

Activation of JNK by TPA promotes apoptosis via PKC pathway in gastric cancer cells

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Abstract

AIM: JNK cascade plays an important role in cell proliferation, differentiation and apoptosis. However, the exact function of JNK cascade for apoptosis induction remains largely unknown. In this study, the role of JNK activation stimulated by TPA in the process of apoptosis induction and its signaling transduction pathway in gastric cancer cells were investigated and determined.

METHODS: Expressions of mRNA and protein were detected by Northern blot and Western blot. Transcription activity was measured by transient transfection and CAT assay. Apoptotic cells were displayed through staining the nucleus with DAPI and were observed under fluorescence microscope. The apoptotic index was determined by counting 1000 cells randomly.

RESULTS: JNK protein was stimulated rapidly by TPA, and reached its highest peak within 3 hr, then decreased in a time-dependent manner, but the expression level of JNK protein induced by TPA was always keeping higher than that in untreated cells. Similar pattern was seen in *c-jun* mRNA level induced by TPA. TPA significantly activated the transcriptional activity of activator protein-1 with a TPA-dose-dependent manner. Furthermore, activation of JNK was mediated through PKC pathway. Treatment of cells with PKC specific inhibitor, Wortmannin, led to repression of JNK even in the presence of TPA. More importantly, all these effects were associated with induction of apoptosis in gastric cancer cells. TPA induced apoptosis obviously in gastric cancer cells. The apoptotic cells became smaller and rounded, and their nuclei became condensation and fragmentation with brightly stained chromatin. However, suppression of JNK by PKC specific inhibitor, Wortmannin, resulted in the decrease of apoptosis induced by TPA in a time-dependent manner, apoptotic index dramatically decreased from 32.56 % to 8.71 %.

CONCLUSION: TPA stimulates JNK cascade, including up-regulation of JNK protein expression level and *c-jun* mRNA expression level, and activation of activator protein-1

transcriptional activity. Activation of JNK is mediated through PKC pathway, which has an association with induction of apoptosis by TPA. Thus, activation of JNK via PKC pathway may represent one of important mechanisms for TPA to induce apoptosis in gastric cancer cells.

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INTRODUCTION

Mitogen-activated protein kinase (MAPK) pathway is an important signal transduction pathway that transduces extracellular signals in intracellular responses and has been implicated in a wide array of physiological processes including cell growth, differentiation, and apoptosis^[1-6]. The extracellular signal-regulated kinase (ERK), the *c-jun* N-terminal kinase (JNK) and the p38 MAPK are the members of MAPK family^[7-9]. Among them, JNK is one of the key mediators of stress signal and inflammatory response evoked by a variety of agents such as UV-irradiation, γ -irradiation, heat shock, and inflammatory cytokine^[10,11]. Once activated by specific kinases, JNK mediates the phosphorylation and activation of the transcription factors, such as *c-Jun*, ATF2, and E1K1^[12,13].

JNK can be strongly activated through stress signal that ultimately leads to apoptosis. Accumulating reports have recently been focused on the potential role of JNK in apoptotic signaling. In PC12 cells, JNK plays a critical role in mediating apoptosis caused by nerve growth factor withdrawal^[14]. The JNK pathway is also required for the induction of apoptosis by ceramide, γ - and UV-irradiation, some chemotherapeutic drugs^[15-17]. Inversely, other reports have shown evidences that activation of JNK is not mechanistically implicated in the apoptotic process. activation of JNK does not correlate with apoptosis induced by the detachment of epithelial cells^[18]. Inhibition of JNK activity by the expression of *c-jun* mutant does not prevent TNF-mediated cell death^[19,20].

The present study investigates the effects of 12-O-tetradecanoyl-Phorbol-13-acetate (TPA) on activation of JNK and its role in apoptosis induction in gastric cancer cells. Our results indicated that TPA activated the expressions of JNK protein, which led to induction of *c-jun* mRNA expression, and was associated with activation of activator protein-1 transcriptional activity. Furthermore, activation of JNK cascade was mediated through protein kinase C (PKC) pathway. All these effects had an association with induction of apoptosis by TPA. Thus, activation of JNK via PKC pathway might represent one of mechanisms for TPA to induce apoptosis in gastric cancer cells.

MATERIALS AND METHODS

Cell line and culture condition

Human gastric cancer cell line, MGC80-3 was established by

Cancer Research Center, Xiamen University^[21]. Cells were maintained in RPMI-1640 medium, supplemented with 10 % FCS, 1 m mol/L glutamine, and 100 u/ml penicillin. The concentration of TPA was 100 ng/ml.

