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Surfactant Protein-A as an Anti-Inflammatory Component in the Amnion: Implications for Human Pregnancy¹

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Abstract

The mechanism of mouse parturition is thought to involve myometrial infiltration by amniotic fluid (AF) macrophages, activated by surfactant protein-A (SP-A). In humans, the concentration of AF SP-A decreases during labor, and no fetal macrophages are found in the myometrium after labor. Therefore, it appears that the mechanisms of labor in mice and humans are different. We investigated a potential role for SP-A in human pregnancy and parturition by examining SP-A expression patterns in AF and amnion. High molecular weight (HMW; >250 kD) oligomeric SP-A was increased in AF with advancing gestation. Interestingly, these oligomers were more abundant in placental amnion before labor at term, while they increased primarily in reflected amnion during labor (p < 0.05). Immunoblotting showed a binding of HMW SP-A in AF to amnion. In C57BL/6 mice, oligomeric SP-A was also readily detected in AF from E15 onwards, but not in amnion. Macrophage density in mice myometrium did not change with advancing gestational age. Microarray analysis of human amnion explants incubated with SP-A revealed a molecular signature of inhibited cytokine-cytokine receptor interaction with down-regulation of IL-1β, CXCL2, and CXCL5 mRNA expression. The findings herein strongly suggest that SP-A signals amniotic anti-inflammatory response via amniotic fluid during pregnancy. We propose that a SP-A interaction among amniotic fluid, placental amnion, and reflected amnion is a unique mechanism for immunoregulation in human pregnancy akin to that established in lung biology. However,

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amniotic fluid SP-A and fetal macrophages by themselves do not seem to be exclusive effectors of parturition in humans.

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Keywords

human; inflammation; mucosa; macrophages; reproductive immunology

Introduction

Human spontaneous labor at term has a prominent pro-inflammatory molecular signature in the chorioamniotic membranes, particularly in the amnion (1,2). This strongly suggests that human parturition has a physiologic inflammatory/immunologic component, and also that the molecules which modulate amniotic pro-inflammatory responses play important roles in parturition. The amnion is a large mucosal surface stretched across the intrauterine surface, which reaches up to $1,876 + -307 \text{ cm}^2$ at term (3), and is the inner layer of the chorioamniotic membranes. The amnion is in direct contact with the amniotic fluid and plays pivotal roles in the induction of labor by producing prostaglandins and in delivery by membrane rupture (4).

Recently, we have demonstrated that the human amnion is biologically compartmentalized into the placental amnion (amnion overlying the placental disc) and the reflected amnion (amnion of the extraplacental chorioamniotic membranes). Of note, the labor-associated proinflammatory response was a more prominent feature in the placental amnion rather than in the reflected amnion (2). Previous studies have shown that inflammation activates amniotic biological cascades associated with human labor. For example, pro-inflammatory cytokines IL-1 and tumor necrosis factor- α (TNF α) stimulated the production of PGE₂ and induced cyclooxygenase-2 expression in the amnion (5,6). IL-1 β and TNF α increased IL-8 expression in the amnion cells (7). Moreover, intra-amniotic infection is the leading identifiable cause of preterm labor and delivery (8). During pregnancy, the biochemical composition of amniotic fluid changes as gestational age advances, and this could have an effect on the biology of the amnion. Conversely, the molecules produced by the amnion could change their amniotic fluid concentrations. In this context, it is reasonable to expect that a certain molecule, whose amniotic fluid concentration changes with progression of gestation and modulates amniotic inflammatory responses, plays an important role in human pregnancy and parturition.

Surfactant protein-A (SP-A) is a C-type lectin encoded by *SFTPA1* and *SFTPA2* (9,10), with a triple helical collagen region on the N-terminus and a carbohydrate recognition domain (CRD) on the C-terminus (11,12). It is expressed by both the lung and the chorioamniotic membranes (13–16). SP-A characteristically shows a marked increase in the amniotic fluid with advancing gestational age, and this is taken as an indicator of fetal lung maturity (17). Because its concentration sharply increases during the third trimester, compared to that of SP-D (18), SP-A is an attractive candidate to control human parturition because it would

link fetal maturity with the onset of labor. Indeed, intra-amniotic administration of SP-A was shown to induce parturition in mice, and treatment with a neutralizing antibody prolonged gestation in that species. Therefore, it has been proposed that activation of fetal amniotic fluid macrophages by SP-A and their subsequent migration into the maternal myometrium are key events for spontaneous parturition in mice. The injection of neutralizing anti-SP-A Ab into the amniotic cavity delayed mouse parturition (19,20). However, fetal macrophages have not been found in the myometrium of women after spontaneous labor at term (21,22), and amniotic fluid SP-A concentration decreases during human spontaneous labor at term (23). Therefore, a puzzling difference in the role of SP-A between humans and mice has emerged.

We hypothesized that decreased amniotic fluid SP-A concentration during human labor at term might be a consequence of sequestration of SP-A by its receptor molecules in the chorioamniotic membranes, particularly within the amnion (23). Since human amnion expresses known receptors of SP-A, such as TLR2 and TLR4, which are critical to immunological responses (24–26), it is very likely that amniotic fluid SP-A modulates the amniotic inflammatory response. It was also shown that the globular heads and the collagen-like tails of SP-A could mediate anti-inflammatory and pro-inflammatory responses by binding to SIRP α (signal-regulatory protein alpha) and to calreticulin/LRP1 (low-density lipoprotein-related protein 1; CD91), respectively, thus conferring dual immunomodulatory roles for SP-A (27). However, the primary immunological impact of SP-A on the human amnion and its biological significance in the amniotic cavity has not been investigated to date. This study was performed to assess immunological and functional aspects of SP-A in human pregnancy and parturition.

Materials and Methods

Patient materials

Samples of human amnion and amniotic fluid were retrieved from the Bank of Biological Materials held at the Perinatology Research Branch, *Eunice Kennedy Shriver* National Institute of Child Health and Human Development, National Institutes of Health. Placental amnion and reflected amnion were obtained using blunt dissection as previously described (2). The study groups were composed of women at term not in labor (TNL; n=28) and term in labor (TIL; n=19). In order to achieve the separation of amnion epithelium and mesoderm, reflected amnion samples obtained from TIL patients were used (n=5). The amnion was rinsed and soaked in PBS for 10 min at room temperature, and the mesodermal layer was meticulously scraped off using a razor blade. Amniotic fluid samples (n=35) were obtained by amniocentesis for clinical indications at different gestational ages. Fetal lung tissue was obtained from an autopsy case of fetal demise *in utero* at the gestational age of 33 weeks, and amniotic fluid and amnion from the same patient were also obtained for comparison. All patients provided written, informed consent, and the collection and use of the samples were approved by the Institutional Review Boards of the participating institutions.

