

Negative feedback regulation of ASK1 by protein phosphatase 5 (PP5) in response to oxidative stress

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Apoptosis signal-regulating kinase 1 (ASK1) is a MAP kinase kinase kinase (MAPKKK) that activates the JNK and p38 MAP kinase cascades and is activated in response to oxidative stress such as hydrogen peroxide (H₂O₂). A yeast two-hybrid screening identified a serine/threonine protein phosphatase 5 (PP5) as a binding partner of ASK1. PP5 directly dephosphorylated an essential phospho-threonine residue within the kinase domain of ASK1 and thereby inactivated ASK1 activity *in vitro* and *in vivo*. The interaction between PP5 and ASK1 was induced by H₂O₂ treatment and was followed by the decrease in ASK1 activity. PP5 inhibited not only H₂O₂-induced sustained activation of ASK1 but also ASK1-dependent apoptosis. Thus, PP5 appears to act as a physiological inhibitor of ASK1–JNK/p38 pathways by negative feedback.

Keywords: apoptosis/ASK1/MAP kinase/oxidative stress/PP5

Introduction

The mitogen-activated protein kinase (MAPK) signaling cascades are well conserved in cells from yeast to human and are composed of three sequentially activating protein kinases which are referred to as MAPK, MAPK kinase (MAPKK) and MAPKK kinase (MAPKKK). Once activated, MAPKKK phosphorylates and thereby activates specific MAPKKs, which then phosphorylates and activates specific MAPKs. Two mammalian MAPKs, c-Jun N-terminal kinase (JNK) and p38 MAPK, are known to be activated by various environmental stresses and regulate diverse cellular functions including cytokine production, differentiation and apoptosis (Nishida and Gotoh, 1993; Xia *et al.*, 1995; Kyriakis and Avruch, 1996; Ichijo, 1999; Widmann *et al.*, 1999; Davis, 2000; Ono and Han, 2000; Matsuzawa and Ichijo, 2001). In addition to being activated by stresses such as oxidative stress, high osmolarity, UV and endoplasmic reticulum stress, the JNK and p38 can also be activated by pro-inflammatory cytokines such as tumor necrosis factor (TNF), Fas-ligand and IL-1. JNK is activated by MAPKKs SEK1 (also known as MKK4) or MKK7, and p38 is activated by MKK3 or MKK6. Numerous candidates for MAPKKKs

that activate SEK1/MKK4, MKK7, MKK3 and/or MKK6 have been reported (Ichijo, 1999; Davis, 2000).

Apoptosis signal-regulating kinase (ASK) 1, a mammalian MAPKKK, activates the JNK and p38 pathways and is activated in response to various cytotoxic stresses, including hydrogen peroxide (H₂O₂), Fas ligation, TNF, serum withdrawal and anti-tumor reagents (Ichijo *et al.*, 1997; Tobiume *et al.*, 1997; Chang *et al.*, 1998; Gotoh and Cooper, 1998; Nishitoh *et al.*, 1998; Saitoh *et al.*, 1998; Wang, T.H. *et al.*, 1998, 1999). Overexpression of ASK1 in epithelial cells in low serum conditions induced apoptosis (Ichijo *et al.*, 1997), and ASK1-deficient cells were resistant to H₂O₂- and TNF-induced apoptosis (Tobiume *et al.*, 2001), indicating that ASK1 plays a pivotal role in stress-induced apoptosis. On the other hand, moderate expression of a constitutively active form of ASK1 induced neuronal differentiation in PC12 cells (Takeda *et al.*, 2000). In addition, low and high expression of exogenous ASK1 in keratinocytes induced differentiation and apoptosis, respectively (Sayama *et al.*, 2000). These results suggest that ASK1 has a broad range of biological activities depending on cell-types, cellular context or the extent of ASK1 activation. The kinase activity of ASK1 is tightly regulated within cells; under non-stressed conditions, ASK1 is inhibited by association with its physiological inhibitor, thioredoxin (Trx). When cells are exposed to H₂O₂ or TNF, reactive oxygen species (ROS)-dependent oxidation of Trx occurs, which results in dissociation of Trx from ASK1 and thereby activation of ASK1 (Saitoh *et al.*, 1998; Liu *et al.*, 2000). Oligomerization-dependent autophosphorylation appears to be the next step required for full activation of ASK1 after the release from Trx (Gotoh and Cooper, 1998; Liu *et al.*, 2000; K.Tobiume, M.Saitoh and H.Ichijo, submitted for publication). On the other hand, mechanisms of how the activated ASK1 returns to an inactive form has not been elucidated.

Many protein phosphatases that directly dephosphorylate and thereby inactivate JNK or p38 have been identified, which include VHR (Ishibashi *et al.*, 1994), CL100 (MKP1) (Charles *et al.*, 1992; Keyse and Emslie, 1992; Alessi *et al.*, 1993; Sun *et al.*, 1993), PAC1 (Rohan *et al.*, 1993; Ward *et al.*, 1994), MKP2 (hVH2, TYP1) (Guan and Butch, 1995; King *et al.*, 1995; Misra-Press *et al.*, 1995), hVH5 (M3/6) (Martell *et al.*, 1995; Theodosiou *et al.*, 1996), Pyst2 (Dowd *et al.*, 1998) and MKP5 (Tanoue *et al.*, 1999). Serine/threonine protein phosphatase (PP) 2C α inactivates the stress-responsive MAPK pathways at the level of either MAPKK or MAPK (Takekawa *et al.*, 1998). Moreover, it was recently shown that PP2C β inactivated a MAPKKK, TAK1, through direct dephosphorylation (Hanada *et al.*, 2001).

Protein phosphatase 5 (PP5) is a member of the serine/threonine protein phosphatase family which includes PP1,

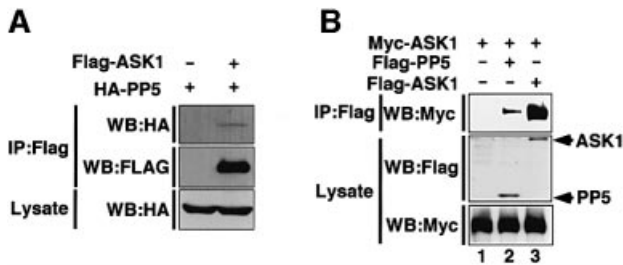


Fig 1. Interaction of PP5 with ASK1 in non-stressed cells. (A and B) 293 cells were transiently co-transfected with the indicated plasmids. Lysates were divided and immunoprecipitated with anti-Flag antibody (M2 gel). Immunoprecipitates were subjected to immunoblot analysis with anti-HA antibody (A, top) or with anti-Myc antibody (B, top). The presence of expressed proteins in the same lysates was verified by the indicated combination of immunoprecipitation (IP) and immunoblotting (WB).

