A pilot study of macrophage responses to silk fibroin particles

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Abstract: Silk fibroin (SF) shows promise for tissue engineering and other biomedical applications due to its excellent biocompatibility, unique biomechanical properties, and controllable biodegradability. The particulate form of SF materials may have many potential uses, including the use as a filler for tissue defects or as a controlled-release agent for drug delivery. However, many past in vivo and in vitro studies evaluating the biocompatibility and biodegradability of SF have involved bulk implants. It is essential to evaluate the inflammatory effects of SF particles before further use. In this study, two different sizes of SF particles were evaluated to assess their impact on the release of tumor necrosis factor (TNF)-α, interleukin (IL)-1β, and IL-6, in comparison with lipopolysaccharide positive control stimulation. The inflammatory processes were characterized using real-time reverse transcription polymerase chain reaction, enzyme-linked immunosorbent assay, and light microscopy evaluations. The results indicated that small silk fibroin particles and large silk fibroin particles, in culture with RAW 264.7 murine macrophage cells for 24 h, caused up-regulation of mRNA coding for TNF-α, which indicated that both size of particles have potential inflammatory effects. There was a statistically significant increase in this up-regulation under small silk fibroin stimulation. However, the immunosorbent assay suggested that there was virtually no observed release of IL-1β, IL-6, or TNF-α, relative to the control group. The results suggest that SF particles of the chosen dimensions may have good biocompatibility in culture with RAW 264.7 murine macrophages. © 2012 Wiley Periodicals, Inc.

Key Words: silk, fibroin, macrophage, biocompatibility, in vitro, inflammation, cytokine


INTRODUCTION
Silk fibroin (SF) shows promise for tissue engineering and other biomedical applications1–3 due to its excellent biocompatibility, unique biomechanical properties, and controllable biodegradability. Such properties have attracted interest of investigators from biomedical fields and other related disciplines. SF is a versatile biomaterial that can be prepared in various forms and shapes, ranging from nanometer to centimeter scales. This versatility allows it to be utilized for a variety of purposes. Although the majority of SF application focuses on tissue engineering involving bone, cartilage, nerve, and blood vessel repair,4–8 there is increased interest regarding the role of particulate SF in drug delivery mechanisms.9–12 The particulate form of SF materials, as injectable fillers, may also have potential use for synthetic repair of tissue defects such as glottic insufficiency caused by unilateral vocal fold paralysis.

However, most of the in vivo and in vitro studies observing biocompatibility and biodegradability of SF have dealt primarily with bulk implants. It is likely that particulate implantation of the same materials may elicit significantly different inflammatory response, compared with bulk implantation.13,14 Observation of implanted materials has suggested that the size, concentration, surface area, and volume dimensions of implanted particulate material play an important role in the tissue-cellular response.15–18 Therefore, it is important that the inflammatory response elicited by particular SF biomaterials could be investigated prior to application on a broader scale.

Monocytes/macrophages serve as an effective study system, due to their active roles in the processes of inflammation and wound healing.19 A variety of investigations have focused on the interactions between biomaterial particles and macrophages, and histological examinations have revealed that macrophages play a central role in the response to the particles.20–22 For this reason, many in vitro studies have aimed at simulating an in vivo response through the examination of macrophages. More specifically,
macrophage cells have been employed extensively for in vitro studies to evaluate the inflammatory responses triggered by various biomaterials. Many cytokines and interleukins, including tumor necrosis factor (TNF)-α, interleukin (IL)-6, and IL-1β, have been widely found in pseudo-membranes of failed total hip arthroplasty cases, as well as in other in vitro studies. These three cytokines, which are proinflammatory cytokines that recruit other inflammatory and immune cells, have been implicated as playing substantial roles in the inflammatory response.

