

Functional roles of C/EBP α and SUMO-modification in lung development

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Abstract. CCAAT enhancer binding protein alpha (C/EBP α) is a transcription factor regulating the core aspects of cell growth and differentiation. The present study investigated the level and functional role of C/EBP α during the development of the rat lung. C/EBP α protein exhibits a dynamic expression pattern. The correlation between the expression of C/EBP α protein and the content of glycogen during lung maturation was analyzed to understand the function of C/EBP α in lung differentiation. The high expression of C/EBP α coincides with the reduction of glycogen in the fetal lung. In addition, the authors identified that changes in the level of C/EBP α are associated with the secretion of pulmonary surfactant. C/EBP α is modified by small ubiquitin-related modifier (SUMO) post-translationally. The results of double immunofluorescence staining and immunoprecipitation demonstrated that SUMO-modified C/EBP α was present in the lung. The sumoylated C/EBP α gradually decreased during lung differentiation and was negatively correlated with pulmonary surfactant secretion, thereby suggesting that the SUMO modification may participate in C/EBP α -mediated lung growth and differentiation. These results indicated that C/EBP α played a role in lung development and provided the insight into the mechanism underlying SUMO-modification.

Introduction

According to the histomorphological characteristics, lung morphogenesis can be divided into five stages in the rat (1). First is the embryonic period, from fertilization to E9.5 (embryonic day 9.5). During this period of organogenesis, the lung primordial branches from the two lung buds that lie on each side of

the future esophagus (2). The segmental branching of the airway is presented by 9.5 days of gestation, which results in the formation of tubular lung, and the following stage is referred to as the pseudoglandular period (E9.5-E16.5). At this stage, the primary lung buds grow ventrally and caudally, and five secondary buds are generated, which form the lobes of the mature lung. Recurrent sprouting and bifurcations of the lung buds result in the formation of pre-acinar airways arising from the process of branching morphogenesis. At the late pseudoglandular stage, the vascular development is quintessential, and all the numbers of pre-acinar airways are completely formed. Structurally, the lung development then enters the next phase, the canalicular period (E16.5-E17.5). Extensive cellular differentiation occurs in the late pseudoglandular and canalicular stages (3), the terminal branches of the bronchial tree are taking shape, and the cuboidal epithelium differentiates into type I and II cells, resulting in the formation of a thin air-blood barrier, as well as the secretion of surfactant. The cell differentiation continues, thereby forming the terminal sacs. These terminal sacs, as the name suggests, leads to the subsequent histological stage, which is known as the sacular period, spanning E17.5 to postnatal day 5 (P5), comprising the phase of rapid expansion (P1-P4). The major characteristics of this stage are increased growth of lung parenchyma, further maturation of the surfactant system, thinning of the interstitial tissue between the airspaces and the production of the last generation of airways by some future alveolar ducts and the outermost periphery alveolar sacs. Together, these changes prepare the lung for respiration after birth. The premature lung in the gestational age <35 weeks remains in the canalicular or sacular phase (4), which are sophisticated metamorphosis stages of lung development. The injury to the developing lung during this period alters subsequent alveolar and vascular development, resulting in simplified alveolar structures, dysmorphic capillary configuration, variable interstitial cellularity and fibroproliferation that are characteristic of the 'new' bronchopulmonary dysplasia (5). The alveolar period (P5-P30) is the last stage in lung development that is responsible for the accomplishment of alveolarization; it includes the phase of alveolarization (P5-P13) and the equilibrium stage (P14-P30).

CCAAT/enhancer binding protein alpha (C/EBP α) is the founding member of a family of basic region/leucine zipper transcription factors. A previous study indicated that C/EBP α participates in regulating the lung differentiation and pulmonary

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maturation (6). The expression of C/EBP α is initiated almost simultaneously to that of cellular differentiation and the emergence of differentiation markers (7). Furthermore, the studies in lung epithelial cell lines have demonstrated that the promoters of several differentiation-dependent genes harbor the C/EBP-binding sites (8-11); the C/EBP α -mediated regulation of these genes has also been proposed. Berg *et al* (12) reported that the abnormal expression of C/EBP α disrupts the lung development. These results indicated a role of C/EBP α in lung development; however, the molecular mechanism is poorly understood.

The post-translational modification is a vital regulatory mechanism underlying proteins exerting pleiotropic effects, thereby improving the structure and function of target proteins. Small ubiquitin-like modifier (SUMO) is a novel protein that can modify the target proteins causing rapid changes in the function and distribution of proteins, subcellular structures and multiprotein complexes (13). The pathway of sumoylation resembles that of ubiquitination, although the enzymes involved in the conjugation of SUMO are different. The SUMO peptide is first processed at the C-terminus by the ATP-dependent heterodimeric SUMO-activating E1 enzyme (Aos1/Uba2). Subsequently, it is transferred to the catalytic cysteine of the E2 conjugating enzyme, Ubc9 (14). The final step involves the transfer of the SUMO moiety from E2 to the specific substrate in the presence of an E3 ligase.

C/EBP α was previously reported to be post-translationally modified by SUMO at a lysine residue (K159) within the 'attenuator domain' of the protein that can negatively affect the transcriptional activity (15-17). Hankey *et al* (18) demonstrated that changes in the sumoylation status of C/EBP α might contribute towards a switch that regulates its transcriptional activity during normal neutrophil development. On the other hand, Sato *et al* (19) reported that the enhancement of C/EBP α -mediated transactivation by BRG1, which is a core subunit of the SWI/SNF chromatin remodeling complex, was inhibited by sumoylation. Furthermore, sumoylation dramatically decreased the expression of the liver-specific albumin gene that harbors the C/EBP α binding site. Notably, the common endodermal origin and the crucial role of C/EBP α in lung and liver suggest the potential transcriptional regulation and that SUMO may have a role in both organs. However, the role of SUMO-modification in the lung has not yet been reported. The C/EBP α studies are primarily focused on the mature lung. The mechanism through which C/EBP α regulates AEC-II (alveolar epithelial cells type II) differentiation and its effect on alveolar maturation in the premature lung have not yet been clarified. The studies on C/EBP α and AEC II differentiation-related constituents, such as pulmonary surfactant proteins, phosphatidylcholine (PC) and glycogen are poorly reported.

In the present study, the authors investigated the level and functional role of C/EBP α during rat lung development. The correlation between the level of C/EBP α and the content of glycogen during lung maturation established a role of C/EBP α in lung differentiation. Furthermore, the changes in the status of C/EBP α were shown to be associated with the secretion of pulmonary surfactant. The SUMO modification of C/EBP α was also found to participate in this phenomenon. These findings indicated that C/EBP α serves a vital role in normal lung development, and provides further insights into the involvement of SUMO.

