Interleukin (IL)-32-mediated CCAAT/Enhancer-binding Protein α (C/EBP α) Phosphorylation by Protein Kinase C δ (PKC δ) Abrogates the Inhibitory Effect of C/EBP α on IL-10 **Production***

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Background: IL-32β promotes IL-10 production in myeloid cells.

Results: IL-32 β -mediated C/EBP α serine 21 phosphorylation by PKC δ induced the dissociation of C/EBP α from IL-10 promoter, thereby promoting IL-10 production.

Conclusion: IL-32 β suppressed the inhibitory effect of C/EBP α on IL-10 production by mediating C/EBP α serine 21 phosphorylation by PKCδ.

Significance: Our data suggest that IL-32 β functions as an intracellular regulator of IL-10 production.

We previously reported that $IL-32\beta$ promotes $IL-10$ produc**tion in myeloid cells. However, the underlying mechanism remains elusive. In this study, we demonstrated that IL-32 abrogated the inhibitory effect of CCAAT/enhancer-binding protein α (C/EBPα) on IL-10 expression in U937 cells. We** observed that the phosphorylation of $C/EBP\alpha$ Ser-21 was inhibited by a PKC δ -specific inhibitor, rottlerin, or IL-32 β knockdown by siRNA and that IL-32 β shifted to the membrane from **the cytosol upon phorbol 12-myristate 13-acetate treatment.** We revealed that IL-32 $\boldsymbol{\beta}$ suppressed the binding of C/EBP $\boldsymbol{\alpha}$ to **IL-10 promoter by using ChIP assay. These data suggest that** PKC δ and IL-32 β may modulate the effect of C/EBP α on IL-10 **expression. We next demonstrated by immunoprecipitation** that IL-32 $\boldsymbol{\beta}$ interacted with PKC $\boldsymbol{\delta}$ and C/EBP $\boldsymbol{\alpha}$, thereby mediating C/EBP α Ser-21 phosphorylation by PKC δ . We showed that IL-32 $\boldsymbol{\beta}$ suppressed the inhibitory effect of C/EBP $\boldsymbol{\alpha}$ on IL-10 **promoter activity. However, the IL-10 promoter activity was** reduced to the basal level by rottlerin treatment. When $C/EBP\alpha$ **serine 21 was mutated to glycine (S21G), the inhibitory effect of** $\mathsf{C}/\mathsf{EBP}\alpha$ S21G on IL-10 promoter activity was not modulated by $IL-32\beta$. Taken together, our results show that $IL-32\beta$ -mediated $\mathsf{C}/\mathsf{EBP}\alpha$ Ser-21 phosphorylation by PKC δ suppressed C/EBP α **binding to IL-10 promoter, which promoted IL-10 production in U937 cells.**

Interleukin 32 (IL-32) was first reported as a proinflammatory cytokine that induces IL-1 β , IL-6, tumor necrosis factor- α (TNF- α), and IL-8 (1). Multiple studies have shown the association of IL-32 with various inflammatory diseases, such as rheumatoid arthritis, inflammatory bowel disease, and chronic obstructive pulmonary disease (2, 3). In addition to those inflammatory diseases, viral and bacterial infections and even cancers induce IL-32 expression. However, the majority of protein is detected in cellular lysates rather than in the supernatants (2, 4– 6). Recently, the interactions of IL-32 with intracellular proteins, such as focal adhesion kinase 1 (FAK 1), paxillin, integrins, protein kinase C δ (PKC δ), protein kinase C ϵ (PKC ϵ), and STAT3, have been demonstrated, which implies that IL-32 may participate in inflammatory responses as an intracellular mediator (7–9).

The CCAAT/enhancer-binding protein $(C/EBP)^3$ family of transcription factors is known to play important roles in cell proliferation and differentiation. The C/EBP family comprises six members: C/EBP α , C/EBP β , C/EBP γ , C/EBP δ , C/EBP ϵ , and $C/EBP\zeta$. Each member, except $C/EBP\zeta$, which lacks a canonical basic region, contains a similar basic region and leucine zipper sequence at its C terminus that mediate DNA binding and dimerization, respectively (10, 11). C/EBP α dimerizes with other members of the C/EBP family or with itself (12) and interacts with other proteins such as transcription factor II B (TFIIB), TATA-binding protein (TBP), retinoblastoma protein (Rb), p300/CREB-binding protein (CBP), p21, and members of the SWI/SNF complex (13–17). C/EBP α is known to up-regulate the expression of granulocytic lineage-specific genes (18– 21), both directly and synergistically with other key genes such as core-binding factor (CBF) complex genes and PU.1 in mye-

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³ The abbreviations used are: C/EBP, CCAAT/enhancer-binding protein; CREB, cAMP-responsive element-binding protein; PMA, phorbol 12-myristate 13-acetate.

by direct interaction with E2F, which induces cell cycle arrest (19, 22).

