

Culture temperature affects
redifferentiation and cartilaginous
extracellular matrix formation in
dedifferentiated human chondrocytes

（培養温度は脱分化したヒト軟骨細胞
において再分化と関節軟骨細胞外基質
形成に影響を与える）

伊藤 明良

Culture temperature affects redifferentiation and cartilaginous extracellular matrix formation in dedifferentiated human chondrocytes

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Running title: Culture temperature in chondrocytes

Abstract

To date, there have been few studies on how temperature affects the phenotype and metabolism of human chondrocytes. Thus, the purpose of this study was to elucidate the effects of culture temperature on chondrocyte redifferentiation and extracellular matrix (ECM) formation using dedifferentiated mature human chondrocytes *in vitro*. Dedifferentiated chondrocytes were cultured in a pellet culture system for up to 21 days. The pellets were randomly divided into three groups with different culture temperature (32°C, 37°C, and 41°C). Chondrocyte redifferentiation and ECM formation were evaluated by wet weight, messenger ribonucleic acid (mRNA), histological, and biochemical analyses. The results showed that the wet weight and the mRNA expressions of collagen type II A1 and cartilage oligomeric matrix protein at 37°C were higher than the corresponding values at 32°C. The histological and biochemical analyses revealed that the syntheses of type II collagen and proteoglycan were promoted at 37°C compared to those at 32°C, whereas they were considerably inhibited at 41°C. In conclusion, the results obtained herein indicated that temperature affects chondrocyte redifferentiation and ECM formation, and modulation of temperature might thus represent an advantageous means to regulate the phenotype and biosynthetic activity of chondrocytes.

Keywords: chondrocyte; temperature; extracellular matrix; differentiation; pellet culture.

Introduction

Articular cartilage (AC) is a hyaline cartilage composed of a dense cartilaginous extracellular matrix (ECM) with sparse distribution of highly specialized cells called chondrocytes. Recently, tissue engineering and cell-based therapies have been explored for AC regeneration,¹ since AC displays a limited capacity for renewal and self-repair.² Autologous chondrocyte implantation (ACI) is a promising cell-based therapy for repairing AC defects.³ However, ACI poses several challenges. Harvested chondrocytes must be expanded to obtain a large number of cells for transplantation, and yet, this process results in the induction of chondrogenic phenotype loss (i.e., dedifferentiation),^{4,5} causing fibro-cartilage-like remodeling. Thus, characterization of the factors regulating the chondrogenic phenotype is desired for inducing redifferentiation and hyaline cartilage for ECM formation. Candidate factors include the microenvironment, such as the presence of growth factors,⁶ scaffolds,⁷ and oxygen tension,⁸ as well as mechanical stimuli.⁹ While these factors have all been well studied, there are conversely few studies that have focused on the role of temperature in chondrocyte redifferentiation.¹⁰ Environmental temperature is known to influence some tissues such as the skin. Interestingly, the temperature within the human knee joint is also influenced by the environmental temperature, with a mean temperature of approximately 32°C, which is 4–5°C lower than the inner body temperature.^{11,12} However, most *in vitro* studies on chondrocytes or AC have been performed using a culture temperature of 37°C, which may not accurately reflect the *in vivo* temperature. In addition, the effect of a high-temperature environment, such as 41°C, remains unclear, although an intermittent heat stimulus (41°C) has been reported to potentially have a positive effect on ECM formation.^{13,14}

To date, there are few studies on how temperature affects the chondrocyte phenotype and metabolism of mature human chondrocytes. We hypothesized that the culture temperature

may influence the ability of dedifferentiated chondrocytes to redifferentiate and produce hyaline-like ECM. Therefore, the purpose of this study was to elucidate the effects of culture temperature (from physiological- to high-temperature) on redifferentiation and ECM formation using dedifferentiated mature human chondrocytes *in vitro*.