Materials and methods

Animals. Sprague-Dawley rats (90-100-days old, weight 250-300 g) were purchased from the Animal Center of Jiangsu University. All rats kept on a 12-h light/dark cycle at a room temperature of 23 \pm 2 $^{\circ}$ C and a relative humidity of 50 \pm 5%, maintained on standard laboratory food and water *ad libitum* throughout the experiment. Rats were mated by 3:1 female: male ratio (15:5). The next morning, the female rats were checked for fertility and recorded as embryonic day 0.5 (E0.5). According to the different stages during the development of rat lung, the authors chose embryonic days 15.5, 17.5 and 19.5, and postnatal days 0.5, 4, 7 and 14 as the observation time-points. Embryos and lungs were isolated from the embryonic and postnatal stages as previously described, and a part of the samples was immediately fixed with 4% paraformaldehyde; the remaining part of the samples was stored at -80 $^{\circ}$ C. The number of animals per group analyzed varies between 5 and 8. The protocols for animal studies were approved by the Laboratory Animal Ethics Committee of the Affiliated Hospital of Jiangsu University (Zhenjiang, China).

Histological analysis and periodic acid-Schiff (PAS) staining. Tissues were fixed with 4% paraformaldehyde in phosphate-buffered saline (PBS) for 24 h at 4 $^{\circ}$ C, washed with PBS, dehydrated by an alcohol gradient and embedded in paraffin. Subsequently, 3 μ m sections were sliced, followed by deparaffinization and rehydration through xylene, ethanol and water. The antigen retrieval was performed in 10 mM citrate buffer (pH 6.0) at a constant pressure of 20 cm H₂O. The tissue sections were stained with either hematoxylin and eosin (H&E) for histological analysis, or PAS reagent (G1281) from Beijing Solarbio Science & Technology Co., Ltd., (Beijing, China) for analyzing the content of glycogen; the content of glycogen in alveolar epithelial cells reflect the maturity of AEC II (20). The sections were observed and images acquired by microscopy at magnification, x400. PAS staining of the lung tissues revealed red- or purple-colored glycogen. The images were analyzed by Image-Pro Plus III (Media Cybernetics, Inc., Rockville, MD, USA) to obtain the mean gray value. A total of 3-5 samples/group were analyzed.

Antibodies. The primary antibodies used were anti-C/EBP α (sc-9314), anti-SP-B (sc-7704) and anti-SP-C (sc-13979 from Santa Cruz Biotechnology, Inc., Dallas, TX, USA), anti-SP-A (LS-C357574) from LifeSpan BioSciences, Inc. (Seattle, WA, USA), anti-SP-D (bs-1583R) from BIOCSS (Beijing, China), and anti-SUMO-1 (4930) and anti- β -actin (#3700) from Cell Signaling Technology, Inc. (Danvers, MA, USA). Biotinylated secondary antibodies (FMS-MS01, FMS-Rb01, FMS-Gt01) were procured from Fcmacs Biotech Co., Ltd. (Nanjing, China). Donkey anti-rabbit IgG/fluorescein isothiocyanate (bs-0295D-FITC) and donkey anti-goat IgG/phycoerythrin (bs-0294D-PE) were obtained from BIOCSS.

Western blot assay. Lung tissues were thawed on ice, washed with cold PBS, sliced into small pieces, lysed in radioimmunoprecipitation assay lysis buffer containing 1 mmol/l protease inhibitor PMSF (Sigma-Aldrich, Merck KGaA, Darmstadt, Germany) and homogenized by Dounce tissue grinders on ice. The extract was centrifuged at 12,000 x g and 4 $^{\circ}$ C for 15 min; the supernatant was collected. Following normalization of protein

concentrations between the samples by BCA kit (Beyotime Institute of Biotechnology, Shanghai, China), 10 ml of the lung lysates were resolved by 12% sodium dodecyl sulfate-polyacrylamide-gel electrophoresis (SDS-PAGE) and transferred to polyvinylidene difluoride membranes. The membranes were blocked in 5% milk-TBST containing 0.1% Tween-20 at 37°C for 2 h, and probed overnight with primary antibodies at 4°C. The following primary antibodies were used: anti-C/EBP α (1:200), anti-SP-A (1:500), anti-SP-B (1:200), anti-SP-C (1:500), anti-SP-D (1:200), anti-SUMO-1 (1:200) and anti- β -actin (1:1,000). The membranes were washed with TBST containing 0.1% Tween-20 and incubated with biotinylated secondary antibodies (1:5,000) for 1 h at 37°C. The immunoreactive bands were visualized by FluorChem FC3 chemiluminescence (ProteinSimple, San Jose, CA, USA) according to the manufacturer's recommendations. The protein expressions were quantified by densitometric analysis using LANE 1D software (Sage, Beijing, China).

PC assay. PC assay kit (ab83377, from Abcam, Cambridge, UK) was used to measure the levels of PC in various samples. Lung tissues were thawed on ice, washed with cold PBS, resuspended in the assay buffer provided by the kit, and homogenized with a Dounce homogenizer on ice. Then, the samples were centrifuged for 5 min at 4°C at 12,000 x g to exclude the insoluble material and collect the supernatant. An equivalent of 2.5 μ g of the samples was loaded per well; PC reaction mixture was added, followed by development mixture according to the protocol of the kit. The colorimetric reaction was measured at 570 nm. The PC assay was carried out three times, following which, the concentration of PC was estimated.

Immunofluorescence. Double immunofluorescence staining was used to detect the localization of sumoylated C/EBP α . The sections were deparaffinized and rehydrated by xylene, ethanol and water. Antigen retrieval was performed in 10 mM citrate buffer (pH 6.0) at a constant pressure of 20 cm H₂O. Subsequently, the sections were blocked with 5% serum in PBS for 20 min at room temperature, followed by incubation with the mixture of anti-C/EBP α (1:100) and anti-SUMO-1 (1:100) antibodies overnight at 4°C. Anti-C/EBP α was detected using a secondary PE-conjugated antibody (1:500) and anti-SUMO-1 was detected by a secondary FITC-conjugated antibody (1:500). The nucleus was counterstained with DAPI for 5 min at room temperature. All washes and antibody dilutions were made in PBS.