It has also been shown that C/EBP α is regulated by phosphorylation and SUMOylation (SUMO $=$ small ubiquitin-like modifier) (23–25). Phosphorylation of Thr-222 and Thr-226 on $\mathsf{C}/\mathsf{EBP}\alpha$ by glycogen synthase kinase 3 (GSK3) causes profound conformational changes, which induce adipogenesis (25). $C/EBP\alpha$ phosphorylation on Ser-21 by extracellular signal-regulated kinase 1/2 (ERK1/2) inhibits granulopoiesis. Phosphorylation of C/EBP α Ser-21 induces conformational changes (26). An oncogenic FLT3 kinase also inhibits the transcriptional activity of C/EBP α by phosphorylating Ser-21 (27).

We previously reported that IL-32 β promotes IL-10 production in myeloid cells (28). In this study, we further verified our results and demonstrated an intracellular mediatory role of IL-32 β for IL-10 production by showing that IL-32 β promoted IL-10 production in a PKC δ -dependent way, where it mediated the phosphorylation of C/EBP α Ser-21 by PKC δ . The phosphorylation of C/EBP α Ser-21 suppressed its inhibitory role in IL-10 induction. Our data suggest that IL-32 β may function as an intracellular regulator of inflammatory response as well as an inducer of inflammation.

EXPERIMENTAL PROCEDURES

Reagents and Cell Culture—The human promyelomonocytic U937 cell line was grown in RPMI 1640 (WelGENE, Daegu, Korea) supplemented with 2 mm L-glutamine, 100 units/ml penicillin, 100 μ g/ml streptomycin, and 10% fetal bovine serum (HyClone, Logan, UT). Human embryonic kidney 293 (HEK293) and lung carcinoma A549 cells were grown in Dulbecco's modified Eagle's medium (DMEM) supplemented with fetal bovine serum (10%) and antibiotics. Phorbol 12-myristate 13-acetate (PMA), lipopolysaccharide (LPS), and polyinosinic: polycytidylic acid (poly(I:C)) were purchased from Sigma. MAPK inhibitor (PD98059), and PKC δ inhibitor (rottlerin) and classical PKC inhibitor (Gö6976) were purchased from Calbiochem.

Measurement of IL-10 Expression Levels and Enzyme-linked Immunosorbent Assay (ELISA)—IL-10 mRNA expression was detected by reverse transcription-polymerase chain reaction (RT-PCR) of total RNAs extracted from U937 cells after the various treatments with 10 nm PMA, 1 μ g/ml LPS, or 10 μ g/ml poly(I:C) for 24 h. U937 cells were also treated with increasing amounts of the PKCδ-specific inhibitor rottlerin for 24 h. The culture media were collected for ELISA. The IL-10 primer set used for PCR was as follows: sense 5--AACCTGCCTAACAT-GCTTCGA-3'; antisense 5'-CTCATGGCTTTGTAGATGC-CT-3'. Human IL-10, $\text{TNF}\alpha$, and IL-8 ELISAs were performed using ELISA kits from R&D systems (Minneapolis, MN).

Immunofluorescence Analyses—For immunofluorescence staining, U937 cells were treated with 50 nm PMA or 1μ g/ml LPS for 40 min; thereafter, 1×10^5 cells were transferred into a 1.5-ml microcentrifuge tube. Cells were fixed with 4% paraformaldehyde and permeabilized with 0.1% Triton X-100 in phosphatebuffered saline (PBS) for 20 min. After blocking with 1% bovine serum albumin (BSA) for 30 min, KU32-52 antibody to IL-32 (1:200) was incubated for 1 h. Fluorescein isothiocyanate (FITC)-conjugated anti-mouse secondary antibody (1:500) was

used for staining. 4',6-Diamidino-2-phenylindole (DAPI) was used for nuclei staining. Fluorescence images were obtained using a Zeiss LSM5 confocal microscope with Plan-Apochromat $63\times/1.4$ oil differential interference contrast objectives.