Methods

Chondrocyte isolation and pellet culture

Human chondrocytes were obtained from the femoral heads (International Cartilage Repair Society grade 0) extracted during bipolar hip arthroplasty performed in two 62- and 89-year-old women (donors A and B, respectively), as previously described.¹⁵ The Ethics Committee of the Faculty of Medicine at Kyoto University approved the procedure, and informed consent was obtained from the donors. The isolated cells from the donors were separately expanded in Dulbecco's modified Eagle medium/Ham's F12 (Nacalai Tesque Inc., Kyoto, Japan) containing 10% fetal bovine serum (Hyclone, Logan, UT), 50 U/mL penicillin (Nacalai Tesque Inc.), and 50 µg/mL streptomycin (Nacalai Tesque Inc.) to dedifferentiate in tissue culture dishes in a CO₂ incubator (5% CO₂ at 37°C with 95% humidity) until passage two or three.

To provide a 3-dimensional environment, a pellet culture system was used.¹⁶ The expanded chondrocytes were trypsinized and subsequently resuspended in a chondrogenic medium (Chondrogenic Differentiation Media BulletKit; Lonza, Walkersville, MD), which was supplemented with 10 ng/mL recombinant human transforming growth factor-beta 3 (R&D Systems, Inc., Minneapolis, MN). Aliquots of 2.5×10^5 cells in 500 µL of the chondrogenic medium were centrifuged at $250 \times g$ for 5 min in 15-mL polypropylene conical tubes. The pelleted cells were randomly divided into three groups with different incubation

temperatures (32°C, 37°C, and 41°C). These temperatures were defined as follows: 32°C, physiological intra-articular temperature; 37°C, conventionally used inner body temperature; and 41°C, the threshold temperature for mammalian cell survival.^{17,18} The pellets from donor A were harvested at 3, 7, 14, and 21 days for analysis. The pellets from donor B cultured at 32°C and 37°C were harvested at 3 and 21 days to confirm the reproducibility of the findings from donor A. Throughout the study, “*n*” indicates the technical replicates of the pellet cultures.

Quantitative real-time reverse transcription-polymerase chain reaction (qRT-PCR)

The generated pellets (*n* = 3 pellets/group) were harvested after 3 and 7 days. The total RNA was extracted and qRT-PCR was performed as previously described.¹⁵ The target genes and reference genes used were as follows: chondrogenesis markers (collagen type II A1 [*COL2A1*], aggrecan [*ACAN*], and cartilage oligomeric matrix protein [*COMP*]); a fibro-cartilage maker (collagen type I A1 [*COL1A1*]); and reference genes (ribosomal protein L13a [*RPL13a*] and tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein [*YWHAZ*]). Their specific primers are listed in Supplementary Table 1.

The data obtained by qRT-PCR were analyzed using the comparative threshold cycle method. Briefly, the quantity of the target genes was normalized to the expression levels of *RPL13a* and *YWHAZ*, which have been proven to be stable under different thermal conditions.¹⁹ The value of the calibration sample (cells cultured at 32°C on day 3) was set to 1, and the values for the other conditions were reported relative to that of the calibration sample.

Histological and immunohistochemical (IHC) analyses

The generated pellets cultured in the three different temperatures obtained on days 7, 14, and 21 were stained with safranin-O/fast green for assessment of proteoglycan deposition, and with picosirius red to visualize the collagen orientation and integrity under a polarizing microscope (Eclipse 80i; Nikon, Tokyo, Japan).²⁰ IHC staining of type II and I collagen was performed to detect each type of collagen deposition according to previously described methods.²¹ To semi-quantify the immunoreactivity of type II collagen at 32°C and 37°C on day 21, the images from each group ($n = 8$ pellets/group) were measured using the ImageJ program (National Institutes of Health, Bethesda, MD) as previously described.¹⁴

Scanning electron microscope (SEM) observation

SEM observation was performed to assess the ultra-microstructure of collagen fibers using a SEM system (H-7650; Hitachi, Tokyo, Japan). The generated pellets on day 21 were fixed and cut into two pieces. The specimen was mounted cut surface up on aluminum stubs. The collagen fibers on the surface, superficial, sub-superficial, transition, and deep regions on the cut surface of the pellets were observed.

Measurement of glycosaminoglycan (GAG) and deoxyribonucleic acid (DNA) content

The total GAG content in the pellets on days 14 and 21 ($n = 6$ pellets/group) was measured using the 1,9-dimethylmethylen blue colorimetric method.²² Moreover, the DNA content in these samples ($n = 6$ pellets/group) was assessed using the Quant-iT™ PicoGreen® assay (Invitrogen Ltd., Paisley, UK) following the manufacturer's instructions. To estimate the GAG productive ability per cell, the GAG/DNA ratio was calculated (GAG content/pellet ÷ DNA content/pellet).

