

Regulation of the Maintenance of Peripheral T-Cell Anergy by TAB1-Mediated p38 α Activation

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In anergic T cells, T-cell receptor (TCR)-mediated responses are functionally inactivated by negative regulatory signals whose mechanisms are poorly understood. Here, we show that CD4⁺ T cells anergized in vivo by superantigen MIs-1^a express a scaffolding protein, transforming growth factor β -activated protein kinase 1-binding protein 1 (TAB1), that negatively regulates TCR signaling through the activation of mitogen-activated protein kinase p38 α . TAB1 was not expressed in naive and activated CD4⁺ T cells. Inhibition of p38 activity in anergic T cells by a chemical inhibitor resulted in the recovery of interleukin 2 (IL-2) and the inhibition of IL-10 secretion. T-cell hybridoma 2B4 cells transduced with TAB1-containing retrovirus (TAB1-2B4 cells) showed activated p38 α , inhibited extracellular signal-regulated kinase (ERK) activity, culminating in reduced IL-2 levels and increased IL-10 production. The use of a p38 inhibitor or cotransfection of a dominant-negative form of p38 in TAB1-2B4 cells resulted in the recovery of ERK activity and IL-2 production. These results imply that TAB1-mediated activation of p38 α in anergic T cells regulates the maintenance of T-cell unresponsiveness both by inhibiting IL-2 production and by promoting IL-10 production.

Anergy is a tolerance mechanism in which lymphocytes are functionally inactivated following an antigen encounter but remain alive for an extended period of time. T cells become anergic after incomplete activation in vitro (33). Such anergic T cells display impaired tyrosine phosphorylation of the T-cell receptor (TCR) ζ chain and recruitment of ZAP-70 (36). Anergic T cells have a defect in the activation of Ras (11) and its downstream mitogen-activated protein kinases (MAPKs), extracellular signal-regulated kinase (ERK) and c-Jun NH₂-terminal kinase (JNK) (10, 19), culminating in the low activity of the AP-1 response element and defects in interleukin 2 (IL-2) synthesis (17). In addition to these defects in TCR signaling, active negative signaling pathways play a role in maintaining T-cell anergy. Activated Rap-1 in anergic T cells antagonizes Ras activity, resulting in reduced Raf-1 protein kinase activity (5). An increase in the level of the cyclic AMP response element-binding protein/cyclic AMP response element modulator complex in anergic T cells results in the repression of IL-2 transcription (28). Furthermore, the presence of some dominant molecules that inhibit anergic T-cell activation has been suggested (41). Searches for anergy-associated genes have identified *Tob* and a gene related to anergy in lymphocytes (*GRAIL*) as genes expressed during the early phase of T-cell anergy induction (2, 43). Forced expression of *Tob* in T cells can repress T-cell proliferation and cytokine transcription. The expression of *GRAIL* in T cells limits cytokine production in a manner that is dependent on E3 ubiquitin ligase activity. Macian et al. used microarrays to identify several sets of anergy-

associated genes that are expressed in a T-cell clone upon ionomycin treatment, which can induce clonal anergy (21). These genes are induced in an *NFAT1*-dependent manner, but their function has yet to be defined.

T cells expressing specific V β chains can be rendered anergic by inoculation of mice with a superantigen (30). These in vivo-anergized T cells display defects in tyrosine phosphorylation of the TCR ζ chain and subsequent recruitment of ZAP-70 (4, 22) similar to T cells anergized in vitro. In vivo-anergized T cells, as opposed to in vitro-anergized T cells, can produce IL-10, a cytokine that inhibits the T-cell response. Such cells can function as regulatory T cells (8, 37). Their unresponsiveness is maintained both by their autonomous anergy and by suppression mediated by inhibitory cytokines (3, 23).

Here, we investigated the MAPK pathways involved in the maintenance of peripheral CD4⁺-T-cell anergy by using TCR V β 8.1-transgenic mice (V β 8.1-tg mice) (47). The activity of p38 kinase was augmented in anergic CD4⁺ T cells, and a scaffolding protein, transforming growth factor β -activated protein kinase 1 (TAK1)-binding protein 1 (TAB1), that can activate p38 α (34) was expressed. The treatment of anergic T cells with an inhibitor of p38 resulted in the partial recovery of ERK activation and IL-2 secretion after TCR stimulation and in the inhibition of IL-10 secretion. The forced expression of TAB1 in a T-cell hybridoma resulted in the activation of p38, culminating in the inhibition of ERK, in the reduction of IL-2 production, and in the up-regulation of IL-10 synthesis after TCR stimulation. These results imply that TAB1 is a key regulatory molecule involved in maintaining T-cell anergy in vivo.

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MATERIALS AND METHODS

Mice. V β 8.1-tg mice, which have been backcrossed with CBA/Ca mice for more than 10 generations (47), were maintained in the Laboratory Animal Center for Research at the Nagasaki University School of Medicine. OT-II mice

were provided by W. R. Heath (The Walter and Eliza Hall Institute of Medical Research, Parkville, Victoria, Australia) as described previously (42). Transgenic mice were used when they were between 8 and 12 weeks old. CBA/J (H-2^k Mls-1^a) mice were obtained from Charles River (Yokohama, Japan). The review board of the Nagasaki University School of Medicine gave approval for the animal studies.

Induction of T-cell anergy and cell culture. T-cell anergy was induced in V β 8.1-tg mice as previously described (48). T-cell-depleted spleen cells (1.5×10^7) were prepared by treatment of CBA/J spleen cells with a combination of anti-CD4 (GK1.5), anti-CD8 (3.155), and anti-Thy-1.2 (30H12) monoclonal antibodies (MAbs) plus complement and were inoculated intravenously into V β 8.1-tg mice. At 12 to 18 days after injection, CD4⁺ T cells were prepared from inguinal, popliteal, brachial, axillary, and mesenteric lymph nodes of untreated or superantigen-inoculated mice. CD4⁺ T cells were purified (92 to 98%) by treatment of lymph node cells with an anti-CD8 MAb (3.155) plus complement, followed by nylon wool column enrichment, or were purified (>98%) by use of BD IMag-anti-mouse CD4 MAb GK1.5 (BD Biosciences Pharmingen, San Diego, Calif.).

CD4⁺ T cells (5×10^4 /well) were treated with SB203580 (1 to 10 μ M) (a gift from SmithKline-Beecham Co.) or PD98059 (10 μ M) (Cell Signaling, Beverly, Mass.) for 2 h at 37°C in 5% CO₂. In some experiments, 150 μ l of medium was removed from each well and replaced with 200 μ l of fresh medium, resulting in a fivefold dilution of the inhibitors at the end of the pretreatment.

After mixing, cells were transferred to wells of a flat-bottom 96-well plate coated with an anti-TCR MAb (H57; 1 to 10 μ g/ml). After 72 h of culturing, proliferation was assessed by using a terminal pulse of 18.5 kBq of [³H]thymidine for 8 h. The proliferative response of CD4⁺ T cells to Mls-1^a or Mls-1^b was evaluated by culturing CD4⁺ T cells (5×10^4) with mitomycin C-treated CBA/J or CBA/Ca spleen cells (0.03×10^5 to 1×10^5). Staining of cells with 5,6-carboxyfluorescein diacetate succinimidyl ester (CFSE) was performed as previously described (20). After culturing, cells were stained with a phycoerythrin-anti-CD4 MAb (eBioscience, San Diego, Calif.) and analyzed by using a FACScan (Becton Dickinson and Co., Mountain View, Calif.).

Th1 and Th2 cells were prepared as previously described (42). Briefly, CD4⁺ T cells (10^5 /ml) purified from OT-II mice were cultured with mitomycin C-treated C57BL/6 spleen cells (4×10^5 /ml) and ovalbumin (amino acids 323 to 339) peptide (10 μ M) for 7 days. Th1 mixtures contained IL-2 (20 U/ml), IL-12 (20 ng/ml), and anti-IL-4 MAb 11B11 (10 μ g/ml). Th2 mixtures contained IL-2 (20 U/ml), IL-4 (100 U/ml), anti-gamma interferon MAb R4-6A2 (15 μ g/ml), and anti-IL-12 MAb C17.8 (10 μ g/ml).

