Bio-functionalized titanium surfaces with the modified silk fibroin carrying titanium binding motif enhanced the ossific differentiation of MC3T3-E1

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Abstract

Silk fibroin (SF) from *Bombyx mori* has superior properties as both a textile and a biomaterial, and has been used to functionalize the surfaces of various medical inorganic materials including titanium (Ti). In this paper, we endowed SF with reversible binding ability to Ti by embedding a titanium binding motif (minTBP-1, RKLPDA). Artificial SF proteins were first created by conjugating gene cassettes for SF motif (AGSGAG) and minTBP-1 motif with different ratios, which have been shown to bind reversibly to Ti surfaces in quartz crystal microbalance analyses. Based on these results, the functionalized SF (TiBP-SF) containing the designed peptide [TS[(AGSGAG)$_3$AS]$_2$RKLPDAS]$_8$ was prepared from the cocoon of transgenic *B. mori*, which accelerates the ossific differentiation of MC3T3-E1 cells when coated on titanium substrates. Thus, TiBP-SF presents an alternative for endowing the surfaces of titanium materials with osseointegration functionality, which would allow the exploration of potential applications in the medical field.

Bio-functionalized titanium surfaces with modified silk fibroin carrying titanium binding motif to enhance the ossific differentiation of MC3T3-E1.

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ABSTRACT. min-TBP-1 modified silk fibroin

Silk fibroin (SF) from *Bombyx mori* has superior properties as both a textile and a biomaterial, and has been used to functionalize the surfaces of various medical inorganic materials including titanium (Ti). In this paper, we endowed SF with reversible binding ability to Ti by embedding a titanium binding motif (minTBP-1, RKLPDA). Artificial SF proteins were first created by conjugating gene cassettes for SF motif (AGSGAG) and minTBP-1 motif with different ratios, which have been shown to bind reversibly to Ti surfaces in quartz crystal microbalance analyses. Based on these results, the functionalized SF (TiBP-SF) containing the designed peptide $[TS[(AGSGAG)_3AS]_2RKLPDAS]_8$ was prepared from the cocoon of transgenic *B. mori* , which accelerates the ossific differentiation of MC3T3-E1 cells when coated on titanium substrates. Thus, TiBP-SF presents an alternative for endowing the surfaces of titanium materials with osseointegration functionality, which would allow the exploration of potential applications in the medical field.

KEYWORDS. Peptide aptamer, transgenic silk fibroin, artificial protein, biomaterial, 13C solid-state NMR

Running title. minTBP-1 functionalized silk fibroin.

INTRODUCTION

"Motif" is the versatile functional unit for constructing artificial biomacromolecules. In some cases, motif can endow non-biomolecules with its assigned functions, while in other cases different motifs can be combined to create artificial proteins having novel combinations of functions (Shiba, 2010). Motif sequences are identified both from natural sequences of proteins and from *in vitro* evolution systems. In the latter cases, a phage display system (Scott & Smith, 1990)) has been employed to create an artificial peptide that binds to biomolecules and non-biological materials such as inorganic materials. The hexapeptidic RKLPDA sequence is one such inorganic material binding peptides. It was originally isolated as a 12-mer peptide binder to a titanium (Ti) surface by using a peptide display system (Sano & Shiba, 2003). Subsequent analyses have revealed that the N-terminal RKLPDA (termed minTBP-1) is both necessary and sufficient for the binding (Sano, Sasaki, & Shiba, 2005). This minTBP-1 motif has been widely employed as an appendix to provide an affinity to Ti in ferritin (Sano, Ajima, et al., 2005), artificial matrix proteins (Kokubun, Kashiwagi, Yoshinari, Inoue, & Shiba, 2008), BMP-2 (Kashiwagi, Tsuji, & Shiba, 2009), antimicrobial peptide (Geng et al., 2018; Yoshinari, Kato, Matsuzaka, Hayakawa, & Shiba, 2010), calcification protein (Tsuji, Oaki, Yoshinari, Kato, & Shiba, 2010), DNA-polymerase (Nishida et al., 2015), carbon nanomaterial-binding peptide (Kokubun, Matsumura, Yudasaka, Iijima, & Shiba, 2018), and self-assembled chemical compounds (Ikemi et al., 2010). Interestingly, it has been revealed that the interaction between minTBP-1 and the surfaces of Ti is governed by reversible interactions (Brandt & Lyubartsev, 2015; Hayashi et al., 2009; Sano, Ajima, et al., 2005; Schneider & Ciacchi, 2012; Skelton, Liang, & Walsh, 2009). In biological systems, weak, specific, reversible binding plays a pivotal role in various biological activities. Indeed, previous studies have shown certain biological activities has been enhanced by adding minTBP-1 to parental biomolecules(Kashiwagi et al., 2009; Kokubun et al., 2008).

Silk fibroin (SF) from *Bombyx mori* possesses many superior and inherent properties as a biomaterial, including advantageous mechanical properties, environmental stability, biocompatibility, low immunogenicity, and biodegradability (Aigner, DeSimone, & Scheibel, 2018; Asakura & Kaplan, 1994; Asakura, Tanaka, & Tanaka, 2019; Koeppel & Holland, 2017; Koh et al., 2015; Velema & Kaplan, 2006; Vepari & Kaplan, 2007). SF fiber has been used in sutures in the surgical field for more than 2,000 years (Thurber, Omenetto, & Kaplan, 2015). Recently, the representative repeating unit of the crystalline fraction (56% of whole SF) (Strydom, Haylett, & Stead, 1977; Zhou et al., 2001; Zhou et al., 2000), AGSGAG sequence, has been used as a motif to create artificial SFs (Asakura et al., 2005; Asakura & Yao, 2002; Yu Suzuki, Aoki, Nakazawa, Knight, & Asakura, 2010; Yao, Ohgo, Sugino, Kishore, & Asakura, 2004). Motif-based artificial protein constructions provide us with opportunities to create artificial biomolecules having novel biological activities

or novel combinations of exiting activities (M. Yang & Asakura, 2005; Mingying Yang, Kawamura, Zhu, Yamauchi, & Asakura, 2009; M. Yang, Muto, Knight, Collins, & Asakura, 2008; M. Yang, Tanaka, et al., 2008; M. Yang, Yamauchi, Kurokawa, & Asakura, 2007). However, the production of these artificial proteins has generally been dependent on microorganisms or cell culture systems, which are obstacles to bulk production.

Recently, a germ line transformation method for the silkworm has been developed using the transposon piggyback (Kojima et al., 2007; Tamura et al., 2000). This enabled the production of functionalized transgenic (TG) SF in a large quantity such as cell-adhesive silks (Asakura et al., 2014; Yanagisawa et al., 2007), colored fluorescent silk (Iizuka et al., 2013), calcium binding silk (Nagano et al., 2011), and silk better suited to artificial blood vessels (Saotome et al., 2015). Such an improvement of the properties of SF for biomaterials has been attained by incorporating suitable DNA sequences. SF consists of a heavy chain (H-chain) and light chain (L-chain) connected by a disulfide bond, as well as a glycoprotein named P25. It is secreted into the posterior silk gland and is thought to be assembled into a high molecular mass elementary unit with a ratio of 6:6:1 (Couble, Chevillard, Moine, Ravel-Chapuis, & Prudhomme, 1985; Takei, Kikuchi, Kikuchi, Mizuno, & Shimura, 1987; Tanaka et al., 1999; Tanaka, Mori, & Mizuno, 1993). The new gene that follows the SF gene is expressed as a new functionalized SF in the posterior silk gland. Subsequently, the SF stored in the middle silk gland is spun out through the anterior silk gland and converted into SF fibers, being coated by silk sericin that emerges in the middle silk gland. After removal of silk sericin by a degumming process, we obtained pure SF. There are many advantages in preparing functionalized SF suitable for biomaterials by the TG technique: (1) the TG silkworms are easy to handle and the larvae are well adapted for artificial rearing; (2) the adult moths cannot fly away and therefore cannot live in the wild; (3) the expression of the recombinant silk proteins can be confirmed visually by fusion of green fluorescent proteins as a marker.

