

# *Actinobacillus actinomycetemcomitans* Y4 capsular polysaccharide induces IL-1 $\beta$ mRNA expression through the JNK pathway in differentiated THP-1 cells

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## Summary

Capsular polysaccharide from *Actinobacillus actinomycetemcomitans* Y4 (Y4 CP) induces bone resorption in a mouse organ culture system and osteoclast formation in mouse bone marrow cultures, as reported in previous studies. We also found that Y4 CP inhibits the release of interleukin (IL)-6 and IL-8 from human gingival fibroblast (HGF). Thus Y4 CP induces various responses in localized tissue and leads to the secretion of several cytokines. However, the effects of Y4 CP on human monocytes/macrophages are still unclear. In this study, THP-1 cells, which are a human monocytic cell line, were stimulated with Y4 CP, and we measured gene expression in inflammatory cytokine and signal transduction pathways. IL-1 $\beta$  and tumour necrosis factor (TNF)- $\alpha$  mRNA were induced from Y4 CP-treated THP-1 cells. IL-1 $\beta$  mRNA expression was increased according to the dose of Y4 CP, and in a time-dependent manner. IL-1 $\beta$  mRNA expression induced by Y4 CP (100  $\mu$ g/ml) was approximately 7- to 10-fold greater than that in the control by real-time PCR analysis. Furthermore, neither PD98059, a specific inhibitor of extracellular signal-regulated kinase nor SB203580, a specific inhibitor of p38 kinase prevented the IL-1 $\beta$  expression induced by Y4 CP. However, JNK Inhibitor II, a specific inhibitor of c-Jun N-terminal kinase (JNK) prevented the IL-1 $\beta$  mRNA expression induced by Y4 CP in a concentration-dependent manner. These results indicate that Y4 CP-mediated JNK pathways play an important role in the regulation of IL-1 $\beta$  mRNA. Therefore, Y4 CP-transduced signals for IL-1 $\beta$  induction in the antibacterial action of macrophages may provide a therapeutic strategy for periodontitis.

**Keywords:** *Actinobacillus actinomycetemcomitans*, IL-1, macrophage, JNK

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## Introduction

Periodontitis is a chronic inflammatory disease characterized by gingival inflammation and alveolar bone resorption. Periodontitis is often caused by infections with Gram-negative bacteria including *Actinobacillus actinomycetemcomitans* [1,2] and *Porphyromonas gingivalis* [3,4].

*A. actinomycetemcomitans* is a gram-negative, capnophilic, fermentative coccobacillus that has been implicated in the aetiology and pathogenesis of several forms of periodontal disease [1]. Clinical, microbiological, and immunological studies have explored the correlation between *A. actinomycetemcomitans* and several types of periodontitis [5,6]. *A. actinomycetemcomitans* produces several tissue-damaging products such as leukotoxin [7,8], lipopolysaccharide (LPS),

capsular polysaccharide [9-11], alkaline and acid phosphatases, an epitheliotoxin, a fibroblast inhibitory factor, and a bone resorption-inducing toxin [5].

Amano *et al.* [12] extracted a serotype-specific capsular polysaccharide-like antigen from whole cells of *A. actinomycetemcomitans* Y4 (serotype b) by autoclaving, purified it by ion-exchange chromatography and gel filtration, and showed that it is a polymer that consists of a repeating disaccharide unit -3)-d-fucopyranosyl-(1,2)-l-rhamnopyranosyl-(1-. Previous studies have shown that *A. actinomycetemcomitans* Y4 capsular polysaccharide (Y4 CP) induces IL-1 from a mouse macrophage cell line [13], bone resorption in a mouse organ culture system and osteoclast formation in mouse bone marrow cultures [14,15], and inhibits the release of IL-6 and IL-8 from human gingival

fibroblasts [16]. On the other hand, the fact that LPS is a bacterial component of gram-negative bacteria was revealed in studies on the details of innate immune responses through gene expression and signal transduction pathways [17,18]. LPS induces mitogen-activated protein kinases (MAPKs), including extracellular signal-regulated kinase (ERK), c-jun NH2-terminal protein kinase (JNK), and p38 mitogen-activated protein kinase (p38). These play key roles in LPS-mediated signal transduction between extracellular membrane stimulation and the cytoplasmic response and nuclear activity in the activation of the gene [19,20].