Western blot analysis. CD4⁺ T cells (3×10^5) were plated on an anti-TCR MAb-coated 96-well plate, centrifuged briefly, and cultured for 5 to 90 min. At the end of the culture period, cells from four wells were combined and resuspended in 50 μ l of lysis buffer A (1% Triton X-100, 10 mM Tris-HCl [pH 7.4], 150 mM NaCl, 1 mM EDTA, 1 mM phenylmethylsulfonyl fluoride). After centrifugation, the lysate was size fractionated by sodium dodecyl sulfate (SDS)-12.5% polyacrylamide gel electrophoresis (PAGE) and transferred to a polyvinylidene difluoride membrane by electrotransfer. The membrane was incubated in blocking buffer (Tris-buffered saline containing 5% skim milk and 0.1% Tween 20) and then probed with anti-phospho-ERK, anti-ERK, anti-phospho-JNK, anti-JNK, anti-phospho-p38, and anti-p38 antibodies (Cell Signaling), anti-p27/kip1 antibody (Transduction Laboratories, San Diego, Calif.), anti-TAB1 antibody (Santa Cruz Biotechnology, Santa Cruz, Calif.), or antiactin antibody (Sigma-Aldrich, St. Louis, Mo.). After washing, the membrane was incubated with horseradish peroxidase (HRP)-conjugated anti-rabbit immunoglobulin antibody or with biotin-conjugated anti-mouse immunoglobulin antibody, followed by HRP-streptavidin. Proteins were visualized by using an ECL kit according to the manufacturer's protocol (Amersham Biosciences Corp., Piscataway, N.J.).

Recombinant TAB1 protein was produced in *Escherichia coli*. A fragment of cDNA corresponding to human TAB1 (amino acids 1 to 312) (TAB1₁₋₃₁₂) was amplified by PCR with human TAB1 cDNA (kindly provided by K. Matsumoto, Nagoya University, Nagoya, Japan) (34), subcloned into pQE32, expressed in *E. coli*, and purified as previously described (44). We confirmed the specificity of the anti-TAB1 antibody by absorption with this recombinant TAB1₁₋₃₁₂ protein.

In vitro kinase assays. Cells were lysed in ice-cold lysis buffer B (1% Triton X-100, 10 mM Tris [pH 7.5], 150 mM NaCl, 2 mM EGTA, 50 mM β -glycerophosphate, 2 mM Na₃VO₄, 10 mM NaF, 1 mM dithiothreitol, 1 mM phenylmethylsulfonyl fluoride, 10 μ g of leupeptin/ml). After centrifugation, the lysate was precipitated with anti-ERK1 or anti-p38 α antibody and protein A-Sepharose, washed three times with lysis buffer B, and washed twice with kinase buffer (20 mM HEPES [pH 7.5], 20 mM MgCl₂, 20 mM MnCl₂, 2 mM dithiothreitol, 25 mM β -glycerophosphate, 100 nM Na₃VO₄). Samples were resuspended in 18 μ l of kinase buffer containing 1 μ g of maltose-binding protein or 1 μ g of ATF2

peptide (Upstate Biotechnology, Lake Placid, N.Y.) in the presence of 20 μ M cold ATP and 0.72 MBq of [γ -³²P]ATP (NEN Life Science Products), and the mixtures were incubated for 30 min at 37°C. The reactions were terminated by the addition of SDS sample buffer and boiling. Phosphorylation of the proteins was detected by autoradiography and quantified by using Image Analyzer BAS-5000 (Fuji Film, Tokyo, Japan). The amounts of precipitated proteins were confirmed by probing the blots with specific antibodies.

Cytokine assays. CD4⁺ T cells (3×10^5) were cultured for 48 h (for IL-2) or 36 h (for IL-10) on plates coated with an anti-TCR MAb. An anti-IL-2 receptor α -chain MAb (3C7) was added to the culture for the IL-2 assay to inhibit the consumption of IL-2 (24). The levels of cytokines in the supernatants were determined by a sandwich enzyme-linked immunosorbent assay (ELISA). IL-2 was measured by using MAb JES6-1A12 as the capture antibody and biotinylated MAb JES6-5H4 as the detecting antibody according to the manufacturer's directions (all MAbs for the ELISA were from BD Biosciences Pharmingen). IL-10 was measured by using MAb JESS-2A5 and biotinylated MAb SXC-1.

RT-PCR. Total cellular RNA was extracted by the acid-guanidium thiocyanate-phenol-chloroform method with Isogen (Nippon-Gene, Tokyo, Japan). RNA (1 μ g) was reverse transcribed by using Moloney murine leukemia virus reverse transcriptase (RT) (Sawaday Technology, Tokyo, Japan). Aliquots of the mixture (0.1 μ g of cDNA) then were amplified by using *Taq* DNA polymerase (Promega, Mannheim, Germany).

The following primer pairs were used for the PCR: IL-2 forward, 5'-AACA GCGCACCACCTTCAA-3'; IL-2 reverse, 5'-TTGAGATGATGCTTTGACA-3'; IL-10 forward, 5'-TCAAACAAAGGACCAGCTGGACAACATACTG-3'; IL-10 reverse, TAB1 forward, 5'-GCTGGACAGTACCTGCTCTCTGCG-3'; TAB1 reverse, 5'-CTCCTGGTTGGCAGTCCAGGCCAT-3'; GATA3 forward, 5'-AGAACC GGCCCTTATCAA-3'; GATA3 reverse, 5'-AGTTTCGC GCAGGATGTC-3'; T-bet forward, 5'-CAACAACCCCTTTGCCAAAG-3'; T-bet reverse, 5'-TCCCAAGCAGTTGACAGT-3'; G3PDH forward, 5'-AC CAGATCCATGCCATCAC-3'; and G3PDH reverse, 5'-TCCACCACCTG TGCTGTA-3'.

Samples were amplified for 25 cycles with the IL-2, T-bet, GATA3, and G3PDH probes (94°C for 60 s, 58°C for 60 s, and 72°C for 60 s); for 36 cycles with the IL-10 probe (94°C for 30 s, 56°C for 50 s, and 72°C for 90 s); and for 18, 24, 30, and 36 cycles with the TAB1 probe (94°C for 30 s, 53°C for 50 s, and 72°C for 90 s). The PCR products were separated by using 2% agarose gels and visualized by staining with ethidium bromide.

Retroviral transduction. The cDNA encoding human TAB1 (a gift from K. Matsumoto, Nagoya University) (34) was subcloned into retroviral vector pMSCVneo (BD Biosciences Clontech, Palo Alto, Calif.). Packaging cell line PT67 was transfected with vector pMSCVneo alone (mock) or with pMSCVneo containing full-length TAB1 cDNA by using DMRIE-C reagent (Invitrogen, Carlsbad, Calif.), and the supernatant was harvested after 36 h of culturing. T-cell hybridoma 2B4 (31) was infected with the retrovirus-containing supernatant in the presence of Polybrene (8 μ g/ml) for 60 to 72 h. The resulting cell line (TAB1-2B4) was cultured in the presence of G418 (500 μ g/ml) to obtain stably transduced cell lines. cDNAs encoding mouse wild-type p38 and dominant-negative p38 (p38DN) were kindly provided by Roger J. Davis (University of Massachusetts) (29) and were subcloned into retroviral vector pMSCVpuro (BD Biosciences Clontech). Supernatant was collected from cell line PT67 transfected with the resulting plasmids and used to produce Mock-2B4 (vector alone) and p38DN-2B4 cell lines. The cell lines were cultured in the presence of G418 (500 μ g/ml) and puromycin (1 μ g/ml) to obtain stably transduced cell lines.