We were interested in the possibility that a combination of the AGSGAG and minTBP-1 motifs could further enhance the functionality of SF when used as a coating agent for titanium substrates. While titanium has already been widely used in dental implants, artificial joints, and vessel stents (Brunette, Tengvall, Textor, & Thomsen, 2001), further improvement of its biocompatibility is still required (Pagel et al., 2016; Schuler et al., 2006; Vidal et al., 2013; Zhang, Zhang, Zhu, Kang, & Neoh, 2008). In the present study, we first synthesized artifice polypeptides containing both AGSGAG and minTBP-1 motifs, and confirmed their reversible interacting ability to titanium, which characterizes the minTBP-1 motif. Then, we constructed the transgenic *B. mori* that spins the cocoon containing the artificial SF. The modified SF prepared from the cocoons enhanced the ossific differentiation of MC3T3-E1 cells grown on the titanium plates coated by the artificial SF. The binding mechanism of silk-TBP on the surface of TiO_2 nanoparticles was examined briefly using the model peptides made from the AGSGAG and minTBP-1 motifs, and ^{13}C solid-state NMR.

MATERIALS AND METHODS

Production of silk-like proteins, TS[(AGSGAG)$_3$AS]$_n$RKLPDAS and TS[(AGSGAG)$_3$AS]$_n$ by *Escherichia coli*.

The recombinant silk-like proteins TS[(AGSGAG)$_3$AS]$_n$RKLPDAS (n = 2 and 4) and TS[(AGSGAG)$_3$AS]$_n$ (n = 2 and 4) were prepared according to the method described previously (Yanagisawa et al., 2007; M. Yang & Asakura, 2005; Mingying Yang et al., 2009; M. Yang, Muto, et al., 2008; M. Yang, Tanaka, et al., 2008; M. Yang et al., 2007). Here, (AGSGAG)$_3$ (Bc3) and RKLPDA represent the crystalline region of SF and minimal sequence of titanium binding peptide, respectively. The construction scheme is shown in Figure 1a. The pUC-Bc3 is the recombinant vector, in which a single unit of DNA sequence coding (AGSGAG)$_3$AS was inserted into pUC118 vector plasmid (Yanagisawa et al., 2007; M. Yang & Asakura, 2005). Because this unit contains *Spe* I and *Nhe* I restriction enzyme recognition sites, and because these enzymes produce complemental sticky ends, directional multiplexing of the unit is possibly though repeated cut-and-paste manipulations (Prince, McGrath, DiGirolamo, & Kaplan, 1995; M. Yang & Asakura, 2005). In this study, we constructed 2-mer and 4-mer of [(AGSGAG)$_3$AS]$_n$ repeats, which were named pUC-(Bc3)$_2$ and pUC-(Bc3)$_4$, respectively. To add RKLPDA motif at the end of repeats of (AGSGAG)$_3$AS, oligonucleotide KY-1775, pCTAGTCGGAAGCTGCCCGACG, and pCTAGCGTCGGGCAGCTTCCGA were annealed and ligated into the *Nhe* I site of pUC-(Bc3)$_2$ and pUC-(Bc3)$_4$ to obtain pUC-(Bc3)$_2$T and pUC-(Bc3)$_4$T. Integrities

of constructed artificial genes were confirmed by sequencing using the Dye Terminator Cycle Sequencing Quick Start Kit (Beckman Coulter, Brea, CA). For the expression in *Escherichia coli*, the inserts of these four plasmids were digested by *Bam* HI and *Hind* III, and were re-cloned into the same sites of the expression vector pET30a (Nobagen Inc Darmstadt, Germany) to obtain pET-(Bc3)$_2$, pET-(Bc3)$_4$, pET-(Bc3)$_2$T, and pET-(Bc3)$_4$T. In pET30a vectors, artificial silk fibroin proteins are expressed as the fusion proteins that carry a vector derived 6xHis-tag sequence at the N-terminal.

Because the pET30a vector is driven by a T7 promoter, the expression vectors were introduced in the *E. coli* strain BL21(DE3)pLysS (Nobagen Inc.), which has T7 polymerase, and the transformed cells were grown in 200–500 mL of LB (Sambrook, Fritsch, & Maniatis, 1989) containing kanamycin and chloramphenicol at 37 °C. At OD$_{660}$ = 0.5, isopropyl β-D-1 thiogalactopyranoside (IPTG) was added to a final concentration of 1 mM to induce protein expression for 4 h. After induction, cells were harvested and frozen and stored at -80 °C until purification. The protein expression was induced at 25 °C by adding IPTG to a final concentration of 1 mM. The recombinant proteins were purified by nickel-chelate chromatography using an ÄKTA explorer 10S/100 (GE Healthcare, Waukesha, WI, USA) column according to the manufacturer's protocol. The optimized conditions for elution were 15 mM imidazole, pH 4.5, and 50 °C. The eluted protein solutions (6–8 mL) were dialyzed against 10 mM NaH$_2$PO$_4$, pH 8.0 and 100 mM NaCl using a Slide-A-Lyzer Dialysis Cassette (3,500 MWCO, Pierce, Waltham, MA), followed by concentration into 500 μL by using Amicon Ultra Centrifugation Filters-4 (Merck Millipore, Billerica, MA). The purified proteins from pET-(Bc3)$_2$, pET-(Bc3)$_4$, pET-(Bc3)$_2$T, and pET-(Bc3)$_4$T were termed (Bc3)$_2$, (Bc3)$_4$, (Bc3)$_2$T, and (Bc3)$_4$T, respectively (Figure 1 a).

QCM analyses.

Quartz Crystal Microbalance (QCM) measurements were carried out on Qsense D300 equipment with a Ti sensor (Q-Sense, Vastra Frolunda) that was intensively cleaned with a UV-ozone ProCleaner system (BioForce Nanosciences, Ames, IA) prior to use. Before measurements, the Ti-sensor and the chamber were equilibrated in 50 mM Tris-HCl (pH 8.0) and 150 mM NaCl (TBS) containing 0.05% polyoxyethylene sorbitan monolaurate (Tween 20) (TBST) at 25 °C. The proteins were prepared in a 2 mL volume at a concentration of 0.5 μM in TBST. The data were obtained and analyzed by Q soft 301 and q-tool (Q-Sense).

Construction of transgenic silk worm that spins the cocoons containing minTBP-1 appended silk fibroin (TiBP-SF).

To introduce the RKLPDA motif (minTBP-1) into the SF heavy chain (HC), a DNA fragment coding [TS[(AGSGAG)$_3$AS]$_2$RKLPDA]$_8$was synthesized (GenScript, Piscataway, NJ, USA) and inserted into the*Bam* HI and *Sal* I sites of plasmid pHC-EGFP (Kojima et al., 2007). The resultant plasmid was digested with *Asc* I and*Fse* I and inserted into the *Asc* I and *Fse* I sites of the plasmid pBac[3xP3-EGFPafm](Horn, Schmid, Pogoda, & Wimmer, 2002) to produce the vector plasmid pBac[HC-TiBP-3xP3EGFP] (Figure 1 b). TG silkworms were produced by injecting the vector plasmid pBac[HC-TiBP-3xP3EGFP] into the eggs of silk worms in order to produce TG silkworms, as reported previously (Tamura et al., 2000). TG silkworms were established using a non-diapausing strain (w1-pnd) and were mated with the diapausing strain (w1) to maintain TG silkworms as diapausing strains. The silkworms were reared on an artificial diet (Nosan Corporation, Yokohama, Japan) or on fresh mulberry leaves at 25 °C. Due to the poor yield and robustness of the silk of TG silkworm strains, they were backcrossed repeatedly with the commercial strains Gunma and 200, which have been shown to have high silk productivity. The resulting hybrid strains were then used for mass rearing of silkworms and collection of silk from the cocoons. The wild-type *B. mori* SF was used for comparison with the TG SF (TiBP-SF). The SF and TiBP-SF were dissolved by lithium bromide and desalinated using PD-10 column (GE Healthcare, Buckinghamshire, UK). Then, the samples were analyzed by 4%–12% gradient SDS-PAGE. The gel was stained with Coomassie Brilliant Blue R250 or transferred to polyvinylidene fluoride (PVDF) membranes. The membrane was treated with an anti-His tag antibody (A190-214A, BETHYL, Montgomery, USA). The immunoreactive bands were visualized using ECL Plus (GE Healthcare, Buckinghamshire, UK) and an LAS-3000 image analyzer (Fujifilm, Tokyo, Japan).