Western blot analysis^[22]

Cells treated by TPA were harvested, and suspended in RIPA buffer (10 m mol/L Tris (pH7.4), 150 m mol/L NaCl, 1 % Triton X-100, 1 % Deoxycholic Acid, 0.1 % SDS, 5 m mol/L EDTA (pH8.0), 1 m mol/L PMSF). Protein concentration was determined by using the Bio-Rad protein assay system according to the manufacturer's direction (Bio-Rad Hercules, CA). Total protein (50 μ g) was subjected to SDS-PAGE and transferred to nitrocellulose membrane for western blot analysis. The membrane was subsequently blocked with 5 % dry milk in TBS-T and then immunoblotted with the corresponding antibody. Protein was detected by using the ECL kit (Amersham Inc.) according to the manufacturer's direction. Before incubation with antibody, the same membrane was stained by ponceau S. (Sigma) to show the amount of protein used in each well. The intensities of bands indicated by Western blot were quantified by densitometer.

RNA preparation and northern blotting

Extraction of total RNA was performed by the guanidine hydrochloride/ultracentrifugation method. Total RNA (20 μ g) was separated on 1 % agarose gel containing formaldehyde, and then transferred to nylon membrane, hybridized with α -³²p-dATP and α -³²p-dCTP labeled *c-jun* probe (Zhong Shang Biotec. CO., China), washed, and autoradiographed as described previously^[23]. The same membrane was probed again with β -actin cDNA to show the amount of RNA used in each well. The intensities of bands indicated by Northern blot were quantified by densitometer.

Transient transfection and CAT assay^[24]

Cells were seeded in a six-well plate and were approximately 70 % confluent at the time of transfection. Cells were transfected by CeLLFECTINtm (Gibco/BRI Life Technologies) by the following procedure: 6 μ l CeLLFECTINtm in 1.0 ml standard medium was added to each well containing 1.0 ml of standard medium supplemented with the -73Col-CAT reporter plasmid (100 ng), β -galactosidase expression vector (400 ng, pCH110, pharmacia). CAT activity was determined using ³H-acetyl-CoA as substrate. β -galactosidase activity was measured to normalize transfection efficiency.

Apoptosis analysis^[25]

Cells were treated with TPA, trypsinized, washed with PBS. The harvested cells were fixed in 3.7 % paraformaldehyde on ice, washed with PBS, then stained with 50 μ g/mL of 4,6-diamidino-2-phenylindole (DAPI, Sigma) containing 100 μ g/mL of DNase-free RNase A to facilitate the examination of nuclei under fluorescence microscope. Apoptotic cells were investigated and counted among 1000 cells randomly. The apoptotic index was a mean of three independent experiments.

RESULTS

Stimulation of JNK expression by TPA

To determine whether TPA activates the JNK expression in gastric cancer cells, MGC80-3 cells were treated with TPA in a time course, and the expression level of JNK protein was detected by western blot analysis. As shown in Figure 1, JNK expressed in MGC80-3 cells, after cells were treated with TPA,

the JNK expression was stimulated markedly. It was noted simultaneously that JNK expression was stimulated rapidly by TPA and reached a highest peak within 3 hr. After 3 hr treatment of TPA, although TPA-stimulated JNK expression decreased in a time course manner, the level of JNK protein was always keeping higher than that in MGC80-3 cells (Figure 1). Thus, the result clearly indicates the activation of JNK by TPA.

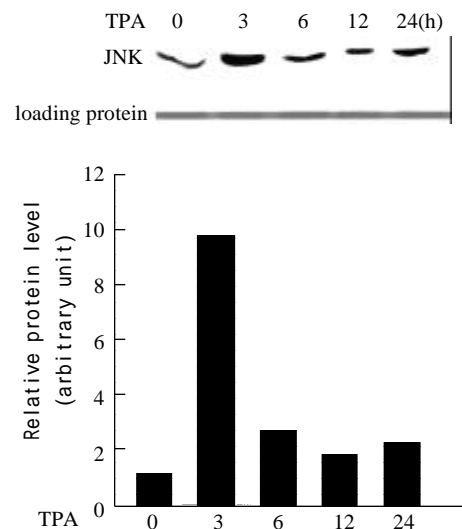


Figure 1 Expression of JNK protein in response to TPA induction. Cells treated with TPA for different time indicated. Expression of JNK protein was detected by Western blot. Amount of protein used in each lane was indicated by staining with ponceau S. The intensity of each band was quantified by densitometer.