PP2A, PP2B, PP2C, PP4 and PP7. PP5 possesses four tetratricopeptide repeat (TPR) domains in its N-terminus (Chen *et al.*, 1994), which are implicated in protein-protein interactions (Blatch and Lassle, 1999). PP5 has been suggested to negatively regulate the functions of p53 and glucocorticoid receptor (GR) (Chen, M.S. *et al.*, 1996; Silverstein *et al.*, 1997; Zuo *et al.*, 1998, 1999; Russell *et al.*, 1999). PP5 interacts with various molecules including CDC16, CDC27 (Ollendorff and Donoghue, 1997), hCRY2 (Zhao and Sancar, 1997) and the HSP90-GR complex (Chen, M.S. *et al.*, 1996; Silverstein *et al.*, 1997; Russell *et al.*, 1999), and the catalytic activity of PP5 was reported to be inhibited or activated *in vitro* by okadaic acid (Chen *et al.*, 1994) and arachidonic acid (Chen and Cohen, 1997; Skinner *et al.*, 1997), respectively. However, physiological substrates of PP5 have not been identified.

Here we report that PP5 directly binds to ASK1 and inhibits ASK1 activity in a negative feedback manner. The interaction of PP5 and ASK1 was induced by the treatment of cells with H₂O₂. PP5 dephosphorylates a critical phospho-threonine residue within the activation loop of ASK1 and thereby inactivated H₂O₂-induced ASK1 activity. Moreover, PP5 inhibited H₂O₂-induced sustained activation of ASK1 and ASK1-dependent apoptosis.

Results

PP5 directly interacts with ASK1

During the course of two-hybrid screening for ASK1 binding proteins, we found that PP5 interacted with ASK1 in yeast (see Materials and methods). We thus examined whether PP5 and ASK1 interact in mammalian cells by a co-immunoprecipitation analysis. When Flag-tagged ASK1 and hemagglutinin (HA)-tagged PP5 were co-transfected in 293 cells, HA-PP5 was co-immunoprecipitated with Flag-ASK1 (Figure 1A). When Myc-ASK1 was co-transfected with Flag-PP5 or Flag-ASK1, PP5-ASK1 and ASK1-ASK1 interactions were clearly observed (Figure 1B). These results indicated that PP5 interacts with ASK1 *in vivo*.

H₂O₂- and TNF-induced interaction of PP5 and ASK1 *in vivo*

Because the observed interaction between PP5 and ASK1 in non-stressed cells was much weaker than the homooligomeric interaction of ASK1 (Figure 1B), we asked whether cell stimulation may alter this interaction. We overexpressed Flag-ASK1 and HA-PP5 in HeLa cells and subjected them to a co-immunoprecipitation analysis after stimulating the cells with certain stresses. Treatment with H₂O₂, one of the most potent activators of ASK1, dramatically increased the association between PP5 and ASK1 (Figure 2A). A slight increase of interaction was also observed by TNF treatment (Figure 2A), which activates ASK1 through a ROS-dependent manner (Saitoh *et al.*, 1998; Liu *et al.*, 2000). We analyzed the dose- and time-dependent effects of H₂O₂ on the interaction of PP5 and ASK1 in HeLa cells. H₂O₂-induced association was observed from 0.5 mM H₂O₂ and increased in a dose-dependent manner (Figure 2B). Similar results were observed in COS7 cells, in which H₂O₂-induced association was observed from as low as 0.05 mM H₂O₂ (Figure 2B). The PP5-ASK1 interaction in HeLa cells was detected within 1 min after treatment with 0.5 mM H₂O₂, peaked at 15 min and decreased thereafter (Figure 2C). More sustained and stronger interaction of PP5 and ASK1 was detected by the treatment with 1 or 5 mM H₂O₂ (Figure 2C). The PP5-ASK1 interaction appeared to be specific since PP2A was unable to bind ASK1 even after treatment of cells with H₂O₂ (Figure 2D). To confirm the observed PP5-ASK1 interaction under more physiological conditions, we examined the endogenous association of PP5 and ASK1 in non-transfected cells. Lysates from H₂O₂-treated A549 cells were immunoprecipitated with normal rabbit IgG or anti-ASK1 polyclonal antibody, and the immunoprecipitates were analyzed by immunoblotting with anti-PP5 antibody. The interaction was clearly induced by H₂O₂ treatment (Figure 2E). TNF-dependent interaction of endogenous PP5 with overexpressed ASK1 was also observed in mouse L929 cells (data not shown).

PP5 has been reported to exist mainly in the nucleus (Chen *et al.*, 1994), whereas ASK1 occurs exclusively in the cytoplasm (see below). To examine a topological rationale to the observed association of PP5 and ASK1, we determined the subcellular localization of PP5. A simple subcellular fractionation using a sucrose-containing buffer divides cellular components into two major fractions: a supernatant which contains mainly cytoplasmic proteins, and a pellet which contains nuclear, cytoskeletal and mitochondrial proteins and large fragments of cellular membranes. Immunoblot analysis revealed that while promyelocyte (PML), a positive control for nuclear protein, was detected only in the pellet, PP5 can be detected not only in the pellet but also in the supernatant (Figure 2F). This result suggests that a substantial amount of endogenous PP5 exists in the cytoplasm. We further confirmed the cytoplasmic localization of PP5 by an immunofluorescence staining of HeLa cells which were transiently transfected with Flag-ASK1 and HA-PP5. HA-PP5 was detected both in the cytoplasm and nucleus, whereas Flag-ASK1 was found mainly in the cytoplasm (Figure 2G). In addition, subcellular localization of PP5 was unaffected by the overexpression of ASK1 or H₂O₂