However, there has still been no report on the effect of Y4 CP on human immunocytes. In this study, we found that Y4 CP affected the gene expression of inflammatory cytokine in macrophages, which play an important role in host defense and inflammation, and examined which signal transduction pathways are used in this gene expression.

## Materials and methods

### Cell culture protocol

THP-1 cells were differentiated to macrophage as follows. THP-1 cells (Dainippon Pharmaceutical Co., Ltd. Japan) were grown in RPMI 1640 supplemented with 10% FCS, 2 mM L-glutamine and  $2 \times 10^{-5}$  M 2-ME in 5% CO<sub>2</sub>-air humidified atmosphere at 37°C. THP-1 cells were treated with 50 nM 1,25-dihydroxy-vitamin D<sub>3</sub> (Calcitriol, Wako, Japan) for 72 h, washed three times with PBS and allowed to rest overnight in RPMI 1640 with 5% FCS [21].

### Microorganisms

*A. actinomycetemcomitans* Y4 (serotype b) were grown in Todd-Hewitt broth (Difco Laboratories, Detroit, MI, USA) supplemented with 1% (wt/vol) yeast extract at 37°C for 3 days in an atmosphere of 5% CO<sub>2</sub> [22]. The organisms were harvested by centrifugation, washed three times with distilled water, and lyophilized.

### Extraction of Y4 CP

The lyophilized cell suspension (300 mg/ml) in saline was autoclaved at 121°C for 15 min [23]. After being autoclaved, the suspension was cooled and centrifuged at  $10\,000 \times g$  for 20 min, and the supernatant was collected. Extraction was repeated on residual whole cells. The supernatants were combined, dialysed extensively with distilled water, and lyophilized.

### Purification of Y4 CP

Serotype antigens were purified according to the method of Amano *et al.* [12]. The autoclaved extracts of

*A. actinomycetemcomitans* Y4 were solubilized with 0.01 M Tris hydrochloride, pH 8.2, to give a final concentration of 100 mg (dry weight) of bacterial extract per ml and dialysed against the buffer at 4°C for 2 days. A 5 ml portion of the antigen suspension was applied to a column of DEAE-Sephadex A-25 (2 × 30 cm; Pharmacia Fine Chemicals, Piscataway, NJ, USA) that had been equilibrated with the buffer and eluted with 200 ml of the buffer followed by a linear gradient of 0 to 1 M NaCl in the buffer at 4°C. Fractions (10 ml each) were monitored for total sugar, protein, and phosphorus. Fractions that showed a positive reaction with anti-*A. actinomycetemcomitans* Y4 serum by immunodiffusion were combined and concentrated in a rotary evaporator. These preparations were dialysed with distilled water at 4°C for 3 days, applied to a column of Sephacryl S-300 (1.5 × 100 cm; Pharmacia), and eluted with distilled water. Fractions that contained the CP with serotype-specific antigens were pooled and lyophilized.

### RT-PCR assay and real-time PCR analysis

For RT-PCR, total cellular RNA was prepared using TRIzol reagent. cDNA was synthesized from total RNA by the extension of random primers with 200 U of Superscript II. PCR of the cDNA was performed using AccuPower PCR PreMix (BIONEER, Daejeon, Korea), which contains specific primers at 20 pmol. The following primers were showed at Table 1. The synthesized PCR products were separated by electrophoresis on 1.5% agarose gel and visualized by ethidium bromide staining. To quantify IL-1β mRNA, real-time PCR was performed using an ABI Prism 7000 (Applied Biosystems, Foster City, CA, USA) with TaqMan Universal PCR master mixture (Applied Biosystems). For PCR, 5 μl of sample was directly added to 45 μl of a RT-PCR mixture prepared from 2× RT-PCR TaqMan Universal PCR master mixture containing each primer at a concentration of 1 μM, 2 mM MgCl<sub>2</sub>, and 100 μM probe. The cycle parameters were as follows: 3 min at 95°C and 40 cycles of 1 min at 95°C, 1 min at 52°C, and 1 min 30 s at 72°C. Cycling was preceded by incubation for 10 min at 95°C to activate AmpliTaq Gold. For reverse transcription, all of these steps were preceded by 30 min of incubation at 48°C. Amplification mixtures were analysed using the ABI Prism detection system. Changes in gene expression were assessed using the comparative Ct method. (<http://docs.appliedbiosystems.com/pebi/docs/04303859.pdf>)

### Inhibitors

PD98059, a specific inhibitor of extracellular signal-regulated kinase (ERK), SB203580, a specific inhibitor of p38 kinase, and JNK inhibitor II, a specific inhibitor of c-Jun N-terminal kinase (JNK), were purchased from Calbiochem-Novabiochem (La Jolla, CA, USA).