Immunoprecipitation. The immunoprecipitation protocol by Sato *et al* (19) was utilized for the detection of sumoylated C/EBP α . In order to detect the electrophoretic mobility of sumoylated C/EBP α , immunoprecipitation assay was carried out. A total of 20 mg lung tissues were thawed on ice, washed with cold PBS, sliced into small pieces and lysed with a buffer containing 1% Triton X-100, 150 mM NaCl, 20 mM Tris (pH 7.5) supplemented with protease inhibitors, followed by grinding using Dounce tissue grinders on ice. The extract was centrifuged at 12,000 x g and 4°C for 15 min; the supernatant was collected. The chromatin was preincubated with 2 μ g anti-C/EBP α antibody for 30 min on ice. Then, 20 μ l Protein A/G PLUS-Agarose beads (Santa Cruz Biotechnology) were added, and the mixture was incubated overnight at 4°C with

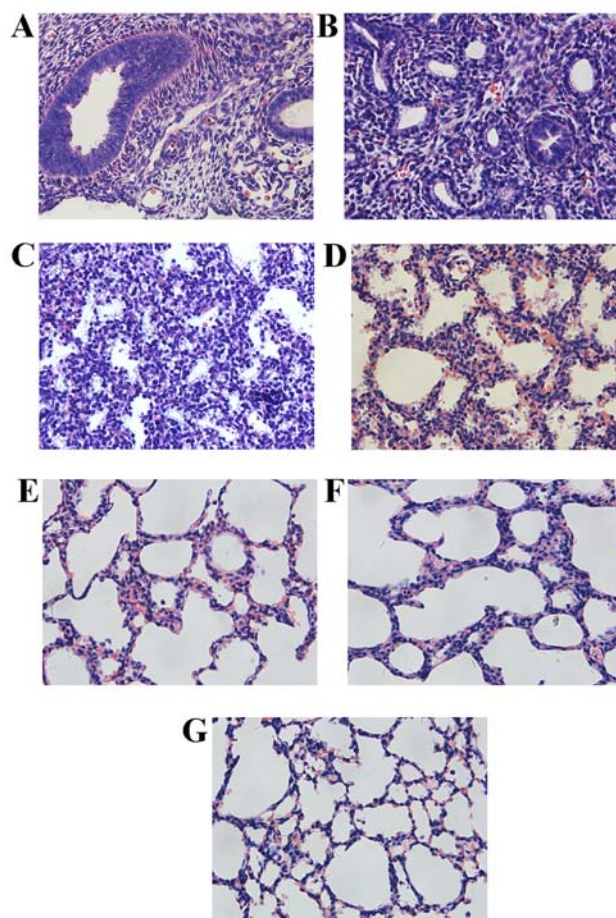


Figure 1. Histology of transverse sections of isolated lungs from several stages of development, fixed in 4% paraformaldehyde, embedded in paraffin and stained with hematoxylin and eosin (H&E). (A) Lung from E15.5, the ring-like pre-acinar airways lined by high columnar epithelium, and surrounded by dense connected tissues. (B) Lung from E17.5, more acinar luminae appeared in this stage, lined with simple cuboidal epithelium. (C) Lung from E19.5, primary alveolar had taken shape and was surrounded by thinner connective tissues that arranged in a streak. (D) Lung from P0.5, a number of primary alveoli increased, with irregular structures. (E) Lung from P4, several ridges protruding into the alveolar space. (F) Lung from P7, air sacs tend to be uniform, and the interstitial tissues thin out. (G) Lung from P14, the basic unit in lung is mature alveoli, with uniform size and separated by razor-thin septum. Magnification, x400.

gentle agitation. The antibody-coated beads were washed with lysis buffer and the immunoprecipitated proteins eluted by heat denaturation for 5 min in 5X Laemmli buffer containing 100 mM β -dithiothreitol. Immunoblotting, to detect the SUMO-1 protein, was performed as previously described.

Statistical analysis. The results are expressed as the mean \pm standard error of the mean of at least three independent experiments. All data were analyzed by the SPSS statistical software (version 17.0; SPSS, Inc., Chicago, IL, USA). Pearson's correlation analysis was used to assess the relationship between indicators. $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Histomorphological variations of lung morphogenesis in different stages. As presented in Fig. 1, the ring-like pre-acinar

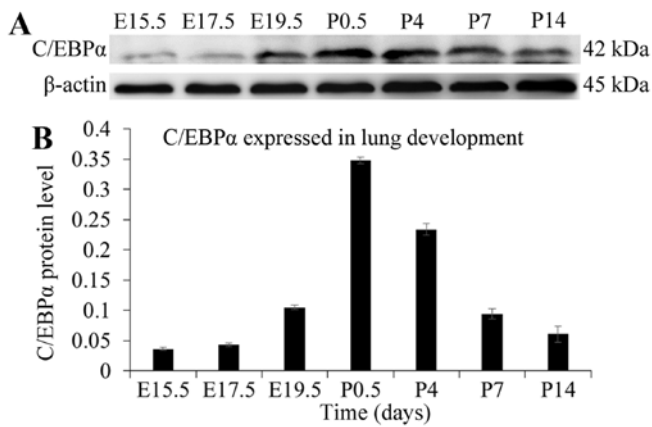


Figure 2. (A and B) C/EBP α protein exhibits a dynamic expression pattern during lung development. Western blot assays were carried out on the total protein lysates of lung tissue. Lung tissues from several different stages of development were lysed and analyzed by immunoblotting with anti-C/EBP α antibody. Data are presented as the mean \pm standard error of the mean. E, embryonic; P, postnatal; C/EBP α , CCAAT enhancer binding protein- α .

airways could be observed at the late pseudoglandular stage at E15.5, lined by high columnar epithelium, surrounded by dense connective tissues. At E17.5, the primitive lung buds were separated completely. A large number of acinar luminae appeared, lined with simple cuboidal epithelium. In addition, the authors also observed the alveolar type I epithelial cells (AEC-I) at this stage. The primary alveolar was shaped at the early canalicular stage at E19.5, surrounded by thin connective tissues arranged in streak-like conformation. At P0.5, the number of primary alveoli was increased with irregularity in the structures. An increasing number of alveolar septa were formed at the late canalicular stage at P4 with some ridges protruding into the alveolar space. At P7, the size of air sacs was predisposed towards uniformity, and the interstitial tissues thinned out. By P14, the basic unit in the lung was mature alveoli, which appeared uniformly sized and separated by a razor-thin septum.

Dynamic expression pattern of C/EBP α protein during lung development. Rat lungs from different developmental stages were investigated for the expression pattern of C/EBP α using the western blot assay. As presented in Fig. 2, little expression of C/EBP α protein could be observed in lung tissues of the late pseudoglandular stage at E15.5. However, after the pseudoglandular-canalicular transition, the amount of C/EBP α protein increased gradually, and the trend was extremely obvious at E19.5 and P0.5. The C/EBP α protein reached its peak level at P0.5, following which, the expression weakened and finally stabilized at P14, resembling the expression pattern in the adult animal (12). In summary, C/EBP α displayed a trend of initial increase followed by a decrease during the lung development. C/EBP α was originally expressed in the late pseudoglandular phase characterized by growth and branching of the lung, which coincided with the cell differentiation and time of emergence of differentiation markers. After the pseudoglandular-canalicular transition, the expression increases and exhibits a widespread pattern, correlating with the extensive cellular differentiation occurring during this period (7).