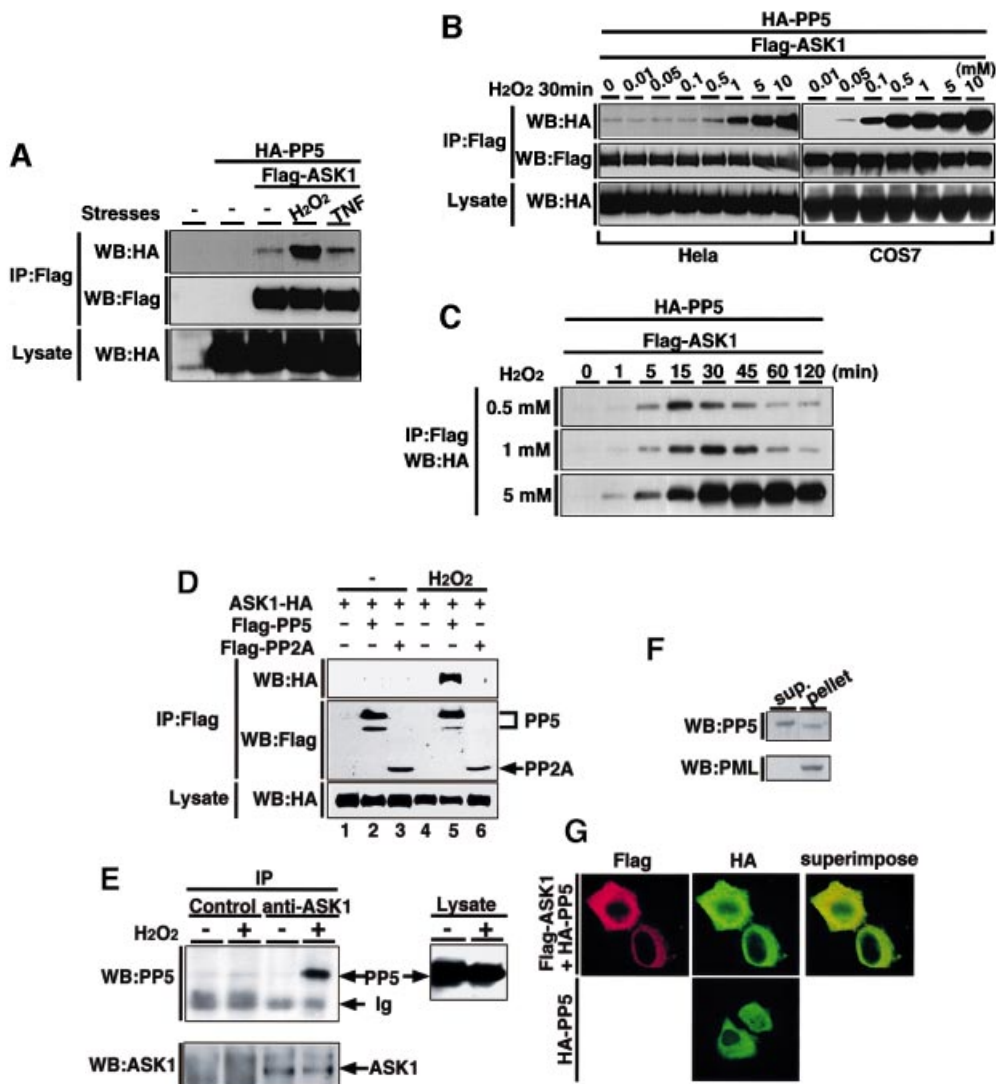


Fig. 2. Oxidative stress enhances the interaction between PP5 and ASK1. (A) H₂O₂- and TNF-induced interaction of PP5 and ASK1. HeLa cells were transiently co-transfected with HA-PP5 and Flag-ASK1. Thirty-six hours later, the cells were treated with 1 mM H₂O₂ or 200 ng/ml TNF for 20 min, and lysates were subjected to co-immunoprecipitation analysis as described in Figure 1A. (B) H₂O₂ dose-dependent interaction of PP5 and ASK1. HeLa cells and COS7 were transfected as in (A), treated with increasing concentrations of H₂O₂ for 30 min and analyzed by co-immunoprecipitation analysis. (C) Time course of the H₂O₂-induced interaction of PP5 and ASK1. HeLa cells were transfected as in (A), treated with indicated concentrations of H₂O₂ for the indicated periods and analyzed by co-immunoprecipitation analysis. (D) Specific interaction of ASK1 with PP5 but not PP2A. 293 cells were transiently co-transfected with ASK1-HA and Flag-PP5 or Flag-PP2A. Cells were treated with 5 mM H₂O₂ for 30 min, and lysates were subjected to co-immunoprecipitation analysis as described in Figure 1A. (E) Interaction of endogenous PP5 and ASK1. Approximately 5 × 10⁷ of A549 cells were treated with 5 mM H₂O₂ for 30 min. Cell lysates were divided and immunoprecipitated with normal rabbit IgG or anti-ASK1 polyclonal antibody (DAV) and were immunoblotted with anti-PP5 monoclonal antibody. The presence of ASK1 and PP5 in the same lysates was verified by immunoblotting (WB). Ig indicates non-specific reactions derived from rabbit IgG. (F) Subcellular localization of endogenous PP5 in A549 cells. A subcellular fractionation was performed as described in Materials and methods, and PP5 was detected by immunoblotting. Anti-PML antibody was used as a positive control for the nuclear protein. (G) Subcellular localization of transfected ASK1 and PP5 in HeLa cells. HeLa cells were transfected with Flag-ASK1 and HA-PP5, or with HA-PP5 alone, and the cells were subjected to an immunofluorescence staining as described in Materials and methods.

treatment (Figure 2G and data not shown). These results indicate that PP5 meets ASK1 mainly in the cytoplasm.