**Table 1.** Primer sequences used for RT-PCR.

Human gene		Sequence	Product size
IL-1 $\beta$	sense	AAA CAG ATG AAG TGC TCC TTC AGG	390 bp
	antisense	TGG AGA ACA CCA CTT GTT GCT CCA	
IL-6	sense	GTG TTG CCT GCT GCC TTC CCT G	320 bp
	antisense	CTC TAG GTA TAC CTC AAA CTC CAA	
TNF- $\alpha$	sense	CAG AGG GAA GAG TTC CCC AG	324 bp
	antisense	TGG AGA ACA CCA CTT GTT GCT CCA	
IL-12p35	sense	CAC TCC AGA CCC AGG AAT GT	293 bp
	antisense	TAC TAA GGC ACA GGG CCATC	
IL-12p40	sense	AAG GAG GCG AGG TTC TAA GC	414 bp
	antisense	TGA TGA AGA AGC TGC TGG TG	
IL-18	sense	GCT TGA ATC TAA ATT ATTATC AGT C	334 bp
	antisense	CAA ATT GCA TCT TAT TAT CAT G	
GAPDH	sense	GTC TTC ACC ACC ATG GAG AAG GCT	393 bp
	antisense	CAT GCC AGT GAG CTT CCC GTT CA	

### Statistical analysis

Data were analysed using the SPSS II software package (SPSS Japan Inc., Tokyo, Japan). The experimental groups were compared by one-way ANOVA with the Tukey HSD-test.