Construction of *Expression Vectors*-IL-32β cDNA was subcloned into pcDNA3.1 $+$ 6 \times Myc vector by using EcoRI and XhoI. cDNAs for PKC α , PKC δ , PKC ϵ , and PKC θ were subcloned into the pcDNA3.1 $+$ 5 \times FLAG vector by using EcoRI and XhoI (7). Human intronless $C/EBP\alpha$ gene was PCR-amplified from THP-1 genomic DNA by using two overlapping primer sets to obtain the entire cDNA; PCR was then conducted again using the two-PCR fragment as templates. The primers sets were as follows: forward (F) 5'-GAATTCATGG-AGTCGGCCGACTTCTAC-3'; reverse (R) 5'-CTCGAGTC-ACGCGCAGTT-GCCCATGGC-3′; forward 2 (F2) 5′-AGAT-CTCGCACTGCGGCCAGACCACCA-3'; reverse 2 (R2) 5'-AGATCTGGAACTGCAGGTGCGGGGCGG-3-. The entire amplified gene was sequenced and cloned into pCS3MT $6\times$ Myc vector using EcoRI and XhoI. A mutant C/EBP α at serine 21 to glycine (S21G) was generated by site-directed mutagenesis. The primer set for mutagenesis was as follows: forward 5'-AGCAGCCACCTGCAGGGCCCCCCGCAC-3'; reverse 5'-GGGCGCGTGCGGGGGGCCCTGCAGGTG-3'. C/EBP α S21G was subcloned into 6 \times Myc pCS3MT vector and confirmed by sequencing. The deletion mutants of $C/EBP\alpha$ and IL-32 β were generated by PCR, and the PCR products were cloned into the pCS3MT $6\times$ Myc vector.

Electroporation—U937 promyelomonocytic cells were transfected with 1.5μ g of IL-32 siRNA or nontargeting siRNA using a NeonTM transfection system (Invitrogen). PKC δ siRNA was transfected in the same way. PKC δ and IL-32 siRNA and nontargeting siRNA were purchased from Dharmacon (Lafayette, CO). The transfected cells were incubated overnight, and 10 nm PMA was then applied for the indicated time, after which cell lysates were prepared for Western blotting.

Western Blotting and Immunoprecipitation—HEK293 cells were co-transfected with pcDNA3.1 + $6\times$ Myc-IL-32 β and $pCDNA3.1 + 5 \times FLAG-PKCs$ (PKC α , PKC δ , PKC ϵ , and PKC θ). The different combinations of IL-32 β , PKC δ , C/EBP α , and mutant C/EBP α S21G were co-transfected into HEK293. After overnight incubation, cells were treated with 10 μ M rottlerin or 10 μ M PD98059 for 1 h for the inhibitor-treated samples. 20 nM PMA was treated for a further 90 min and then lysed in 50 mm HEPES, pH 7.5; 150 mm NaCl; 5% glycerol; 20 mm β -glycerophosphate; 1% Nonidet P-40; 0.5% Triton X-100; and 1 mm EDTA.Western blotting was performed using the Myc tag antibody (Millipore-Upstate, Bedford, MA), pC/EBP α (serine 21) antibody (Cell Signaling, Danvers, MA), and C/EBP α antibody (Santa Cruz Biotechnology). U937 cells were lysed after PMA treatment for the indicated time. Treatment with inhibitors was performed 1 h before PMA treatment. The optical density of the bands was measured by using Image Studio Lite Software after scanning the blots (LI-COR Biosciences, Lincoln, NE). For immunoprecipitation, cell lysates were mixed with 1μ g of Myc antibody, 3 μ g of C/EBP α antibody, or 3 μ g of goat polyclonal IL-32 antibody (R&D systems) for 1 h, respectively, and then pulled down using 35 μ l of protein G-agarose bead (Kirkegaard & Perry Laboratories, Gaithersburg, MD).

FIGURE 1. The involvement of IL-32 β in IL-10 production in U937 cells. A, IL-32 β was knocked down in U937 cells by transfection with 1.5 µg of IL-32 siRNA or nontargeting siRNA (*NT*), and the cells were then treated with 10 nM PMA for 24 h. The culture media were collected for ELISA, and cell lysates were prepared for Western blotting. *con*, control. The silencing of IL-32 was confirmed by Western blotting. *B–D*, ELISAs for IL-10 (*B*), TNF-- (*C*), and IL-8 (*D*) were performed. All values are mean S.E. (*n* 3). *, *p* 0.03 (*B*). *siIL-32*, IL-32 siRNA; *siCtrl*, siRNA control.