PP5 inhibits ASK1 activity *in vivo*

To explore a potential effect of PP5 on ASK1 *in vivo*, we determined the kinase activity of ASK1 by an immunocomplex kinase assay after transfection of ASK1 with or without PP5 in HeLa cells. Co-expression of PP5 partially inhibited the basal kinase activity of ASK1 (Figure 3A, lane 3). This partial inhibition may reflect the weak

interaction of PP5 and ASK1 in non-stressed cells (Figure 1B and see below). ASK1 activates JNK and p38 pathways but not the extracellular signal-regulated kinase (ERK) pathway (Ichijo *et al.*, 1997). We thus tested whether PP5 specifically inhibits ASK1-dependent JNK and p38 pathways. Figure 3B shows that expression of PP5 reduced ASK1-induced activation of JNK in a dose-dependent manner. ASK1-induced p38 activation was also reduced by PP5 (data not shown). On the other hand, PP5 had little effect on an MEKK1-induced activation of JNK (Figure

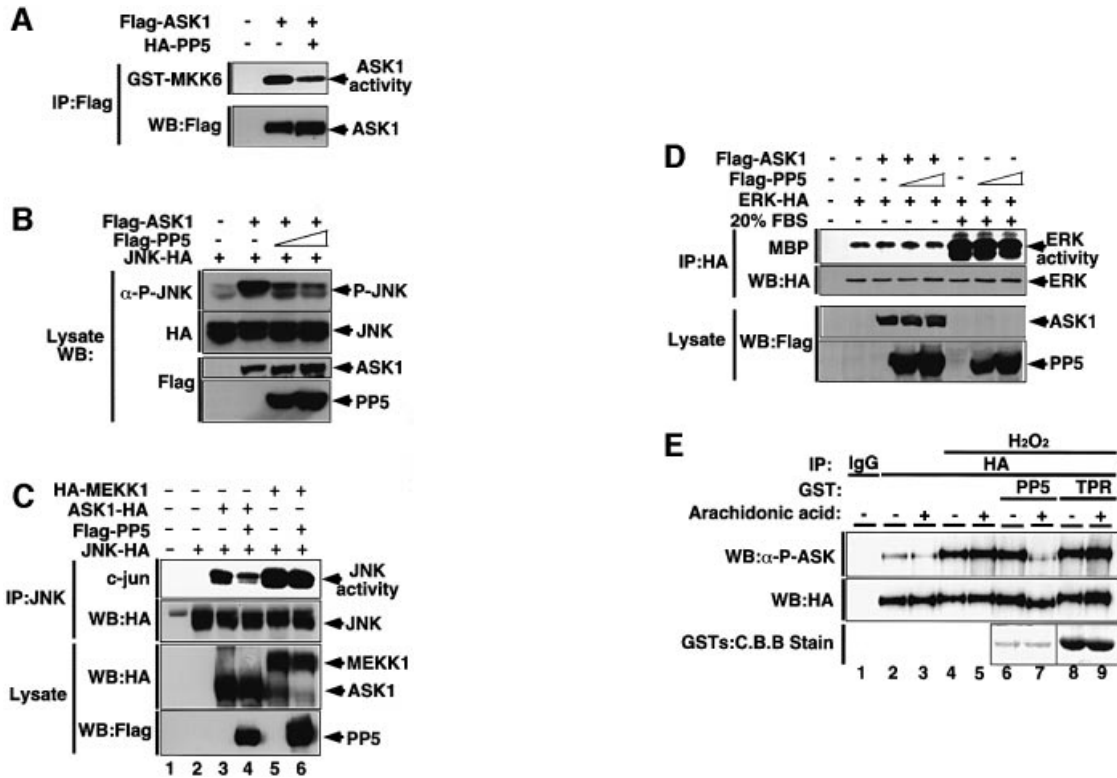


Fig. 3. PP5 dephosphorylates and inactivates ASK1 *in vivo* and *in vitro*. (A) PP5 reduces ASK1 activity *in vivo*. HeLa cells were transfected with the indicated plasmids. Thirty-six h later, immunocomplex kinase assay for ASK1 was performed as described in Materials and methods. ASK1 activity was measured using GST-MKK6 as a substrate (top). Consistent expression of Flag-ASK1 was confirmed by immunoblotting (bottom). (B) PP5 reduces ASK1-induced activation of JNK. 293 cells were transfected with the indicated plasmids. JNK activity was measured by immunoblotting using phospho-specific antibody to SAPK/JNK (Thr183/Tyr185). Expression of JNK-HA, Flag-ASK1 and Flag-PP5 was confirmed by immunoblotting using the indicated antibodies. (C) PP5 does not inhibit MEKK1-induced activation of JNK. 293 cells were transfected with the indicated plasmids. Lysates were immunoprecipitated with anti-JNK antibody, and JNK activity was measured by immunocomplex kinase assay using GST-c-jun as a substrate. Expression of JNK-HA, ASK1-HA, HA-MEKK1 and Flag-PP5 was confirmed by immunoblotting using the indicated antibodies. (D) PP5 does not inhibit the ERK pathway. 293 cells were transfected with the indicated plasmids. Cells were then stimulated with 20% FBS for 20 min, and ERK activity was measured by immunocomplex kinase assay using MBP as a substrate (top). Expression of transfected plasmids were confirmed by immunoblotting using the indicated antibodies. (E) PP5 directly dephosphorylates a critical phospho-threonine residue of ASK1. PAE cells stably transfected with ASK1-HA were treated with 1 mM H₂O₂ for 30 min. ASK1 was immunoprecipitated with anti-HA, incubated with recombinant full-length PP5 or with truncated PP5 (TPR domain only) for 20 min in the presence (+) or absence (-) of arachidonic acid. The samples were subjected to immunoblotting analysis with anti-phospho-ASK1 antibody (P-ASK). The presence of ASK1 and GST fusion proteins was verified by immunoblotting (WB) and staining with Coomassie Brilliant Blue dye (C.B.B stain), respectively.

3C) or a serum-induced activation of ERK (Figure 3D), indicating that PP5 selectively inhibits ASK1 and its downstream targets, JNK and p38, *in vivo*.