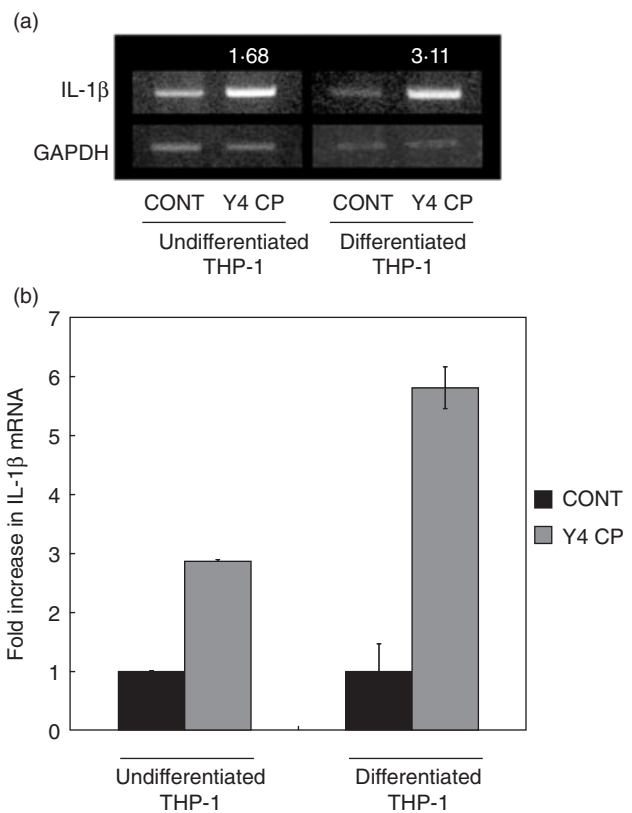
### Results

#### Y4 CP-induced IL-1 $\beta$ mRNA expression in undifferentiated and differentiated THP-1 cells

The reactivity of Y4 CP in undifferentiated and differentiated THP-1 cells was evaluated by the induction of IL-1 $\beta$  mRNA which is secreted from monocyte/macrophage. Both cell groups were treated with or without Y4 CP (100  $\mu$ g/ml) for 4 h. After treatment, total RNA was harvested, and IL-1 $\beta$  mRNA expression was analysed by RT-PCR. The densities of PCR products were expressed numerically using NIH image (N.I.H. USA). The densities of undifferentiated and differentiated THP-1 cells were standardized by GAPDH of each group. The ratio of undifferentiated or differentiated *versus* control was calculated by the density of each. The ratios of undifferentiated and differentiated THP-1 cells were 1.68 and 3.11, respectively (Fig. 1a). Furthermore, quantitative analysis was performed by real-time PCR. While IL-1 $\beta$  mRNA expression of undifferentiated THP-1 cells was approximately 3-fold that in the control, IL-1 $\beta$  mRNA expression in differentiated THP-1 cells was approximately 6-fold that in the control upon Y4 CP stimulation (Fig. 1b).

#### Inflammatory cytokine mRNA expression compared to that of IL-1 $\beta$

Monocyte and macrophage are well known to secrete not only IL-1 $\beta$  but also other inflammatory cytokines. Therefore, mRNA expression of TNF- $\alpha$ , IL-6, IL-12 and IL-18 from differentiated THP-1 cells that had been stimulated



**Fig. 1.** Y4 CP-induced IL-1 $\beta$  mRNA expression in undifferentiated and differentiated THP-1 cells. To confirm the reactivity of Y4 CP, undifferentiated and differentiated THP-1 cells were examined with regard to IL-1 $\beta$  mRNA expression. Both cell groups were cultured with Y4 CP (100  $\mu$ g/ml) or medium alone (control) for 4 h. After treatment, total RNA was prepared. IL-1 $\beta$  mRNA expression was analysed by RT-PCR. The densities of PCR products were expressed numerically using NIH image (N.I.H., USA). The densities of undifferentiated and differentiated THP-1 cells were standardized by GAPDH of each group. (a) The ratio of undifferentiated or differentiated *versus* control was calculated by the density of each. (b) IL-1 $\beta$  mRNA expression was analysed by real-time PCR.

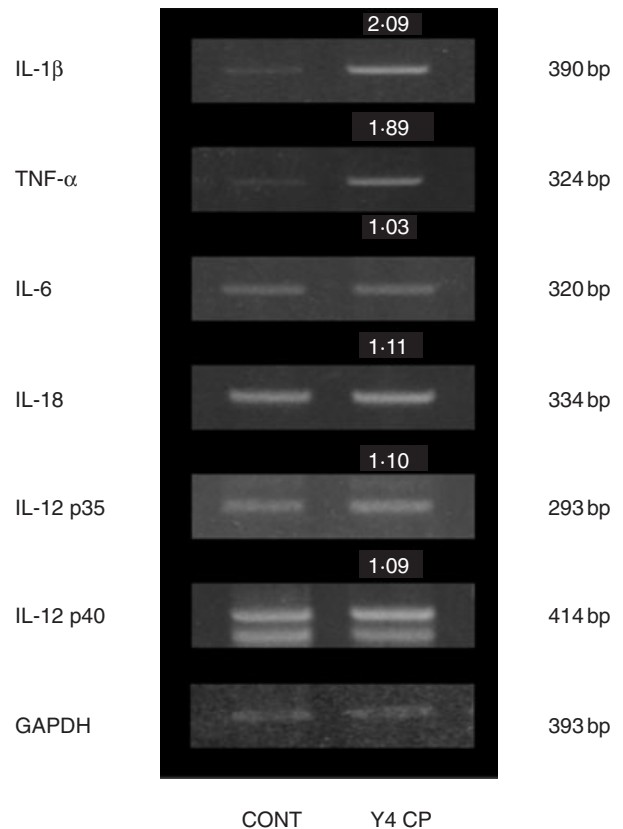
with Y4 CP was examined by RT-PCR. As shown in Fig. 2, the densities of PCR products were calculated using the same formula as in Fig. 1: the ratios of IL-1 $\beta$ , TNF- $\alpha$ , IL-6, IL-12 p35, IL-12 p40 and IL-18 were 2.09, 1.89, 1.03, 1.10, 1.09, and 1.11, respectively. IL-1 $\beta$  and TNF- $\alpha$  mRNA are both strongly expressed by Y4 CP.