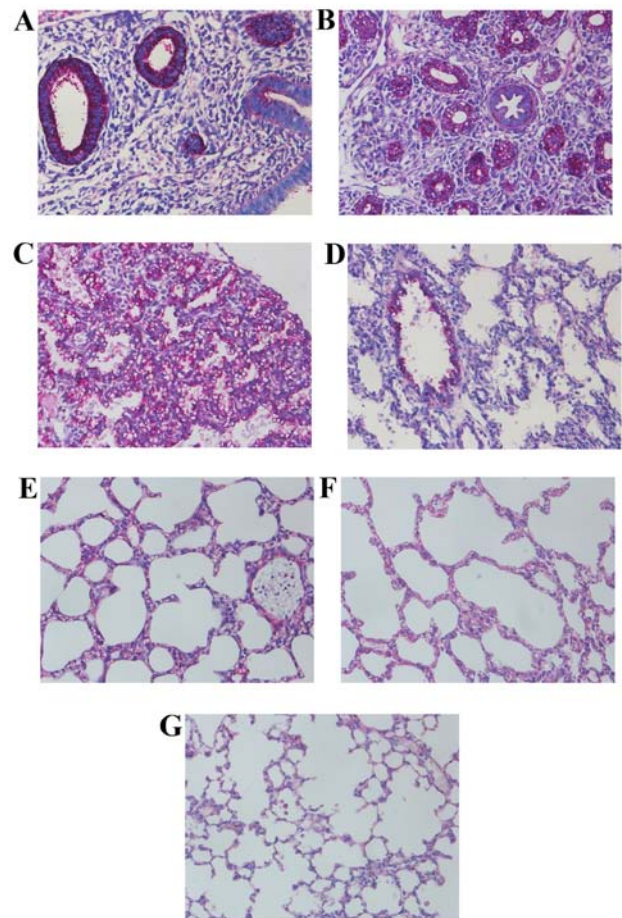


Figure 3. Periodic-acid and schiff staining of lung tissues from several different stages of development. Red/purple color identifies mucin and carbohydrate macromolecules such as glycogen. (A-G) Lungs from E15.5, E17.5, E19.5, P0.5, P4, P7 and P14, respectively. The content of glycogen was decreased during lung differentiation. E, embryonic; P, postnatal. Magnification, x400.

Correlation between glycogen content and C/EBP α protein expression. The epithelial content of glycogen reflects the maturity of AEC-II (20). The intracellular glycogen is transformed into the pulmonary surfactant phospholipids; with AEC-II differentiation, the content is reduced gradually. As indicated by PAS staining of lung tissues in Fig. 3, the glycogen was identified by red or purple color. Glycogen occurs in the cytoplasm; the content was high at the late pseudoglandular stage at E15.5, followed by a decrease as the lung develops. This tendency is noticeable, especially at the canalicular and early saccular stages, which correspond to a wide range of cell differentiations during this stage. Finally, the glycogen content stabilized at the alveolar stage. To investigate the functional role of C/EBP α in AEC-II differentiation, the authors analyzed the link between the changes in the status of C/EBP α and the content of glycogen during lung maturation. As observed in Fig. 4, the level of glycogen exhibited a pattern approximately opposite to the expression of C/EBP α protein at the embryo and early postnatal stages, establishing a positive correlation between the C/EBP α expression and lung differentiation. At the late development period of the lung, primary and mature alveolar became the basic structural unit of lung tissue, and the pulmonary function improved gradually. The reduced glycogen consumption corresponds to the relatively low level

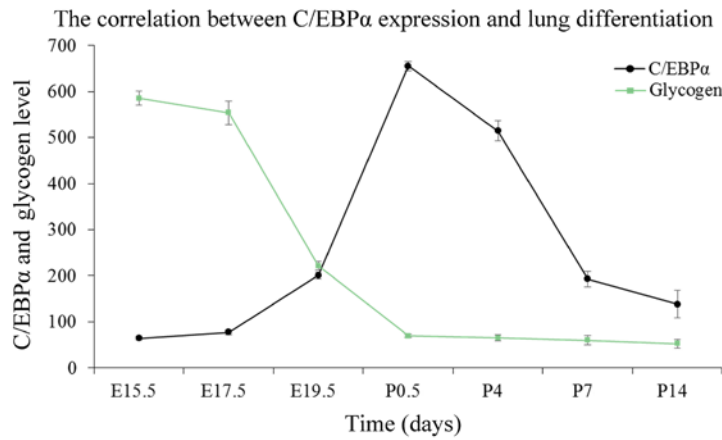


Figure 4. C/EBPα protein expression is positively correlated with lung differentiation. The amount of glycogen showed a pattern approximately opposite to the expression of C/EBPα protein at the embryo and early postnatal stages, corresponding to the lung differentiation. During this period, both stabilized at the late stage of lung development. C/EBPα, CCAAT enhancer binding protein-α; E, embryonic; P, postnatal.