ASK1 is a direct substrate for PP5

PP5 negatively regulates GR- and/or p53-signaling pathways (Chen, M.S. *et al.*, 1996; Silverstein *et al.*, 1997; Zuo *et al.*, 1998, 1999; Russell *et al.*, 1999); however, a direct substrate for PP5 has not been identified. Moreover, although the above results indicate that PP5 interacts with and inactivates ASK1 *in vivo*, the mechanism of inactivation of ASK1 is unknown. We thus examined whether PP5 can directly dephosphorylate and thereby inactivate ASK1 *in vitro*. To this end, immunoprecipitated ASK1-HA was incubated with recombinant glutathione *S*-transferase (GST) fusion proteins of PP5 (GST-PP5) or the TPR domain only (GST-TPR), and the phosphorylation status of ASK1 was monitored by an immunoblot analysis using a phospho-specific antibody to a critical phospho-threonine residue (Thr845) within the activation loop of ASK1

(K.Tobiume, M.Saitoh and H.Ichijo, submitted for publication). Phosphorylation of Thr845 of ASK1, which represents an activation status of ASK1 (K.Tobiume, M.Saitoh and H.Ichijo, submitted for publication), was induced by H₂O₂ treatment (Figure 3E, top panel, compare lanes 2 and 4). GST-PP5 but not GST-TPR, dephosphorylated the Thr845 of ASK1 only in the presence of arachidonic acid, a specific activator of PP5 (Figure 3E, top panel, lane 7). Moreover, ASK1 incubated with active PP5 was found to migrate faster on SDS-PAGE (Figure 3E, middle panel, lane 7), also suggesting a dephosphorylation of ASK1. These results indicate that PP5 can directly dephosphorylate at least Thr845 of ASK1 and thereby inactivate ASK1.

PP5 inhibits ASK1 in a negative feedback manner *in vivo*

Trx has been identified as a physiological inhibitor of ASK1 under non-stressed conditions (Saitoh *et al.*, 1998). It was of interest to compare the modes of inhibitory action

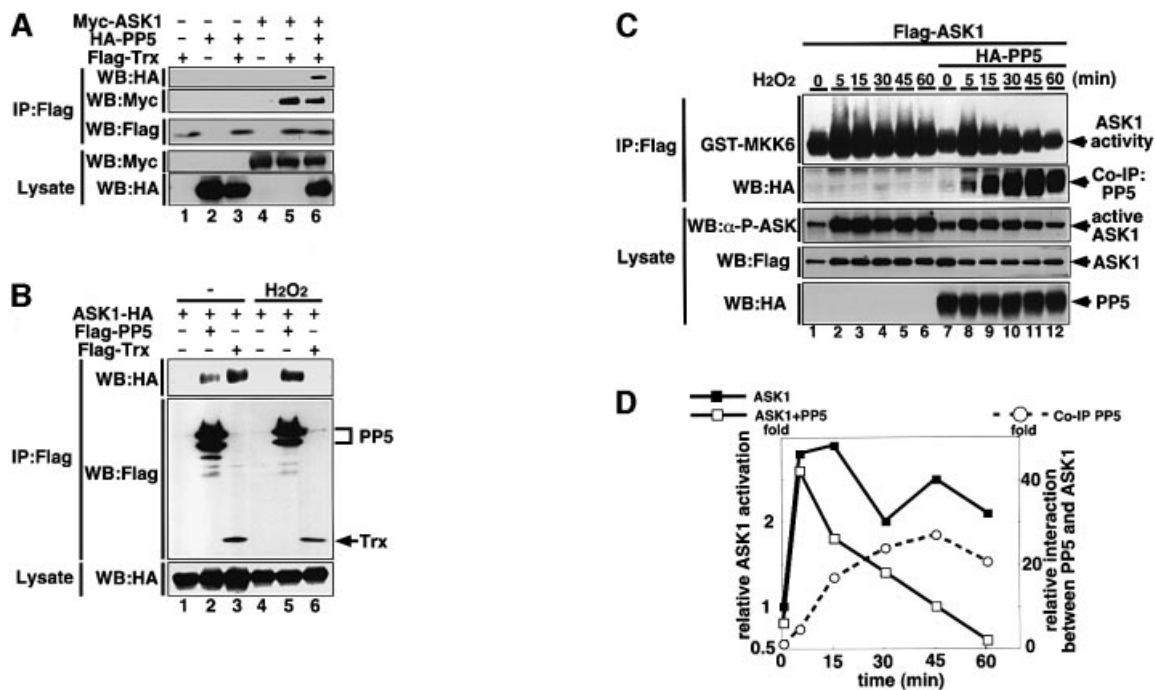


Fig. 4. PP5 inhibits H₂O₂-induced sustained activation of ASK1 *in vivo*. (A) PP5 can participate in the complex of ASK1 and Trx. 293 cells were transiently co-transfected with the indicated plasmids. Lysates were divided and immunoprecipitated with anti-Flag antibody (M2 gel). Immunoprecipitates were subjected to immunoblot analysis with anti-HA antibody (top panel) or with anti-Myc antibody (second panel). The presence of expressed proteins in the same lysates was verified by the indicated combination of immunoprecipitation (IP) and immunoblotting (WB). (B) Opposing effects of H₂O₂ on the interactions of ASK1 with PP5 and Trx. 293 cells were transfected with the indicated plasmids, treated with 5 mM H₂O₂ for 30 min and analyzed by co-immunoprecipitation analysis. (C) PP5 inhibits H₂O₂-induced sustained activation of ASK1. Flag-ASK1 was transiently transfected with or without HA-PP5 into HeLa cells. Thirty-six h later, the cells were treated with 5 mM H₂O₂ for the indicated periods. Lysates were divided, immunoprecipitated or immunoblotted, and the kinase activity of ASK1 (top panel), phosphorylation status of Thr845 of ASK1 (third panel) and co-immunoprecipitated HA-PP5 (second panel) were analyzed. The presence of HA-PP5 and Flag-ASK1 in the same lysates was verified by immunoblotting (WB). (D) The intensity of GST-MKK6 phosphorylation (scale in the left) and the amount of co-immunoprecipitated PP5 (scale in the right) in (C) were quantified and represented by a graph. Relative values of activation and interaction were calculated by dividing the intensities of phosphorylated GST-MKK6 or co-immunoprecipitated HA-PP5 (Co-IP PP5) at different time points by those at time zero.