**Quantitative analysis of IL-1 $\beta$  mRNA expression in differentiated THP-1 cells after Y4 CP stimulation**

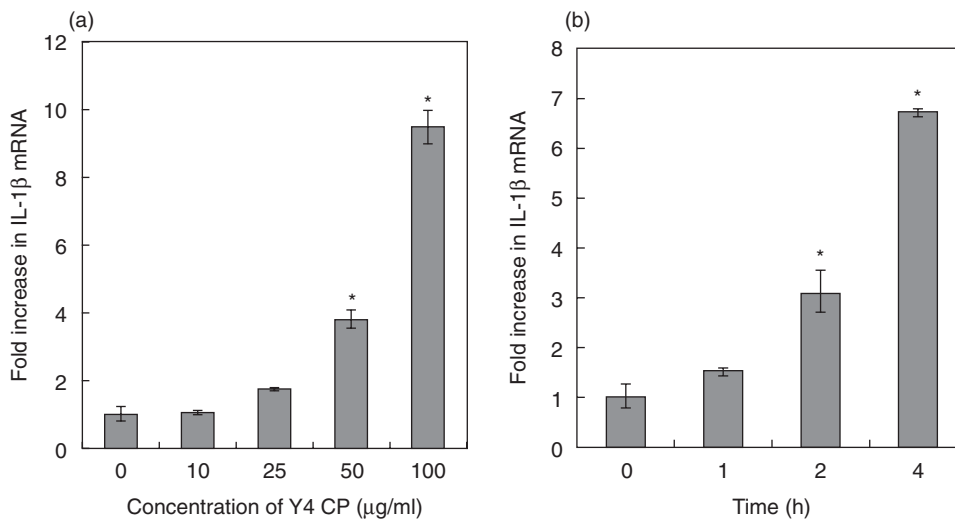
The optimal dose and duration of culture are very important for measuring mRNA expression. We treated differentiated THP-1 cells with various concentrations (0, 10, 25, 50, 100  $\mu$ g/ml) of Y4 CP for various durations (0, 1, 2, 4 h). After stimulation, the expression of IL-1 $\beta$  mRNA in differentiated THP-1 cells was evaluated by real-time PCR. Treatment of differentiated THP-1 cells with Y4 CP caused an increase in the expression of IL-1 $\beta$  mRNA in a dose and time-dependent manner (Fig. 3a). When differentiated THP-1 cells were stimulated with 100  $\mu$ g/ml Y4 CP for 4 h, Y4 CP induced approximately 7- to 10-fold greater IL-1 $\beta$  mRNA expression than that in the control.

**Signal pathways in IL-1 $\beta$  mRNA expression**

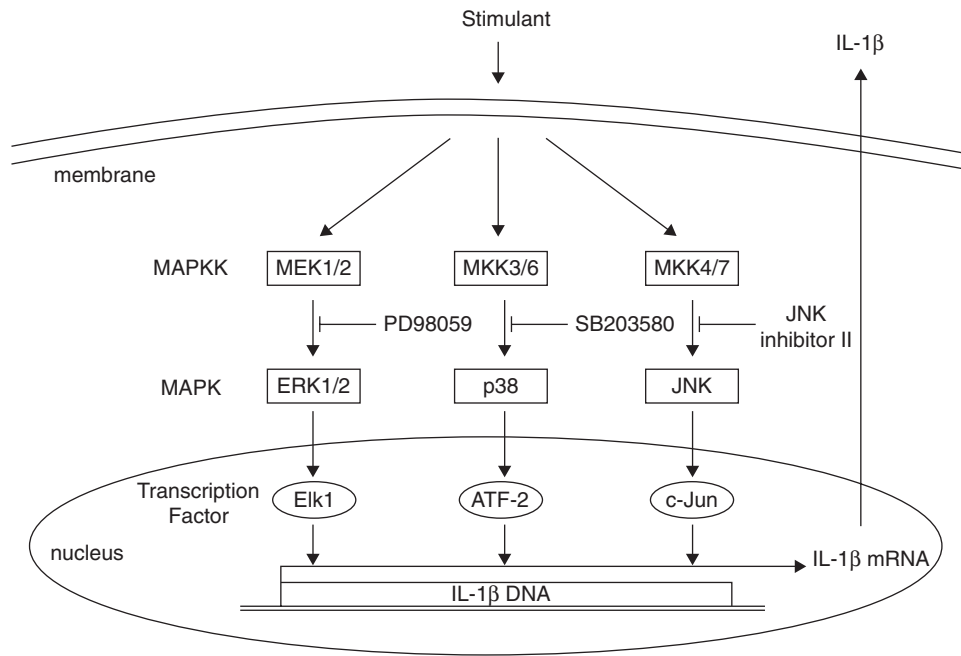
Various members of the MAPK family may modulate the expression of IL-1 $\beta$  in stimulated monocytes/macrophages. To investigate which MAPK pathway is involved in the expression of IL-1 $\beta$  mRNA when differentiated THP-1 cells are stimulated with Y4 CP, we used inhibitors of several MAPKs (Fig. 4). Differentiated THP-1 cells were treated with PD98059 (1–10  $\mu$ M), SB203580 (1–10  $\mu$ M), JNK Inhibitor II