IL-10 Reporter Plasmid and Luciferase Assay—IL-10 promoter region (from -1338 to $+24$) was subcloned into pGL3 Basic using XhoI and HindIII sites. The primer set was as follows: sense 5'-CTCGAGTGTGGAAGGGGAAGGTGA-AGG-3'; antisense 5'-AAGCTTACAGAGCAGTGCTGAGC-TGTG-3'. HEK293 cells were co-transfected with the different combinations of pGL3-IL-10 promoter, $C/EBP\alpha$, mutant $C/EBP\alpha$ S21G, and IL-32 β by using Lipofectamine® 2000 reagent (Invitrogen). After overnight incubation, the samples were treated with 10 μ M rottlerin or 10 μ M PD98059, after which they were treated with 20 nM PMA for a further 24 h. Luciferase assays were performed using Dual-Luciferase® reporter assay system (Promega, Madison, WI).

Chromatin Immunoprecipitation (ChIP) Assay—We used the commercially available ChIP assay kit (Millipore) according to the manufacturer's instructions. Briefly, U937 cells were electroporated with IL-32 siRNA or nontargeting RNA. After overnight incubation, cells were treated with PMA for 90 min. The cells were fixed with 1% formaldehyde, lysed in kit lysis buffer, and sonicated with 5 pulses for 5 s each. After centrifugation at 13,000 rpm for 20 min, the supernatants were precleared with 45 μ l of protein A-agarose/salmon sperm DNA (50% slurry) for 60 min. After a brief centrifugation, the supernatants were mixed with 3 μ g of C/EBP α antibody and maintained overnight. Sixty microliters of protein A-agarose/ salmon sperm DNA (50% slurry) was added to each sample, and the pulled-down DNA fragments were eluted. PCR amplification using the eluted DNA as the template was performed for 35 cycles at an annealing temperature of 61 °C. The primers for PCR amplification of the IL-10 promoter were as follows: sense (from -577 to -557) 5'-CTTTGAGGATATTTAGCC-CAC-

FIGURE 2. PMA-dependent shift of IL-32 β localization in U937 cells. Immunofluorescence staining was performed to localize the endogenous IL-32 β in U937 cells. Cells were treated with 50 nm PMA or 1 μ g/ml LPS for 40 min. Cells were fixed, permeabilized, and labeled for nuclei with DAPI and endogenous IL-32 with FITC. Fluorescence signals were analyzed by confocal microscopy. The same amount of mock IgG (1 μ g) as KU32-52 antibody was used for control.

FIGURE 3. The interaction of PKC δ with IL-32 β and its involvement in IL-10 production. A, IL-32 β -interacting PKC isoforms were screened by immunoprecipitation. HEK293 cells were co-transfected with 6×Myc-tagged IL-32 β and each of 5×FLAG-tagged PKC isoforms (α , δ , ϵ , and θ). After overnight incubation, PMA (20 nm) treatment was performed for 90 min. Immunoprecipitation (*IP*) was carried out with 1 µg of Myc tag antibody. The expression level of each gene was determined by Western blotting (WB) with 30 µg of whole cell lysates (WCL). *B*, after co-transfection of HEK293 cells with IL-32*B* and PKC δ , cells were treated with 10 μ M rottlerin (Rott) or Gö6976 (6976), a classical PKC inhibitor, for 1 h before PMA (20 nM) treatment. Immunoprecipitation was performed with 3 μ g of PKC δ antibody. U937 cells were treated with increasing doses of rottlerin 1 h before the PMA (10 nm) treatment; thereafter, the cells were incubated for 24 h. C–*E*, culture media were collected for IL-10 (C), TNF- α (*D*), and IL-8 (*E*) ELISA. Nontreated cells were included as a control (*con*). All values are mean \pm S.E. (*n* = 3). *, *p* < 0.05; 0 *versus* 1 μ*M* rottlerin, **, *p* < 0.02; 1 *versus* 10 μ*M* rottlerin (*B*), ***, *p* < 0.01; 0 *versus* 10 μ*M* rottlerin (*C*).

3-; antisense (from 296 to 276) 5--GGGCTACCTCTCTT-AGAATAA-3'. Primers for Bcl-2 and lactoferrin ChIP were: Bcl-2 sense (from -395 to -375) 5'-CATTCTTTTTAGCCG-TGTTAC-3'; antisense (from -238 to -218) 5'-GATCTTTA-TTTCATGAGGCAC-3'; lactoferrin sense (from -211 to -192) 5'-CAGCCAGGCAGAACCAGGAC-3'; antisense (from -23 to -4) 5'-GCCACTTGCGCCTGCCCTGC-3'. We measured the optical intensity of the bands by using GelQuant.NET software provided by BiochemLabSolutions. The ChIP products were also analyzed by quantitative real-time PCR by using iQ^{TM} SYBR® Green supermix (Bio-Rad).