Statistical analysis

JMP 11 software (SAS Institute, Cary, NC) was used for the statistical analyses. Descriptive statistics were calculated as means and 95% confidence intervals. Statistical significance for the donor A experiments was determined using the paired *t*-test for the semi-quantitative evaluation of type II collagen, or by one-way analysis of variance using the post-hoc multiple comparison Tukey-Kramer test for other experiments. For donor B, statistical significance was determined using the paired *t*-test. The correlation between GAG content and wet weight was examined using Pearson's correlation coefficient. The differences observed were considered to be significant if the *P* value was < 0.05 .

Results

The results from donors A and B showed similar trends. Therefore, only the results from donor A are described below, whereas the results from donor B, which were used to confirm the reproducibility of the findings from donor A, are described in Supplementary Fig. 1.

Wet weight measuring

Representative pellets generated at the three different temperatures are shown in Fig. 1(a). The pellets generated at 32°C and 37°C showed a ball-like shape, whereas that at 41°C showed a disc-like shape. Fig. 1(b) shows the wet weight changes over time. Although the wet weight was heavier at 32°C than at the other temperatures on day 3, the heaviest pellets were those cultured in 37°C obtained on days 14 and 21. On the other hand, at 41°C, the wet weight was significantly lighter than at the other temperature, and did not change over time.

Gene expression analysis

Gene expression analysis related to the cartilaginous ECM was performed on days 3 and 7 to analyze the early effects of temperature (Fig. 2). The gene expressions analyzed were found to be significantly down-regulated at 41°C. Conversely, the expressions of *COL2A1*, *COL1A1*, *ACAN*, and *COMP* were all up-regulated on day 7 compared to those on day 3 at 32°C and 37°C. The expressions of *COL2A1* and *COMP* were significantly more up-regulated at 37°C than at 32°C on days 3 and 7, whereas no significant differences were observed for the expressions of *COL1A1* and *ACAN* in this early phase.

Histological and IHC analyses

Representative images are shown in Fig. 3. Safranin-O staining, type II collagen IHC staining, and picrosirius red staining revealed a progressive deposition at 32°C and 37°C, but not at 41°C. IHC staining of type I collagen showed early and intense deposition at 32°C and 37°C, but not at 41°C. Picrosirius red staining at 32°C was observed not only in the superficial region but also in the deep region (Fig. 3 [white arrow]) on days 14 and 21, while that at 37°C was observed mainly in the superficial region. To clarify the differences in type II collagen deposition between 32°C and 37°C on day 21, semi-quantitative evaluation was performed. The mean intensity per pixel (Fig. 4a) and mean percentage of type II collagen positive area (Fig. 4b) were found to be significantly higher at 37°C than at 32°C.

SEM observation

Fig. 5 shows representative images of the surfaces and cut surfaces of the pellets on day 21. While the surface of the pellets at 32°C and 37°C appeared even and smooth, these characteristics were not observed at 41°C. In the superficial region, dense and layered collagen fiber formations parallel to the surface were observed at 32°C and 37°C, whereas no

collagen formations were observed at 41°C. In the sub-superficial region, the collagen fibers were randomly oriented, and they appeared sparse through the deep region. The collagen fibers in the transition and deep regions at 32°C appeared denser than at 37°C.

Measurement of GAG and DNA content

The GAG content per pellet at 41°C was significantly lower than that at the other temperature environments (Fig. 6a). The GAG content in the pellet generated at 37°C tended to be higher than that obtained at 32°C. Moreover, the DNA content per pellet at 41°C was significantly lower than that at the other temperature environments (Fig. 6b). When the GAG content was normalized according to the DNA content, the value was found to be significantly higher at 37°C compared to at 32°C on day 21 (Fig. 6c). The GAG content and the wet weight had a strong positive correlation ($R^2 = 0.91$, $P < 0.01$, $n = 36$) (Supplementary Fig. 2).