Primary amnion cell and amnion explant cultures

Human amnion cells were obtained from the reflected amnion of TNL patients and incubated for 6 h with Dulbecco's Modified Eagle Medium (DMEM, Mediatech, Inc) containing 2.5% (v/v) amniotic fluid and 10% FBS. Human reflected amnion explants (2.5 cm \times 2.5 cm) obtained from TNL patients were incubated for 6 h with DMEM with 2.5% amniotic fluid pre-adsorbed with rabbit polyclonal neutralizing anti-SP-A Ab (Chemicon International Inc.) or normal rabbit sera, and harvested after washing with PBS. For the analysis of the extent of extracellular SP-A binding in the amnion, the amnion tissues were

treated with 2% collagenase (Worthington Biochemical Corporation) at 37°C for 30 min and used for protein isolation.

Animal experiments

This study was approved by the Animal Investigation Committee of Wayne State University, Detroit, Michigan. Timed pregnant female C57BL/6 mice (n=16; Charles River Laboratories International Inc.) were maintained in an animal facility under pathogen-free conditions. Mice were sacrificed on E13, E15, E17, and E19. Fetal, uterine, placental tissues, and amniotic fluid were collected.

Immunoblotting

Proteins were extracted from liquid nitrogen-pulverized amnion, fetal lung tissue, or cell pellets using RIPA lysis buffer (Sigma) containing a proteinase inhibitor cocktail (Roche). Protein lysates were electrophoresed in 4-15% gradient SDS-PAGE (Bio-Rad) under reducing or non-reducing conditions, and then electro-blotted onto polyvinylidene difluoride membranes (GE Healthcare Bio-Sciences Corporation). After blocking for 1 h with 5% nonfat dry milk in Tris-buffered saline (TBS) containing 0.1% Tween 20 (0.1% TBS-T), the membranes were incubated with primary Abs specific to SP-A (rabbit polyclonal, Chemicon International Inc.; goat polyclonal, Chemicon International Inc.; mouse monoclonal, AntibodyShop), CD91 (mouse monoclonal, 5A6, Abcam Inc.), calreticulin (mouse monoclonal, FMC75, Abcam Inc.), SIRPa1 (rabbit polyclonal, Upstate Biotechnology Inc.), TLR4 (mouse monoclonal, Santa Cruz Biotechnology, Inc.), MD2 (rabbit polyclonal, Abcam Inc.), TLR2 (rabbit polyclonal, Abcam Inc.), HPRT (rabbit polyclonal, Santa Cruz Biotechnology, Inc.), β -actin (mouse monoclonal, Sigma-Aldrich), and GAPDH (mouse monoclonal, Sigma-Aldrich) at 4°C overnight. HRP-conjugated anti-mouse IgG, anti-rabbit IgG, or anti-goat IgG (Santa Cruz Biotechnology, Inc.) was used as a secondary Ab. Signals were detected by chemiluminescence.

Immunoprecipitation

Amnion proteins were obtained using lysis buffer (pH 8.0) containing 150 mM NaCl, 50 mM Tris-HCl, 1% NP-40, and a proteinase inhibitor cocktail. A total of 125 micrograms of protein lysates were pre-incubated with protein G agarose (Upstate Biotechnology Inc.) for 2 h on a rotary shaker, and then specific primary Abs and CaCl₂ (final concentration: 1 mM) were added for overnight incubation at 4°C. After washing, immune complexes harvested with protein G agarose were eluted and electrophoresed in SDS-PAGE under non-reducing conditions for further analysis. Immunoblottings were carried out as described above.

Immunofluorescent staining

Double label immunofluorescent staining on human amnion tissue was conducted using a panel of Abs to SP-A (rabbit polyclonal, Chemicon International Inc.), CD68 (mouse monoclonal, Chemicon International Inc.), CD91 (mouse monoclonal, Zymed), and TLR4 (mouse monoclonal, Santa Cruz Biotechnology, Inc.). Five-micrometer-thick frozen tissue sections of reflected amnion tissues (TIL; *n*=3) or intact (whole mount) placental amnion tissues (TNL; *n*=3) were fixed with 4% (w/v) paraformaldehyde or 85% (v/v) ethanol, permeabilized with 0.25% Triton X-100 and incubated with 5% (w/v) BSA in PBS for 30 min at room temperature. Sections or tissues were incubated with a primary Ab in 1% (w/v) BSA in PBS for 1 h, followed by incubation with Alexa 488 goat anti-mouse IgG and Alexa 594 donkey anti-rabbit IgG (Invitrogen) in 1% (w/v) BSA for 30 min and mounted in ProLong® Gold antifade reagent with DAPI (Invitrogen). The stained sections were examined using a Leica TCS SP5 spectral confocal system (Leica Microsystems).

Assessment of SP-A oligomerization in amniotic fluid

To assess whether immunoreactive bands corresponding to MMW and HMW SP-A represent oligomeric SP-A, 2.5 microliters of amniotic fluid obtained by amniocentesis for amnioreduction in a twin pregnancy at the gestational age of 29 weeks and 10 nanograms of SP-A isolated from bronchoalveolar lavage fluid of a patient with pulmonary alveolar proteinosis (a generous gift from Dr. Whitsett, University of Cincinnati College of Medicine) were resolved in 4–15% gradient SDS-PAGE under non-reducing conditions. The pieces corresponding to 50-65 kD, 150-250 kD, and more than 250 kD were cut from the gel and homogenized using liquid nitrogen. The proteins in gel homogenates were isolated using Laemmli Sample Buffer (Bio-Rad). Isolated proteins were electrophoresed again in 4-15% gradient SDS-PAGE under reducing conditions, and immunoblotting was performed with anti-SP-A Ab (mouse monoclonal, AntibodyShop). For two-dimensional gel electrophoresis, protein samples extracted with ReadyPrep Rehydration/Sample Buffer (Bio-Rad) were rehydrated with 7 cm IPG strip (pH 4-7, Bio-Rad) at room temperature for 12 h followed by isoelectric focusing (IEF) using a PROTEAN IEF Cell (Bio-Rad). Voltages were applied at 250V for 20 min, followed by 4,000V for 2 h and 4,000V for 10,000V-h. The focused IPG strips were equilibrated and run on 4–15% gradient Mini-PROTEAN Ready Gel with IPG comb (Bio-Rad) followed by immunoblotting with mouse monoclonal anti-SP-A Ab.