Statistical Analysis—Statistical analysis was performed using the unpaired two-tailed Student's *t* test. Differences were considered statistically significant at $p < 0.05$.

RESULTS

IL-32 Promotes IL-10 Production in U937 Cells upon PMA $Stimulation$ —We demonstrated previously that IL-32 β promoted the production of the anti-inflammatory cytokine IL-10 in myeloid cells (28). To verify that IL-32 β is involved in IL-10 production, we knocked down IL-32by using siRNAin U937 cells (Fig. 1*A*). Only the production of IL-10 was decreased upon PMA treatment after IL-32 β silencing, whereas the levels of TNF- α and IL-8 were not changed (Fig. 1). The ectopic introduction of IL-32 β into U937 cells further increased IL-10 production (data not shown). Therefore, these data confirmed that IL-32 β up-regulated the production of IL-10 by PMA treatment in U937 cells.

*IL-32β Associates with PMA-activated PKC*δ—We localized IL-32 β in U937 cells by using confocal microscopy. Under normal conditions, IL-32 β was distributed in the cytosol, but it shifted to the membrane upon PMA stimulation. LPS treatment did not result in the movement of IL-32 β in U937 cells (Fig. 2). This finding suggests that IL-32 β played an indirect mediatory role rather than a role as a direct transcriptional activator for IL-10 expression. Because the effect of IL-32 β on IL-10 production was PMAdependent, and its membrane localization upon PMA stimulation resembled the movement of PKC, we screened the relevant PKCs by immunoprecipitation, as we had previously performed with IL-32 α (7). IL-32 β specifically interacted with PMA-activated $PKC\delta$ because the interaction was inhibited by a $PKC\delta$ -specific inhibitor, rottlerin, but was not inhibited by a classical PKC inhibitor, Gö6976 (Fig. 3, *A* and *B*). Meanwhile, several studies demonstrated that PKC δ is involved in IL-10 production (29-31). Thus, we examined whether $PKC\delta$ was involved in IL-10 production in U937 cells as well. When U937 cells were treated with increasing doses of rottlerin, the production of IL-10 was decreased in a rottlerin dose-dependent manner (Fig. 3*C*). The TNF- α level was also affected by rottlerin, but IL-8 was not (Fig. 3, *D* and *E*). These data suggest that IL-32 β may be inter-related with PKC δ for IL-10 production.

Serine 21 of C/EBP- *Is Phosphorylated by PKC in an IL-32 involved Way in U937 cells*—The human promyelomonocytic U937 cells undergo monocytic differentiation by PMA stimu-

FIGURE 4. **The involvement of IL-32** β **in C/EBP** α **phosphorylation by** PKC δ . *A*, U937 cells were treated with 10 nm PMA for the indicated time. The phosphorylation of C/EBP α at serine 21 was detected by phospho-C/EBP α (*S21*)-specific antibody. Quantitation is shown below the blots, normalized to 0 min (for 10, 30, and 60 min). *B*, U937 cells were treated with 10 μ M rottlerin (*Rott*) or 10 μ M PD98059 (*PD*) for 1 h; thereafter, they were treated with 10 nm PMA for a further 60 min. Quantitation is shown below the blots, normalized to nontreated control for the others. C , U937 cells were transfected with 2 μ g of PKC δ siRNA and nontargeting siRNA (NT). Cells were treated with 10 nm PMA for 30 min after overnight incubation, and then Western blotting was performed. *D*, 1.5 μ g of IL-32 siRNA and nontargeting siRNA (NT) was introduced into U937 cells. After overnight incubation, cells were treated with 10 nm PMA for the indicated time. IL-32 β band (25 kDa) was detected with KU32-52 antibody.