between PP5 and Trx. We first examined whether PP5 can participate in the complex with ASK1 and Trx. HA-PP5 was co-immunoprecipitated with Flag-Trx in 293 cells only in the presence of ASK1 (Figure 4A), indicating that a Trx-ASK1-PP5 ternary complex can be formed in non-stressed conditions. This also suggests that PP5 may play a role in keeping ASK1 inactive together with Trx under non-stressed conditions; however, since the interaction between PP5 and ASK1 was much stronger in H₂O₂-treated cells (Figure 2), the effects of PP5 on ASK1 are likely to be exerted mainly in stressed conditions. We thus analyzed the stoichiometry of interaction between PP5-ASK1 and Trx-ASK1 in the cells treated with or without H₂O₂ (Figure 4B). Although the interaction of PP5 and ASK1 was much weaker than that of Trx and ASK1 in non-stressed cells (Figure 4B, lanes 2 and 3), H₂O₂ clearly induced the dissociation of Trx from ASK1, and reciprocally induced the association of PP5 with ASK1 (Figure 4B, lanes 5 and 6). These results suggest that these two ASK1 inhibitors may play different roles in ASK1 inhibition; PP5 appears to mainly target and inactivate the activated form of ASK1. We next examined the kinetics of the PP5-ASK1 interaction and that of activation and phosphorylation states of ASK1 in H₂O₂-stimulated cells (Figure 4C). Without co-transfection of PP5 (Figure 4C, lanes 1-6), ASK1 activity (Figure 4C, top

panel; Figure 4D) and activating phosphorylation of ASK1 (Figure 4C, third panel) were induced by H₂O₂ within 5 min and sustained for at least 60 min. In contrast, when PP5 was co-transfected (Figure 4C, lanes 7-12), H₂O₂-induced activation as well as phosphorylation of ASK1 peaked at 5 min and decreased thereafter. Reciprocally, PP5 started to bind to ASK1 after 5 min, and the interaction was increased with time (Figure 4C, second panel; Figure 4D). This inverse correlation between PP5-ASK1 complex formation and activation/phosphorylation of ASK1 strongly suggests that PP5 specifically targets the activated form of ASK1 by negative feedback.

PP5 inhibits ASK1-dependent apoptosis

We have recently shown by deleting ASK1 in mice that H₂O₂-induced sustained activations of JNK and p38 are lost in ASK1^{-/-} embryonic fibroblasts, and that ASK1^{-/-} cells are resistant to H₂O₂-induced apoptosis (Tobiume *et al.*, 2001). Thus, H₂O₂-induced sustained activation of JNK/p38, which resulted from sustained activation of ASK1, strongly correlated with apoptosis. These findings suggest that duration of ASK1 activation may directly link to the determination of cell fate (survival or apoptosis). Since PP5 inhibited only sustained but not transient ASK1 activity induced by H₂O₂ (Figure 4C and D), we examined

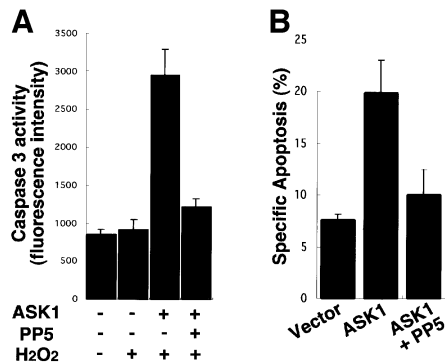


Fig. 5. PP5 inhibits ASK1-dependent apoptosis. (A) PP5 inhibits ASK1-dependent activation of caspase-3-like protease. Indicated plasmids were transiently transfected into 293 cells and caspase-3-like protease activity was measured as described in Materials and methods. Results are the means of duplicate determinations \pm SE from one of more than three representative experiments. (B) PP5 inhibits ASK1-dependent cell death. The indicated plasmids were transiently transfected into HeLa cells with pEGFP, and apoptotic cell death was determined by a morphological analysis as described in Materials and methods. Results are the means of duplicate determinations \pm SE from one of two representative experiments.

whether PP5 inhibits H₂O₂-induced ASK1-dependent apoptosis. ASK1 was transfected into 293 cells with or without PP5, and H₂O₂-induced apoptosis was assessed by caspase-3 activity. While ASK1 enhanced H₂O₂-induced caspase-3 activation, co-expressed PP5 suppressed the ASK1-dependent apoptosis (Figure 5A). The inhibitory effect of PP5 on ASK1-dependent apoptosis was also confirmed in HeLa cells as determined by cell morphology (Figure 5B). Collectively, PP5 negatively regulates H₂O₂-induced sustained activation of ASK1–JNK/p38 pathways, and thereby inhibits ASK1-dependent apoptosis by negative feedback.

Discussion

Transient and persistent activations of MAPK are known to lead to different cell fates (Marshall, 1995); early and transient activation of ERK induces proliferation of PC12 cells, whereas prolonged activation of ERK induces neuronal differentiation. Early/transient and late/sustained activations of JNK induced by TNF (Guo *et al.*, 1998; Roulston *et al.*, 1998), UV-C or gamma-radiation (Chen, Y.R. *et al.*, 1996) have been reported to correlate with survival and apoptosis, respectively. However, the mechanism by which duration of MAPK activation is regulated has not been fully elucidated. In the present study, we found that PP5 directly interacts with and inactivates activated ASK1 in a negative feedback manner and thereby inhibits ASK1-dependent sustained activations of JNK/p38 and apoptosis. Such a negative feedback system may be useful for cells to determine their fates (survival or apoptosis) in response to exposed stresses depending on their dose or duration. The ratio of expression levels between ASK1 and PP5 may be an important determinant of cellular sensitivity to oxidative stresses.

In this study, we found that PP5 specifically targets active form(s) of ASK1. However, which part of active configuration of ASK1 is recognized by PP5 is unknown.

Although H₂O₂-induced phosphorylation of Thr845 returned to the basal level after 60 min (Figure 4C, third panel, compare lanes 7 and 12), ASK1 still bound a substantial amount of PP5 (Figure 4C, second panel, compare lanes 7 and 12). These results suggest that phosphorylation of Thr845 itself is unlikely to give rise to a site recognized by PP5. A fine mapping of interaction sites between PP5 and ASK1 may answer this question.