Liquid chromatography-tandem mass spectrometry (LC-MS/MS)

To further characterize the oligomeric nature of HMW SP-A in amniotic fluid and the specificity of immunoreactive bands, 25 microliters of amniotic fluid were immunoprecipitated with rabbit polyclonal anti-SP-A Ab (Chemicon International Inc.) as previously described. Amniotic fluid immunoprecipitates and purified SP-A were resolved in 4-15% gradient SDS-PAGE under non-reducing conditions, the region corresponding to the HMW SP-A immunoreactive band (>250 kD) was cut from the gel, and proteins were extracted as described earlier. Isolated proteins were electrophoresed in 4-15% gradient SDS-PAGE under reducing conditions. Both immunoblotting for SP-A and Coomassie blue staining were performed. After destaining, slices corresponding to SP-A bands (32 kD and 64 kD) were cut and analyzed by LC-MS/MS. Gel slices were cut into 1 mm³ cubes and each was washed with water for 15 min; the liquid was replaced with 50 mM ammonium bicarbonate (AB):acetonitrile (ACN) (1:1, v:v). Liquid was removed and gel plugs were shrunk with ACN. The gel pieces were rehydrated with AB, and after 5 min an equal volume of ACN was added. After 15 min, the liquid was removed, and ACN was added. After shrinking the gel plugs, the ACN was removed and the gel pieces dried in a vacuum centrifuge. The gel plugs were allowed to swell in 10 mM DTT/AB and then to reduce for 1 h at 55°C. After reaching room temperature, the excess liquid was replaced with 55 mM iodoacetamide/AB and incubated for 1 h in the dark. The liquid was removed and the gel plugs were washed twice for 15 min with AB:ACN (1:1). The gel pieces were shrunk with ACN and dried as explained above. In-gel digestion was performed overnight at 37°C after the addition of sequencing grade trypsin (Sigma) in 25 mM AB. Peptides were extracted from the gel plugs the next day using two rounds of the following protocol: 1) Water was added to the gel pieces which were then incubated for 15 min, 2) An equal volume of ACN was added, sonicated for 2 min, and incubated for 15 min, 3) Excess liquid was transferred to an Eppendorf tube and the gel pieces were shrunk with ACN, and 4) After 15 minutes, excess ACN was transferred to an Eppendorf tube. Volatile buffers and salts of the peptides were removed during two rounds of drying and resolubilizing in water. Peptides were resuspended in 5% ACN, 0.1% formic acid, and 0.005% TFA, and separated by reverse phase chromatography (Magic C18 column, Michrom BioResources, Inc.). Peptides were ionized with the ADVANCE ion source (Michrom BioResources, Inc) and introduced into an LTQ XL mass spectrometer (Thermo Fisher Scientific Inc.). Abundant species were

fragmented with collision-induced dissociation (CID). Data analyses were performed using BioWorks (Thermo Scientific) and Scaffold (Proteome Software, Inc.) software that incorporated SEQUEST and ProteinProphet algorithms. A human protein database (IPI ver. 3.49) was used for the analysis.

Microarray analysis

To study the effects of SP-A treatment in the amnion, microarray experiments were conducted. Human reflected amnion explants $(2.5 \times 2.5 \text{ cm})$ obtained from TNL patients (n=6) were kept in DMEM media containing 0.5% FBS for 1 h and were treated with purified SP-A. Vehicle (distilled water) and SP-A were negative for endotoxin to <0.005 EU/ml for the water and <0.005 EU/µg for SP-A. The amnion explants were treated with SP-A at a final concentration of 1 µg/ml for 12 h, and total RNA was isolated using TRI reagent (Ambion). After purification of total RNA samples using an RNeasy Mini Kit, 100 nanograms of total RNA were amplified and biotin-labeled with the Illumina TotalPrep RNA Amplification kit (Ambion). Labeled cRNAs were hybridized to Illumina's HumanHT-12 v3 expression BeadChips. BeadChips were imaged using a BeadArray Reader, and raw data was obtained with BeadStudio Software (Illumina). After logarithmic transformation, the bead-level expression intensities were normalized using the quantile normalization algorithm (28). A paired moderated t-test using the *limma* package (29) of Bioconductor was used to gauge the effect of treatment on gene expression levels. A combined criteria was used to identify differentially expressed probes which included a false discovery rate (FDR) (30) adjusted p value of less than 0.25 and a fold change greater than 1.5 fold. The differentially expressed genes were further analyzed to identify enriched Gene Ontology terms, such as molecular functions and biological processes, using the GOstats package (31) of Bioconductor. Moreover, pathway analysis was performed using an overrepresentation approach on the Kyoto Encyclopedia of Genes and Genomes (KEGG) human signaling and metabolic pathways database (www.genome.jp/kegg/). For the signaling pathways available in KEGG, the signaling pathway impact analysis (SPIA) algorithm (32,33) was applied to make full use of the pathways topology and the directed gene-gene interactions to assess pathway significance. This later analysis was performed using the SPIA package of Bioconductor. The complete microarray data set is available in a MIAME (Minimal Information About a Microarray Experiment)-compliant format in the ArrayExpress database (www.ebi.ac.uk/microarray-as/ae) [entry ID: E-TABM-741]).

Real-time Reverse Transcription-Polymerase Chain Reaction (qRT-PCR)

To validate the microarray analysis results, expression of select cytokine and chemokine genes were analyzed using TaqMan gene expression assays for IL-1 β (HS00174097_m1), CXCL2 (HS00236966_m1), CXCL5 (HS00171085_m1), and CCL3 (HS00234142_m1). Amniotic total RNA was reverse transcribed using ImPromIITM reverse transcriptase and oligo(dT) primers (Promega Corporation), and subsequently used for PCR. PCR was done using an ABI 7500 Fast Real Time PCR System (Applied Biosystems Inc.). The human ribosomal protein, large, P0 (RPLP0; Applied Biosystems Inc.) was used for standardization of PCR results.

Amniotic fluid cytology preparation

Human amniotic fluid samples obtained by amniocentesis were added to PreservCyt transport medium (Cytyc Corporation), and the slides for cytologic examination were prepared using a ThinPrep 2000 processor (Cytyc Corporation). Immunohistochemical staining was performed for CD68 using a mouse monoclonal anti-CD68 Ab (Dako).

Flow cytometry

The amniotic fluid samples obtained from five patients were analyzed. Cord blood samples from the same patients were available in two cases and used as positive controls for CD14 staining (data not shown). To stain cord blood, whole blood (100 μ l) anticoagulated with EDTA was incubated for 30 min at 4 °C with APC-conjugated isotype control or APC-conjugated mouse anti-CD14 (BD Biosciences). Red blood cells were lysed by incubation with 1X FACS Lysing Solution (BD Biosciences), and the cells were pelleted by centrifugation at 300×g for 5 min, and washed twice with PBS. Stained cells were finally suspended in 1% paraformaldehyde in PBS for flow cytometric acquisition. Amniotic fluid was centrifuged at 300xg for 5 min, washed with PBS, resuspended in BD staining buffer (BD Biosciences) and stained as whole blood. All acquisition and analyses were performed with a LSRII flow cytometer (BD Biosciences).

Measurement of macrophage density in the mouse myometrium

To evaluate intra-myometrial macrophage density, immunostaining was performed using a rat monoclonal anti-F4/80 Ag (Serotec). The numbers of F4/80 positive macrophages were counted in the myometrial sections containing the mid-plane of the placenta from four pregnant mice at each gestational age (E13, E15, E17, and E19). Eight high-power fields corresponding to every 45° angle point starting from the mesometrial border in a clockwise direction, covering the entire myometrial circumference, were evaluated in each section. Image-Pro Plus 6.2 software (Media Cybernetics, Inc.) was used for the analysis.