**Fig. 2.** Expression of cytokine mRNAs in differentiated THP-1 cells stimulated with Y4 CP. Differentiated THP-1 cells were treated with Y4 CP (100  $\mu$ g/ml) or medium alone (control) for 4 h. After treatment, total RNA was prepared and inflammatory cytokine mRNA expression was examined by RT-PCR. The densities of PCR products were calculated using the same formula as in Fig. 1.



**Fig. 3.** (a) Effect of Y4 CP on IL-1 $\beta$  mRNA expression in differentiated THP-1 cells. Differentiated THP-1 cells were cultured in the presence of different concentrations of Y4 CP (10–100  $\mu$ g/ml) or without Y4 CP as a control. (b) Effect of the duration of culture (1–4 h) with Y4 CP (100  $\mu$ g/ml) on IL-1 $\beta$  mRNA expression in differentiated THP-1 cells. IL-1 $\beta$  mRNA levels were determined by real-time PCR. The results are expressed as ratios of the levels in the Control. Values shown are means  $\pm$  standard deviations of triplicate assays. \* $P$  < 0.01 versus control.



**Fig. 4.** Proposed model for signal transduction pathways of MAP kinase in the regulation of IL-1 $\beta$  expression.

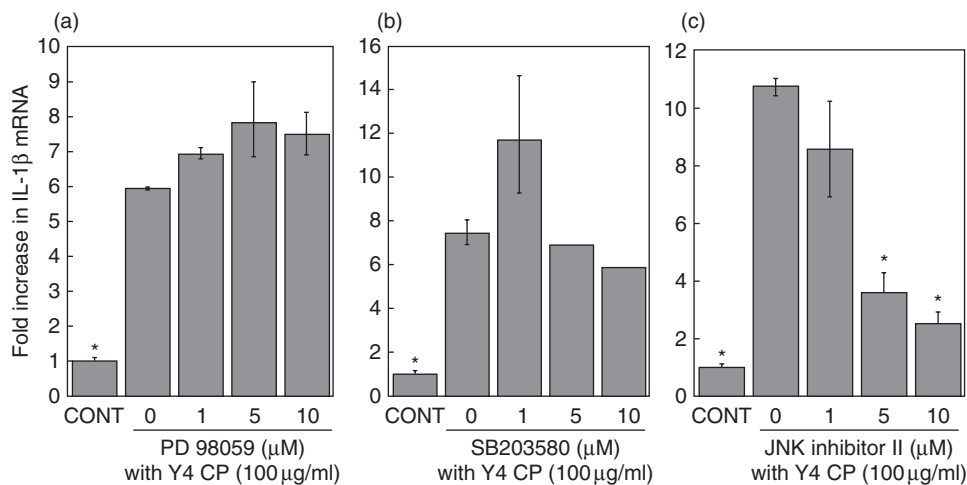
(1–10  $\mu$ M), or vehicle (DMSO) for 30 min, respectively, and then stimulated with Y4 CP (100  $\mu$ g/ml) for 4 h.