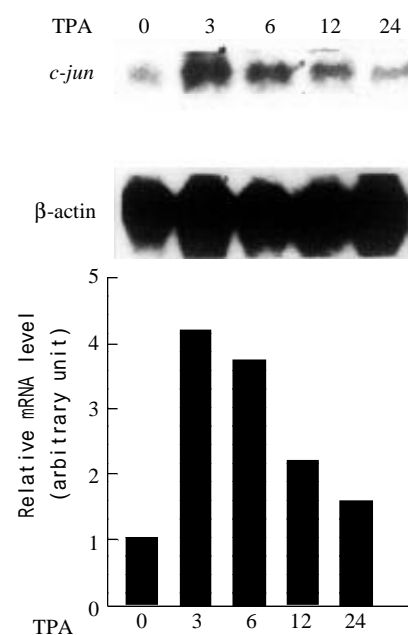


Figure 2 Expression of *c-jun* mRNA in response to TPA induction. Cells treated with TPA for different time indicated. Expression of *c-jun* mRNA was revealed by Northern blot. The same membrane was probed again with β -actin cDNA to show the amount of RNA used in each well. The intensity of each band was quantified by densitometer.

Effect of TPA on *c-jun* mRNA expression

Since activation of the JNK cascade is known to induce *c-jun* expression in several cell systems^[26,27], we examine whether *c-jun* expression in mRNA level is also induced by TPA in gastric cancer cells. Northern blot analysis, using *c-jun* as a probe, revealed that MGC80-3 cells expressed *c-jun* mRNA at lower level, after treatment of TPA, expression level of *c-jun* mRNA was upregulated in a time-dependent manner (Figure 2). More interestingly, the induction patterns between JNK and *c-jun* by TPA were very similar. *c-jun* mRNA also reached its maximal peak induced by TPA within 3 hr, followed by a decrease with extension of TPA treatment in a time-course. Taken together, the data with Figure 1 demonstrate that JNK cascade indeed plays a role in TPA signaling in gastric cancer cells.

Transcriptional activity of activator protein-1 in response to TPA

Activator protein-1 is composed of products of the proto-oncogenes *c-jun* and *c-fos*^[28]. The result mentioned above implies that TPA may affect transcriptional activity of activator protein-1 due to the activation of *c-jun*. To further determine whether TPA activates the transcriptional activity of activator protein-1, we measured transcriptional activity of activator protein-1 in MGC80-3 cells treated with various concentrations of TPA by the methods of transient transfection and CAT assay. When the reporter -73Col-CAT expression vector that contains an AP-1 binding site^[29] was transfected into MGC80-3 cells, a strong induction of reporter transcriptional activity was observed when cells were treated by TPA. This induction of reporter transcriptional activity was TPA-dose-dependent obviously (Figure 3). Thereby, TPA had a capability of activating the transcriptional activity of activator protein-1.

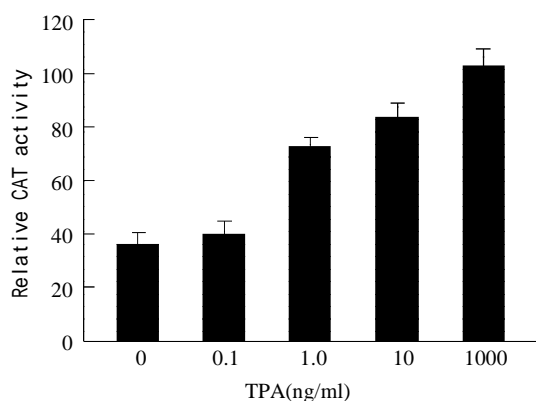


Figure 3 Effect of TPA on transcriptional activity of activator protein-1. Cells were treated with various concentrations of TPA indicated. Various expression vectors were transfected into cells as described in materials and methods. Transcriptional activity was measured by the method of transient transfection and CAT assay. Data shown represent mean of duplicate experiments (\pm SE).

TPA activates JNK cascade through PKC pathway

Since the target of TPA is protein kinase C (PKC)^[30,31], we examine whether stimulation of JNK cascade by TPA is through PKC pathway in gastric cancer cells. When MGC80-3 cells were pretreated with PKC specific inhibitor, Wortmannin^[32,33] for 2 hr, and then treated with TPA for another 3 hr, there was a significant inhibition observed in TPA-stimulated JNK protein expression (Figure 4, lane 4). As controls, JNK expression was clearly detected in MGC80-3 cells (Figure 4, lane 1) and TPA-

stimulated JNK expression was seen obviously in the TPA-treated cells (Figure 4, lane 3), and Wortmannin alone slightly repressed JNK expression (Figure 4, lane 2), respectively. The similar result was observed in MGC80-3 cells treated with another PKC inhibitor, PKC inhibitor peptide^[34] (data not shown). These results suggest that activation of JNK by TPA is mediated through PKC pathway.