Reporter gene assay. A luciferase reporter plasmid containing the mouse IL-10 promoter (-1536 to +64) was a kind gift from S. Smale (University of California at Los Angeles) (7). The reporter gene assay was performed by using a dual luciferase reporter assay system (Promega). Jurkat cells (10^7) were transiently transfected with vector pCMV, which contained no insert or human TAB1 cDNA (10 μ g), in addition to internal control phRL-TK (3 μ g) and luciferase reporter construct pGL2B driven by the mouse IL-10 promoter (-1536 to +64) (5 μ g) by electroporation (220 V, 65 ms) with a Gene Pulser (Bio-Rad, Hercules, Calif.). After 24 h of culturing, cells were treated with phorbol myristate acetate (PMA) (100 ng/ml) and ionomycin (2 μ M) for 18 h, and cell lysates were analyzed for luciferase activity according to the manufacturer's instructions (Promega).

RESULTS

Activation of p38 MAPK in anergic T cells. T-cell anergy was induced in the majority of T cells in V β 8.1-tg mice after inoculation of T-cell-depleted CBA/J (Mls-1^a) spleen cells. Both

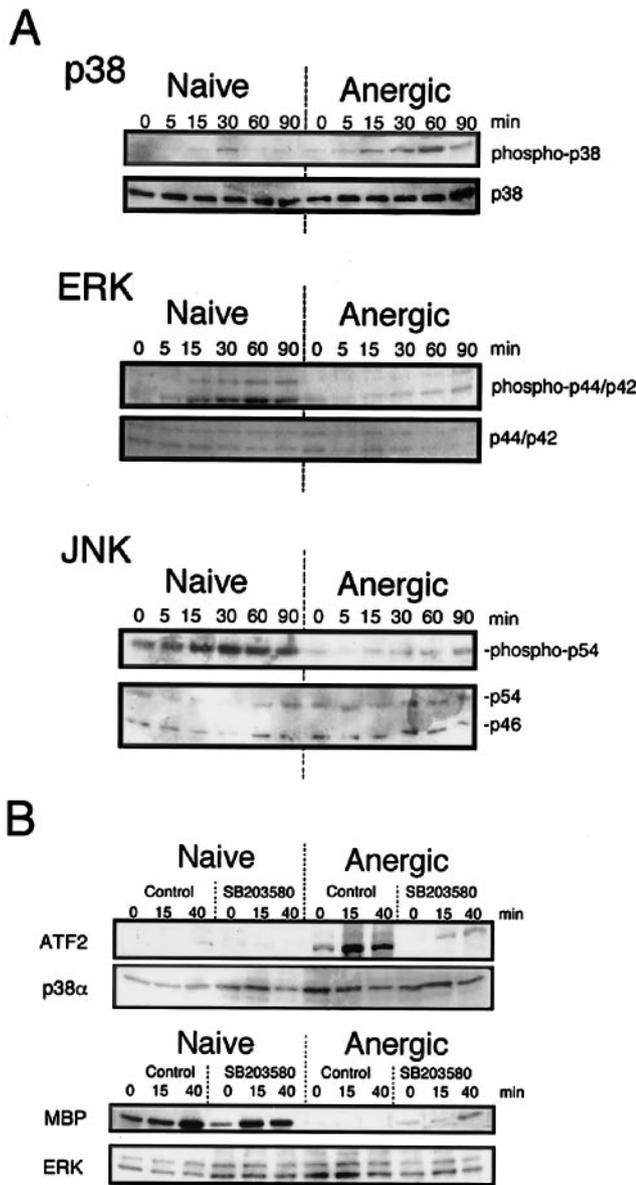


FIG. 1. Anergic CD4⁺ T cells show decreased activation of ERK and JNK and increased activation of p38 MAPK in response to TCR stimulation. (A) CD4⁺ T cells were purified from untreated Vβ8.1-tg mice (naive) or Mls-1^a-treated Vβ8.1-tg mice (anergic) by treatment of lymph node cells with anti-CD8 MAb and complement, followed by nylon wool column enrichment. Cells were cultured for 3 h at 37°C in complete medium to decrease basal levels of MAPK activity. Cells then were placed in anti-TCR MAb (10 μg/ml)-coated plates for various times and lysed, and the levels of phosphorylated and total MAPKs were determined by immunoblotting with specific antibodies. Representative results from three independent experiments are shown. (B) In vitro immune complex kinase assay of p38α and ERK1. Naive or anergic CD4⁺ T cells were prepared as described for panel A and placed in anti-TCR MAb (10 μg/ml)-coated plates in the presence or absence of SB203580 for various times. Cell lysates were immunoprecipitated with anti-p38α or anti-ERK1 antibody, and in vitro kinase assays of the immunoprecipitates were performed with ATF2 and maltose-binding protein (MBP) as substrates for p38α and ERK1, respectively. Proteins were separated by SDS-PAGE and analyzed by autoradiography. The quantities of p38α and ERK1 were assessed by probing the immunoblots with antibodies specific for p38α or ERK1. Representative results from three independent experiments are shown.

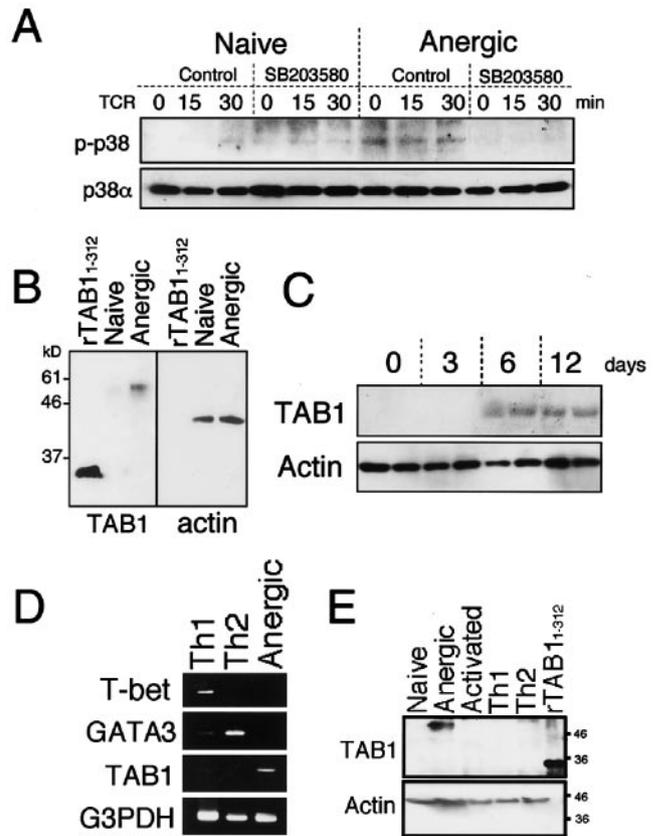


FIG. 2. Anergic CD4⁺ T cells express TAB1. (A) Naive and anergic CD4⁺ T cells were purified by using BD IMag-anti-mouse CD4 MAb, cultured with or without SB203580 (10 μM) for 2 h before stimulation, and plated in wells coated with anti-TCR MAb (10 μg/ml) for 0, 15, or 30 min. Cells were harvested and lysed, and the levels of phospho-p38 and total p38α were determined by immunoblotting with specific antibodies. Representative results from three independent experiments are shown. (B) Lysates from naive and anergic CD4⁺ T cells (4 × 10⁶/lane) were separated by SDS-12.5% PAGE and examined for the presence of TAB1 by immunoblotting with specific antibodies. After stripping, the membrane was immunoblotted with antiactin antibody. rTAB1₁₋₃₁₂, N-terminal 312-amino-acid fragment of recombinant TAB1. (C) Expression of TAB1 during the induction of T-cell anergy. CD4⁺ T cells were purified from individual mice on the indicated days after Mls-1^a inoculation. Lysates from two samples per group were separated by SDS-12.5% PAGE and examined for the expression of TAB1 by immunoblotting. Representative results from two independent experiments are shown. (D) CD4⁺ T cells from OT-II mice were cultured for 7 days in Th1 and Th2 mixtures. Anergic CD4⁺ T cells were prepared from Vβ8.1-tg mice 14 days after Mls-1^a inoculation (anergic), RNA was extracted from each cell type, and RT-PCR analysis of T-bet, GATA3, TAB1, and G3PDH was performed as described in Materials and Methods. (E) CD4⁺ T cells were prepared from Vβ8.1-tg mice (naive). Anergic, Th1, and Th2 cells were prepared as described for panel D. CD4⁺ T cells from Vβ8.1-tg mice were stimulated with anti-TCR MAb (0.5 μg/ml) and soluble anti-CD28 MAb (1 μg/ml) for 24 h in vitro (activated). Total cell lysates were separated by SDS-12.5% PAGE, blotted, and probed with anti-TAB1 antibody. A recombinant TAB1 fragment (rTAB1₁₋₃₁₂) was used as a positive control.