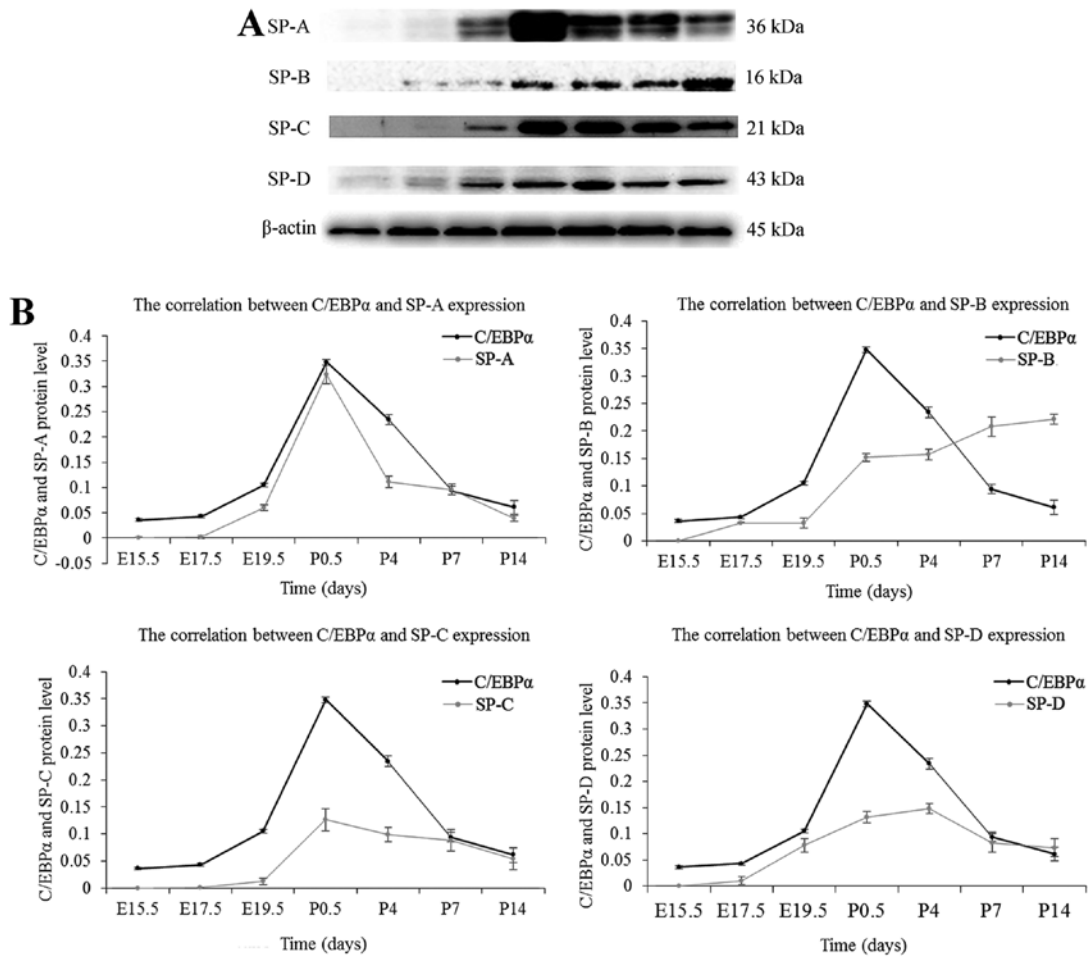


Figure 5. C/EBPα serves a role in the secretion of pulmonary surfactant proteins. (A) Western blot assays were carried out on total protein lysates of lung tissue. Lung tissues from several different stages of development were lysed and analyzed by immunoblotting with anti-SP-A, SP-B, SP-C and SP-D antibodies, respectively. (B) Positive correlation between C/EBPα and SPs expression, except SP-B. The amount of SP-B increases continually after birth. Data are presented as the mean ± standard error of the mean. C/EBPα, CCAAT enhancer binding protein-α; E, embryonic; P, postnatal.

of C/EBPα. This suggested that C/EBPα might facilitate cell differentiation throughout the lung development phase.

Correlation between pulmonary surfactant proteins and C/EBPα expression. Lung surfactant contains four associated

proteins, surfactant protein (SP)-A, SP-B, SP-C and SP-D. In the present study, the authors investigated the secretion of pulmonary surfactant proteins during the whole development process of lung, using the western blot assay. As presented in Fig. 5, little expression of SP-A could be seen at the late

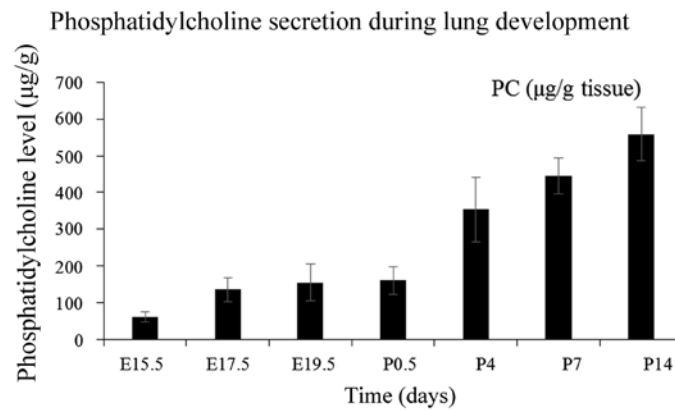


Figure 6. PC assay kit was used to measure the PC levels in lung samples from several development stages. Samples were homogenized with a Dounce homogenizer, centrifuged, supernatants collected. The PC reaction mixture and development mixture were added respectively, and colorimetric assay was used to detect the concentration of PC. Data are presented as the mean \pm standard error of the mean. PC, phosphatidylcholine; E, embryonic; P, postnatal.

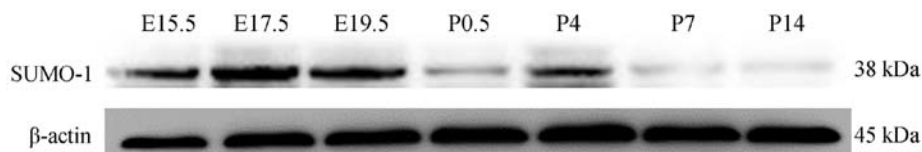


Figure 7. SUMO-1 protein was expressed at several lung development stages. Western blot analysis was carried out on total protein lysates of lung tissue. Lung tissues from several different stages of development were lysed and analyzed by immunoblotting with the anti-SUMO-1 antibody. SUMO, small ubiquitin-like modifier.

pseudoglandular stage at E15.5, whereas SP-B, SP-C, and SP-D were first detected at E17.5; the expression of SPs was increased gradually with the development of fetal lung. The secretion of SP-A, SP-C and SP-D was maximal at the postnatal age between 12 h and 4 days, then decreased gradually and stabilized at P14. The expression of SP-B continued to rise after birth, and the increase was superior to the embryonic period. SP-B secretion increased slowly at P14 and stabilized gradually. The expression patterns of SPs were approximately similar to that of the C/EBP α protein except for SP-B. The results suggested a positive correlation between C/EBP α expression and the secretion of SPs; SPs are differentiation-dependent genes, and these results were consistent with the finding that C/EBP α promotes lung differentiation. However, the function of C/EBP α on SP-B during the postnatal days was not clarified.