Discussion

Temperature, which can be manipulated easily in the cell culture process, and possibly also in the clinical treatment, may be one of the key microenvironmental parameters regulating the chondrogenic phenotype and ECM formation. We investigated the effects of three different culture temperatures (32°C, 37°C, and 41°C) on the ability of dedifferentiated mature human chondrocytes to redifferentiate and form ECM *in vitro*. To the best of our knowledge, this is the first report on the effect *in vivo* intra-articular temperature on human chondrocyte metabolism *in vitro*. Our results demonstrated that the wet weight measured up to day 21 showed time-dependent increases at 32°C and 37°C, suggesting ECM accumulation (Fig. 1b). However, on the other hand, the wet weight at 41°C did not change over time.

Moreover, the ECM-related genes (*COL2A1*, *COL1A1*, *ACAN*, and *COMP*) at 41°C were significantly down-regulated compared to at 32°C and 37°C (Fig. 2a–d). These results from the mRNA expression analysis are consistent with those of the safranin-O staining and the IHC staining of type II and I collagens (Fig. 3), as well as with the results of the GAG quantification (Fig. 6a), which all indicated that ECM formation was dramatically inhibited at 41°C. To elucidate this phenomenon further, we observed the generated pellets at 41°C using SEM to clarify the ultra-microstructure of the collagen, and found that the collagen fiber content in these pellets was very low (Fig. 5). In addition, the results of the DNA quantification suggested that there were significant decreases in the cell number within the pellets at 41°C, by day 14 of culture (Fig. 6b).

Peltonen et al.²³ reported that collagen cannot fold into a triple-helix conformation at a temperature of approximately 40°C. Therefore, the reasons for the inhibition of ECM formation at 41°C may be explained by a combination of cell loss, inhibition of ECM-related mRNA expression, and perhaps also by collagen misfolding. Thus, while intermittent heat stimuli may have a positive effect on ECM formation^{13,14}, prolonged exposure to heat stimuli may have the opposite effect, and we should hence consider heat stimuli as a thermal dose combining both temperature and duration.²⁴

Interestingly, in this study, the wet weight showed time-dependent increases at 32°C and 37°C (Fig. 1b), indicating that a cooler environment (32°C) can resemble ECM produced at 37°C, although the wet weight of samples in the 37°C was significantly higher than samples in the 32°C group on days 14 and 21. The wet weight of AC is known to be mainly composed of water (60–85%), type II collagen (15–22%), and proteoglycan (4–7%).²⁵ As proteoglycan traps water, the most important factor affecting the wet weight is thought to be the proteoglycan content. Herein, we observed a strong correlation between the proteoglycan

content and the wet weight (Supplementary Fig. 2); and therefore, the wet weight was thought to be heavier at 37°C due to the higher proteoglycan content (Figs. 3 and 6).

Compared to 32°C, from the aspect of the differentiation state, the culture temperature of 37°C appeared to enhance redifferentiation of the pellet, which comprised dedifferentiated chondrocytes. Dedifferentiated chondrocytes exhibit increasing fibro-cartilage marker type I collagen and decreasing hyaline-cartilage marker type II collagen.^{4,5} In this study, the *COL2A1* mRNA expression (Fig. 2a) and type II collagen protein synthesis (Fig. 4) were higher at 37°C compared to at 32°C. As for the proteoglycan synthesis (Figs. 3 and 6c), the culture temperature of 37°C was also associated with a higher synthesis rate compared to 32°C in the late phase (day 21). Therefore, in our experimental condition, the culture temperature of 37°C was able to induce redifferentiation at a higher rate than 32°C, likely by promoting type II collagen synthesis in the early phase and proteoglycan synthesis in the late phase. Taken together, these results suggest that the cells implanted in a patient through ACI are likely to be affected by the intra-articular temperature, and that their growth would be promoted by regulating the temperature at approximately 37°C.

The findings from the picrosirius red staining observed under a polarizing microscope showed that integrated collagen fibers were observed in the deep region from day 14 at 32°C (Fig. 3 [white arrow]), whereas it was mainly observed in the superficial region at 37°C. These collagen architectural differences according to the culture temperature are consistent with the findings of previous studies that used immature porcine chondrocytes.¹⁶ Furthermore, upon SEM observation, the collagen fibers in the transition and deep regions also seemed to be denser at 32°C than at 37°C (Fig. 5). In addition, we noted that the compressive response of the pellets was altered by the culture temperature (Supplementary Fig. 3). Although the detailed mechanism of this phenomenon remains unclear, these findings suggest that thermal

environment may affect the function of an articular cartilage. To verify this possibility, further investigations are needed.