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Surface modification of the Ti surface with SF or TiBP-SF.

Pure wrought Ti disks (ø7 mm × 1 mm) were purchased from Furuuchi Chemistry (Tokyo, Japan). The surface of each Ti disk was polished with a sequence of 320-, 600-, and 1200-grit silicon carbide papers (RIKEN, Saitama, Japan) under a stream of water. The polished Ti surface was used after being rinsed with ultrapure water (MilliQ®), ultrasonically cleaned with acetone, and then dried at room temperature. For coating, the disk was immersed in 5 ml 2 mass% SF or TiBP-SF aqueous solution at 20 °C for 24 h. After being air-dried at 20 °C for 1 h, the disk was rinsed with 70% ethanol and then air-dried again at 20 °C for 24 h.

Cell culture.

Osteoblast-like cells from rat calvaria (MC3T3-E1) cells were cultured in sterile cell culture plates containing α-minimal essential medium (α-MEM) with 10% heat-inactivated fetal bovine serum (FBS) (Biowest, Riverside, MO, USA), 2 mM L-glutamine and 10 U/mL penicillin/streptomycin (Gibco, Grand Island, NY, USA) in a humidified 95% air/5% CO_2 incubator at 37 °C. When the cells reached sub-confluence, they were harvested and sub-cultured. The cell suspension (5.0×10^4 cells) was slowly dripped onto the substrates (non-coated, SF-coated and TiBP-SF coated) in 48-well plates. Three replicates of each sample were tested during the culture period. Plates were shaken using gentle agitation for 10 min and were then incubated for 3 h at 37 °C. Then, 0.5 mL of medium was added along the side of each well.

Scanning electron microscopy observations.

At the end of the various incubation times, the non-attached cells on the different substrates were removed from the substrates by rinsing twice with 0.1 M sodium cacodylate buffered solution, dehydrated in a graded series of ethanol (50%, 70%, and 90%), and then placed in 100% ethanol. The dehydrated samples were then immersed in tert-butyl alcohol under vacuum (FDU-1200, EYELA, Tokyo, Japan). The freeze-dried samples were mounted on aluminum stubs and were then sputter-coated with a gold-palladium alloy. Scanning electron microscope (SEM, S-2150, Hitachi, Tokyo, Japan) observations of each sample were then performed at numerous magnifications at 10 kV.

ALP assay

Next, 5×10^4 MC3T3-E1 cells/mL in α-MEM were seeded in 6-well plates overnight. The medium in each well was removed the next day and replaced with osteoblast differentiation-inducing medium (OIM) (MK430, Takara Bio Inc., Tokyo, Japan), and then incubated at 37 °C for 3 weeks in a 5% CO_2 incubator. Day 0 is the date the OIM medium was added. ALP activity was determined using an ALP assay kit (ab83369, Abcam, Cambridge, MA, USA) following the manufacturer's instructions. In brief, untreated control and NaF-treated MC3T3-E1 cells were homogenized in assay buffer after washing with cold PBS and centrifuged at 15,000 rpm for 15 min at 4 °C. For each measurement, the supernatants were added in triplicate in 96-well plates and each well was brought to a total volume of 80 µL with assay buffer, followed by the addition of 20 µL stop-solution for a background control. Standard curves were generated as detailed in the manufacturer's manual. The ALP activity of each sample was calculated based on a comparison between the standard and sample curves and is reported as Unit/mL (U/mL).

Cell differentiation, Alizarin Red S staining, and quantification.

Here, 5×10^4 MC3T3-E1cells/ml in α-MEM were seeded overnight with SF or TiBP-SF. The medium was removed the next day, replaced with OIM, and incubated at 37 °C for 3 weeks in a 5% CO_2 incubator. Day 0 is the date the differentiation medium was added to cells. The entire medium was replaced every 3 days. Alizarin red S staining was performed to analyze calcium deposits. Cells were washed twice with Hanks balanced salt solution (HBSS) (Gibco 14025, Thermo Fisher Scientific, Waltham, MA, USA) and then fixed with 4% paraformaldehyde for 15 min. The cells were stained with 1% Alizarin Red S solution for 3 min at 37 °C and rinsed with distilled water to exclude non-specific staining. For quantitation, cells stained with Alizarin red were de-stained with 10% cetylpyridinium chloride (Sigma-Aldrich, Saint Louis, MO, USA) for 15 min, after which the extracted stain was transferred to 96-well plates, and the absorbance at 562 nm

was measured using a microplate reader. After 2 and 3 weeks of osteogenic induction, the level of OC in the supernatant was directly measured using an ELISA kit (USCN Life Science, Wu Han, China) according to the manufacturer's instructions and a microplate reader at 450 nm. The assays were performed in triplicate.

Quantitative RT-PCR analysis.

At weeks 2 and 3 after incubation with OIM, total RNAs were isolated using an miRNeasy Mini kit (Qiagen, Hilden, Germany). First-strand cDNAs were synthesized from 1 μg of each total RNA using Super Script VILO Master Mix (Life Technologies, Carlsbad, CA, USA). One μL of each cDNA was subjected to real-time RT-PCR using TaqMan Gene Expression Assays (Applied Biosystems, Foster City, CA, USA) for Runx2 (Mm00501584_m1), Osterix (Mm04209856_m1), ALP (Mm04209856_m1), Collagen Type 1 (Col I) (Mm04209856_m1), OPN (Mm00436767_m1) and OC (Mm03413826_m1), and Pre-Developed TaqMan Assay Reagents (Applied Biosystems) for β-actin (Mm02619580_g1) as an internal control. Two-step PCR cycling was carried out as follows: 2 min at 50 °C for 1 cycle, 10 min at 95 °C for 1 cycle, then 15 s at 95 °C and 1 min at 60 °C for 40 cycles. Two independent measurements were averaged, and relative mRNA expression levels were calculated as a ratio to β-actin expression of each sample.

Statistical analysis.

All quantitative data are expressed as means ± SD. Statistical analyses were performed using the statistics program SPSS 20. Results were analyzed by one-way ANOVA, followed by post hoc tests for pair-wise comparisons (Tukey's test). Differences were considered to be significant for $p < 0.05$.

Synthesis of the Model Peptides of the AGSGAG and minTBP-1 motifs.

To examine the binding mechanism of silk-TBP on the surface of TiO_2 nonoparticles, the model peptides of the AGSGAG and minTBP-1 motifs, i.e., RKLPDA (minTBP-1) , $(AGSGAG)_2$, AGSGAGGRKLPDAG-GAGSGAG (Silk-TBP), and AGSG[1-^{13}C]AGGRKLPD[3-^{13}C]AGGAGSGAG (^{13}C Silk-TBP) were synthesized in a stepwise fashion on Fmoc-amino acid-PEG-PS resin by a Pioneer Peptide Synthesis System (PE Biosystems, USA) using Fmoc solid phase method in our laboratory (Asakura et al., 2005; Asakura & Yao, 2002; Yu Suzuki et al., 2010; Yao et al., 2004). The purity was more than 95%, as verified by high-performance liquid chromatography and ^{13}C solution NMR. The [1-^{13}C]Ala (99.9%^{13}C enrichment) and [3-^{13}C]Ala (99.9% ^{13}C enrichment) were purchased from Cambridge Isotopes Laboratories, Andover, MA, USA.