Correlation between the secretion of pulmonary surfactant phospholipids and C/EBP α expression. The pulmonary surfactant phospholipids consist primarily of PC, synthesized in AEC-II, stored in the lamellar body, and excreted to the alveolar surface, thereby reducing the surface tension, increasing the compliance and preventing the alveolar collapse. The authors investigated the secretion of PC through lung tissues from various developmental stages. The augmented production of PC by fetal lungs does not commence until the gestation is ~80% complete (21). As presented in Fig. 6, the amount of PC at the late pseudoglandular stage is low. After a slow increasing period before birth, the content seems to increase rapidly at postnatal days and almost stabilizes during the alveolar period. The secretion pattern of PC was in accordance with the expression profile of C/EBP α in utero; however, the after-birth tendency was opposite without a significant correlation. Chen *et al* (22) reported that C/EBP α exerts a positive effect

on adipocyte differentiation and restrains the accumulation of lipid. A slow increase of PC was observed at the embryonic period with a high expression of C/EBP α . Conversely, C/EBP α expression was reduced after birth, but the content of PC increased rapidly. Thus, it is possible that C/EBP α accelerates the secretion of AEC-II and metabolism of PC and other extracellular lipids concurrently. The secretion and cyclic utilization of PC are primarily regulated by SPs (23). As previously noted, C/EBP α exerts a positive effect on the expression of SP genes in the embryonic period, and thus, it may be speculated that C/EBP α can indirectly promote the synthesis of PC.

SUMO-modification of C/EBP α occurs during lung development. C/EBP α is considered to serve a vital role in lung morphogenesis and cytodifferentiation (12). The modification of C/EBP α by SUMO post-translationally can alter the protein function (24). However, explicit data on the sumoylated C/EBP α in lung development is absent, and therefore, rat lung tissues were used to study the sumoylation of C/EBP α . As presented in Fig. 7, SUMO-1 protein expressed during lung maturation. Next, C/EBP α and SUMO-1 were analyzed using immunofluorescence. As presented in Fig. 8, C/EBP α was partially colocalized with SUMO-1 in the AECs at P14, suggesting that sumoylation of C/EBP α occurred in the lung. Then, the expression of sumoylated C/EBP α during lung development was assessed. As shown in Fig. 9A, a shifted band attributable to modified C/EBP α was detected at different time-points during the development of rat lung. The shifted band seemed to be sumoylated C/EBP α , as its mobility was in accordance with that of the confirmed sumoylated C/EBP α in the hepatocytes (19). To confirm whether this shifted band was indeed SUMO-1-modified C/EBP α , the immunoprecipitation assay was carried out. As shown in Fig. 9B, the sumoylated C/EBP α was precipitated

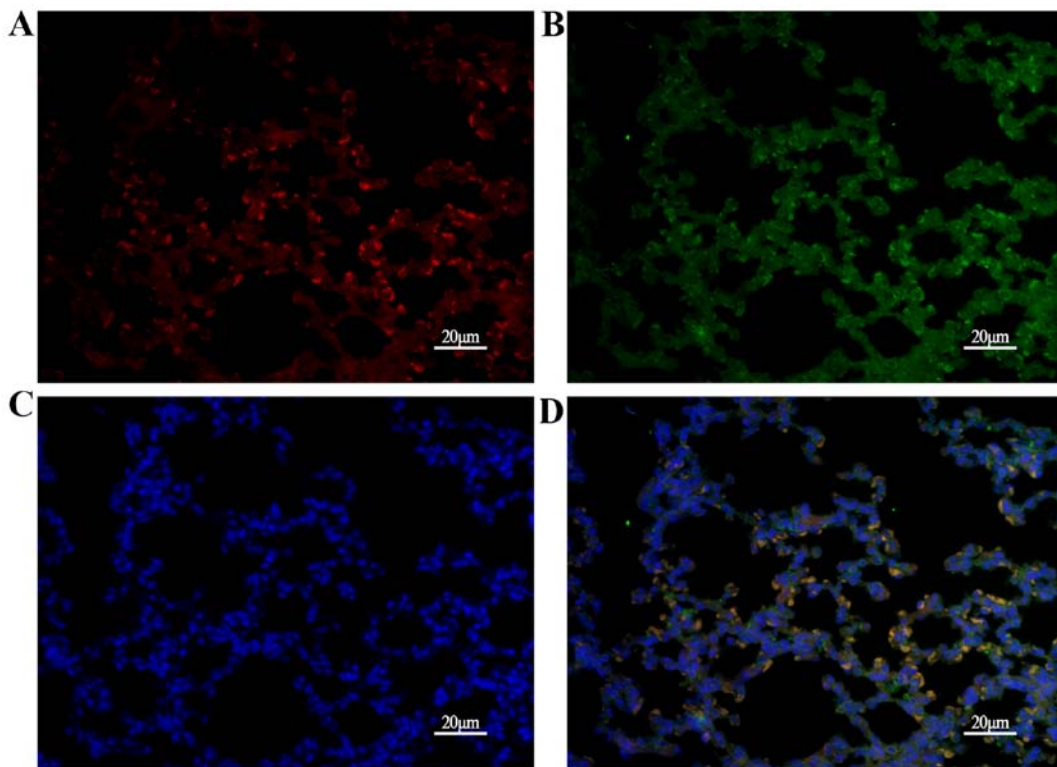


Figure 8. C/EBP α was partially colocalized with SUMO-1 in the alveolar epithelial cells. (A) Expression of C/EBP α at P14 was detected by immunofluorescence with phycoerythrin (red fluorescence). (B) Expression of SUMO-1 was detected by immunofluorescent fluorescein isothiocyanate (green fluorescence). (C) The nucleus was detected by DAPI (blue fluorescence). (D) Merged image. The yellow color indicates co-expression of C/EBP α and SUMO-1. C/EBP α , CCAAT enhancer binding protein- α ; SUMO, small ubiquitin-like modifier.

with the anti-C/EBP α antibody and immunoreacted with the anti-SUMO-1 antibody. The mobility of the SUMO-1-modified C/EBP α was found to be identical with that of the shifted band described above. Therefore, it was confirmed that this shifted C/EBP α band was generated by sumoylation.

Correlation between sumoylated C/EBP α and lung development. The above experiments suggested that the C/EBP α protein was sumoylated in the lung. C/EBP α is an established key regulator of lung development (25,26), and thus, the function of the sumoylated C/EBP α protein needs to be addressed. As shown in Fig. 9, sumoylated C/EBP α exhibits a developmental expression pattern during lung development, which was scarcely detected at the late pseudoglandular stage at E15.5; the expression increased and was maximal at E17.5, following which, the sumoylated C/EBP α decreased until the saccular period. At the alveolar stage, it increased slightly and finally stabilized. The decreased sumoylation of C/EBP α occurred at the lung development stage, which was characterized by extensive cell differentiation and secretion of pulmonary surfactant. The results demonstrated that the amount of glycogen exhibited a pattern approximately similar to the expression of sumoylated C/EBP α protein at the embryo and early postnatal stages. A negative correlation between sumoylation status of C/EBP α and lung differentiation was observed in Fig. 10A. In addition, as shown in Fig. 10B, the correlations of pulmonary surfactant secretion with C/EBP α and sumoylated C/EBP α were inverse. These data suggested that sumoylation may exert a suppressive effect on C/EBP α -mediated AEC-II differentiation and secretion.