PP5 interacts not only with ASK1 but also with GR complex (Chen, M.S. *et al.*, 1996; Silverstein *et al.*, 1997; Russell *et al.*, 1999), CDC16, CDC27 (Ollendorff and Donoghue, 1997) and hCRY2 (Zhao and Sancar, 1997). Thus, it is formally possible that the anti-apoptotic activity of PP5 observed in this study may not be solely due to its inhibitory action on ASK1. Interestingly, antisense oligonucleotide-mediated inhibition of PP5 has been reported to activate transcriptional activity of p53, a potent inducer of apoptosis, suggesting that PP5 inhibits p53 function *in vivo* (Zuo *et al.*, 1998). On the other hand, p53 is reported to be activated by JNK or p38 via phosphorylation or stabilization of p53 protein (Fuchs *et al.*, 1998; Bulavin *et al.*, 1999; Huang *et al.*, 1999; Potapova *et al.*, 2000; She *et al.*, 2000). We do not know yet whether ASK1-induced apoptosis requires JNK/p38-dependent p53 modification. Such a study will disclose an exact mechanism of how oxidative stress induces apoptosis through the ASK1–MAPK cascade.

Materials and methods

Yeast two-hybrid system

To analyze the regulatory mechanisms of ASK1, we employed the yeast two-hybrid system to search for proteins that bind to ASK1 by using LexA DNA binding domain–ASK1–K709R (kinase inactive mutant form of ASK1) as a bait (Saitoh *et al.*, 1998). We identified several positive clones which were closely related to ASK1, and designated this gene as ASK2 [K. Takeda and H. Ichijo, manuscript in preparation; also called MAPKKK6 (Wang, X.S. *et al.*, 1998)]. A human fibroblast cDNA library in the pJG4-5 prey plasmid (a gift from Roger Brent) was screened for proteins that interact with mouse ASK2/MAPKKK6 using the EGY48 yeast reporter strain and the pSH18-34 reporter plasmid as described (Kawabata *et al.*, 1995). The bait plasmid expressing ASK2 protein was constructed in-frame with the LexA DNA-binding domain of the pEG202 bait plasmid. Plasmids of positive clones were recovered, and the cDNA inserts were sequenced. DNA sequencing analysis of the ASK2-interacting clones revealed that one of them encoded a full-length cDNA of PP5. To assay the interaction between ASK1 and PP5, ASK1 and PP5 constructs were co-transformed along with the pSH18-34 reporter plasmid into EGY48 yeast strain. Transformants were tested on 5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside (X-gal; Calbiochem) containing plates. PP5 interacted with not only ASK2 but also ASK1 in yeast (data not shown).

Cell culture, expression vectors and transfections

HeLa, A549 and COS7 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal calf serum (FCS). DMEM containing a higher concentration of glucose (4.5 mg/ml) was used for 293 cells. Porcine aortic endothelial (PAE) cells stably transfected with ASK1-HA were cultured in HAM's F12 medium supplemented with 10% FCS and 200 μ g/ml G418. All cells were cultured with 100 units/ml penicillin G in a 5% CO₂ atmosphere at 37°C. A Flag tag was inserted at the N-termini of human ASK1 (Flag-ASK1), rat PP2A α catalytic subunit (a gift from Dr Hitoshi Nakagama; Flag-PP2A), human Trx (Flag-Trx) and human PP5 (Flag-PP5) in pcDNA3.0 (Invitrogen). An HA tag was inserted at the N-termini of human PP5 (HA-PP5) and human MEKK1 (a gift from Dr Gary Johnson; HA-MEKK1), and at the C-termini of human ASK1 (ASK1-HA), mouse JNK3-1 (JNK-HA) and *Xenopus* MAPK (ERK-HA) in pcDNA3.0. Six copies of the Myc tag was inserted at the N-terminus of human ASK1 (Myc-ASK1)

in pcDNA3.1. Transfection was performed with Fugene 6 (Roche) according to the manufacturer's instructions.

Antibodies and reagents

Monoclonal antibodies to the HA tag (clone 3F10 and 12CA5) were purchased from Roche Molecular Biochemicals. Monoclonal antibodies to PP5 and to JNK3 (MAPKp49) were purchased from Transduction Laboratories. Rabbit polyclonal antisera to ASK1 antibody (DAV) (Saitoh *et al.*, 1998) and phospho-specific polyclonal antibody to ASK1 (P-ASK; K.Tobiome, M.Saitoh and H.Ichijo, submitted for publication) was as described. Phospho-specific polyclonal antibody to SAPK/JNK (Thr183/Tyr185) was purchased from New England Biolabs. Anti-PML monoclonal antibody (clone PG-M3) and normal rabbit IgG were purchased from Santa Cruz Biotechnology Inc. The anti-Flag monoclonal antibodies (M2, biotinylated-M2 and M2 gel), myelin basic protein (MBP) and arachidonic acid were purchased from Sigma. Human recombinant TNF- α was purchased from Pepro Tech EC Ltd. The anti-Myc monoclonal antibody (clone 9E10) was purchased from Calbiochem.

Co-immunoprecipitation assay and immunoblotting

Cells were lysed in a lysis buffer containing 150 mM NaCl, 20 mM Tris-HCl pH 7.5, 5 mM EGTA, 1% Triton X-100, 1% deoxycholate, 12 mM β -glycerophosphate, 50 mM NaF, 1 mM sodium orthovanadate, 1 mM dithiothreitol (DTT), 1 mM phenylmethylsulfonyl fluoride (PMSF), 1.5% aprotinin. Cell extracts were clarified by centrifugation, and the supernatants were immunoprecipitated with anti-Flag antibody gel (M2 gel) or anti-ASK1 antibody (DAV) using protein A-Sepharose (Zymed). The beads were washed twice with the washing buffer A (1% Triton X-100, 500 mM NaCl, 20 mM Tris-HCl pH 7.5, 5 mM EGTA, 1 mM DTT), and twice with the washing buffer B (150 mM NaCl, 20 mM Tris-HCl pH 7.5, 5 mM EGTA, 1 mM DTT). The beads were subjected to SDS-PAGE followed by electroblotting onto PVDF membranes. After blocking with 5% skim milk in TBS-T (150 mM NaCl, 50 mM Tris-HCl pH 8.0, 0.05% Tween 20) for 1 h, the membranes were probed with antibodies. The antibody-antigen complexes were detected using the enhanced chemiluminescence (ECL) system (Amersham Pharmacia Biotech). Aliquots of whole-cell lysates were subjected to immunoblotting analysis to confirm appropriate expression of proteins. The amount of co-immunoprecipitated HA-PP5 were quantitated by Quantity One® (PDI), and were presented in fold amount of co-immunoprecipitated HA-PP5 compared with untreated cells.