lation. A tissue-specific transcription factor, $C/EBP\alpha$, blocks 12-*O*-tetradecanoylphorbol-13-acetate-induced monocytic differentiation of bipotential myeloid cells when ectopically expressed (32) and induces growth arrest by repressing proliferation genes (10). Hence, we investigated the status of C/EBP α in U937 cells. C/EBP α was expressed in U937 cells and was phosphorylated at serine 21 by PMA stimulation (Fig. 4*A*). The phosphorylation of $C/EBP\alpha$ Ser-21 was inhibited by the treatment of rottlerin, a $PKC\delta$ -specific inhibitor, which means that $C/EBP\alpha$ Ser-21 could be phosphorylated by PKC δ as well. C/EBP α Ser-21 phosphorylation was also inhibited by PD98059, an ERK1/2 inhibitor, to a lesser extent than rottlerin, as consistent with the findings of a previous study (26) (Fig. 4*B*). To further verify that PKC δ is involved in C/EBP α Ser-21 phosphorylation, PKC δ was knocked down with 80% efficiency compared with GAPDH by using $PKC\delta$ -specific siRNA. As shown in Fig. 4C, phospho-C/EBP α at Ser-21 was significantly decreased by PKC δ silencing, which means that PKC δ is also involved in $C/EBP\alpha$ Ser-21 phosphorylation. We then examined whether IL-32 β might affect the phosphorylation of $C/EBP\alpha$. When IL-32 β was knocked down by siRNA, the phosphorylation of Ser-21 of C/EBP α was not induced upon PMA treatment (Fig. $4D$), which suggests that IL-32 β involved is in $C/EBP\alpha$ Ser-21 phosphorylation. To analyze the relevance of $C/EBP\alpha$ to IL-10 expression, we performed ChIP for the putative C/EBP α -binding site on IL-10 promoter after silencing of IL-32 β . We observed that the binding of C/EBP α to IL-10 promoter was increased by IL-32 β knockdown. However, C/EBP α binding to Bcl-2 or lactoferrin promoter was not affected by IL-32 β silencing. We further verified the ChIP results by quantitative real-time PCR (Fig. 5). This finding implies that IL-32 β suppressed C/EBP α binding to IL-10 promoter, thereby relieving the IL-10 promoter from $C/EBP\alpha$ repression. ERK1/2 is

Consensus binding sequence of C/EBP: TTGCGCAAT

FIGURE 5. **The suppression of C/EBP** α **binding to IL-10 promoter by IL-32** β **. ChIP was performed with U937 cells after silencing of IL-32** β **. U937** cells were transfected with 1.5 μ g of IL-32 siRNA and nontargeting siRNA (NT). After an overnight incubation following transfection, 10 nm PMA was treated for 90 min. ChIP was carried out with 3 μ g of C/EBP α antibody. Quantitation is shown below the PCR data, normalized to control (*con*) nontargeting siRNA for the others. The consensus binding sequence of C/EBP and a putative $C/EBP\alpha$ -binding site and its location on IL-10 promoter are presented. ChIP results from Bcl-2 and lactoferrin promoters are presented as controls. IL-10 ChIP products were also measured by quantitative real-time PCR. The results are given as -fold change of the IL-10/lactoferrin ChIP ratio as compared with the value of nontreated nontargeting siRNA. All values are mean \pm S.E. $(n = 3)$.

known to directly phosphorylate C/EBP α Ser-21 through an FXFP docking site. However, we speculated that $PKC\delta$ might phosphorylate $C/EBP\alpha$ Ser-21 in a IL-32 β -mediated way, which inhibited the binding of C/EBP α to IL-10 promoter.

*IL-32β Interacts with PKCδ and C/EBP*α *and Mediates* $C/EBP\alpha$ Ser-21 Phosphorylation—We next investigated how IL-32 β mediated C/EBP α Ser-21 phosphorylation by PKC δ . IL-32 β association with PMA-activated PKC δ was shown in Fig. 3, A and B. We then examined whether $C/EBP\alpha$ interacted with these molecules together. We cloned human $C/EBP\alpha$ $cDNA$ into $6\times$ Myc-tagged expressing vector and then performed immunoprecipitations after co-transfection of HEK293 with different combinations of IL-32 β , C/EBP α , and PKC δ . IL-32 β co-immunoprecipitated with C/EBP α and PKC δ , and this interaction was inhibited by rottlerin, but not by PD98059, an ERK1/2 inhibitor (Fig. 6, *A* and *B*). The interaction of $C/EBP\alpha$ with IL-32 β was also confirmed in U937 cells. The interaction was inhibited by the treatment of rottlerin, but not by PD98059 (Fig. $6C$). We then proved that IL-32 β mediated the phosphorylation of C/EBP α Ser-21 by PKC δ . We transfected HEK293 cells with $6\times$ Myc-tagged C/EBP α with or without IL-32 β and then immunoprecipitated C/EBP α with Myc antibody. $C/EBP\alpha$ Ser-21 was not phosphorylated in the absence of IL-32 β . The phosphorylation of Ser-21 was inhibited by rottlerin, but the phospho-C/EBP α Ser-21 was still detected even in the presence of a ERK1/2 inhibitor, PD98059 (Fig. 6*D*), which means that PKC δ also phosphorylated C/EBP α Ser-21 in a IL-32 β -mediated way. When C/EBP α serine 21 was mutated to glycine, it was no longer phosphorylated (Fig. 6*E*). However, the interaction of C/EBP α with PKC δ and IL-32 β was not inhibited by the mutation of serine 21 to glycine (S21G). We reconfirmed the interaction of IL-32 β with mutant C/EBP α S21G in A549 lung carcinoma cells, which means that the phos-