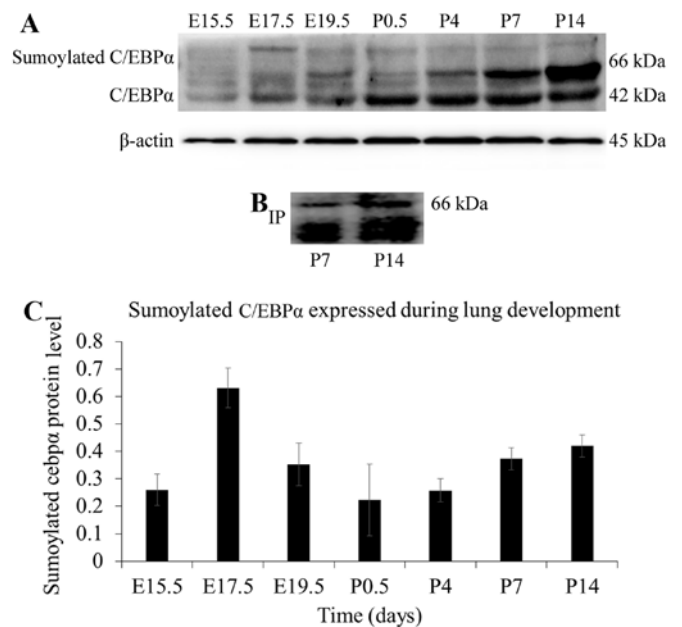


Figure 9. SUMO-modification of C/EBP α occurs during lung development. (A) Lung tissues from different development stages were lysed with mild lysis buffer and analyzed by immunoblotting with a specific antibody to C/EBP α . (B) IP, whole tissue lysates of P7 and P14 rat lungs were immunoprecipitated using anti-C/EBP α antibody and detected with anti-SUMO-1 antibody. (C) Sumoylated C/EBP α exhibits a developmental expression pattern during lung development. Data are presented as the mean \pm standard error of the mean. SUMO, small ubiquitin-like modifier; C/EBP α , CCAAT enhancer binding protein- α ; IP, immunoprecipitation; P, postnatal.

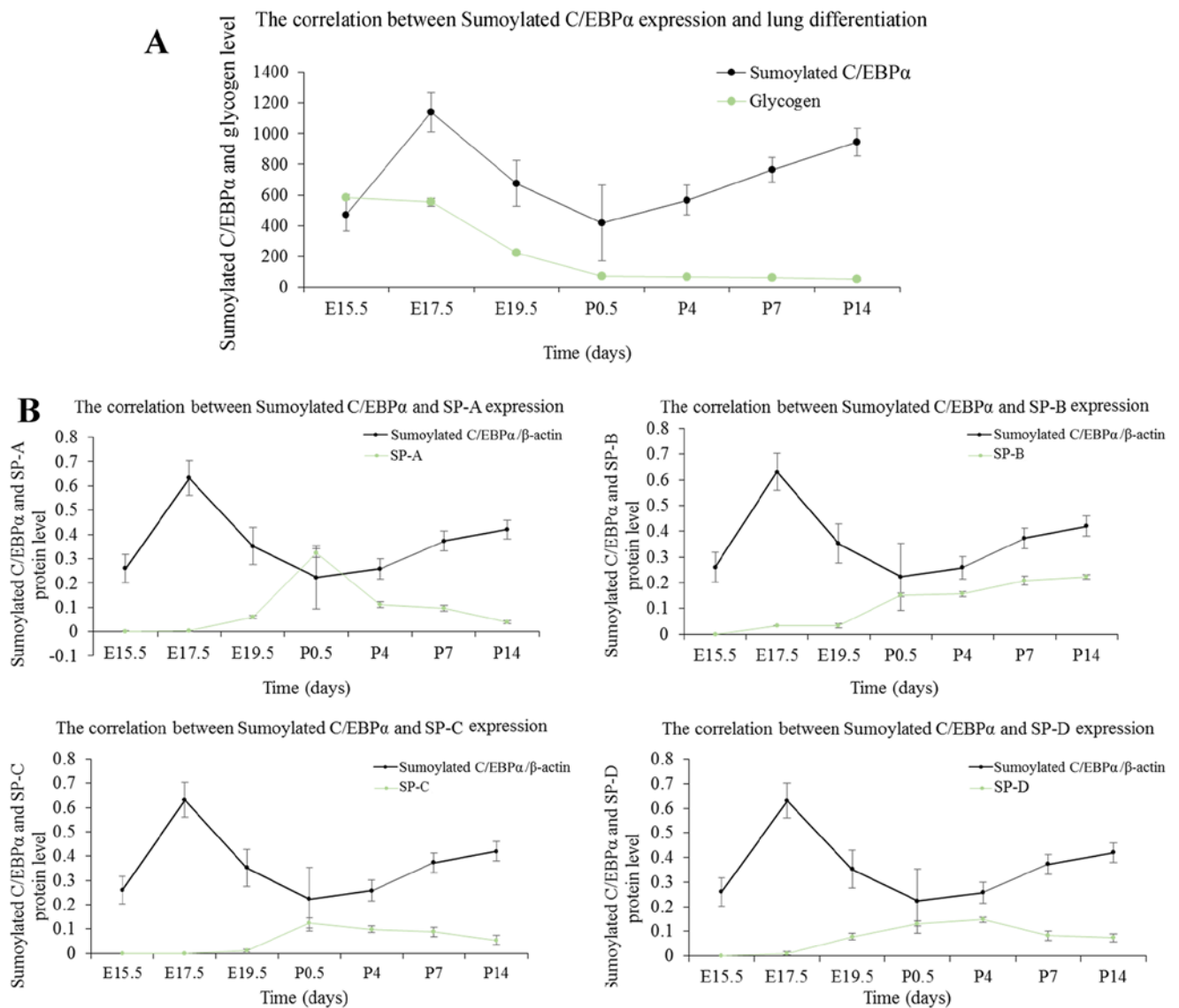


Figure 10. Sumoylated C/EBP α protein may play a role in lung differentiation and the secretion of pulmonary surfactant proteins. (A) Negative correlation between sumoylated C/EBP α and lung differentiation was observed during the period of lung differentiation, both stabilized at the late stage of lung development. (B) Sumoylated C/EBP α and SPs were negatively correlated, except SP-B. The expression of SP-B increased continually after birth. C/EBP α , CCAAT enhancer binding protein- α ; E, embryonic; P, postnatal.