Statistical analysis

The Mann-Whitney U test was used to compare the relative densities of high molecular weight (HMW) SP-A oligomers in the reflected amnion between TIL and TNL cases. Comparisons of protein expressions between different anatomical structures (epithelium vs. mesoderm, placental amnion vs. reflected amnion) and qRT-PCR data obtained from amnion explants according to SP-A treatment were analyzed using the Wilcoxon signed-rank test. The Kruskall-Wallis test was used for the analysis of myometrial macrophage counts at each mouse gestational age. SPSS version 15.0 (SPSS Inc.) was employed for statistical analysis. A p value of <0.05 was considered statistically significant. The methods used for microarray analysis have been described above in the appropriate section.

Results

Oligomeric SP-A in the amniotic fluid and reflected amnion increases with advancing gestational age and labor

Upon immunoblotting for SP-A using amniotic fluid, amnion, and fetal lung samples (all obtained from a single fetal demise *in utero* case at the gestational age of 33 weeks) both in reducing and non-reducing gel electrophoresis conditions, anti-SP-A Abs generated from different species (rabbit polyclonal, goat polyclonal, and mouse monoclonal as described above) showed different immunoreactive patterns. The immunoblotting results clearly indicated that SP-A is largely present in oligomers *in vivo*. The major SP-A bands corresponded to non-reducible dimers and supra-trimeric oligomers. Under non-reducing conditions (Fig. 1*A*), the rabbit polyclonal anti-SP-A Ab most effectively detected HMW (>250 kD) SP-A oligomers and non-reducible dimers. On the other hand, both goat polyclonal and mouse monoclonal anti-SP-A Abs effectively recognized medium molecular weight (MMW; 75 kD-250 kD) SP-A oligomers. Under reducing conditions (Fig. 1*B*), non-reducible SP-A dimers with a molecular weight of 64 kD were readily detected by both rabbit and goat polyclonal Abs. Monomeric SP-A polypeptides were detected by all of the Abs, particularly by the mouse monoclonal Ab.

In order to confirm that the amniotic fluid proteins in the HMW and MMW bands detected by anti-SP-A antibodies were genuine SP-A oligomers, proteins were isolated from gel slices corresponding to SP-A in HMW and MMW regions electrophoresed under nonreducing conditions. For this experiment, amniotic fluid obtained from a twin pregnancy at 29 weeks of gestation was used. Isolated proteins were electrophoresed again under reducing conditions, and analyzed by immunoblotting. During immunoblotting using mouse monoclonal anti-SP-A Ab, monomeric SP-A polypeptides were detected in both MMW and HMW SP-A. The 50–65 kD fraction mainly demonstrated a 64 kD immunoreactive band, which was consistent with non-reducible dimers previously described (Fig. 2A) (34). We also performed LC-MS/MS analysis of amniotic fluid SP-A from immunoprecipitates using rabbit polyclonal antibody and purified SP-A (Fig. 2, B and C), and confirmed that HMW SP-A is an oligometric form composed of SP-A monomers and non-reducible dimers. Mass spectra of amniotic fluid SP-A is shown in Figure 2C and identified peptides are summarized in Table I. Furthermore, two-dimensional electrophoresis of the HMW fraction of amniotic fluid SPA (n=3), followed by immunoblotting with mouse monoclonal anti-SP-A Ab, revealed the same isomer patterns as that of purified SP-A (Fig. 2, D, E, F, and G). The immunoblotting data were quite consistent with the observation made by He (35) that SPA multiple isomers (mass 20.5-26 kD, pI of 4.5-5.4; mass 26-32 kD, pI of 4.5-5.4; mass 32–42 kD, pI of 4.5–5.4) are found in human bronchoalveolar lavage fluid. However, isomers with molecular mass ranging between 50 kD-75 kD were not as prominent or absent in amniotic fluid HMW SP-A compared with purified pulmonary SP-A.

Evaluation of amniotic fluid SP-A throughout the gestational period, using both rabbit polyclonal and goat polyclonal Abs, revealed that HMW SP-A oligomers, not MMW SP-A oligomers, increased with advancing gestational age and became a predominant form in the amniotic fluid in third-trimester cases (Fig. 3, *A* and *B*). Oligomeric SP-A bands were also readily detected in the human reflected amnion. A comparison of reflected amnion samples obtained from TNL and TIL patients showed a significant increase in HMW SP-A oligomers after labor on densitometric analysis (p < 0.05; Fig. 3, *C*, *D*, and *E*).

To compare amniotic fluid SP-A profiles of humans and mice, samples of amniotic fluid and fetal membranes obtained from timed-pregnant C57BL/6 mice (E13, E15, E17, and E19) were analyzed. As in humans, non-reducible dimers and supratrimeric oligomers of SP-A were also readily detected in mouse amniotic fluid from the gestational age of E15 and gradually increased (Fig. 3*F*). HMW SP-A oligomers were evident from E17. However, in contrast to humans, supratrimeric SP-A was not detected in the fetal (extraembryonic) membranes of the placenta (Fig. 3*G*).

Binding of amniotic fluid SP-A to the amnion as a distinct feature of human pregnancy

The next line of investigation was conducted to determine whether SP-A oligomers in the amnion are, at least in part, originating from the amniotic fluid as there was a reciprocal relationship between amniotic fluid SP-A concentration and HMW oligomeric SP-A in the reflected amnions of TIL patients (23). Immunoprecipitation of amnion lysates obtained from TNL and TIL patients using Abs against SP-A, SIRP α , calreticulin, MD2, and TLR2, and subsequent immunoblotting for SP-A, showed co-immunoprecipitation of SP-A with these known receptor molecules tested (Fig. 4, *A* and *B*). Immunofluorescent staining of the placental amnion for SP-A, CD91, and TLR4 demonstrated distinct co-localizations of SP-A with CD91 (Fig. 4*C*) or TLR4 (Fig. 4*D*) in the epithelium.

Furthermore, incubation of primary amnion cells obtained from TNL patients with amniotic fluid depleted HMW oligomeric SP-A in amniotic fluid in a time-dependent manner, with a reciprocal increase of HMW oligomeric SP-A in amnion cells (Fig. 5A). The HMW oligomeric SP-A was not detected in the amnion incubated with amniotic fluid in the

presence of neutralizing anti-SP-A Ab (Fig. 5*B*), showing that HMW oligomeric SP-A in the amnion largely originates from amniotic fluid. Interestingly, this was not the case with MMW SP-A oligomers as they did not change after incubation with amnion cells (Fig. 5*C*). The extracellular binding nature of oligomeric SP-A was also supported by a drastic decrease in the intensity of HMW oligomeric SP-A bands on the immunoblots after collagenase treatment of the amnion (Fig. 5*D*).