Immunocomplex kinase assay

GST constructs for MKK6 kinase negative mutant (GST-MKK6) and c-jun (1-79) (GST-c-jun) were prepared in pGEX-4T-1 vector (Amersham Pharmacia Biotech) by PCR. Cells were lysed and immunoprecipitated as described above. The beads were incubated with 1 μ g of GST-MKK6 as a substrate for ASK1 for 20 min at 30°C in a final volume of 30 μ l of kinase buffer (20 mM Tris-HCl pH 8.0, 20 mM MgCl₂ and 0.3 μ Ci of [γ -³²P]ATP). GST-c-jun and MBP were used as substrates for JNK and ERK, respectively. Kinase reactions were stopped by adding SDS sample buffer. The beads were resolved on SDS-PAGE followed by immunoblotting analysis with ECL. Phosphorylation of GST-MKK6 or MBP was analyzed by a Fuji BAS2000 Image Analyzer. Aliquots of whole-cell lysates were subjected to immunoblotting analysis to confirm appropriate expression of proteins.

In vitro phosphatase assay

GST constructs for full-length PP5 (GST-PP5) and TPR domain of PP5 (GST-TPR) were prepared in pGEX-4T-1 vector (Amersham Pharmacia Biotech). GST fusion proteins were purified as described (Skinner *et al.*, 1997) using glutathione-Sepharose beads (Amersham Pharmacia Biotech). GST fusion proteins were eluted with a buffer containing 50 mM Tris-HCl pH 8.0, 4 mM MnCl₂, 0.1% 2-mercaptoethanol, 10 mM glutathione and stored at -20°C in 50% glycerol. PAE cells stably transfected with ASK1-HA were treated with 1 mM H₂O₂ for 30 min, and cell lysate was prepared by lysis buffer. Aliquots of the lysates were immunoprecipitated with anti-HA antibodies (12CA5) and protein A-Sepharose beads (Zymed), and the beads were washed twice with washing buffer B. The beads were then added with GST-PP5 or GST-TPR in the presence or absence of 100 μ M arachidonic acid, and reaction mixtures were incubated at 30°C for 20 min. Phosphatase reaction was stopped by adding SDS sample buffer. The beads were then resolved on SDS-PAGE followed by electroblotting onto PVDF membranes and analyzed by immunoblotting with ECL.

Measurement of caspase-3 activity

293 cells were transiently transfected with vector alone, Flag-ASK1, or Flag-ASK1 plus Flag-PP5. Forty-eight h later, the cells were washed with phosphate-buffered saline (PBS) and refed with serum-free medium. Cells were stimulated with or without 0.1 mM H₂O₂ plus 25 mM aminotriazole (catalase inhibitor) for 4 h. Caspase-3-like activity was measured by a CPP32/caspase-3 fluorometric protease assay kit (MBL) in which a fluorogenic synthetic peptide DEVD-7-amino-4-trifluoromethyl coumarine (AFC) was used as a substrate. The fluorescence of the released AFC was measured with an excitation wavelength of 360 nm and an emission wavelength of 530 nm. To confirm appropriate expressions of transfected plasmids, the same lysates of duplicate wells were combined and verified by immunoblot analysis.

Cell death assay

HeLa cells were transiently transfected with vector alone, Flag-ASK1, or Flag-ASK1 plus Flag-PP5 along with pEGFP C1 (Clontech). Thirty-six h later, the cells were washed with PBS and refed with serum-free medium. Cells were then treated with or without 0.5 mM H₂O₂ plus 25 mM aminotriazole (catalase inhibitor) for 8 h. Cell morphology and green fluorescence protein (GFP) signals were analyzed using a microscope. The ratio of apoptotic cells was determined by dividing the number of GFP-positive cells showing apoptotic morphologies, e.g. membrane blebbing and cell shrinkage, etc., by the total GFP-positive cell number (~500 cells). Specific apoptosis was calculated as the percentage of apoptotic cells in each experimental condition minus the percentage of apoptotic cells of the vector control.

Subcellular fractionation

A549 cells were washed three times in ice-cold PBS, scraped and harvested by centrifugation at 2000 *g* for 10 min at 4°C. The pellet was resuspended in a buffer containing 250 mM sucrose, 20 mM HEPES-KOH pH 7.5, 10 mM KCl, 1.5 mM MgCl₂, 1 mM EDTA, 1 mM EGTA, 1 mM PMSF, 0.5 mM DTT. After incubating on ice for 10 min, cells were homogenized with 20 strokes of a Dounce homogenizer (Wheaton). The homogenate was centrifuged at 2000 *g* for 10 min at 4°C and the supernatant was collected as the cytoplasmic fraction (sup.). The pellet was washed in ice-cold PBS and lysed in a lysis buffer containing 150 mM NaCl, 20 mM Tris-HCl, pH 7.5, 5 mM EGTA, 1% Triton X-100, 1% deoxycholate, 12 mM β -glycerophosphate, 1 mM DTT, 1 mM PMSF, 1.5% aprotinin. Each subcellular fraction was resolved on SDS-PAGE and analyzed by immunoblotting using anti-PP5 antibody. Anti-PML antibody was used as a positive control of the nuclear protein.

Immunofluorescence staining of cells

HeLa cells were plated onto Lab-Tek Chamber Slide (Nalge Nunc International), transiently transfected with HA-PP5 and Flag-ASK1, or with HA-PP5 alone, and cultured overnight. After a brief wash at room temperature (RT) in PBS, the cells were fixed in 4% paraformaldehyde in PBS for 30 min at RT. After several washes in PBS, the cells were permeabilized and blocked in PBS containing 0.2% Triton X-100 and 3% bovine serum albumin (BSA) for 30 min at RT. They were then incubated with anti-HA (clone 3F10; rat) and anti-Flag (clone M2; mouse) antibodies in PBS containing 0.2% Triton X-100 and 3% BSA (dilution buffer) for 1 h at RT, followed by three washes with PBS. Samples were incubated for 45 min with fluorescein isothiocyanate-conjugated anti-rat IgG and Texas Red® dye-conjugated anti-mouse IgG (Jackson ImmunoResearch Laboratories, Inc.) in the dilution buffer, washed three times with PBS, and the slides were mounted in 90% glycerol. Samples were then analyzed and recorded by an Olympus Fluoview microscope.

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