Pretreatment of THP-1 with SB203580 led to a slight decrease in the expression of IL-1 $\beta$  mRNA. However, significant differences were not observed upon pretreatment with SB203580 or PD98059. On the other hand, JNK Inhibitor II prevented the up-regulation of IL-1 $\beta$  mRNA expression in Y4 CP-stimulated THP-1; indeed, the expression of IL-1 $\beta$  mRNA was inhibited by 76% with 10  $\mu$ M JNK Inhibitor II.

These findings suggest that the JNK pathway is probably involved in mediating the response to Y4 CP (Fig. 5).

### Discussion

Periodontitis is initiated by oral microbacteria such as *P. gingivalis* or *A. actinomycetemcomitans* that induce an inflammatory cascade, which stimulates host-mediated tissue destruction. Recent advances in the understanding of



**Fig. 5.** Effects of protein kinase inhibitors, i.e. inhibitors of ERK (PD98059)(a), p38 (SB203580)(b) and JNK (SP600125)(c), on Y4 CP-induced IL-1 $\beta$  mRNA expression. Differentiated THP-1 cells were pretreated with PD98059, SB203580, or JNK Inhibitor II (1, 5, 10  $\mu$ M) for 30 min. Cells were then cultured with Y4 CP (100  $\mu$ g/ml) for an additional 4 h. Values shown are means  $\pm$  standard deviations of triplicate assays. \* $P$  < 0.01 versus Y4 CP stimulation without inhibitor.



inflammation have provided insight into the mechanisms involved in periodontal tissue destruction. Several mediators appear to be involved, including a variety of cytokines produced by several different cell types. The primary mediators, such as IL-1 and TNF, have been shown to contribute to several events that are essential for the initiation of an inflammatory response and, ultimately, tissue destruction [24–26]. They can induce the up-regulation of adhesion molecules on leucocytes and endothelial cells and stimulate the production of chemokines, which are needed to recruit circulating leucocytes. IL-1 also induces the expression of other mediators that amplify or sustain the inflammatory response, such as prostaglandins, and the production of lytic enzymes, such as matrix metalloproteinases; they also can enhance bacterial killing and phagocytic activity [27]. Furthermore, IL-1 is synergistic in its capacity to enhance bone resorption [28]. Although the periodontium has a high capacity for repair following injury, in some situations cytokines may limit repair by inducing apoptosis of matrix-producing cells [29]. Moreover, many studies have reported that gingival crevicular fluid (GCF) IL-1 levels are significantly elevated in all forms of periodontitis, compared to health or gingivitis [30–39]. Ishihara *et al.* [30] reported that the degree of periodontitis, classified according to alveolar bone resorption, was correlated with the total amounts of IL-1 $\alpha$  and IL-1 $\beta$  in GCF and the level of an IL-1 activity index. Thus, IL-1 is a significant and integral component of the host response to periodontal infection.

*A. actinomycetemcomitans* is a major pathogenic bacterium that is responsible for aggressive periodontitis (localized juvenile periodontitis). *A. actinomycetemcomitans* LPS has been shown to play a role in cellular and humoral immunity and inflammatory bone resorption *in vitro*. For example, LPS from *A. actinomycetemcomitans* induced IL-1 and prostaglandin E<sub>2</sub> production from calvarial organ cultures, and IL-1 is responsible for the induction of bone resorption [40] and osteoclast formation in mouse bone marrow cultures [41]. LPS from *A. actinomycetemcomitans* also induces IL-6, which is related to the proliferation and maturation of plasma cells, and is produced by monocytes [42] and human gingival fibroblasts [43]. However, the biological activities of *A. actinomycetemcomitans* CP are completely different from those of *A. actinomycetemcomitans* LPS. For instance, while *A. actinomycetemcomitans* LPS induces IL-6 production, its CP reduces the production of IL-6 by gingival fibroblasts. Therefore, it seems there are different signalling pathways among *A. actinomycetemcomitans* LPS and CP in immunocytes.