proliferation and IL-2 production in response to Mls-1^a or anti-TCR MAb were impaired in CD4⁺ T cells from Mls-1^a-inoculated Vβ8.1-tg mice (4, 48) (see Fig. 3A). CD4⁺ T cells from naive Vβ8.1-tg mice (naive CD4⁺ T cells) and from mice

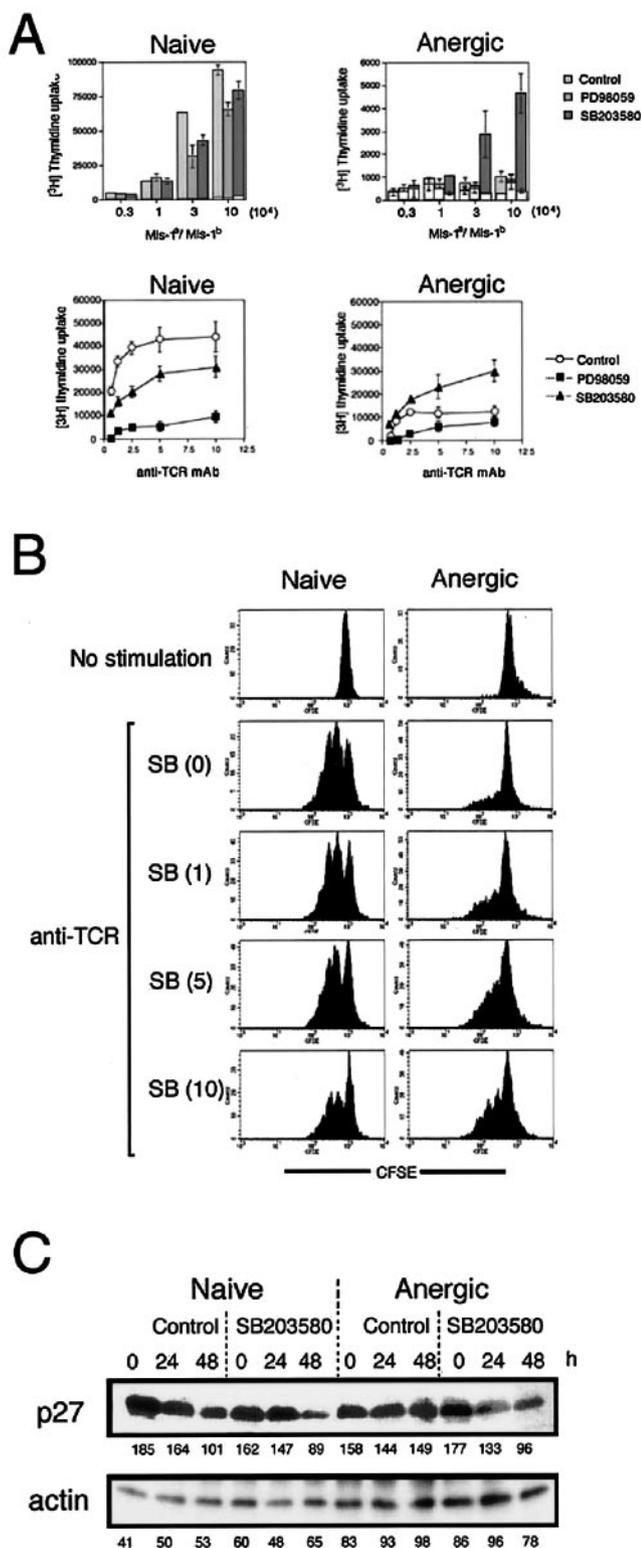


FIG. 3. T-cell proliferation is inhibited by p38 MAPK activity in anergic CD4⁺ T cells. (A) Naive and anergic CD4⁺ T cells were pretreated with dimethyl sulfoxide (control), PD98059 (10 μM), or SB203580 (10 μM) for 2 h and were cocultured with mitomycin C-treated T-cell-depleted CBA/J (Mls-1^a) or CBA/Ca (Mls-1^b) spleen cells in the presence or absence of a lower concentration (2 μM) of the inhibitors (top) or stimulated with anti-TCR MAb (0.625 to 10 μg/ml) in the same manner (bottom). Cultures were maintained for 3 days,

that were injected with T-cell-depleted CBA/J spleen cells (anergic CD4⁺ T cells) expressed comparable levels of TCR and CD4 (4, 48). The proportions of CD25⁺ cells in the naive and anergic CD4⁺-T-cell populations were ~6% and ~13%, and those of CD44^{low} CD45RB^{high} cells were ~82% and ~35%, respectively.

To determine the activities of MAPKs, naive and anergic CD4⁺ T cells were cultured for 3 h to decrease the basal level of MAPK activation and were stimulated with plate-bound anti-TCR MAb. The activation status of MAPKs in these T cells was evaluated by immunoblotting with phospho-specific antibodies (Fig. 1A). Phosphorylation of p38 after TCR stimulation was enhanced in anergic CD4⁺ T cells over that in naive CD4⁺ T cells, which showed limited phosphorylation. In contrast, the phosphorylation of ERK and JNK in anergic CD4⁺ T cells was severely reduced, consistent with the previous report of T cells that were anergized in vitro (10, 19) or in vivo (25).

Next, we evaluated the activity of p38α and ERK by using an in vitro kinase assay (Fig. 1B). The kinase activity of p38α in anergic CD4⁺ T cells was detectable after 3 h in resting cultures and was enhanced by stimulation with anti-TCR MAb. In contrast, the activity of ERK was hardly detectable in anergic CD4⁺ T cells but was apparent in naive CD4⁺ T cells. These results indicated the marked shift in the activation status of MAPKs between naive and anergic CD4⁺ T cells.

TAB1 expression and TAB1-mediated phosphorylation of p38α. The enhanced activation of p38α in anergic CD4⁺ T cells suggested to us that the expression of the signaling molecule upstream of p38α might be altered. There are at least two separate mechanisms that can activate p38α. One is dependent on a prototypical kinase cascade in which p38α is phosphorylated by MKK3/6; the other is dependent on TAB1-mediated p38α phosphorylation (13). One way to distinguish between them is to examine the sensitivity of their own phosphorylation to an inhibitor of p38α and β, SB203580. The former pathway is typically insensitive to SB203580 treatment, while the latter is sensitive (13). Therefore, we examined whether the phosphorylation of p38 in anergic CD4⁺ T cells is inhibited by SB203580 by immunoblotting (Fig. 2). SB203580 inhibited the level of phospho-p38 in anergic but not in naive CD4⁺ T cells, suggesting that the phosphorylation of p38 in anergic T cells was dependent on its own activation. Therefore,

and proliferation was assessed by measuring [³H]thymidine uptake. The responses of T cells to Mls-1^a (CBA/J) and to Mls-1^b (CBA/Ca) are represented by shaded and open columns, respectively. Representative results from 10 independent experiments are shown. Error bars indicate standard deviations. (B) Naive and anergic CD4⁺ T cells were labeled with CFSE, left untreated or treated with various doses of SB203580 (SB) for 2 h, and cultured in the presence of one-fifth the indicated concentrations of SB203580 for 3 days in wells of anti-TCR MAb-coated plates. Cells then were stained with phycoerythrin-anti-CD4 MAb and analyzed by flow cytometry. Representative results from three independent experiments are shown. (C) Naive and anergic CD4⁺ T cells were stimulated with anti-TCR MAb (10 μg/ml) for the indicated times in the presence or absence of SB203580, and the expression of p27/kip1 was analyzed by Western blotting. The expression of actin was used as a control. Arbitrary densitometric units for these bands were analyzed by using NIH Image software. Representative results from three independent experiments are shown.