In this investigation, we provide initial pilot results for an in vitro study of biocompatibility in SF particles. Two different sizes of SF particles were selected to investigate their role in stimulating the release of TNF-α, IL-1β, and IL-6, in comparison with lipopolysaccharide (LPS) positive control stimulation. The inflammatory processes were characterized using real-time reverse transcription polymerase chain reaction (RT-PCR), enzyme-linked immunosorbent assay (ELISA), and light microscopy evaluations.

**MATERIALS AND METHODS**

**Preparation of SF particles**

SF particles were prepared according to previously established methods. *Bombyx mori* silk cocoons were degummed twice in 0.5% (w/w) aqueous NaHCO₃ solution at 95°C for 30 min. After being washed thoroughly with distilled water and then dried, the degummed silk was dissolved in 9.5M LiBr aqueous solution for about 1 h at 45°C with gentle stirring. After being filtered, the resulting fibroin solution was dialyzed against deionized water for 4 days at room temperature with a 14 kDa cutoff dialysis membrane to remove the salt. The dialyzed SF solution was centrifuged against deionized water for 4 days at 6000 rpm for 10 min. Then, the clarified solution in the supernatant was collected and concentrated by reverse dialysis to a final concentration of 10–45 µm (small silk fibroin particles; SFS) and 45–125 µm (large silk fibroin particles; SFL). The size and morphology of the particles were confirmed by scanning electron microscopy (SUPERSCAN SSX-550; Shimadzu, Japan) (Fig. 1).

All the SF particles were treated with 70% ethanol solution for at least 48 h to sterilize and remove bound endotoxins.33 Particles were rinsed thoroughly with distilled water, then suspended and sonicated in sterile phosphate buffered saline (PBS) before use. All procedures were carried out with extreme care and performed under clean and sterile conditions (instruments, containers, etc.) to avoid any contamination.

**Macrophage culture**

A RAW 264.7 murine macrophage cell line obtained from the Cell Bank of the Chinese Academy of Sciences (Shanghai, China) was used in this study. We selected this specific cell line because of their reproducibility and the ease of culturing that they permit. This is also one of the cell lines frequently used by other authors to evaluate particle responses.14,26 The cells were expanded and maintained in 75 cm² cell culture flasks (Corning Glass Works, Corning, NY). They were suspended in Dulbecco’s Modified Eagle Medium (DMEM, Sigma), containing 10% fetal bovine serum (FBS, Gibco), and...
cultured in an incubator with a moist atmosphere consisting of 5% CO₂ to 95% air at 37°C. Media contained 100 U/mL of penicillin and 100 μg/mL of streptomycin.

After formation of the cell monolayer (80% cell confluence), the culture medium was carefully removed and the flasks were washed twice with PBS solution. An additional 15 mL of PBS was added to the flask, and the cells that remained on the flask surface were harvested with a cell scraper (Corning Glass Works).

The medium containing the cells was then pipetted and transferred to 20-mL sterile tubes, which were centrifuged at 1000 rpm for 10 min to separate the cells from the medium. After centrifugation, the supernatant was discarded and 10 mL of culture medium was added to each tube containing the cells. The total number of cells was counted in a hemocytometer, yielding a subcultivation ratio of 1:8. The surplus cells were preserved in complete growth medium supplemented with 5% (v/v) dimethylsulfoxide (DMSO, Sigma) in liquid nitrogen.

**Macrophage stimulation and assay**

RAW 264.7 macrophages were seeded in 2.5 mL cell cultures in different wells of a 6-well plate, delivering approximately 2 × 10⁵ cells per well. The macrophage cultures were plated for 24 h, washed with PBS, then subject to new culture media prior to stimulation.

Once prepared for experimental use, the cells were challenged with SFL particles (5 mg/well, n = 6) and SFS particles (5 mg/well, n = 6), respectively. For experimental purposes, the SFL replicates correspond to group A, whereas the SFS particles comprise group B. We chose an SF dosage of 5 mg/well, because at this dosage, the cells were thinly covered by the particles. Our results depict data gathered after 24 h incubations only, since previous studies have shown that the maximum cytokine production—by cultured macrophages in response to particles—occurs after 15–20 h of incubation. And our preliminary experiment found that longer observation times without replacement of the culture media may result in over proliferation and death of the cells.