FIGURE 6. **IL-32** β **-mediated phosphorylation of C/EBP** α **Ser-21 by PKC** δ **. A and B, IL-32** β **was associated with C/EBP** α **and PKC** δ **, and then mediated the** phosphorylation of C/EBP α serine 21 by PKC δ . HEK293 cells were co-transfected with 6×Myc-tagged IL-32 β and C/EBP α and with 5×FLAG-tagged PKC δ . After overnight incubation, cells were treated with 10 μ*M* rottlerin (*Rott*) or 10 μ*M* PD98059 (PD) for 1 h for the inhibitor-treated samples. 20 nM PMA was treated for 90 min. Three micrograms of KU32-52 antibody (Α) or C/EBPα antibody (Β) was used. *IP*, immunoprecipitation; *WCL*, whole cell lysate. C, the expression levels of the transfected genes were determined by Western blotting with 30 μ g of whole cell lysates (*WCL*). Immunoprecipitations (*IP*) were performed with U937 cells after 10 μ M rottlerin or PD98059 treatment for 1 h and treatment with 20 nM PMA for a further 90 min. 3 μ g of goat polyclonal IL-32 antibody was used. After co-transfection of HEK293 cells with 6×Myc-tagged IL-32 β and C/EBP α , cells were treated with inhibitors and PMA in the same way as in *A. D*, cell lysates were immunoprecipitated with 1 μ g of Myc tag antibody; the phosphorylated C/EBP α at serine 21 was detected using phospho-C/EBP α Ser-21-specific antibody. *E* and F, a mutant C/EBP α S21G was co-transfected with IL-32 β and PKC δ into HEK293 cells, and then immunoprecipitations were performed with Myc tag antibody (*E*) or KU32-52 antibody (*F*). G, A549 lung carcinoma cells were co-transfected with a mutant C/EBP α S21G, IL-32*β*, and PKC δ . Each plasmid was also transfected as controls. Immunoprecipitation was carried out with Myc tag antibody. The expression levels of the introduced genes were determined by Western blotting with 30 μ g of whole cell lysates (*WCL*).

FIGURE 7. Interaction domain mapping of C/EBP α and IL-32 β . A, functional domains of C/EBP α and its deletion mutants are schematically illustrated. TAD; transactivation domain, *RD*; regulatory domain, *DBD*; DNA-binding domain, *bLZ*; basic leucine zipper domain. HEK293 cells were co-transfected with 6×Myctagged C/EBP α mutants and 5 \times FLAG-tagged IL-32 β . After overnight incubation, cells were treated with 20 nm PMA for 90 min. Immunoprecipitation was performed with 1 μ g of FLAG antibody. *B*, a schematic illustration of IL-32*β* deletion mutants is presented. HEK293 cells were co-transfected with 5×FLAGtagged C/EBP α and 6 \times Myc-tagged IL-32 β mutants. Immunoprecipitation was performed as in A with 1 μ g of FLAG antibody. The pulled down D1 is marked by arrow. The expression levels of the transfected genes were determined by Western blotting with 30 µg of whole cell lysates (*WCL*). Immunoglobulin G light chain (*IgG L*) is indicated by an *arrow*.

phorylation of C/EBP α Ser-21 did not influence its association with IL-32 β and PKC δ (Fig. 6, *F* and *G*). We also characterized the interaction domains of C/EBP α and IL-32 β . IL-32 β inter-

acted with the full-length $C/EBP\alpha$ *(FL)* as well as F2 and F4 (Fig. 7*A*). F2 contains the DNA-binding domain and the basic leucine zipper domain, and F4 contains only the basic leucine zip-

FIGURE 8. **The release of C/EBP** α **block on IL-10 expression by IL-32** β **.** A, HEK293 cells were co-transfected with pGL3-IL-10 promoter, IL-32 β , or C/EBP α . After overnight incubation, cells were treated with 20 nm PMA for 24 h, and then firefly luciferase assay was performed. All values are mean \pm S.E. *, **, $p < 0.001$. B, luciferase assay was performed in the same way as in A after co-transfection of HEK293 cells with pGL3-IL-10 promoter, IL-32β, C/EBPα, or mutant C/EBPα S21G. For inhibitor-treated samples, treatment with 10 μ m rottlerin (*Rott*) and 10 μ m PD98059 (*PD*) was performed 1 h before PMA (20 nm) treatment for 24 h. All values are mean \pm S.E. *, **, p $<$ 0.001. C, HEK293 cells were co-transfected with a mutant C/EBP α S21G, IL-32 β , and pGL3-IL-10 promoter, and then luciferase assay was performed in the same way as in *B*. All values are mean \pm S.E. ($n = 3$). *, $p < 0.001$.