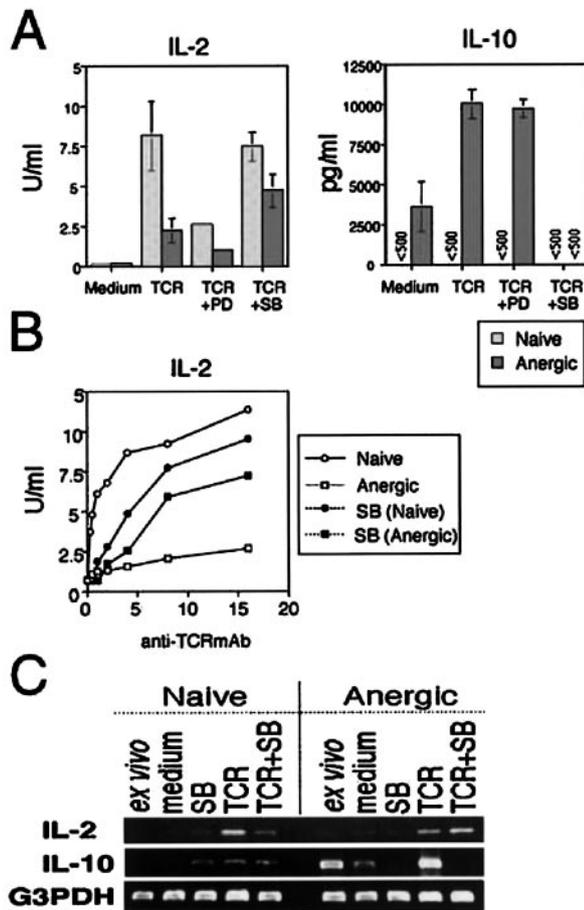


FIG. 4. The production of IL-2 is inhibited by p38 activity in anergic CD4⁺ T cells, while IL-10 production is p38 dependent. (A) CD4⁺ T cells from untreated (naive) or CBA/J-inoculated (anergic) Vβ8.1-tg mice were cultured in plates coated with anti-TCR MAb in the presence or absence of SB203580 (SB) or PD98059 (PD). Culture supernatants were collected 48 h (for IL-2) or 36 h (for IL-10) later, and the levels of IL-2 and IL-10 were determined by an ELISA. An anti-IL-2 receptor α-chain MAb (3C7) was added to the culture for the IL-2 assay to inhibit the consumption of IL-2. Representative results from three independent experiments are shown. Error bars indicate standard deviations. (B) Naive and anergic CD4⁺ T cells were cultured in plates coated with various doses of anti-TCR MAb in the presence or absence of SB203580, and the levels of IL-2 in supernatants were determined as described for panel A. Representative results from three independent experiments are shown. (C) Expression of mRNAs for IL-2, IL-10, and G3PDH in naive or anergic CD4⁺ T cells. Naive or anergic CD4⁺ T cells were plated on anti-TCR MAb (10 μg/ml)-coated plates in the presence or absence of SB203580 as described in the legend to Fig. 1A for 18 h, and total RNA was purified from each well. cDNA samples (0.1 μg) from RNA were amplified for 25 cycles for IL-2 and G3PDH or for 36 cycles for IL-10 by RT-PCR. Representative results from three independent experiments are shown.

we examined whether TAB1 is expressed in these CD4⁺ T cells by RT-PCR (data not shown) and immunoblotting (Fig. 2B). Anergic CD4⁺ T cells showed clear expression of TAB1, in contrast to little expression in naive CD4⁺ T cells. Thus, we concluded that naive CD4⁺ T cells do not express significant levels of TAB1. The study of expression kinetics indicated that the expression of TAB1 in CD4⁺ T cells became detectable 6

days after superantigen inoculation, corresponding to the induction of T-cell anergy (Fig. 2C).

We also examined the expression of TAB1 in T cells that were activated *in vitro* or that were cultured with Th1 and Th2 mixtures (Fig. 2D and E). CD4⁺ T cells cultured with Th1 and Th2 mixtures preferentially expressed T-bet and GATA3, respectively, transcription factors regulating Th1 and Th2 differentiation. The expression of TAB1 was, however, barely detectable by RT-PCR and Western blot analysis. Also, CD4⁺ T cells activated by stimulation with anti-TCR MAb plus anti-CD28 MAb for 24 h did not express TAB1. These results indicated that TAB1 is specifically expressed in anergic CD4⁺ T cells and can induce the phosphorylation of p38α.

Effect of MAPK inhibitors on anergic T cells. To investigate the role of the activated p38 pathway in the maintenance of anergy, we examined whether SB203580 can modulate the anergic state of CD4⁺ T cells. Treatment of anergic T cells with SB203580 inhibited the activity of p38 but also slightly increased ERK activity (Fig. 1B). The proliferative responses of naive and anergic CD4⁺ T cells were evaluated in the presence of MEK1 inhibitor PD98059 or SB203580. These cells were pretreated with PD98059 (10 μM) or SB203580 (10 μM) for 2 h and were stimulated with Mls-1^a or plate-bound anti-TCR MAb in the presence of a lower concentration of the inhibitors (2 μM). Both inhibitors suppressed the proliferative responses of naive CD4⁺ T cells to TCR occupancy. However, SB203580 augmented the proliferative responses of anergic CD4⁺ T cells in response to Mls-1^a and anti-TCR MAb (Fig. 3A and B). We obtained similar results with another p38 kinase inhibitor, FR167653 (32) (data not shown).

To monitor the responses of CD4⁺ T cells at the population level, we labeled the cells with fluorescent dye CFSE and examined their proliferation by measuring the diminution of the CFSE level (Fig. 3B) (20). Proliferation of anergic CD4⁺ T cells in response to anti-TCR MAb was augmented by SB203580 in a dose-dependent manner. We also examined the level of a p27/kip1 cyclin-dependent kinase inhibitor which was reported to regulate the blockade of the cell cycle in anergic CD4⁺ T cells (6). The level of p27/kip1 was reduced 48 h after stimulation of naive but not anergic CD4⁺ T cells. In the presence of SB203580, however, the expression of p27/kip1 was reduced 48 h after TCR stimulation in anergic CD4⁺ T cells (Fig. 3C), suggesting that the cells produced IL-2 and thus induced the degradation of p27/kip1 (Fig. 4).

Cytokine production by anergic CD4⁺ T cells. We next examined the production of cytokines in the presence of MAPK inhibitors (Fig. 4). PD98059 inhibited the production of IL-2 in response to anti-TCR MAb in both naive and anergic CD4⁺ T cells. SB203580 partially inhibited the production of IL-2 by naive CD4⁺ T cells but also enhanced IL-2 production by anergic CD4⁺ T cells (Fig. 4A). We also evaluated the production of IL-10, since *in vivo*-anergized CD4⁺ T cells could produce IL-10 (8, 23, 37). Although naive CD4⁺ T cells produced barely detectable levels of IL-10 in our assay, anergic CD4⁺ T cells secreted IL-10 without stimulation and produced higher levels of IL-10 after culturing with anti-TCR MAb. SB203580 completely inhibited the production of IL-10 by anergic CD4⁺ T cells, indicating that the production of IL-10 by anergic CD4⁺ T cells is p38 dependent. To determine whether the level of cytokine production was regulated at the