Preparation of TiO_2 particles with ^{13}C Silk-TBP on the surface.

The nanoparticles of TiO_2 (Aeroxide TiO_2P25), which were kindly provided by NIPPON AEROSIL co., JP, had an average particle size of 21 nm and a surface area of 50 ± 15 m²/g. The TiO_2 nanoparticles were suspended in water and sonicated twice for 30 min with a 50% duty cycle. The suspension was then centrifuged at 10,000 rpm for 15 min, and the precipitated TiO_2 nanoparticles were washed in distilled water. This process was repeated, and the particles were dried overnight. The dried particles were suspended in 20 mM phosphate buffer at pH = 7.0 and sonicated again for 10 min with a 50% duty cycle. The peptides (TBP, $(AGSGAG)_2$, and Silk-TBP) were dissolved in 20 mM phosphate buffer and mixed with the suspended TiO_2 nanoparticle solutions, respectively. The weight ratio of the peptide and suspended TiO_2 nanoparticle was 1:1 for each case. Finally, the state of dispersion of the mixture after standing overnight was observed.

Solid-state ^{13}C CP/MAS NMR Measurement .

The ^{13}C Silk-TBP (20 mg) was dissolved in 20 mM phosphate buffer at pH = 7.0 (10 ml) and mixed with the TiO_2 nanoparticles (50 mg). The solution was stirred for 4 h at 100 rpm to reach equilibrium. Then equilibrated. Then the suspension was centrifuged at 10,000 rpm for 10 min and excess peptide dissolved in the supernatant was removed. This process was repeated. Thus, TiO_2 particles with ^{13}C Silk-TBP adsorbed on the surface were obtained after drying. The ^{13}C CP/MAS NMR spectra of the ^{13}C Silk-TBP and ^{13}C Silk-TBP adsorbed on the surface of TiO_2 nanoparticles were recorded on a Bruker Avance 400 NMR spectrometer with an operating frequency of 100.0 MHz for ^{13}C at a sample spinning rate of 8 kHz in a 4 mm diameter ZrO_2 rotor. The numbers of scans were 1 K for ^{13}C Silk-TBP and 40 K for ^{13}C Silk-TBP adsorbed

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on the TiO_2 nanoparticles. The pulse delay was 5 s and contact time was 2 ms. The ^{13}C chemical shifts were referenced to TMS using adamantane as a secondary standard (^{13}CH peak at 28.8 ppm).

RESULTS AND DISCUSSION

QCM analyses

Designer proteins (Bc3)$_2$, (Bc3)$_4$, (Bc3)$_2$T, and (Bc3)$_4$T (Figure 1 a) were synthesized and purified as described in MATERIALS AND METHODS. Using QCM, the mode of interactions between the crated artificial proteins and titanium surface was assessed. In the QCM assay, binding of molecules on the surface of the sensor decreases the resonance frequency of the sensor and detachment of molecules from the surface is observed as the resonance frequency increases. Steps of QCM measurements are schematically illustrated in Figure 2 a. First, the Ti sensor was equilibrated with water for 30–60 min and the measurement was started (time = 0). After 5 min, TBST (a solution buffer of the proteins) was injected 3 times (time = 5, 10, and 15 min). Then, protein solution was injected into the measurement chamber (time = 20 min). After 30 min, the sensor was washed with TBST 3 times (time = 50, 55, and 60 min), followed by washing with water 3 times (time = 65, 70, and 75 min). Results obtained from (Bc3)$_2$ and (Bc3)$_2$T are shown in Figure 2 b. Injections of the protein solutions rapidly decreased the resonance frequency of the sensor, indicating that both (Bc3)$_2$ and (Bc3)$_2$T have bound to the surfaces of Ti. Because TBST washing only slightly increased the frequency, the binding of these proteins to Ti is stable under the conditions. When the measurement sensor was washed with water (time = 65 min), an increase of frequencies was observed, which was due to buffer change (not detachment of proteins from the sensor). When the rates of frequency increase in water were compared between (Bc3)$_2$ and (Bc3)$_2$T (time = 65–80 min), it can be recognized that (Bc3)$_2$T was detaching faster than B(Bc3)$_2$. Similar results were obtained from (Bc3)$_4$ and (Bc3)$_4$T (Figure 2 c).

Construction of transgenic *Bombyx mori* that spins the cocoons containing the functionalized silk fibroin (TiBP-SF).

The QCM analyses indicated that the artificial proteins composed of SF core motifs and minTBP-1 motif have the abilities to bind to (and detach from) the titanium surface. To evaluate the effect of the appended minTBP-1 motif on the nourishing ability of SF for cells, we decided to construct the large-scale production system for SF-minTBP-1 proteins. For this experiment, we selected the (Bc3)$_2$T = [TS[(AGSGSG)$_3$AS]$_2$RKLPDA] as a core unit. This was because in the QCM experiments, (Bc3)$_2$T bound in larger quantities and detached faster (especially in water) than (Bc3)$_4$T, suggesting the properties of minTBP-1 was more manifested in this formula.

We have already established the transgenic *B. mori* system, which has enabled us to prepare large amounts of engineered SF (Asakura et al., 2014; Iizuka et al., 2013; Kojima et al., 2007; Nagano et al., 2011; Saotome et al., 2015; Tamura et al., 2000; Yanagisawa et al., 2007). In this study, a DNA fragment coding [TS[(AGSGSG)$_3$AS]$_2$RKLPDA]$_8$((Bc3)$_2$T) was inserted into the plasmid pHC-EGFR(Kojima et al., 2007) to construct pBac[HC-TiBP-3xP3EGFP] (Figure 1 b), which was injected into the eggs of the silkworm, and transgenic silkworms were selected as described previously (Kojima et al., 2007; Tamura et al., 2000; Yanagisawa et al., 2007). The established transgenic silkworms spin cocoons that are apparently indistinguishable from wild-type samples (data not shown). From the cocoons, SF proteins were prepared following standard procedures described in MATERIALS AND METHODS, and the obtained fibroin proteins (named TiBP-SF) were analyzed by SDS-PAGE followed by CBB staining (Figure 3 a). From the TiBP-SF sample, in addition to SF H-chain and SF L-chain proteins, an additional protein band was observed that was not detected from the wild-type SF sample. Western blotting using an anti-His tag antibody confirmed that the protein was derived from a transgenic gene (Figure 3 b).

The apparent molecular weight of the appendix protein (75 K) was larger than that of the estimated molecular weight (52 kD). Similar abnormal mobilities of the modified SFs in SDS-PAGE have also been reported in other experiments (Asakura et al., 2014; Iizuka et al., 2013; Kojima et al., 2007; Saotome et al., 2015; Teule et al., 2012; Yanagisawa et al., 2007), which may be due to either skewed amino acid compositions of the proteins or particular secondary structures. Because the ratio of minTBP-1 motif to SF motif should be

extremely small (approximately 0.1 to 0.9 % in a similar system (Yanagisawa et al., 2007)) compared to the model artificial proteins used in the experiments of Figure 2, we were interested in the effect of minTBP-1 on the binding ability of SF to titanium. As shown in Figure 4 a, QCM analyses indicated that TiBP-SF decreased the amounts of protein absorbed on the titanium sensor, which agreed well with the results obtained from $(Bc3)_2T$ and $(Bc3)_4T$. This means that with a decreased ratio of minTBP-1/silk motif, the minTBP-1 motif masks the binding ability of sill peptide to the material surface. Thus, doping SF with minTBP-1 dramatically changed the binding properties of SF. This notion was further confirmed by the observation that the Silk model peptide $(AGSGAG)_2$ was not able to disperse Ti particles in aqueous solution, whereas Silk-TBP model peptide AGSGAGGRKLPDAGGAGSGAG can endow the Ti particle with dispersibility similar to the mimTBP-1 peptide (Figure 4 b).