Discussion

In the present study, the authors demonstrated that C/EBP α exhibits a dynamic expression pattern during lung development. To investigate the functional roles of C/EBP α in the lung, the authors analyzed its correlation to the differentiation and secretion of AEC-II, the accepted stem cell of the pulmonary epithelium (27). It was identified that the changes in the status of C/EBP α were associated with AEC-II differentiation as reflected by the content of glycogen during lung maturation. The expression pattern of C/EBP α protein is almost identical to that of SPs, and exerts a positive effect on AEC-II differentiation and secretion. In addition, SUMO was demonstrated to modify C/EBP α in the lung tissue, and sumoylation may have a negative effect on C/EBP α -mediated lung maturation.

During previous years, the physiological roles of C/EBP α have begun to be identified. C/EBP α is relatively highly expressed in the lung; is a key regulator of lung differentiation (25,26) and

directly activates the transcription of several lung-specific and differentiation-dependent genes (10,11). C/EBP α expression is initiated in close temporal proximity to the emergence of AEC-II and the original formation of pulmonary surfactant system. The result has shown a positive correlation between the C/EBP α expression and AEC-II differentiation and secretion. The expression patterns of SPs, except SP-B, are almost similar to that of the C/EBP α protein. The promoter of SP-A and SP-D gene harbors the C/EBP binding sites (8,9). The current observations are in agreement with those described previously; C/EBP α was positively correlated with the expression trend of the two hydrophilic proteins, SP-A and SP-D. SP-B and SP-C are small hydrophobic proteins; their promoter contains the TTF-1 binding sites (28). TTF-1 is the synergy transcription factor of C/EBP α . C/EBP α , SP-A and SP-D occur primarily in AEC-II, and to a lesser extent in bronchioalveolar epithelial (Clara) cells (7,29,30). On the other hand, the expression of the SP-C gene is restricted to AEC-II (31), whereas that of SP-B occurs in

both AEC-II and bronchiolar epithelial cells (29). As a result, a part of SP-B is not regulated by C/EBP α . In summary, C/EBP α exhibits a dynamic expression pattern during lung development; the expression of C/EBP α is related to the glycogen content and the secretion of SPs. Altogether, these results suggested that the role of C/EBP α in lung maturation is exercised by the regulation of pulmonary surfactant.

The authors demonstrated that C/EBP α could be modified by SUMO-1 protein in the lung; the sumoylated C/EBP α displays a negative correlation with AEC-II differentiation and secretion. This suggests that SUMO-modification may be involved in the regulation of C/EBP α -induced lung maturation. SUMO-1 modifies C/EBP α at lysine residue 159 within a conserved domain that negatively modulates the transcriptional activity (17). Sumoylation confers new protein-protein interaction properties of the transcription factors, which in turn can significantly alter the transcriptional activity. Histone deacetylases (HDACs) are typically correlated with the repression of transcription and can be recruited to sumoylated transcription factor complexes. C/EBP α promotes the transcription of several genes expressed in a tissue-specific and differentiation-dependent manner (32,33); HDAC3 interacts with sumoylated C/EBP α to negatively regulate the liver X receptor alpha expression (34). C/EBP α associates with the SWI/SNF complex and p21 can increase its stability and transcription activity; however, the sumoylation of C/EBP α inhibits the binding with these factors (19,35,36). Taken together, the presented results suggested that sumoylation might suppress the C/EBP α -mediated transactivation of SP genes and restrain PC synthesis indirectly. SPs are differentiation-dependent genes, and these results were consistent with the finding in our study that the sumoylation of C/EBP α expression is converse to lung differentiation. Therefore, additional studies on the effect of ectopic sumoylation of C/EBP α in lung are essential.

Furthermore, the results showed that the level of C/EBP α sumoylation was low in the developing lung, less than half of the total level of C/EBP α , similar to that reported for other sumoylated proteins (37-40). However, the physiological effects of sumoylation were abstruse. Intriguingly, the descending expression of sumoylated C/EBP α corresponded to the increasing C/EBP α at canalicular and saccular stages, the sophisticated metamorphosis period during the lung development. Injury during this phase will inhibit the expression of C/EBP α , resulting in abnormal lung differentiation and pulmonary surfactant secretion (41). In addition, a transient increase in the expression of SUMO-1 was observed at P4. The lung tissue was in a period of expansion during the postnatal days 1-3, which led to the improvement of pulmonary morphology and functions. At P4, the differentiation of AEC-II approached equilibrium, and the proliferation served as the major hallmark of expression. Sato *et al* (19) found that sumoylation of C/EBP α blocked its inhibitory effect on cell proliferation by disrupting the formation of a proliferation-inhibition complex owing to the failure of interaction between sumoylated C/EBP α and other proteins. This phenomenon has not yet been reported in the lung. However, the common endodermal origin and the crucial role of C/EBP α in the lung and liver suggested a common role of SUMO in both organs. This feature may provide a valuable insight into further investigations. The current results found that SUMO-modification participates in normal C/EBP α -mediated

lung development, and offers a possibility that it might continue to partake in abnormal ways; thus, sumoylated C/EBP α may be a putative therapeutic target in lung injury.

Two isoforms were generated by the C/EBP α gene, with the resultant p42 kDa (full length) and p30 kDa (truncated) C/EBP α proteins, and the N-terminus is abbreviated in the p30 kDa protein. The heterodimerization between the two isoforms can restrain the ability of p42 C/EBP α to transactivate the target genes (24). The p30 C/EBP α can be sumoylated, although it is less sensitive to the changes in sumoylation than the p42 isoform (18). p30 C/EBP α enhances the sumoylation of p42 C/EBP α via the upregulation of Ubc9, which results in decreased transcriptional activity of the latter (15). Changes in the ratio of p42:p30 have been confirmed to contribute to tipping the scales from normal myelopoiesis to pre-leukemic or even leukemic myelopoiesis (42). As indicated in the results, there is a shifted band with electrophoretic mobility just lower to that of the sumoylated C/EBP α . We also observed a band at the consistent location by the immunoprecipitation assay. The shifted band could be speculated as the sumoylated p30 C/EBP α . Since similar investigations in the lung are absent, further studies are essential.

In conclusion, the authors found that C/EBP α exhibits a dynamic expression pattern during lung development and exhibits a positive effect on AEC-II differentiation and secretion. SUMO post-translationally modifies C/EBP α that occurs in the lung tissue; its expression during development corresponded to that during lung differentiation and several differentiation-dependent genes expression. Furthermore, these results suggested that sumoylation may act as a negative regulator of the C/EBP α -mediated transactivation in the lung.

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