For our positive control (n = 6; group C), LPS (Sigma, L6529) was added to the cells at a terminal concentration of 1 μg/mL. The negative control (n = 6; group D) consisted of cells cultured in standard medium alone. All stimulations, with the exception of LPS positive controls, were carried out in the presence of 5 μg/mL Polymyxin B to inhibit any LPS contamination.

Cells were examined using light microscopy in order to observe their morphology, which was assessed in terms of cell attachment to the plate. After exposure to the stimulation for 24 h, the supernatant of the cell culture media was collected and centrifuged to remove particulate matter. The solution was aliquoted and stored at -20°C for later evaluation of cytokine levels. The cells in each well were washed with PBS to remove the particles and the growth media, followed by the addition of 1 mL of TRIzol® Reagent (Invitrogen, Carlsbad, CA) in order to lyse the cells. We pipetted the cells up and down several times to homogenize the samples, and then stored them at -80°C to prepare them for mRNA extraction.

**Real-time RT-PCR assays of the stimulated macrophages**

RNA was extracted from RAW 264.7 cells using TRIzol® Reagent, following the manufacturer’s protocol. Quantity and purity of RNA were determined by absorbance values indicated on a spectrophotometer (BioMate 3S, Thermo Scientific) at 260 and 280 nm. For reverse transcription, 1.6 μg of RNA was reversely transcribed with Oligo (dT)₁₅ (Fermentas), and complementary DNA was amplified according to the manufacturer’s instructions. Primers were designed and synthesized by Sangon Biotech Shanghai Co. Ltd, Shanghai, China. Real-time RT-PCR was performed to detect gene expression of IL-1β, IL-6, and TNF-α, using the ABI Prism 7500 (Applied Biosystems, Foster City, CA) real-time system in triplicate for all samples.

The forward and reverse primers for IL-1β are 5'-TGAAATGCGACCCTTTTGACAG-3' and 5'-CCACAGCCACAATGAGTGATAC-3', respectively; for IL-6, forward primer 5'-ACAAAGGCAGTCTCTTCAGAG-3' and reverse primer 5'-AAGATGAATTTGGATGGTCTTGG-3'; and for TNF-α, forward primer 5'-GGCCTATGCTCACGCTCTTCTC-3' and reverse primer 5'-CAGTGGTGTTGTTGCTAGA-3'. The housekeeping gene, β-actin, was selected as a reference gene in order to control the intersample variation in RNA isolation and integrity. This was done by using a pair of primers: forward primer 5'-GAGACCTCTCAACACCCAGC-3' and reverse primer 5'-ATGTCAGGCCAGATTTCC-3'.

The amplification conditions were 95°C for 2 min followed by 40 cycles of 95°C for 20 s, 57.5°C for 30 s, and 72°C for 32 s. For each assay, 0.25 μM (terminal concentration) of both forward and reverse primers and approximately 32 ng of complementary DNA template were added to each reaction of the GoTaq qPCR Master Mix (Promega) for a total reaction volume of 20 μL.

At the end of each reaction, a cycle threshold (CT) was manually setup at the level that reflected the best kinetic PCR parameters, followed by observation of melting curves and subsequent analysis. Ct is defined as the PCR cycle number during which an increase in fluorescence beyond a threshold occurs. Amplification products were quantified by comparison of experimental Ct levels. The expression data for each gene product were normalized against the Ct level of the β-actin housekeeping gene. The resulting transcript Ct levels are reported as mean relative changes (± range) compared to untreated controls. The ΔΔCt method of relative quantification was adapted and optimized to estimate the quality of the four genes previously indicated.