Intra- and inter-amniotic compartmentalization of oligomeric SP-A binding in humans

The human amnion is composed of amnion epithelium and mesoderm (connective tissue). Myofibroblasts and macrophages are major cellular subsets of the mesodermal layer (36). Upon immunofluorescent staining of the chorioamniotic membranes, SP-A immunofluorescence was found in both amniotic epithelial and mesodermal cells (Fig. 6*A*). Immunoblotting of amnion epithelium and mesoderm obtained from TIL patients also demonstrated more abundant oligomeric SP-A in the mesoderm compared to the epithelium (Fig. 6*B*). Expression patterns of known SP-A receptor molecules were also different between the two layers. Interestingly, TLR2 and MD2 expressions were higher in the mesoderm, while the expressions of other receptors, such as CD91, calreticulin, SIRP α , and TLR4 were higher in the epithelium (Fig. 6*B*). Differential expressions of all these proteins between the epithelium and mesoderm were significant upon densitometric analysis (p < 0.05 for all).

Previously, we showed that a prominent increase in IL-1 β mRNA expression associated with labor at term was a more pronounced feature of the placental amnion rather than the reflected amnion (2). Therefore, further analyses of the distribution patterns of SP-A and receptors of SP-A (CD91, calreticulin, SIRP α , TLR2, TLR4, and MD2) between the placental amnion and the reflected amnion were performed by serial immunoblottings using a panel of specific Abs. In TNL cases (*n*=6), HMW oligomeric SP-A was predominantly localized in the placental amnion compared to the reflected amnion, but this difference was not noted in TIL cases (*n*=6), largely due to a decrease and an increase in SP-A in the placental amnion and the reflected amnion, respectively (Fig. 6*C*). This difference was significant upon densitometric analysis of HMW SP-A oligomers in placental amnion and reflected amnion between TNL and TIL cases (*p* < 0.01). The median RA/PA SP-A density ratio was 0.27 (range: 0.13–0.77) in TNL cases, while it was 1.86 (range: 0.4–3.27) in TIL cases. Among the receptors screened – CD91, calreticulin, SIRP α , TLR2, TLR4, and MD2 –, the expression of CD91 (*p* = 0.05) and SIRP α (*p* = 0.07) tended to be higher in the placental amnion (Fig. 6*C*).

Biological effects of SP-A in the amnion

As the findings indicated significant binding of amniotic fluid SP-A to the amnion, we examined the effects of SP-A in the amnion *in vitro*. Primary reflected amnion explants obtained from TNL patients were treated with SP-A for 12 h at the concentration of 1 µg/ml. Fig. 7*A* shows a "volcano plot" summarizing the differential expression of the genes between SP-A-treated and non-treated groups. Sixty-five probes corresponding to 60 unique genes met our significance criteria (Table II). More than 100 biological processes, including *angiogenesis* and *response to stimulus*, were significantly enriched in the list of differentially expressed genes. The SPIA identified the Cytokine-cytokine receptor interaction pathway as being significantly impacted, more specifically, inhibited, by the SP-A treatment. qRT-PCR analyses confirmed a significant decrease in mRNA expression of IL-1β (p < 0.05), CXCL2 (p < 0.05), and CXCL5 (p < 0.05) following SP-A treatment. CCL3 mRNA expression was not significantly different (Fig. 7*B*).

Macrophages in the amniotic fluid and myometrium: Observations in humans and mice

It has been proposed that amniotic fluid fetal macrophages play a key role in mouse parturition (19). Therefore, we determined whether macrophages were readily detectable in the amniotic fluid of humans as a substantial number of amniotic fluid macrophages were expected if their myometrial infiltration was a key phenomenon associated with labor. Cytologic preparations of human amniotic fluid samples (*n*=19) obtained at various clinical settings for different gestational periods, were analyzed by immunostaining for CD68. Macrophages were detected very rarely in seven amniotic fluid samples tested (Fig. 8A), and they were not found in the rest of the samples. The demographics of the patients are summarized in Table III. Flow cytometric analyses of amniotic fluid cells obtained from five additional patients were further performed to assess proportion of CD14 positive macrophages. CD14 positive cells were not detected in three cases (gestational age; 36, 38, and 39 weeks not in labor). In the remaining two cases (gestational age; 19 and 37 weeks), the proportions of CD14 positive cells were 0.007 % (isotype: 0.003 %) and 0.004 % (isotype: 0.003 %), respectively (Fig. 8*B*).

The density of myometrial macrophages in C57BL/6 mice was also analyzed to see if it increased with advancing gestational age in contrast to humans. Microscopic examination of macrophages in the mouse myometrium was done among the pregnant mice across gestational periods (E13, E15, E17, and E19). F4/80 positive macrophages were found mainly in the endometrium and sparsely in the myometrium (Fig. 8*C*). However, the density of macrophages in the myometrium did not change significantly across gestational periods (Fig. 8*D*). F4/80 positive macrophages were not found in the amnion or visceral yolk sac comprising mouse fetal membranes.

Discussion

The novel findings of this study are that: 1) there is a redistribution of HMW SP-A oligomers between the "placental amnion" and the "reflected amnion" with labor at term in humans. This is largely due to the increased binding of amniotic fluid SP-A to the reflected amnion; 2) amniotic binding of supratrimeric SP-A oligomers is a unique feature of human pregnancy as it is not observed in C57BL/6 mice; 3) macrophages are very rare in human amniotic fluid; 4) there is no significant macrophage influx into the myometrium in association with labor at term in C57BL/6 mice; and 5) SP-A treated human amnion demonstrates a molecular signature consistent with anti-inflammatory properties (inhibition of cytokine-cytokine receptor interaction). Therefore, amniotic fluid SP-A per se has modest anti-inflammatory effects on the amnion.

Our data clearly show that HMW SP-A oligomers (rather than monomers, dimers, or MMW oligomers) primarily increase in amniotic fluid with advancing gestational age. This finding indicates that the effects conferred by amniotic fluid SP-A as gestational age advances would be influenced more by HMW oligomers. Supratrimeric SP-A oligomerization is related to the stability of collagen triple helix and ligand aggregation (37). Oligomeric SP-A distribution in the human amnion was found to be more abundant in the mesodermal layer containing fetal tissue macrophages and myofibroblasts, and it is intriguing that the receptors we screened were more abundant in the epithelial layer (36,38). The amniotic SP-A distribution pattern in humans is in stark contrast with that observed in C57BL/6 mice. The difference in the amniotic distribution of oligomeric SPA between humans and mice is likely due to the differences in the anatomical constitution of their respective fetal membranes. Mouse fetal membranes are primarily composed of amnion and yolk sac, and the very thin amnion proper lacks sizable amounts of stroma and macrophages as well (39), although it is possible that anti-SP-A oligomers in the mouse amnion. We also wish to

bring to attention the fact that SP-A genes are duplicated in humans producing two isoforms, which are able to form higher order SP-A oligomers (40). This may not be the case in mice, which have a single SP-A gene (9). The polymorphism of SP-A genes has been shown to be associated with susceptibility to certain diseases (41), and the relationship among SP-A genetic polymorphism, the degree of higher order SP-A oligomerization, and susceptibility to pregnancy complications would be worthy of further studies.