Osteoblast differentiation on the TiBP-SF-coated titanium surfaces.

The above data demonstrated that TiBP-SF and SF have distinct behaviors in terms of binding to the titanium surface. However, it was not clear whether the altered properties of TiBP-SF can improve the performance of titanium for medical devices. To address this question, we performed biological assessments using an osteoblastic cell line (MC3T3-E1), which has often been used to evaluate osteoblast differentiation. After coating titanium plates with either SF or TiBP-SF (as described in MATERIALS and METHODS), MC3T3-E1 was grown on the plates for 14 and 21 days in the differentiation-inducing medium (OIM). The morphologies of the cells on the plates were then observed by SEM. As shown in Figure 5 a, the enhanced differentiation of MC3T3-E1, which is characterized by an elongated sheet-like cell structures with multiple filopodia-like intercellular connections protruding from their leading edges, was evident after 21 days when the titanium plates were coated with TiBP-SF. Without coating (or SF-coating), these characteristic morphologies were not evident.

The formation of filopodia from MC3T3-E1 after 21 days of growth on the TiBP-SF coated Ti plate suggested the cells were differentiated into a hydroxyapatite-mineralizing state, which was further confirmed by the increased alkaline phosphatase (ALP) activity and osteocalcin (OC) production. As shown in Figure 5 b, both the ALP activity and OC production from the cells grown on the TiBP-SF coated Ti substrate were the highest among the 3 groups after 21 days of growth. Moreover, the ALP activity of the SF group was lower than that of the TiBP-SF group (p < 0.05), indicating that TiBP-SF was more beneficial than SF for inducing cell differentiation and promoting cell mineralization. The mineralization of MC3T3-E1 cells was also evaluated by alizarin red staining (Figure 5 c). Cells grown on TiBP-SF displayed higher mineralization capacities than those grown on SF (p < 0.01), as well as the control after culture for 21 days. The calcified deposits were uniform and clustered in all locations. To evaluate the cell differentiation state in more detail, the expression levels of osteoblast-related genes (including Runx2, Osterix, OC, OPN, ALP, and Col I) were evaluated by real-time PCR analyses after culture for 14 and 21 days (Figure 5 d). Runx2 and Osterix are master transcription factors involved in regulating the process of MC3T3-E1cell differentiation. After culture for 14 days, the expression levels of both the Runx2 and Osterix genes in cells cultured on TiBP-SF were up-regulated more than those cultured on SF or on the control. Although the expression level of the Runx2 gene decreased in all groups after culture for 21 days, the TiBP-SF group remained much higher than the SF group. A similar trend was also found for Osterix expression. The early markers of osteoblasts (such as the transcription of ALP and Col I) were also up-regulated after culture for 21 days on TiBP-SF compared with the other groups. Furthermore, culture on TiBP-SF up-regulated the mRNA expression levels of OC and OPN at most time intervals. Thus, TiBP-SF was shown to have an ability for enhancing the differentiation of an osteoblast model cell line *in vitro* .

Conformations of [13]C Silk-TBP and[13]C Silk-TBP adsorbed on the surface of TiO_2 nanoparticles in the solid state.

To obtain further insights into the binding mechanisms of the SF-minTBP-1 proteins, we synthesized the [13]C -labeled model peptide for solid state NMR analyses. Figure 6 shows[13]C CP/MAS NMR spectra (10-70 ppm) of (a) [13]C Silk-TBP (AGSG[1-^{13}C]AGGRKLPD[3-^{13}C]AGGAGSGAG) and (b) [13]C Silk-TBP adsorbed on the surface of TiO_2 nanoparticles together with the assignment (Asakura et al., 2005; Asakura & Yao, 2002;

Yu Suzuki et al., 2010; Y. Suzuki, Shindo, & Asakura, 2016; Wishart, 2011; M. Yang & Asakura, 2005; Yao et al., 2004). The chemical shifts and the assignments are summarized in Table 1. According to the chemical shifts of the peaks, Ala Cβ and Ala Cα carbons, the conformation of the ^{13}C Silk-TBP sample in a solid state is considered to be random coil. This is due to the sample preparation by freeze-dry treatment (Asakura et al., 2005; Yao et al., 2004). After adsorption treatment of the peptides on the surface of the TiO$_2$ nanoparticles, a small^{13}C labeled methyl peak of [3-^{13}C]Ala^{13} residue in the partial sequence (RKLPD[3-^{13}C]A) in the peptide could be still observed. Interestingly, the Ala^{13}Cβ chemical shift changed from a random coil (16.3 ppm) to an α-helix (14.4 ppm) (Figure 6). Namely, the Ala^{13} residue adsorbed on the TiO$_2$ particle took an α-helix form. The α-helix peak also became remarkably sharper compared to the Ala Cβ random coil peak of the peptide before adsorption. This could indicate that the interaction between the minTBP-1 and the surfaces of TiO$_2$ is governed by reversible interactions. In addition, relatively sharp peaks at 21.0 and 25.6 ppm could be observed, which are assigned to the side chain carbons of natural abundant Leu and Lys residues in the sequence RKLPDA. Namely, the peak at 21.0 ppm is assigned to the Leu Cδ$_1$,Cδ$_2$, and Lys Cγ carbons, and the peak at 25.6 ppm to the Lys Cδ carbon (according to the chemical shifts). Thus, the Leu and Lys residues are considered to be attached to the surface of TiO$_2$ particles, although the backbone peaks of these residues could not be observed.

CONCLUSIONS

By using a transgenic *B. mori* system, we synthesized the modified silk fibroin TiBP-SF, which is doped with an artificial protein composed of silk fibroin motif (AGSGAG) and titanium binding motif (RKLPDAS) in the [TS[(AGSGAG)$_3$AS]$_2$RKLPDAS]$_8$formula. The TiBP-SF enhanced the ossific differentiation of osteoblast-like cells when coated on the surfaces of a titanium plate. Accordingly, TiBP-SF has potential applications in the medical field.

Acknowledgments

Figure captions

Figure 1. (a) Construction flow of the recombinant plasmids from genetic blocks coding for the crystalline region of silk fibroin ((AGSGAG)$_3$AS, blue arrow) and for the minimal titanium binding motif (RKLPDA , orange arrow). B, N, S, and H represent*Bam* HI, *Nhe* I, *Spe* I, and *Hind* III, respectively. From the resultants plasmids, the His-tagged proteins, (Bc3) $_2$, (Bc3) $_4$, (Bc3)$_2$T, and (Bc3) $_4$T were produced in BL21(DE3)pLysS and purified by nickel-chelate chromatography, whose domain structures are schematically illustrated by the colored ellipses (indicated in the inner box). (b) The structure of the vector for constructing the transgenic silkworm. The gene for [TS[(AGSGAG)$_3$AS]$_2$RKLPDA]$_8$is under the control of Fibroin H promoter. (c) The structure of proteins translated from pBac[HC-TiBP-3xP3EGFP], which has the N-terminal domain (NTD), the C-terminal domain (CTD), and C-terminal His-tag, which were derived from the parental vector, pBac[3xP3-EGFPafm] (Horn et al., 2002). Arrows indicate right and left arms of the transposon piggyBac. EGFP was used for screening the transgenic silkworm.