Y4 CP induced IL-1 $\beta$  mRNA expression in both undifferentiated and differentiated THP-1 cells. Differentiated THP-1 cells showed significantly increased IL-1 $\beta$  mRNA expression compared to undifferentiated THP-1 cells (Fig. 1). The cell markers CD14 and CD11a, which are involved in cell signalling in response to a range of bacterial pathogen-associated molecular patterns, are both increased upon

treatment of these cells with vitamin D<sub>3</sub> [44]. In our experiments, increased IL-1 $\beta$  expression might help explain the enhanced sensitivity of cells to Y4 CP. We previously reported that Y4 CP could increase osteoclast formation in mouse bone marrow culture systems, and concluded that IL-1 secreted by bone marrow cells after Y4 CP stimulation might induce osteoclast formation. This previous study supports the notion that Y4 CP affects immunocytes such as monocyte/macrophage in bone marrow cells. However, IL-1 $\beta$  production was not able to be detected though mRNA expression was observed in this experiment. We speculate that Y4 CP might induce the IL-1 $\beta$  mRNA expression only and other Y4 component such as leukotoxin will activate the caspase-1 activity, and it will help to produce the mature IL-1 production [45].

A recent study has shown that the induction of the IL-1 $\beta$  gene in mouse calvarial bone cells stimulated with *P. gingivalis* fimbria is regulated by transcriptional factor activation protein-1 [46]. In contrast, in *A. actinomycetemcomitans*, IL-1 $\beta$  expression was only measured upon LPS stimulation [47,48], and not with other *A. actinomycetemcomitans* components. Our previous studies have only considered cytokine production *in vitro*. Therefore, there is no evidence concerning signalling molecules. The MAPK pathway is one of the major modulators of cytokine mRNA expression; consequently MAPK pathways were examined under our experimental conditions. JNK inhibitor II, a specific inhibitor of JNK, significantly and additively suppressed IL-1 $\beta$  mRNA expression along with Y4 CP. On the other hand, PD98059 and SB203580, specific inhibitors of ERK1/2 and p38 kinase, had no effect on IL-1 $\beta$  expression by Y4 CP. These results suggest that at least a JNK pathway is essential for IL-1 $\beta$  expression after the stimulation of THP-1 cells with Y4 CP.

Many reports deal with IL-1 $\beta$  expression induced by LPS in monocyte/macrophage and related signalling pathways. LPS induces MAPKs, including ERK, JNK and p38. These molecules play key roles in LPS-mediated signal transduction between extracellular membrane stimulation and the cytoplasmic response and nuclear activity in the activation of the gene [19,20]. Specifically, ERK activation involves cytokine induction and regulation during responses to bacterial products [49–51]. In addition to responding to numerous physiological and stress stimuli [52–55], JNK is considered to play roles in regulating the expression of various stress-induced proteins and inflammatory cytokines [55,56]. p38 is activated in response to stress signals such as LPS, osmotic stress, and pro-inflammatory cytokines [51,57–59]. Previous studies have shown that the p38 pathway plays a critical role in LPS-stimulated cytokine release [49,59], including IL-1 and TNF induction in monocytes [20]. In this study, the JNK pathway was shown to be important not only for LPS- but also for Y4 CP-stimulated IL-1 $\beta$  expression. Many of the downstream targets of the JNK pathway are transcription factors, including c-Jun, ATF-2, and Elk-1 [60]. These transcription factors regulate various genes that encode

inflammatory mediators. In addition, a JNK pathway inhibitor also blocked gene transcription and reduced protein production. For example, the immunosuppressant dexamethasone reduced the LPS-induced translation of TNF- $\alpha$  mRNA by selectively inhibiting the JNK pathway [61]. The JNK-to-c-jun pathway is important for cell apoptosis and the cell cycle, and Y4 CP also induced osteoblast apoptosis [62]. Therefore, further studies are warranted to examine Y4 CP-induced apoptosis in macrophage through a JNK pathway. Recently, it is found that TLRs are important for the recognition of various bacterial components. Previous our group indicated that Y4 CP induces to make osteoclast formation in C3H/HeJ mouse bone marrow cells [15]. It is thought that the recognition of Y4 CP does not need TLR4. However, the rest of TLRs necessity for Y4 CP recognition of THP-1 cells is still unknown. Therefore, TLRs and Y4 CP relationship will discover for future project.

In conclusion, the present results showed that *A. actinomycetemcomitans* Y4 CP induces IL-1 $\beta$  gene expression in macrophage. Our results also suggest that the bioactivity of Y4 CP is at least mediated by the activation of JNK. The current data provide new insight into the induction of immune responses by Y4 CP, and we believe that these findings may be useful for further research into infectious diseases such as periodontitis.

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