The next question under investigation was how SP-A-induced changes within the amnions observed in this study could be explained in the context of human labor. The reduced SP-A distribution in the placental amnion with labor is consistent with the over-expression of IL-1 β mRNA in the amnion. IL-1 β mRNA expression with labor was higher in the placental amnion rather than in the reflected amnion (2). Therefore, it may be argued that the decreased SP-A in the placental amnion might be a part of the mechanism involved in upregulation of IL-1ß mRNA. However, increased amniotic fluid SP-A by itself does not seem to be a prerequisite for the initiation of human parturition for two reasons: 1) We found clear examples of women not in labor with barely detectable HMW SP-A oligomers in the placental amnion, and also, we observed cases of women in labor with barely detectable HMW SP-A oligomers in the reflected amnion; and 2) SP-A knock-out mice undergo spontaneous labor at term. Since SP-A is a lectin, it is possible that there is functional redundancy, and that SP-A is not essential for the onset of labor (42). It is interesting that SP-A also down-regulates CXCL2 and CXCL5 mRNA expression. CXCL2 and CXCL5 are potent neutrophil chemokines, and the pattern of regulation by SP-A is not quite consistent with the enhanced neutrophil chemotaxis reported in other studies (43,44). It should be noted that TLR2 and TLR4 show intra-amniotic (epithelium vs. mesoderm) differences in their expressions. The reasons why TLR2 and TLR4 show reciprocal expression patterns between the amnion epithelium and mesoderm needs further investigation, but it strongly suggests that the functional requirements of the amnion epithelium facing the amniotic cavity and the mesoderm facing the chorion (endometrial cavity side) are different.

Although the importance of SP-A-induced amniotic fluid macrophage activation was proposed in mice, the rarity of macrophages in human amniotic fluid along the continuum of gestational age does not support a mechanistic relationship between the initiation of labor and the trafficking of amniotic fluid fetal macrophages. Previous clinical studies have also shown that amniotic fluid macrophages increase under a pathological condition, such as fetal open neural tube defects, and are rare or absent in normal pregnancies (45,46). We were not able to demonstrate an increase in myometrial macrophage density or a directional migratory pattern of macrophages in the myometrium or the fetal membranes of C57BL/6 mice. Therefore, although fetal or maternal macrophage infiltration has been described as one of the major changes in the myometrium associated with labor both in humans and other animal species (19,47,48), a more thorough evaluation and systematic analysis would be necessary to confirm if such macrophage migration precedes labor. There are preexisting tissue macrophages in the utero-placental compartment, and activation of these macrophages would be more effective and biologically relevant if a functional involvement of macrophages is necessary for the initiation of labor. It is noteworthy that Mackler et al (49) reported a decline in the number of C3/HeN mouse myometrial macrophages before birth compared to day 15 of pregnancy.

SP-A was shown to be a signal for a pro-inflammatory response in mouse parturition (19), whereas several studies have shown its anti-inflammatory, protective role in pulmonary inflammation. *SFTPA* knockout mice showed defects in pulmonary bacterial clearance and bacterial uptake by alveolar macrophages while pro-inflammatory cytokines, $TNF\alpha$ and IL-6, were increased upon intra-tracheal administration of group B Streptococcus and Haemophilus influenzae when compared to wild types of animals (50). The analysis of

alveolar macrophages of Sprague-Dawley rats, CD14-null 129J, or C3H/HeJ mice demonstrated CD14-independent inhibition of LPS-induced TNF α , IL-1 α , and IL-1 β mRNA expressions by SP-A (51). It is important to note that alveolar macrophages are a prototypic example of alternatively activated macrophages, and SP-A exerts anti-inflammatory responses of these macrophages characterized by decreased NADPH oxidase activity and reactive oxygen intermediates production (52). TLR2 and TLR4 signaling are also dampened in SP-A-treated macrophages with decreased phosphorylation of I κ B α (53).

In summary, sequestration of oligomeric SP-A to the reflected amnion seems to explain, in part, a decrease in amniotic fluid SP-A concentration during labor (Fig. 9). The amnion represents an extra-pulmonary source of amniotic fluid SP-A (15,16). The present study reveals a novel example of how a collectin molecule in the amniotic fluid can orchestrate its biological effect on the chorioamniotic membranes. Amniotic fluid and the amnion comprise an environment similar to the case of the lung (54) in the context of SP-A interaction among the amniotic fluid, placental amnion, and reflected amnion. We propose that amniotic SP-A protects the integrity of the amnion and amniotic cavity from pro-inflammatory stimuli during pregnancy, just as it protects the lung from alveolar inflammation. Yet, SP-A and fetal macrophages in the amniotic fluid per se do not seem to be molecular or cellular prerequisites for parturition in humans.

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Abbreviations used in this paper

AF	amniotic fluid
CRT	calreticulin
FDR	false discovery rate
Ε	epithelial layer
GP	goat polyclonal
HMW	high molecular weight
KEGG	Kyoto Encyclopedia of Genes and Genomes
Μ	mesodermal layer
MM	mouse monoclonal
MMW	medium molecular weight
PA	placental amnion
RA	reflected amnion
RP	rabbit polyclonal
qRT-PCR	quantitative RT-PCR
SPIA	signaling pathway impact analysis
SP-A	surfactant protein-A
TIL	term in labor
TNL	term not in labor

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Figure 1.

Immunoreactive patterns of SP-A in human amniotic fluid, amnion, and fetal lung samples to Abs generated from different species (rabbit polyclonal, RP; goat polyclonal, GP; mouse monoclonal, MM) and specificity of immunoreactive bands. *A*, Two microliters of amniotic fluid and 10 micrograms of total proteins from the amnion and the lung obtained from the same patient were electrophoresed in 4–15% gradient SDS-PAGE under non-reducing conditions, and subsequently electro-blotted. Under non-reducing conditions, the RP anti-SP-A Ab detects HMW (>250 kD) SP-A oligomers and dimers most effectively, whereas GP and MM anti-SP-A Abs effectively recognize MMW (75 kD–250 kD) SP-A oligomers. *B*, Under reducing conditions, non-reducible SP-A dimers (64 kD) are readily detected by both RP and GP Abs. Monomeric SP-A polypeptides are detected by all Abs, but, most effectively, by the MM Ab.



Figure 2.