Figure 2. (a) Schematic drawings representing the changes in frequency ([?]f; left axis) of titanium sensor upon (i) buffer injection, (ii) sample injection, (iii) buffer wash and (iv) water wash in QCM analyses. (b) Changes in resonance frequency with 0.5 μm (Bc3)$_2$ (blue) and (Bc3) $_2$T (green). (c) Changes in resonance frequency with 0.5 μm (Bc3) $_4$(blue) and (Bc3) $_4$T (green).

Figure 3. (a) SDS-PAGE and Coomassie Brilliant Blue R250 staining for the functionalized SF, TiBP-SF, and its parental SF. The arrow indicates the artificial protein containing [TS[(AGSGAG)$_3$AS]$_2$RKLPDA]$_8$(Figure 1 c). The open circle, the closed circle, and the triangle indicate Fibroin H-chain protein, Fibroin L-chain protein, and Fibro hexamerin protein (Inoue et al., 2000), respectively. (b) Western blotting using anti-His antibody, which recognizes the C-terminal His-tag of the

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artificial SF.

Figure 4. (a) Changes in resonance frequency of titanium sensor with 1 mg/mL of SF (blue) and TiBP-SF (green). Details of experimental procedures are described in Figure 1 caption and MATERIALS AND METHODS. (b) Dispersion of TiO_2 particles (21 nm) with TBP (RKLPDA) peptide, Silk-TBP (AGS-GAGGRKLPDAGGAGSGAG) peptide, Silk ($(AGSGAG)_2$) peptide and no peptide (control). Details of experiments are described in MATERIALS AND METHODS.

Figure 5. (a) SEM images (magnification of 2000×) of non-coated (cont), SF-coated and TiBP-SF-coated polished Ti surfaces after cultivating MC3T3-E1 cells for 14 or 21 days. The bars indicate 10 μm. (b) Effects of TiBP-SF coating on the alkaline phosphatase (ALP) activity and the expression level of osteocalcin (OC) of MC3T3-E1 cells after 14 and 21 days incubation. Values are expressed as means ± SD (n = 3). Asterisks indicate P < 0.05 in the t-test. (c) Effect of TiBP-SF coating on the mineralization in MC3T3-E1 cells. Cells were stained with Alizarin Red (left) and quantified as described in the MATERIALS AND METHODS after 14 and 21 days incubation. Asterisks indicate P < 0.05 in the t-test. (d) Expression levels of mRNA for Runx2, Osterix, ALP) Collagen Type I (Col I), Osteopontin (OPN) and OC, which were quantified by quantitative RT-PCR, in MC3T3-E1 cells after 14 and 21 days incubation on non-coated (control), SF-coated, and TiBP-SF-coated polished Ti surfaces. The relative mRNA expression levels were calculated as a ratio to β-actin expression of each sample. Values represent means ± SD from three independent experiments. Asterisks indicate P < 0.05 in the t-test.

Figure 6. ^{13}C CP/MAS NMR spectra (10–70 ppm) of (a) ^{13}C Silk-TBP (AGSG[1-^{13}C]AGGRKLPD[3-^{13}C]AGGAGSGAG) and (b) ^{13}C Silk-TBP adsorbed on the surface of TiO_2 nanoparticles together with the assignments.

Table 1. ^{13}C chemical shifts and assignments of ^{13}C CP/MAS NMR spectra of (a) ^{13}C

Silk-TBP and (b) ^{13}C Silk-TBP adsorbed on the surface of TiO_2 nanoparticles.

Chemical Shift		Assignment	
(a)	(b)		
(b) 14.5 ppm			Ala ^{13}Cb a-helix
(a) 16.2 ppm	Ala ^{13}Cb random coil		
(b) 21 ppm			Leu Cd_1,d_2 Lys Cg
(b) 25.5 ppm			Lys Cd
(a) 42 ppm (broad)	Gly Ca		
(b) 42–43 ppm (broad)			Gly Ca
(a) 49 ppm (broad)	Ala Ca random coil	-	

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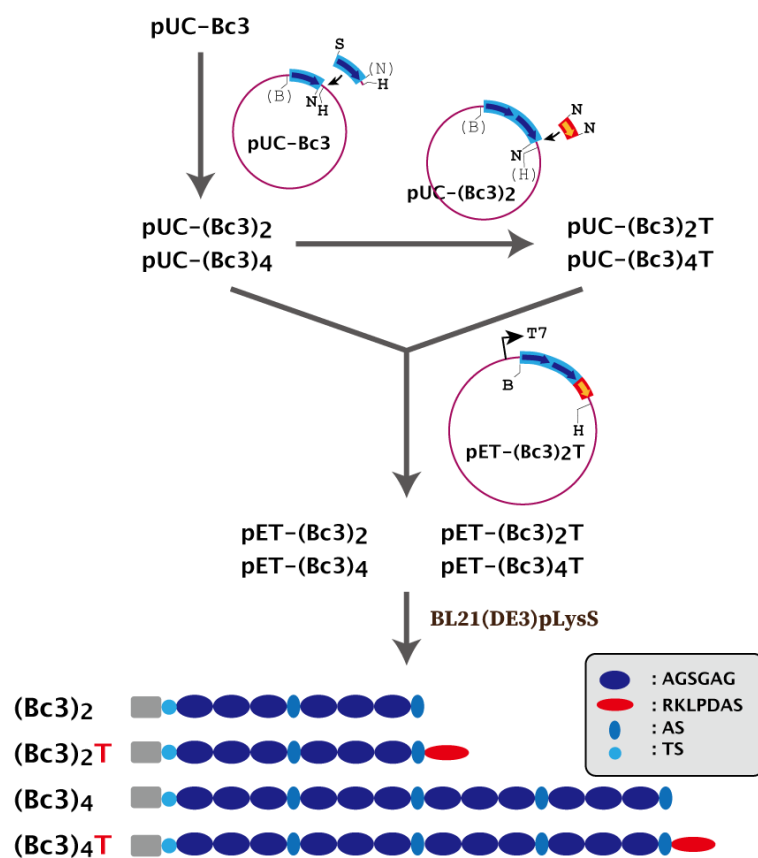
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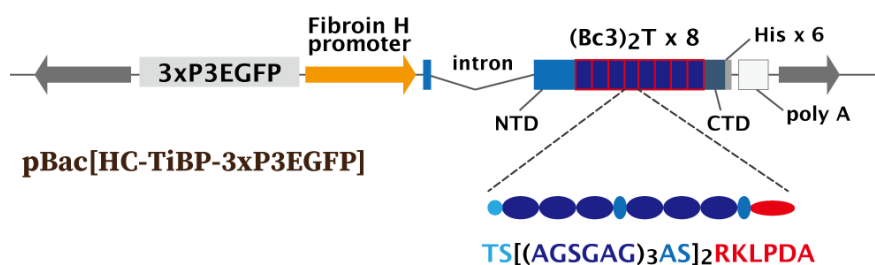