**Determination of macrophage cytokine release**

Concentrations of IL-1β, IL-6, and TNF-α in the supernatants of the cell culture were quantified using ELISA kits (BioTNT, Shanghai, China) according to the manufacturer’s protocol. These were ELISA kits that are murine specific. Cytokine concentration was calculated by comparison with standard curves of known values of recombinant murine IL-1β, IL-6, and TNF-α. Measurements of optical densities were performed with ELISA reader (DNM-9602; Beijing Prolong New Technology Co., Ltd., Beijing, China) at 450 nm wavelength. All samples and standards were processed in duplicate wells and the results were expressed in picograms per
milliliter. The cytokines released in the blank control were subtracted from all other samples as a baseline.

**Statistical analysis**
Statistical analysis was performed using a commercially available statistical software prism 5.0 (GraphPad Software Inc., San Diego, CA). All data were expressed as mean ± standard deviation (SD) for \( n = 6 \). Statistics were produced using one-way analysis of variance (ANOVA) tests as well as Tukey’s HSD post-hoc comparison test. The value of \( p < 0.05 \) was considered statistically significant.

**RESULTS**

**Light microscopy observations of RAW 264.7 cell activation**
After stimulation for 24 h, the challenged macrophages were viewed under an inverted phase contrast microscope (Nikon Ti-E, Japan), showing activation of the cells. All three stimulated sample groups (SFL, SFS, and LPS) exhibited morphological changes, which included enlarged cell sizes, increased vacuolization, and the presence of pseudopod-like structures. The morphological changes of the RAW 264.7 cell line in groups A and B were almost the same, having slight enlargement and polymorphism of cell bodies, with a few cells having bubble-like constitution in the plasma [Fig. 2(A) and (B)]. In group C, however, the similar morphological changes were dramatically characterized by alveolar macrophage changes [Fig. 2(C)]. In contrast, group D showed almost no changes of cell morphology except proliferation of the cells [Fig. 2(D)].

**Induction of mRNAs of IL-1β, IL-6, and TNF-α of the stimulated macrophages**
The mRNA expression of proinflammatory cytokines IL-1β, IL-6, and TNF-α were evaluated by real-time RT-PCR (Fig. 3). LPS stimulation caused a marked increase in the expression of mRNAs indicative of IL-1β, IL-6, and TNF-α (\( p < 0.001 \) vs. control). Despite this result, IL-1β and IL-6 mRNAs for the SFL and SFS stimulated groups were observed to be relatively similar to the control group (\( p > 0.05 \), respectively). However, macrophages stimulated by SFL and SFS both showed increased expression of TNF-α-suggestive mRNA when compared with the control cells (\( p < 0.001 \), respectively), but much less than the positive control (\( p < 0.001 \)). Furthermore, we observed higher levels of such mRNA in the SFS-stimulated group, compared with the SFL group (\( p < 0.001 \)). These results indicated that the SFS particles may have higher inflammatory-inducing nature.
Concentrations of IL-1β, IL-6, and TNF-α in the supernatants of the cell culture after stimulations

We examined the pro-inflammatory IL-1β, IL-6, and TNF-α cytokines using ELISA. The stimulation of the macrophages with LPS for 24 h increased levels of all the three cytokines noticeably, compared with the other three groups ($p < 0.001$, respectively) [Fig. 4(A)–(C)]. In contrast, SFL and SFS administration at a concentration of 5 mg/well created no obvious difference in cytokine concentrations, compared with the negative control cells ($p > 0.05$, respectively). Additionally, there was no significant difference between the particle populations with respect to expression and secretion of the cytokines being observed ($p > 0.05$). These results suggest that neither SFL nor SFS will provoke inflammatory responses from RAW 264.7 macrophages.

**DISCUSSION**

The purpose of this investigation was to evaluate the immunostimulating properties of two different sizes of SF particles. Inflammatory response was gauged by morphological changes; production of IL-1β, IL-6, and TNF-α cytokines;

![Figure 3](image-url)

**FIGURE 3.** IL-1β, IL-6, and TNF-α expression of the stimulated macrophages. Each value in real-time RT-PCR graphs (A, B and C) represents the mean ± SD of 6 wells of cells in each group. # $p < 0.001$ vs. control group; *** $p < 0.001$ vs. SFS group.