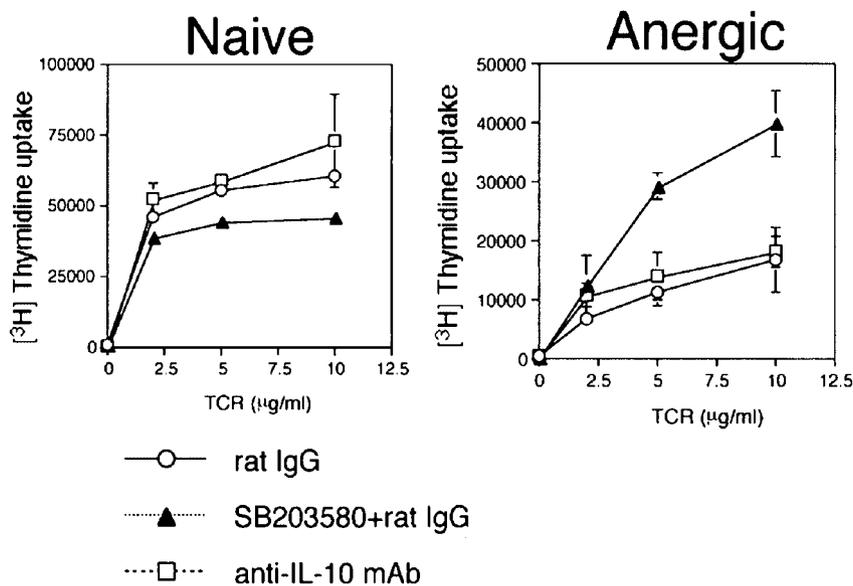


FIG. 5. The maintenance of unresponsiveness in anergic $CD4^+$ T cells is independent of IL-10 production. Naive and anergic $CD4^+$ T cells were cultured in plates coated with anti-TCR MAb (0 to 10 $\mu\text{g/ml}$). The cultures were maintained in the presence of control rat immunoglobulin G (IgG; 20 $\mu\text{g/ml}$), anti-IL-10 MAb (20 $\mu\text{g/ml}$), or SB203580 (2 μM) plus control rat IgG for 72 h, and proliferation was assessed by measuring the incorporation of [^3H]thymidine. Cells cultured in the presence of SB203580 were incubated with the inhibitor at a final concentration of 10 μM for 2 h before TCR stimulation. Representative results from three independent experiments are shown. Error bars indicate standard deviations.

level of mRNA expression, RT-PCR analysis of IL-2 and IL-10 was performed (Fig. 4C). Anergic $CD4^+$ T cells expressed a high level of IL-10 mRNA *ex vivo* and maintained this high level upon stimulation *in vitro* with anti-TCR MAb. The expression of IL-10 mRNA by anergic $CD4^+$ T cells was clearly inhibited by SB203580.

One possible explanation for the increased proliferation and IL-2 production of anergic $CD4^+$ T cells in the presence of SB203580 was a reduction in IL-10 production, which might otherwise have suppressed the response of these cells. To exclude this possibility, we attempted to determine whether the addition of neutralizing anti-IL-10 MAb could induce the proliferation of anergic $CD4^+$ T cells (Fig. 5). Anti-IL-10 MAb (20 $\mu\text{g/ml}$) had little effect on the proliferative response of anergic $CD4^+$ T cells. The concentration of anti-IL-10 MAb used in this assay was sufficient to neutralize the activity of IL-10, since this concentration of anti-IL-10 MAb was able to completely block the inhibitory effect of recombinant IL-10 on the costimulatory activity of anti-CD28 MAb on the $CD4^+$ -T-cell proliferative response to anti-TCR MAb (data not shown) (1). Thus, the effect of SB203580 on the proliferation of anergic $CD4^+$ T cells was not simply due to the inhibition of IL-10 production by these cells.

TAB1 negatively regulates T-cell activation. We next examined the effect of the forced expression of TAB1 on T-cell function. Our initial attempt to transfer DNA into anergic $CD4^+$ T cells by using a retrovirus was unsuccessful, since anergic T cells did not replicate DNA. Thus, we introduced TAB1 cDNA into T-cell hybridoma 2B4 (24), which does not express endogenous TAB1 (Fig. 6A), and generated stable cell lines containing vector alone (Mock-2B4) or TAB1 (TAB1-2B4). As expected, p38 α kinase was constitutively active and

phospho-p38 was detected in TAB1-2B4 cells (Fig. 6B and C). Treatment of TAB1-2B4 cells with SB203580 resulted in down-regulation of the phospho-p38 level, indicating that the phosphorylation of p38 depends on its own activity. RT-PCR of IL-2 and IL-10 mRNAs indicated that TAB1-2B4 cells expressed lower levels of IL-2 mRNA both before and after TCR stimulation compared with Mock-2B4 cells (Fig. 6D). IL-10 mRNA was detectable in TAB1-2B4 cells but not in Mock-2B4 cells. The production of IL-2 in the supernatant was reduced in TAB1-2B4 cells after stimulation with anti-TCR MAb as well as PMA and ionomycin (Fig. 6E). The IL-10 reporter assay showed that the expression of TAB1 induced IL-10 promoter activity in Jurkat cells and that this effect was augmented by stimulation with PMA and ionomycin (Fig. 6F).

The effect of TAB1 on the activation of ERK was investigated by immunoblotting (Fig. 7). We used PMA to activate ERK kinase in 2B4 cells, since the phosphorylation of ERK in 2B4 cells after stimulation with anti-TCR MAb was barely detectable. The stimulation of TAB1-2B4 cells with various doses of PMA induced phospho-ERK to levels lower than those in Mock-2B4 cells (Fig. 7A). However, these cells showed increased levels of phospho-ERK in the presence of SB203580 (Fig. 7B). To confirm that this recovery of ERK activation in the presence of SB203580 resulted from the inhibition of p38, we introduced p38DN into Mock-2B4 or TAB1-2B4 cells and generated stable cell lines containing vector alone (Mock/Mock-2B4 and TAB1/Mock-2B4) or p38DN (Mock/p38DN-2B4 and TAB1/p38DN-2B4). In TAB1-2B4 cells, the levels of ERK protein expression were up-regulated for unknown reasons, but it was clear that PMA-induced ERK activation was up-regulated in TAB1/p38DN-2B4 cells compared with TAB1/Mock-2B4 cells. We also examined the acti-

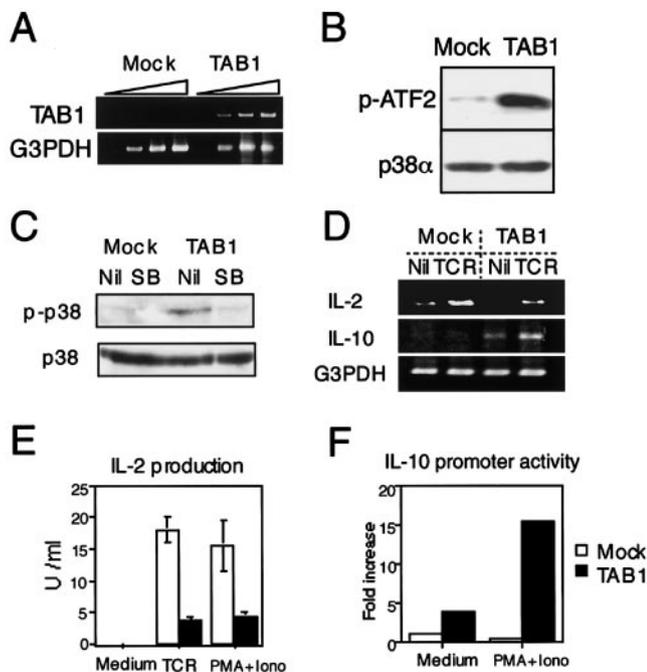


FIG. 6. TAB1 inhibits IL-2 and increases IL-10 production. (A) T-cell hybridoma 2B4 was transduced with retrovirus that contained the vector alone (Mock) or human TAB1 (TAB1), and stable cell lines were selected in the presence of G418 (500 μ g/ml). RNA was prepared from each cell line and was subjected to RT-PCR analysis for TAB1 mRNA expression. PCR was performed for 18, 24, 30, and 36 cycles (as indicated by the triangles over the lanes). (B) p38 α was immunoprecipitated with specific antibodies from lysates of Mock-2B4 cells (Mock) or TAB1-2B4 cells (TAB1) and was subjected to an *in vitro* kinase assay with ATF2₁₉₋₉₆ peptide as a substrate. The quantity of p38 α was assessed by immunoblotting with anti-p38 α antibody. (C) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were incubated with (SB) or without (Nil) SB203580 (10 μ M) for 2 h, lysed, and subjected to immunoblotting with anti-phospho-p38 and anti-p38 α antibodies. (D) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were cultured for 18 h in plates coated (TCR) or not coated (Nil) with anti-TCR MAb (10 μ g/ml) and were subjected to RT-PCR analysis for IL-2, IL-10, and G3PDH mRNAs. (E) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were cultured with anti-TCR MAb (TCR) or with PMA (20 ng/ml) and ionomycin (Iono; 1 μ M) for 24 h. The levels of IL-2 in the supernatants were determined by an ELISA. Error bars indicate standard deviations. (F) Jurkat cells were transiently transfected with pGL2B luciferase reporter constructs driven by the mouse IL-10 promoter (-1536 to +64), with vector pCMV, which contained either no insert (Mock) or human TAB1 cDNA (TAB1), and with internal control phRL-TK. Cells were cultured with PMA (100 ng/ml) and ionomycin (2 μ M) for 18 h, and luciferase activity was measured.