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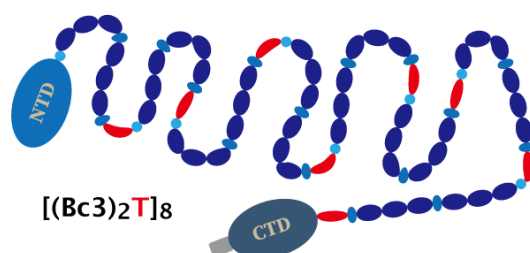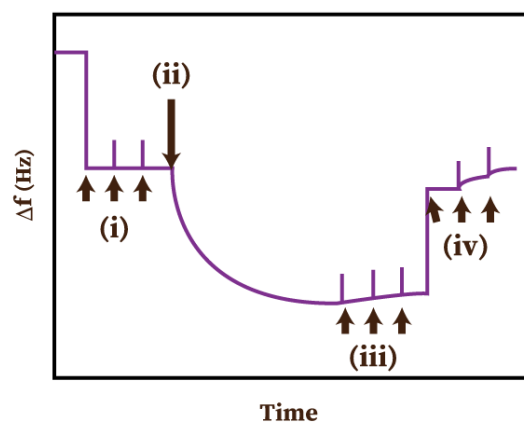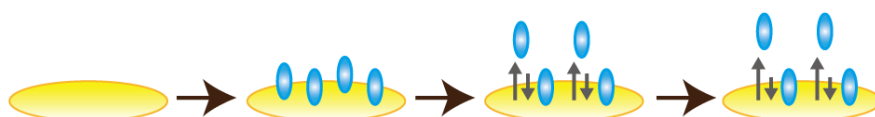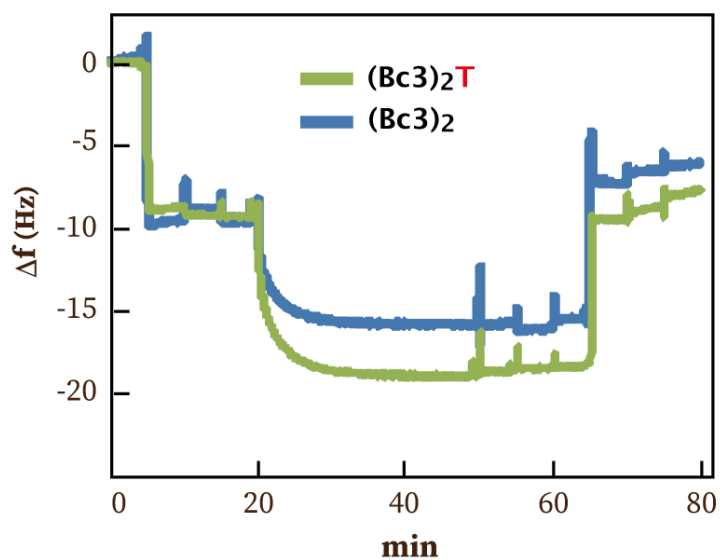
(a)

pUC–Bc3

pUC–Bc3

pUC–(Bc3)2

pUC–(Bc3)2
pUC–(Bc3)4

pUC–(Bc3)2T
pUC–(Bc3)4T

pET–(Bc3)2T

T7

pET–(Bc3)2
pET–(Bc3)4

pET–(Bc3)2T
pET–(Bc3)4T

BL21(DE3)pLysS

(Bc3)2

(Bc3)2T

(Bc3)4

(Bc3)4T

: AGSGAG
: RKLPDAS
: AS
: TS

(b)

3xP3EGFP

Fibroin H promoter

intron

(Bc3)2T x 8

His x 6

poly A

NTD

CTD

pBac[HC-TiBP-3xP3EGFP]

TS[(AGSGAG)3AS]2RKLPDA

(c)

NTD

CTD

[(Bc3)2T]8

(a)

Δf (Hz)

(ii)

(i)

(iii)

(iv)

Time

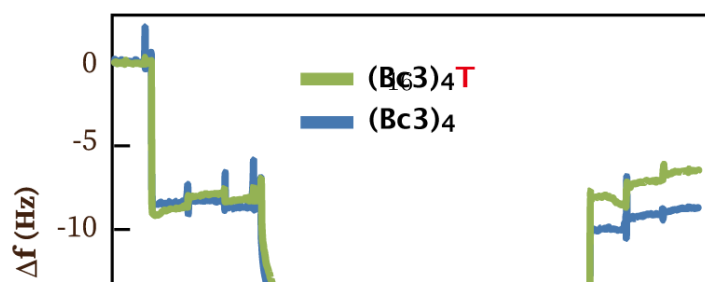(i) buffer injection $\longrightarrow$ (ii) sample injection

$\longrightarrow$ (iii) buffer wash $\longrightarrow$ (iv) water wash

(b)

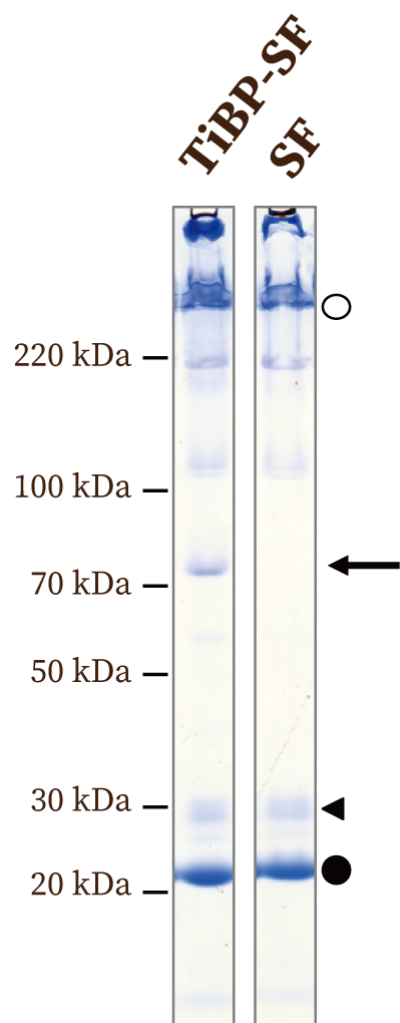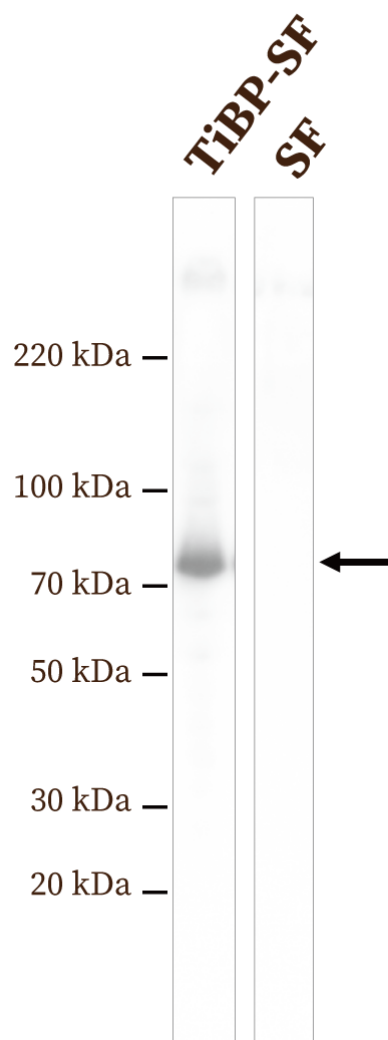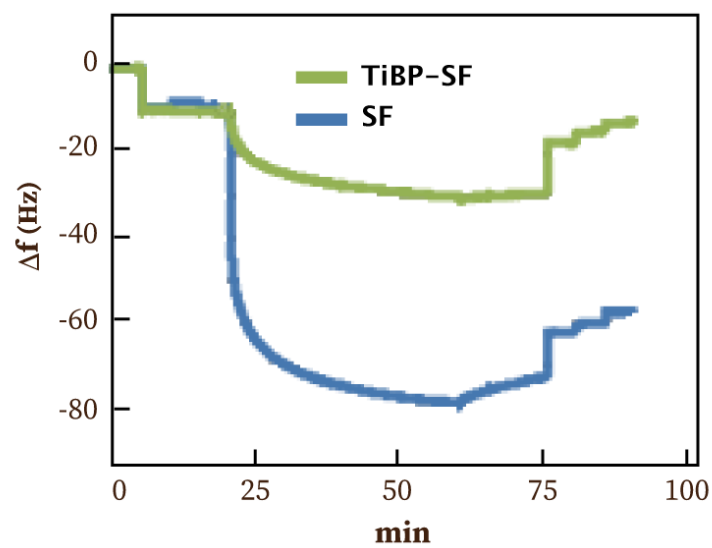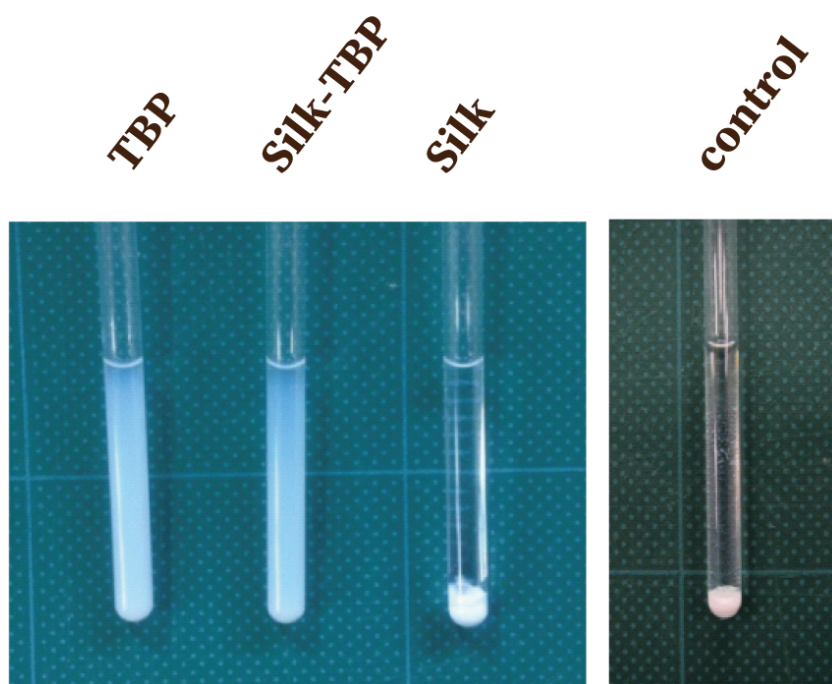Δf (Hz)