Characterization of SP-A in the amniotic fluid. A, Amniotic fluid (AF) proteins in slices corresponding to HMW and MMW bands on a non-reducing SDS-PAGE gel were extracted and resolved again by SDS-PAGE under reducing conditions. AF obtained from a twin pregnancy at 29 weeks, and fetal lung tissue from an autopsy case with gestational age of 33 weeks were compared. Immunoblotting with mouse monoclonal anti-SP-A Ab recognizes monomeric SP-A polypeptides in both HMW and MMW fractions, while the 50–65 kD fraction demonstrates largely non-reducible dimers (64 kD). B, Immunoblotting using MM anti-SP-A Ab of HMW SP-A fraction in the same AF (29 weeks) immunoprecipitates with RP anti-SP-A Ab and purified SP-A demonstrates an identical pattern showing monomeric SP-A (32 kD) and non-reducible SP-A dimer (64 kD). C, Mass spectra of monomeric SP-A shown in Figure 2B (*) on LC-MS/MS analysis. LC-MS/MS analysis also confirmed SP-A in monomeric and dimeric SP-A (arrows) in purified SP-A (data not shown). D, E, F, and G, Two-dimensional electrophoresis and SP-A immunoblotting profiles of amniotic fluid samples and purified SP-A. HMW SP-A fraction of AF proteins from three cases (D, 29 weeks; E, 38 weeks; F, 37 weeks). G, Purified SP-A. Electrophoresis was done under reducing conditions followed by immunoblotting with MM anti-SP-A Ab demonstrates the same isomer patterns as that of purified SP-A. However, the upper bands representing nonreducible dimers were hardly detected in all of the AF samples compared with purified SP-A.



Figure 3.

Expression patterns of SP-A in the amniotic fluid and the amnions of humans and C57BL/6 mice. A and B, Immunoblotting analysis of amniotic fluid samples obtained across gestational periods shows a clear correlation between gestational age and the HMW oligomeric SP-A expression level detected by the RP anti-SP-A Ab. Two microliters of amniotic fluid samples were electrophoresed in 4-15% gradient SDS-PAGE under a nonreducing condition. C and D, SP-A immunoblotting of the reflected amnion samples shows more prominent HMW (>250 kD) oligomeric SP-A in the cases with labor (TNL: term not in labor, TIL: term in labor). The difference is less prominent for MMW (75 kD-250 kD) SP-A which was detected using the GP anti-SP-A Ab. Ten micrograms of amnion samples were electrophoresed in 4–15% gradient SDS-PAGE under a non-reducing condition. E, Densitometric comparison of HMW SP-A oligomers shows a significant increase of HMW SP-A oligomers in the reflected amnion of TIL cases compared to TNL cases. F, Immunoblotting of C57BL/6 mouse amniotic fluid using the RP anti-SP-A Ab also shows increasing oligomeric SP-A as gestation progresses. HMW SP-A oligomers are readily detectable in E17. Two microliters of amniotic fluid samples were electrophoresed in 4-15% gradient SDS-PAGE under a non-reducing condition. G. In contrast to the findings for the human amnion, mouse fetal membranes do not show oligomeric forms of SP-A. Ten

micrograms of fetal membrane samples were electrophoresed in 4–15% gradient SDS-PAGE under a non-reducing condition.



C CD91 SP-A DAPI



D TLR4 SP-A DAPI



Figure 4.

Binding of SP-A to SP-A receptors in the human amnion. *A*, Human amnion protein lysates (125µg) were immunoprecipitated using specific Abs to SP-A and known SP-A receptors, such as SIRP α , calreticulin (CRT), MD2, and TLR2, and subsequently probed with anti-SP-A Ab. For isotype control experiments to assess specificities of immunoprecipitative reaction, equal amount of proteins were pooled from the same TNL and TIL cases and used. The blot demonstrates co-immunoprecipitation of oligomeric SP-A to SIRP α , CRT, MD2, and TLR2. The species of the primary Ab host used for immunoprecipitation reaction are shown at the bottom of the gel. SP-A immunoblottings were done using a rabbit polyclonal antibody. *B*, The result of immunoprecipitation. The same pooled proteins (TNL + TIL) were used. *C* and *D*, Immunofluorescent images of the amnion show distinct co-localizations of SP-A with CD91 (*C*) or TLR4 (*D*) in the amnion epithelium. Immunofluorescent staining was performed on placental amnion whole mount samples obtained from three TNL cases.



Figure 5.

Binding of amniotic fluid SP-A oligomers to the human amnion. *A*, Primary amnion cells incubated with amniotic fluid show an increase in HMW SP-A oligomers in a time-dependent manner (left), while they are being depleted in the conditioned amniotic fluid (right). *B*, The addition of a neutralizing Ab in the amniotic fluid inhibits HMW SPA binding to the amnion explants. *C*, The MMW SP-A oligomers are not detected in the amnion cells after incubation with amniotic fluid (left), nor were they changed in the conditioned amniotic fluid (right). *D*, The SP-A oligomers in the placental amnion (PA) and the reflected amnion (RA) were decreased after treatment with collagenase.



Figure 6.

Distribution of SP-A in the human amnion. A, Immunofluorescent staining for SP-A and CD68, a macrophage marker, demonstrates SP-A immunoreactivity in both the amniotic epithelial and mesodermal layers. In the epithelial layer, the signals are detected in both apical portion and basal portion of the cells (original magnification X630). B, Serial immunoblotting for SP-A and its potential receptors in the amnion epithelial layer (E) and the mesodermal layer (M). SP-A is more abundant in the mesodermal layer (p < 0.05). Interestingly, the majority of the potential SP-A receptors was more abundant in the amnion epithelial layer than in the mesodermal layer (p < 0.05), with the exception of TLR2 and MD2, whose expression is clearly higher in the mesodermal layer (p < 0.05). C, Immunoblotting shows more abundant SP-A in the placental amnion (PA) before labor, and SP-A distribution between the placental amnion and the reflected amnion (RA) becomes similar in labor due to an increase in the reflected amnion and a decrease in the placental amnion. The difference in the distribution of HMW SP-A between TNL and TIL cases was significant on densitometric analysis of HMW SP-A oligometric in PA and RA (p < 0.01). The median RA/PA SP-A density ratio was 0.27 (range: 0.13-0.77) in TNL cases, while it was 1.86 (range: 0.4–3.27) in TIL cases. Among the potential receptors screened, the expression of CD91 (p = 0.05) and SIRP α (p = 0.07) tended to be higher in the PA than in the RA. TLR2 (p = 0.14) and TLR4 (p = 0.14) expressions were not significantly different between PA and RA.



Figure 7.

Effects of SP-A treatment in the human amnion explants. *A*, Human reflected amnion explants obtained from TNL patients were treated with SP-A (1 µg/ml) for 12 h. A volcano plot showing the relationship between their fold changes (log₂ of, *x* axis) and FDR-corrected *p* values ($-\log_{10}$ of, *y* axis). Positive values on the *x*-axis indicate genes whose expression increased after SP-A treatment, while negative values indicate genes down-regulated after SP-A treatment. *B*, Confirmative qRT-PCR data showing changes in the expression of select cytokines and chemokines. IL-1 β , CXCL2, and CXCL5 mRNA expressions decreased significantly after treatment with SP-A (*p* < 0.05 for all comparisons). CCL3 mRNA expression was not significantly different. * p<0.05 by Wilcoxon signed rank test



Figure 8.