![Figure 4](image-url)

**FIGURE 4.** ELISA analysis of IL-1β, IL-6 and TNF-α secretion. Each value represents the mean ± SD of 6 wells containing supernatants of the cell culture in each group. # $p < 0.001$ vs. control group.
and transcription of the matching mRNA by RAW264.7 macrophages, following exposure to these biomaterials.

We chose to assay for IL-1β, IL-6, and TNF-α, which are proinflammatory cytokines that recruit other inflammatory and immune cells; activating these compounds upon arrival at implant sites. These three cytokines are thus suggested to play a valuable role in the inflammatory response process, making them ideal indicators of such activity. Obviously, additional or other cytokines could have been measured, but we limited our selection to these three for our pilot study. These are also the three cytokines selected for evaluation by many other investigators as described in the Introduction section.

Noticeable presences of pseudopod-like structures and bubble-like constitution in the plasma were observed in macrophages responding to both SFL and SFS 24-h stimulation. However, there was a more pronounced response to LPS stimulation. Treatment with LPS caused morphological changes characterized by larger nuclei, prominent nucleoli and relatively prominent cytoplasm with increased granularity, and enhanced secretion of TNF-α and other proinflammatory cytokines by macrophages as previously reported. In this study, the functional consequence accompanied with these morphological changes in SFL and SFS groups is up-regulated transcription of TNF-α mRNA, which encode TNF-α, a pleiotropic molecule that plays a central role in inflammation. We observed higher expression of TNF-α mRNA with SFS stimulation compared with that in the SFL dose. With respect to translation of mRNA, although these were statistically insignificant, similar differences in TNF-α secretion also existed. The results indicated that both SFL and SFS have potential inflammatory activity and furthermore demonstrated that SFS may cause more inflammation.

There were no statistically significant differences observed between the SFS and SFL samples for the presence of either the IL-1β or IL-6 mRNAs, compared with the negative control (Fig. 3). Similarly, there were no significant differences in measured secretion levels for either of these corresponding cytokines (IL-1β or IL-6), when observing the SFS and SFL samples relative to the control (Fig. 4). These results suggest that TNF-α may be the most abundant inflammatory mediator resulting from stimulation by SF particles. A study by Kaplan and co-workers also observed that insoluble, irregularly shaped fibroin particles of approximately 10–200 μm size induced significant TNF-α production under an experimental setup similar to our SF particle. SF particles have certain anti-tumor activity effect14 under an experimental setup similar to our SF particle. The data presented in this study demonstrate that SFS particles induced more significant macrophage activations characterized by TNF-α mRNA transcription than those observed during SFL stimulation. These results are likely due to the larger surface area–volume ratio inherent in SFS based on equal particle mass. This is in agreement with the previously published work which demonstrated that the total surface area or volume of the particles appeared to be an important factor in determining the inflammatory response during particle-induced cell activation.15,24

The inconsistency between transcription level of TNF-α mRNA and translation level of TNF-α may contribute to different efficiency of gene expression in different stimulated groups, inappropriate choice of testing time points, and the small sample size of the ELISA test. These limitations will remain unconfirmed until further experimentation can be performed under a better protocol design.

CONCLUSION

SFS and SFL in culture with RAW 264.7 murine macrophage cells for 24 h can cause obvious up-regulation of TNF-α mRNA—observed to a higher degree in the SFS sample group—which indicated that both sizes of SF material could provoke inflammatory activity mediated by TNF-α. Despite the presence of precursory TNF-α mRNA, there was no statistically significant difference in the release of IL-1β, IL-6, or TNF-α compared with the control group. These results indicated that SF particles of the tested dimensions may have limited inflammatory effect, thus suggesting good biocompatibility. Based on these findings, there seems to be a promising future for SF particle application in a variety of medical fields. However, further evaluation should be performed prior to any attempt at in vivo application of these particles.

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