(Bc3)$_2$T
(Bc3)$_2$

0   20   40   60   80

min

(c)

Δf (Hz)

(Bc3)$_4$T
(Bc3)$_4$

16

(a)

(b)

TiBP-SF

SF

TiBP-SF

SF

220 kDa —

100 kDa —

70 kDa —

50 kDa —

30 kDa —

20 kDa —

○

←

◄

●

220 kDa —

100 kDa —

70 kDa —

50 kDa —

30 kDa —

20 kDa —

←

(a)

(b)

18

(a)

cont   SF   TiBP-SF

14 days

21 days

(b)

(c)

(d)

(a)

Ala C$^\beta$ (random coil)

Ala C$^\alpha$ (random coil)

Gly C$^\alpha$

(b)

Leu C$^{\delta1,\delta2}$ Lys C$^\gamma$

Ala C$^\beta$ (α-helix)

Lys C$^\delta$

Gly C$^\alpha$

65 60 55 50 45 40 35 30 25 20 15 10

ppm from TMS
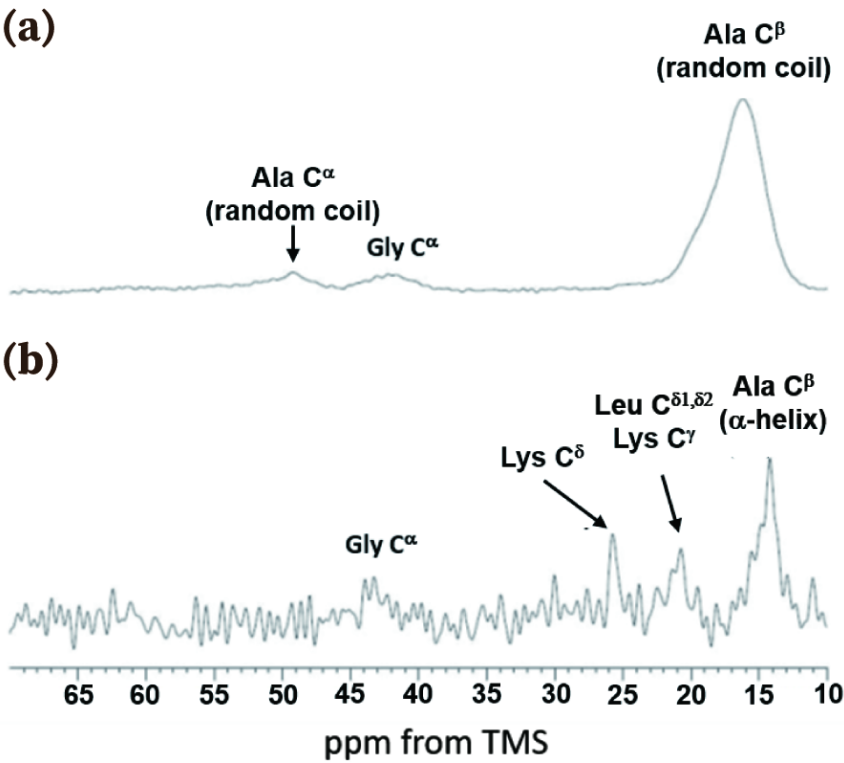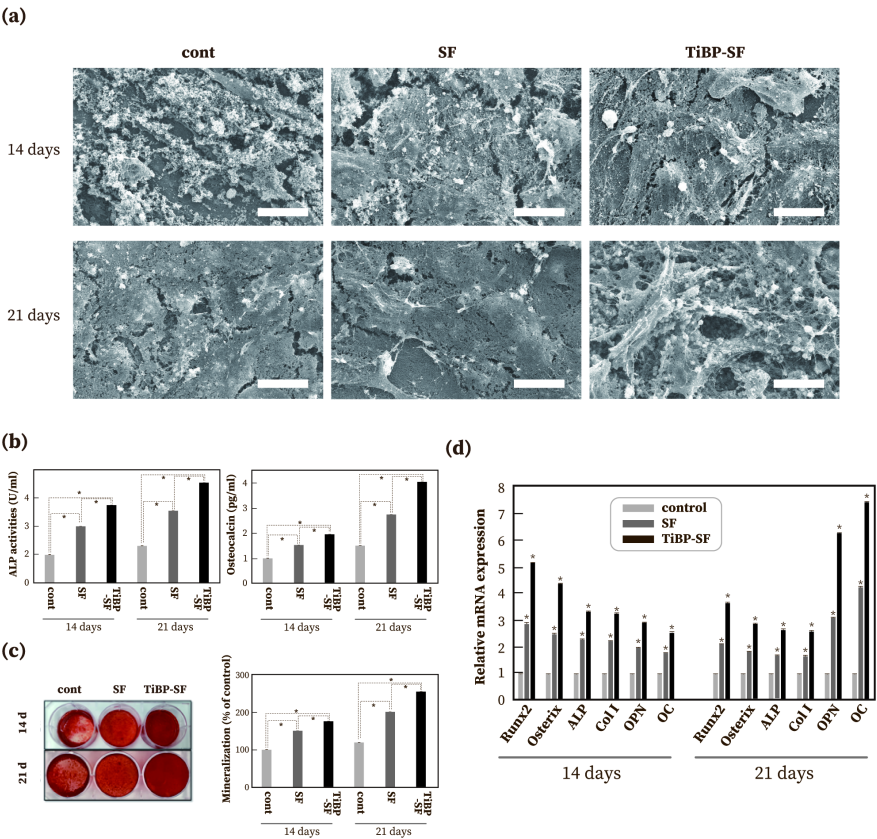
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Table 1.docx available at https://authorea.com/users/354975/articles/478415-bio-functionalized-titanium-surfaces-with-the-modified-silk-fibroin-carrying-titanium-binding-motif-enhanced-the-ossific-differentiation-of-mc3t3-e1