Macrophages in the human amniotic fluid (A and B) and in the myometrium of C57BL/6 mice (C and D). A, Amniotic fluid cytologic specimens show CD68 immuno-positive macrophages among scattered epithelial cells. Case 4: a case of a patient presenting with intrauterine fetal demise at the gestational age of 21.4 weeks, Case 15: a case of a TNL patient at the gestational age of 38.7 weeks (original magnification X200; hematoxylin was used for counterstaining). B, A histogram from flow cytometric analysis of amniotic fluid cells does not show CD14 positive macrophages. The amniotic fluid was obtained from a TNL patient at the gestational age of 38 weeks. Green: CD14, Red: isotype. C, F4/80 immuno-positive macrophages (arrows) in the C57BL/6 myometrium obtained at various gestational ages: E13, E15, E17, and E19 original magnification X200; hematoxylin was used for counterstaining). D, The density of macrophages in the mouse myometrium did not change significantly across gestation. The numbers of F4/80 positive macrophages were counted in myometrial sections containing the mid-plane of the placenta from four pregnant mice at each gestational age (E13, E15, E17, and E19). Eight high-power fields corresponding to every 45° angle point starting from the mesometrial border in a clockwise direction, covering the entire myometrial circumference, were evaluated in each section. Image-Pro Plus 6.2 software (Media Cybernetics, Inc.) was used for the analysis.



Figure 9.

A schematic representation of SP-A dynamics among the amniotic fluid, placental amnion (amnion overlying the placental disc), and the reflected amnion (amnion of the extraplacental chorioamniotic membranes). There is a redistribution of SP-A in the amniotic fluid and the amnion with labor. HMW SP-A oligomers are more abundant in the placental amnion compared to the reflected amnion before labor. With labor at term, increased binding of HMW SP-A oligomers to the reflected amnion results in a decrease of amniotic fluid SP-A. Decreased amniotic fluid SP-A concentration during labor (23) is consistent with this model.

Table I

Peptides from Surfactant Protein-A

Average m/z	Number of peptides	Modification(s)	aa residues	Sequence
2329.390	1	Oxidation (M)	95–115	GPPGLPAHLDEELQATLHDFR
1490.220	2		123–137	GALSLQGSIMTVGEK
1506.575	22	Oxidation (M)	123–137	GALSLQGSIMTVGEK
2201.185	4	Carbamidomethyl (C)	138–157	VFSSNGQSITFDAIQEACAR
1447.344	95		167–179	NPEENEAIASFVK
2234.765	2		180–199	KYNTYAYVGLTEGPSPGDFR
2106.885	4		181–199	YNTYAYVGLTEGPSPGDFR
2132.020	1	Carbamidomethyl (C), Oxidation (M)	220-236	GKEQCVEMYTDGQWNDR
781.364	7	Carbamidomethyl (C)	243-248	LTICEF

Table II

List of differentially expressed genes in amnion explant culture after SP-A treatment

Gene	Fold change	Adjusted <i>p</i> value
CYR61	-1.97	0.001
LOC648517	-1.89	0.006
TCEA3*	1.65	0.019
KLF2	-1.59	0.022
FOSB	-1.88	0.031
SLC22A3	-1.73	0.031
EGR1	-1.71	0.031
FZD8	-1.69	0.031
AMOT*	1.59	0.031
HBEGF	-1.52	0.031
TWIST1	-1.50	0.031
CTGF	-2.12	0.031
SERPINE1	-1.69	0.033
FRMD6	-1.60	0.041
ADRB2	-1.50	0.048
GUCY1A3*	1.51	0.049
ANPEP	-1.56	0.049
GSTO2*	1.50	0.049
MYC	-1.51	0.049
PDK4	-1.54	0.051
LRIG1	-1.61	0.057
MAP1B*	1.58	0.060
GFPT2	-1.61	0.062
LOX	-1.55	0.062
AKR1C2	-1.65	0.065
KCNMB4*	1.51	0.068
ANKRD35*	1.53	0.068
ERN1	-1.58	0.075
EVI2B	-1.51	0.075
TH^{*}	-1.57	0.075
CRLF1	-1.59	0.078
SFRP2*	1.98	0.083
CYP1B1	-1.50	0.085
NAMPT	-1.53	0.095
HSPA6	-1.94	0.096
ST3GAL1	-1.52	0.097
GPR68	-1.57	0.097
GABRP*	2.08	0.097

Gene	Fold change	Adjusted p value
FAM183A*	1.54	0.105
ADFP	-1.72	0.113
RASD1	-1.58	0.119
ECM1	-1.50	0.130
AGR3*	1.57	0.140
SILV*	1.52	0.142
HBA1	-1.81	0.144
CXCL2	-2.12	0.147
CDH10*	1.57	0.151
ANGPTL4	-1.61	0.152
CLEC1B*	1.79	0.163
HBG1	-1.79	0.165
PHLDA1	-1.63	0.180
LGALS7B*	1.51	0.182
ESM1	-1.52	0.199
HBG2	-1.61	0.213
IL1B	-2.26	0.218
IL1RL1	-1.55	0.222
CXCL5	-1.72	0.240
FOS	-1.58	0.243
CCL3	-1.66	0.246
COL1A1	-1.58	0.249

* Genes up-regulated after SP-A treatment

number	Gestational age at amniocentesis (weeks)	Diagnosis at amniocentesis	CD68 (+) macrophages	Gestational age at delivery (weeks)	Placental pathology
1	18.4	Short cervix	+	19.4	Acute chorioamnionitis
2	19.6	Genetic amniocentesis	I	41.9	
3	21.2	Short cervix	Ι	26.6	Acute chorioamnionitis
4	21.4	Intrauterine fetal death	+	38.1	
5	23.0	Genetic amniocentesis	+	38.3	
9	24.4	Short cervix	+	35.1	Acute chorioamnionitis
7	24.9	Genetic amniocentesis	I	25.0	
8	28.3	Twins - Advanced cervical dilatation	Ι	33.3	
6	28.3	Genetic amniocentesis	I	40.1	
0	32.0	Preterm contractions	I	39.7	
1	35.1	Twins - Preterm contractions - Mild preeclampsia	I	35.1	
2	35.1	Twins - Preterm contractions - Mild preeclampsia	I	35.1	
3	35.9	Mild preeclampsia	I	35.9	
4	38.0	Term in labor	I	38.0	
5	38.7	Term not in labor	+	38.7	
9	39.1	Term not in labor	I	39.9	
7	39.1	Term not in labor	Ι	39.1	
8	39.1	Term not in labor	+	39.1	
6	41.9	Term not in labor	+	41.9	Acute chorioamnionitis

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Table III

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Clinicopathologic information of the cases included in the cytological preparation of their amniotic fluid