vation of JNK in these cells, since the activation of JNK was also impaired in anergic CD4⁺ T cells. The phosphorylation of JNK after PMA treatment was reduced in TAB1-2B4 cells compared with Mock-2B4 cells. However, unlike that of ERK, the phosphorylation of JNK was not up-regulated in TAB1/p38DN-2B4 cells (Fig. 7C), suggesting that the reduction in JNK phosphorylation in TAB1-2B4 cells was independent of p38. Further study is under way to determine the molecular basis of TAB1-mediated JNK inactivation.

Parallel to the levels of ERK activation, TAB1-2B4 cells produced less IL-2 mRNA after stimulation with PMA plus ionomycin and more IL-2 mRNA in the presence of SB203580

or p38DN (Fig. 7D and E). These results indicated that the forced expression of TAB1 in T cells resulted in the constitutive activation of p38 α and the inhibition of ERK activation, culminating in the reduced expression of IL-2 after T-cell activation and the induction of IL-10 expression. This response pattern was similar to that of anergic CD4⁺ T cells, which showed reduced levels of IL-2 production and proliferation even in response to various doses of PMA (1 to 25 ng/ml) and ionomycin (2 μ M) (data not shown). Finally, the study of TCR-stimulated IL-2 production in these hybridoma cell lines indicated that the inhibition of IL-2 production by the introduction of TAB1 was mediated by the p38 pathway (Fig. 7F).

DISCUSSION

T-cell anergy induced by superantigen is maintained primarily by intrinsic mechanisms and is independent of regulatory T cells (3, 23). In fact, Mls-1^a-induced anergy could be maintained without CD25⁺ regulatory T cells (data not shown). In this study, we investigated the molecular basis of this anergy and showed that TAB1 is expressed in CD4⁺ T cells anergized *in vivo* and is not expressed in naive or activated CD4⁺ T cells implying that TAB1 is a unique functional molecule expressed in anergic T cells. TAB1 was originally identified as an activator of the MAPK kinase kinase, TAK1, that induces MAPK activation cascade in transforming growth factor β signaling pathway (34). It is a scaffolding protein that can also bind to p38 α and induces its phosphorylation. This TAB1-mediated phosphorylation of p38 α requires its intrinsic kinase activity, and is thus inhibited by SB203580 (13). We speculate that TAB1-mediated signaling in anergic CD4⁺ T cells could regulate the distal TCR signaling events, such as the activation of MAPKs, as well as the production of cytokines after TCR engagement. In fact, forced expression of TAB1 in a T-cell hybridoma induced the activation of p38 α MAPK prior to T-cell activation, promoted IL-10 synthesis and was inhibitory to IL-2 expression after activation with PMA and ionomycin. In anergic CD4⁺ T cells and TAB1-2B4 cells, p38 α MAPK is activated prior to TCR stimulation, which can be inhibited by SB203580, indicating that phosphorylation is dependent on its own kinase activity.

We propose two possibilities for the SB203580-sensitive phosphorylation of p38. One is that the activation of p38 α itself is TAB1-mediated autophosphorylation, as was proposed by Ge et al. (13). The second possibility is that TAB1 mediates feedback control of the MAPK pathway by p38 α (9). Further studies are under way to distinguish between these possibilities. In either cases, these studies imply that the maintenance of Mls-1^a-induced T-cell anergy is regulated at the level of MAPK signaling. In fact, the production IL-2 by anergic CD4⁺ T cells in response to various concentrations of PMA and ionomycin was reduced when compared with control (data not shown), although some of the previous studies showed that anergic T cells responded normally to PMA and ionomycin (33). These differences might reflect the different degrees of T-cell anergy that can be induced *in vivo* (40, 45).

Our study showed that the effect of SB203580 treatment on MAPKs signaling and cytokine secretion differed significantly between naive and anergic CD4⁺ T cells. In naive CD4⁺ T cells, SB203580 had little effect on ERK activation and slightly

