdtABC genes: possible activity of GroELlike molecule. Oral Microbiol Immunol 2003; 18:339-49.
 41. Bartold PM, Walsh LJ, Narayanan AS. Molecular and cell biology of the gingiva. Periodontol 2000 2000;24:28-55.
 42. McCulloch CA, Lekic P, McKee MD. Role of physical forces in regulating the form and function of the periodontal ligament. Periodontol 2000 2000;24:56-72.
 43. McCulloch CA, Bordin S. Role of fibroblast subpopulations in periodontal physiology and pathology. J Periodontal Res 1991; 26:144-54.
 44. Ivanovski S, Li H, Haase HR, Bartold PM. Expression of bone associated macromolecules by gingival and periodontal ligament fibroblasts. J Periodontal Res 2001; 36:131-41.
 45. Christersson LA, Wikesjo UM, Albin B, Zambon JJ, Genco RJ. Tissue localization of *Actinobacillus actinomycetemcomitans* in human periodontitis. II. Correlation between immunofluorescence and culture techniques. J Periodontol 1987;58:540-5.
 46. Christersson LA, Albin B, Zambon JJ, Wikesjo UM, Genco RJ. Tissue localization of *Actinobacillus actinomycetemcomitans* in human periodontitis. I. Light, immunofluorescence and electron microscopic studies. J Periodontol 1987;58: 529-39.
 47. Belibasakis GN, Mattsson A, Wang Y, Chen C, Johansson A. Cell cycle arrest of human gingival fibroblasts and periodontal ligament cells by *Actinobacillus actinomycetemcomitans*: involvement of the cytolethal distending toxin. APMIS. 2004 Oct; 112(10):674-85.
 48. Kanno F, Korostoff J, Volgina A, DiRienzo

- JM. Resistance of human periodontal ligament fibroblasts to the cytolethal distending toxin of *Actinobacillus actinomycetemcomitans*. J Periodontol. 2005 Jul; 76(7):1189-201.
49. Belibasakis GN, Johansson A, Wang Y et al. The cytolethal distending toxin induces receptor activator of NF-kappaB ligand expression in human gingival fibroblasts and periodontal ligament cells. Infect Immun. 2005 Jan;73(1):342-51.
50. Nestic D, Hsu Y, Stebbins CE. Assembly and function of a bacterial genotoxin. Nature. 2004 May 27;429(6990):429-33.

*Actinobacillus actinomycetemcomitans*의 cytolethal distending toxin subunit CdtA 유전자 클로닝과 단백질 발현

고선영¹, 정동근¹, 유소현¹, 김형섭^{1,2}

1. 전북대학교 치과대학 치주과학교실
2. 전북대학교 구강생체과학연구소

Cytolethal distending toxin(CDT)은 세포 주기 중 G2에서 M 기로의 전환을 막아 세포의 증식을 억제할 수 있는 세균 단백 독소의 일종이다. 구강 미생물 중 유일하게 *Actinobacillus actinomycetemcomitans* (*A. actinomycetemcomitans*)만이 이 CDT를 생성 할 수 있는 것으로 알려져 있다. *A. actinomycetemcomitans*는 localized aggressive periodontitis (LAP)의 원인균으로 여겨지며 비 운동성의 그람 음성 구간균이고 37°C, 5% CO₂ 하에 성장이 왕성하다.

*A. actinomycetemcomitans*의 CDT는 3개의 인접한 유전자인 *cdtA*, *cdtB*, *cdtC*에 의해 형성되며 각각의 유전자에 대한 단백질의 기능은 아직 완전히 밝혀지지 않았다. 현재까지 연구에 의하면 *cdtA*는 CDT의 세포부착과 관련이 있는 것으로 여겨지며 이 유전자의 기능 이상 시 CDT의 독성 효과가 현저히 감소한다고 알려져 있다. 따라서 본 연구는 *A. actinomycetemcomitans*의 *cdtA* 유전자를 클로닝, 단백질 발현하여 향후 치주질환의 발병 과정에서 CdtA의 역할을 규명하고 질환의 예방 및 치료법에 도움을 주고자 하였다.

A. actinomycetemcomitans Y4 균주를 *cdtA* 유전자 클로닝을 위해 사용하였다. *A. actinomycetemcomitans*의 genomic DNA는 genomic DNA 추출 kit를 사용하여 분리하고 *cdtA*에 특이적인 primer를 이용하여 PCR을 통해 *cdtA* 유전자를 증폭하였다. 증폭된 *cdtA* 유전자를 T-vector에 클로닝 하였으며, 클로닝 된 *cdtA* 유전자는 단백질 발현을 위해 pRSET A vector에 서브클로닝 한 후 발현 균주인 BL21(DE3)를 이용하여 발현시켰다. 발현 후 Ni-NTA AP conjugate를 이용한 Western blot을 통해 pRSET-CDTA를 확인하였다.