per domain. Therefore, these data mean that IL-32 β binds to the basic leucine zipper region of C/EBP α . C/EBP α bound to the N-terminal region of IL-32 β (Fig. 7*B*). Most isoforms of IL-32 contain the same C terminus (D2), but the N terminus (D1) is diverse because of alternative splicing. Meanwhile, the two mutants of IL-32 β had no effect on IL-10 production, which means that only the intact form of IL-32 β is functional (data not shown).

IL-32β Suppressed the Inhibitory Effect of C/EBPα on IL-10 $Expression$ —We further investigated the effect of $C/EBP\alpha$ on IL-10 expression using IL-10 promoter reporter assay. We observed that IL-10 promoter was not activated under C/EBP α expression without IL-32 β . However, in the presence of IL-32 β , the promoter was activated (Fig. 8*A*). The promoter activity was inhibited by rottlerin, but not by PD98059, which means that ERK1/2 was not involved in IL-10 expression in this system. The activity of $C/EBP\alpha$ is repressed by the phosphorylation of Ser-21 (26); as is consistent with this study, a mutant C/EBP α S21G, which could not be phosphorylated by PKC δ , repressed IL-10 promoter activity even under co-expression with IL-32 β (Fig. 8*B*). When the cells were treated with rottlerin under mutant C/EBP α S21G expression, the promoter activity was further reduced, but was not modulated by PD98059 (Fig. 8*C*). These data therefore suggest that IL-32 β -mediated phosphorylation of C/EBP α Ser-21 by PKC δ suppressed the inhibitory effect of $C/EBP\alpha$ on IL-10 production.

DISCUSSION

The initiation of the proinflammatory response is a prerequisite for the proper immune responses upon viral or bacterial infections. Although the proinflammatory reaction is essential, it should be resolved to prevent damage to the host. IL-32 has been studied for its proinflammatory function because multiple studies have shown its association with inflammatory diseases. IL-32 β is known to induce TNF- α , IL-8, IL-6, and macrophage inflammatory protein-1 α (MIP-1 α), all of which are proinflammatory mediators. However, we showed that IL-32 β induced an anti-inflammatory cytokine IL-10 in myeloid cells. In this study, we questioned the mechanism of action of IL-32 β because IL-32 β protein was detected only with cellular lysates, not in the culture medium. In this study, we solved our previous question by revealing the mechanism by which IL-32 β promotes IL-10 production. Recently, several studies have shown the intracellular mediatory function of IL-32 (7, 8, 33).We demonstrated that IL-32 β mediated the phosphorylation of serine 21 on C/EBP α by PMA-activated PKC δ , which resulted in IL-10 production. Phosphorylation of C/EBP α at Ser-21 inhibits its transcriptional activity through induction of conformational changes. The involvement of $PKC\delta$ in IL-10 production is known, but there is little evidence of the effect of C/EBP α on IL-10 production. Although a previous study showed that $C/EBP\alpha$ induced IL-10 production (34), the effect was synergistic with cAMP-responsive element-binding protein/activating transcription factor (CREB/ATF). C/EBP α not only activates the granulopoiesis-related genes, but also represses cell cyclerelated genes, which means that C/EBP α acts as a transcriptional activator as well as a transcriptional inhibitor. There are a couple of models explaining the inhibitory function of C/EBP α . C/EBP α may inhibit S-phage genes indirectly through an association with E2F, or by direct binding to E2F sites. Alternatively, C/EBP α may directly inhibit S-phage genes by binding to the C/EBP-binding site on promoters of those genes (10). On the grounds of these studies, we examined the effect of IL-32 β on C/EBP α and showed that IL-32 β inhibited C/EBP α binding to IL-10 promoter by mediating Ser-21 phosphorylation by PKC δ . However, whether the inhibition of the IL-10 gene by $C/EBP\alpha$ results from the binding of $C/EBP\alpha$ to IL-10 promoter or the interaction of other inhibitory proteins with $C/EBP\alpha$ located on the IL-10 promoter remains unclear. In any event, according to our results, IL-32 β has an intracellular mediatory function in addition to its proinflammatory function.

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