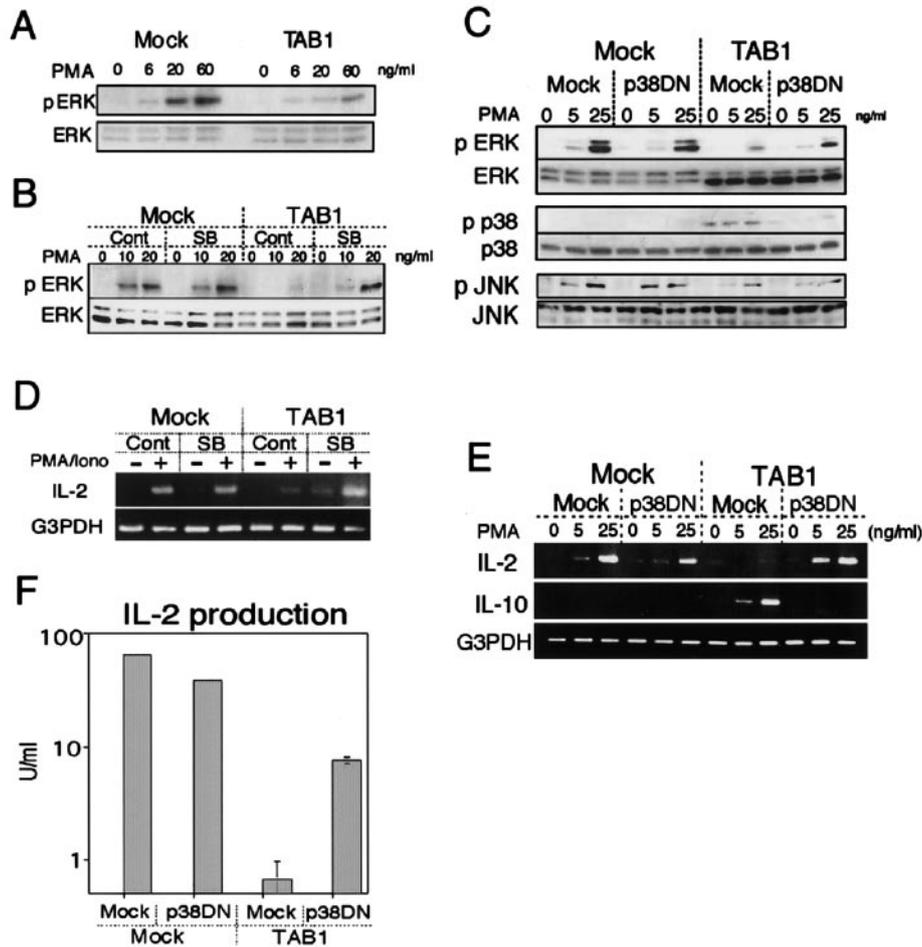


FIG. 7. The inhibition of ERK kinase activity by TAB1 is dependent on p38. (A) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were cultured with various doses of PMA (0 to 60 ng/ml) for 10 min, and the levels of phosphorylated ERK and total ERK were determined by immunoblotting with specific antibodies. (B) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were cultured in the presence (SB) or absence (Cont) of SB203580 (10 μ M) for 2 h and subsequently stimulated with PMA (0, 10, or 20 ng/ml) for 10 min. Cells were lysed, and the levels of phosphorylated ERK and total ERK were determined by immunoblotting with specific antibodies. (C) Mock-2B4 cells were transduced with a retroviral supernatant containing empty vector (Mock/Mock) or p38DN (Mock/p38DN) to establish stable cell lines. TAB1-2B4 cells were also transduced to establish cell lines expressing empty vector (TAB1/Mock) or p38DN cDNA (TAB1/p38DN). These cell lines were stimulated with PMA (0, 5, or 25 ng/ml) for 10 min and lysed, and the levels of phosphorylated MAPks and total MAPks were determined by immunoblotting with specific antibodies. (D) Mock-2B4 cells (Mock) and TAB1-2B4 cells (TAB1) were treated or not treated with SB203580 (10 μ M) for 2 h and stimulated with PMA (20 ng/ml) and ionomycin (Iono; 2 μ M) for 8 h in the presence (SB) or absence (Cont) of SB203580, respectively. Cells were subjected to RT-PCR analysis for IL-2 and G3PDH mRNAs. Representative results from two independent experiments are shown. (E) Mock/Mock, Mock/p38DN, TAB1/Mock, and TAB1/p38DN cells were stimulated with PMA (0, 5, or 25 ng/ml) and ionomycin (2 μ M) for 8 h. Cells were subjected to RT-PCR analysis for IL-2, IL-10, and G3PDH mRNAs. Representative results from two independent experiments are shown. (F) Mock/Mock, Mock/p38DN, TAB1/Mock, and TAB1/p38DN cells were stimulated with anti-TCR MAb (H57; 10 μ g/ml) for 24 h. The levels of IL-2 in culture supernatants were determined by an ELISA. IL-2 was not detectable in cultures without anti-TCR MAb.

inhibited IL-2 production, while in anergic CD4⁺ T cells, it potentiated ERK activity and IL-2 production. Although the effect of SB203580 on other molecules, such as c-Raf and MAPK kinase kinase of the ERK pathway, has been described (16, 26), we believe that the effect of SB203580 on anergic T cells is specific to p38. Activation of c-Raf by SB203580 cannot explain its differential effects on naive and anergic CD4⁺ T cells. In addition, our study with a T-cell hybridoma indicated that the expression of TAB1 and thus the activation of p38 α is critical for the effect of SB203580 on ERK activation, because SB203580 had little effect on ERK activation in Mock-2B4

cells but potentiated ERK in TAB1-2B4 cells (Fig. 6). Furthermore, the effect of SB203580 on TAB1-2B4 cells was recapitulated by the expression of p38DN. Therefore, we concluded that the activation of the ERK pathway in anergic CD4⁺ T cells by SB203580 was caused by the release of this pathway from the negative regulatory effects of p38 α . Thus, the activation of p38 α has different effects on naive and anergic CD4⁺ T cells.

One possibility is that the constitutive activation of p38 α prior to TCR occupancy has a negative regulatory effect on the ERK pathway. The phosphorylation of p38 α in naive CD4⁺ T

cells is mediated by MAPK kinases (MKK3 and MKK6), which are activated after TCR stimulation and thus are not inhibitory to ERK, while p38 α is activated prior to TCR occupancy by TAB1-mediated mechanisms in anergic T cells. The pretreatment of anergic CD4⁺ T cells with SB203580 may release the ERK pathway from the inhibitory effect of p38 α . Alternatively, a third protein that associates with p38 α may dictate the differential outcome of p38 α activation in naive and anergic T cells. TAB1 is a 504-amino-acid protein that has a putative protein phosphatase 2C-like domain and p38 α - and TAK1-binding domains (14) and can form a complex with TAK1, TAB2, and TRAF6 (39). It is possible that the complex of TAB1 and p38 α in anergic T cells associates with a third protein, which has an inhibitory effect on the ERK pathway. Similar cross talk between two MAPK pathways has been reported in other biological systems. Singh et al. reported that p38 α negatively regulates the expression of a low-density lipoprotein receptor through an ERK signaling cascade in hepatoma cell line HepG2 through the use of a pharmacological inhibitor as well as constitutively active MKK6 (35). Others also showed that this type of inhibitory effect of p38 α on the ERK pathway could be mediated through an upstream kinase of the ERK cascades (46) or by a direct interaction between p38 and ERK (49).

CD4⁺ T cells anergized in vivo produce IL-10, a cytokine which inhibits the T-cell response and could induce anergy in peripheral T cells (15, 23, 38). Repeated inoculation of superantigen SEB in vivo induced unresponsiveness of specific T cells and IL-10 production (18). Influenza hemagglutinin-specific T cells that were rendered anergic in vivo also expressed high levels of IL-10 mRNA (8). Consistent with these studies, anergic CD4⁺ T cells in V β 8.1-tg mice produced IL-10. Anergic CD4⁺ T cells secreted IL-10 ex vivo prior to TCR stimulation and increased levels of IL-10 after TCR occupancy. The SB203580 treatment completely inhibited the production of IL-10 in anergic CD4⁺ T cells, indicating that its production depends on the activation of p38 (Fig. 3A and C). TAB1-2B4 cells also showed increased IL-10 mRNA production, similar to p38-dependent IL-10 production in human T cells (18, 27) and monocytes (12). However, the effect of the p38 inhibitor on the proliferation of anergic CD4⁺ T cells was independent of their IL-10 secretion, because neutralization of IL-10 by anti-IL-10 MAb in vitro did not rescue the proliferation of anergic CD4⁺ T cells (Fig. 4). This is consistent with the previous study, showing that a treatment of mice with neutralizing anti-IL-10 MAb prior to the challenge with superantigen did not rescue IL-2 production (37).

T-cell unresponsiveness induced by superantigen inoculation in vivo is maintained both by the autonomous anergy of cells and by cytokine-mediated suppression (3, 23). The expression of TAB1 and thus the activation of p38 α regulate the secretion of IL-2 and IL-10 in distinct ways; one is the inhibition of ERK activation culminating in the inhibition of IL-2 production, and the other is the promotion of IL-10 production, both of which contribute to the maintenance of CD4⁺ T-cell unresponsiveness. Therefore, TAB1 is a key signaling protein in the maintenance of T-cell unresponsiveness. Modulation of TAB1 activity could alter the function of anergic and regulatory T cells.

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AUTHOR'S CORRECTION

Regulation of the Maintenance of Peripheral T-Cell Anergy by TAB1-Mediated p38 α Activation

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Volume 24, no. 16, p. 6957–6966, 2004. In this paper we identified an ~56-kDa protein expressed in anergic CD4⁺ T cells by Western blotting using anti-TAB1 peptide antibody (N-19; Santa Cruz Biotechnology). This molecule was not present in naïve CD4⁺ T cells, and we concluded that TAB1 is expressed in anergic and not in naïve CD4⁺ T cells. However, our recent work indicated that this is not correct. After publication of the paper, we generated monoclonal antibodies (MAbs) specific for recombinant human TAB1 protein to confirm the expression of TAB1 in naïve and anergic CD4⁺ T cells. Contrary to our expectations, these MAbs identified an ~70-kDa molecule in naïve CD4⁺ T cells at levels similar to those in anergic CD4⁺ T cells. Subsequent experiments have shown that the ~70-kDa protein is the TAB1 molecule, because these antibodies detect a molecule of the same size in COS7 cells transfected with the expression vector containing human TAB1 cDNA and not with the control vector and, furthermore, this band is specifically absent in cells lacking the TAB1 gene.

Therefore, the ~56-kDa molecule that we identified in anergic CD4⁺ T cells by anti-TAB1 antibody (N-19) was not TAB1. The expression of TAB1 was shown in Fig. 2 of the paper, and we retract these data. We apologize for releasing and misinterpreting the immature data. While we are unable to conclude that the activation of p38 α in anergic CD4⁺ T cells is regulated by up-regulation of TAB1 protein, the main conclusions that anergic CD4⁺ T cells show enhanced p38 mitogen-activated protein kinase activity and that signals involving TAB1 could be a critical regulatory point in maintaining CD4⁺ T-cell anergy by activating p38 remain unchanged.