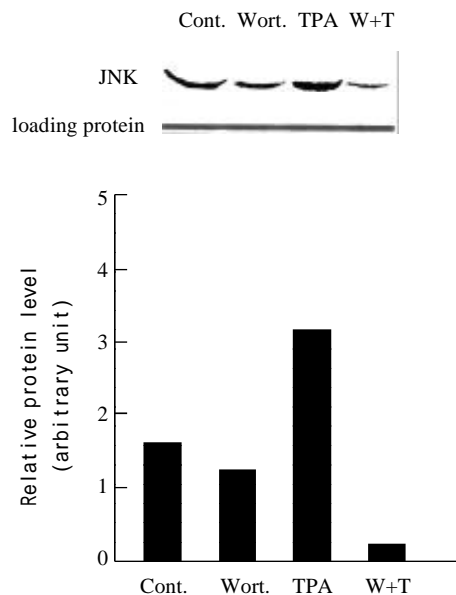


Figure 4 Effect of various agents, including Wortmannin (shown by Wort. or W), a PKC specific inhibitor, on expression of JNK protein. Cells were treated with Wortmannin for 2 hr, following by TPA induction for another 3 hr. Protein was detected by Western blot. Amount of protein used in each lane was indicated by staining with ponceau S. The intensity of each band was quantified by densitometer.

TPA-induced apoptosis is correlated with JNK expression through PKC pathway

To determine the effects of TPA on JNK and PKC functions, and their relationship with induction of apoptosis in gastric cancer cells, we analyzed apoptotic rate in MGC80-3 cells treated with relevant agents. Compared to the untreated MGC80-3 cells, treatment of cells with TPA caused the classical morphological characteristics of apoptosis, cells became smaller and rounded, and their nuclei became condensation and fragmentation with brightly stained chromatin (Figure 5A). However, when treatment of cells with PKC specific inhibitor, Wortmannin, subsequently with TPA, the apoptotic cells were decreased obviously, cells displayed a normal nuclear morphology similar to that of untreated cells (Figure 5A). In addition, the number of apoptotic cells was increased by the induction of TPA in a time-dependent manner. The highest apoptotic rate reached 32.56 % after treating cells with TPA for 48 hr, but only 5.46 % in untreated MGC80-3 cells (Figure 5B).

Since TPA-activated JNK cascade was mediated through PKC pathway in MGC80-3 cells (Figure 4), we analyzed whether this pathway was an association with apoptosis induced by TPA. Compared to TPA that induced apoptosis obviously, pretreated cells with PKC specific inhibitor, Wortmannin, significantly inhibited apoptosis induced by TPA, apoptotic rate was rather low, the highest apoptotic rate was only 8.71 %, although those cells were treated subsequently by TPA for another 48 h (Figure 5B). Accordingly, all these data further suggest that induction

of apoptosis by TPA is through PKC pathway, which may associate with the activation of JNK cascade.

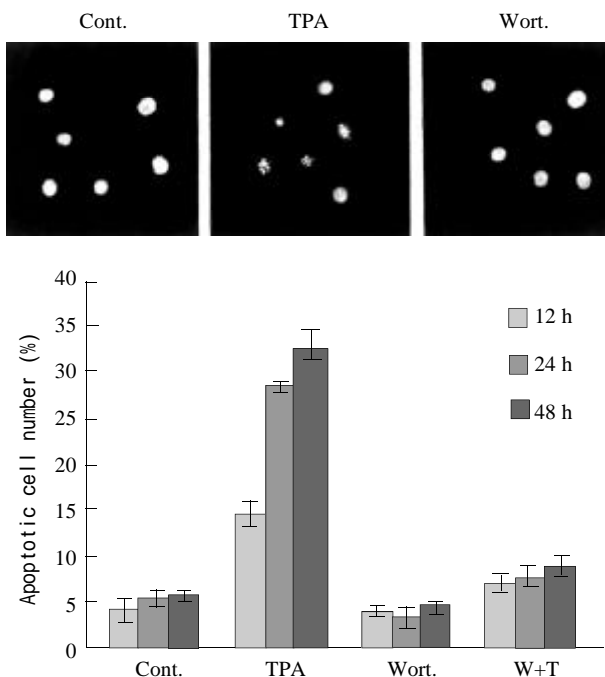


Figure 5 Effect of various agents, including TPA and Wortmannin, on apoptosis induction. Cells were treated with TPA and/or Wortmannin for different time indicated, and then the apoptotic rate was analyzed as described in material and methods. (A) The morphological changes in nucleus in response to different agents. (B) Apoptosis index.

DISCUSSION

The dynamic balance among the extracellular signal-regulated kinase (ERK), the *c-jun* N-terminal kinase (JNK) and the p38 Mitogen-activated protein kinase (p38 MAPK) activities critically determines the cellular fate in response to stimuli of proliferation, differentiation and apoptosis^[35,36]. The role of ERK signaling in the processes of cell differentiation and proliferation has been clearly recognized^[37,38], while requirement of JNK and p38 MAPK activations for apoptosis remains controversy. In some cases, JNK and p38 MAPK activity is strictly required for apoptosis^[39-41]. However, in other circumstances, they are not associated with the induction of apoptosis^[42]. In this study, we demonstrate that TPA stimulates JNK cascade through PKC pathway, which is associated with induction of apoptosis in gastric cancer cells.

JNK cascade plays an important role in cell proliferation, differentiation and apoptosis^[43,44]. In HeLa cells, the activation of JNK was very rapid and sustained for at least 24 hr post-irradiation^[45]. We also found that JNK activity was activated by TPA in gastric cancer cells, MGC80-3. This activation occurred very rapidly within 3 hr of the TPA addition (Figure 1), suggesting that JNK protein may be one of the TPA primary response targets. More importantly, JNK expression level activated by TPA was always keeping higher than that in untreated MGC80-3 cells. This evidence also suggests that the JNK pathway is a major one for mediating TPA effect, which should be functional in gastric cancer cells.

JNK cascade involves many factors. Activated transcription factor 2 (ATF-2) and *c-jun* are activated mainly by JNK^[43]. Transcriptional regulation of *c-jun* expression is mainly mediated by a TPA-response element in its promoter, which binds to *c-jun*/ATF-2 heterodimers^[46]. Increase of *c-jun* mRNA

by various stimuli is highly regulated by phosphorylation of *c-jun* or ATF-2 that bind to the activator protein-1 binding sites present in the *c-jun* promoter^[47]. Our observation showed a possible interrelation between JNK and *c-jun*, which were activated rapidly by TPA and reach its maximal peak within 3 hr, and then decreased with time-dependent manner (Figure 1, 2). Activator protein-1 is a transcriptional factor mainly composed of the *c-jun/c-jun* homodimer or *c-jun/c-fos* heterodimer. The fact that TPA also activated the transcriptional activity of activator protein-1 significantly in MGC80-3 cells (Figure 3) suggests that signaling that causes the transcriptional activity of activator protein-1 is functional in gastric cancer cells. Thus, induction of *c-jun* by TPA in MGC80-3 cells is likely due to the activation of JNK, which led to the activation of activator protein-1 transcriptional activity. This is a JNK cascade stimulated by TPA indeed.

How to function for JNK activation by TPA in gastric cancer cells is still largely unknown. MGC80-3 cells underwent apoptosis in response to TPA, as evidenced by typical morphological changes (Figure 5A, B), which is consistent with our previous observation^[48,49]. Our results demonstrated that apoptosis induction by TPA was associated with JNK activation via PKC pathway. When PKC specific inhibitor, Wortmannin was used to treat MGC80-3 cells prior to TPA treatment, it would prevent TPA further to activate JNK expression (Figure 4). In this case, apoptotic cells were much fewer than that in TPA-treated cells, apoptotic index dramatically decreased from 32.56 % to 8.71 % (Figure 5A, B). Thus, these data not only confirm that TPA-induced apoptosis is associated with JNK activation through PKC pathway, but also indicate that JNK cascade is required for apoptosis induction by TPA, which may be one mechanism for TPA to inhibit the growth of gastric cancer cells.

In summary, the present study demonstrates that the activation of JNK cascade is mediated through a signaling transduction pathway that involves PKC-dependent pathway, which leads to activation of JNK, induction of the *c-jun* expression and activation of activator protein-1 transcriptional activity in response to TPA. The activation of JNK cascade is closely associated with induction of apoptosis by TPA in gastric cancer cells.

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