Our study has a few limitations. First, the detailed signaling cascade responsible for inducing the differences in chondrocyte metabolism according to the different culture temperatures remains unclear. Second, we investigated only three typical culture temperature conditions. The intra-knee joint temperature in active osteoarthritis and rheumatoid arthritis has been reported to be 34–36°C,¹¹ and the temperature is altered by the patients' physical activity level.²⁶ Thus, further investigations are needed to apply multilevel temperature conditions in order to gain an understanding of the precise effects of thermal environment on chondrocytes. Third, we only analyzed cells obtained from two individuals. Therefore, in order to generalize our findings, larger studies are warranted in the future.

In conclusion, the culture temperature of 37°C, which mimics the inner body temperature, was found to promote redifferentiation and ECM formation better than 32°C, which mimics the *in vivo* intra-articular temperature, whereas that of 41°C drastically inhibited ECM formation. Therefore, modulation of thermal environment might represent an advantageous means to regulate the phenotype and biosynthetic activity of chondrocytes. In addition, *in vitro* experiments should consider the culture temperature, since this markedly influences the chondrocyte metabolism and phenotype.

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Figure legends

Figure 1: Macroscopic observations and wet weight.

(a) Representative images of the generated pellets. Scale bar = 1 mm. (b) Wet weight changes of the pellets. The wet weight was heavier at 37°C than at 32°C on days 14 and 21. Values are presented as means $\pm$ 95% confidence intervals ($n = 9$ pellets/group; $*P < 0.05$, 32°C vs. 37°C; $**P < 0.01$, 32°C vs. 37°C; $\ddagger P < 0.01$, 32°C vs. 41°C; $\S\S P < 0.01$, 37°C vs. 41°C).

Figure 2: Gene expression analysis.

Relative mRNA expressions of (a) collagen type II A1 (*COL2A1*), (b) collagen type I A1 (*COL1A1*), (c) aggrecan (*ACAN*), and (d) cartilage oligomeric matrix protein (*COMP*) are shown. These genes were up-regulated from days 3 to 7 at 32°C and 37°C, but not at 41°C. *COL2A1* and *COMP* at 37°C were significantly higher than at 32°C on days 3 and 7. There were no significant differences in the *COL1A1* and *ACAN* gene expressions between 32°C and 37°C. The gene expressions at 41°C were all significantly down-regulated. Values are presented as means $\pm$ 95% confidence intervals ($n = 3$ pellets/group; $*P < 0.05$, 32°C vs. 37°C; $**P < 0.01$, 32°C vs. 37°C; $\dagger P < 0.05$, 32°C vs. 41°C; $\ddagger P < 0.01$, 32°C vs. 41°C; $\S P < 0.05$, 37°C vs. 41°C; $\S\S P < 0.01$, 37°C vs. 41°C).

Figure 3: Histological and immunohistochemical analyses.

Representative images of the histological and immunohistochemical findings are shown. The staining intensities of safranin-O and type II and I collagen increased over time at 32°C and 37°C, but not at 41°C. Picrosirius red staining demonstrated integrated collagen fibers in

the deep region of the pellet at 32°C on days 14 and 21 (white arrow). Scale bar = 500 μm; magnification, ×100.

Figure 4: Semi-quantitative evaluation of type II collagen immunohistochemical staining.

(a) The mean type II collagen intensity per pixel and (b) the mean percentage of the type II collagen positive area were calculated on day 21. Both values were higher at 37°C than at 32°C. Values are presented as means ± 95% confidence intervals ($n = 8$ pellets/group; $**P < 0.01$).

Figure 5: Scanning electron microscope observations.

Representative images of the surface and cut surface of the generated pellets on day 21 are shown. The surfaces at 32°C and 37°C, but not at 41°C, appeared to be even and smooth. In the cut surface, dense and layered collagen fibers parallel to the surface were observed in the superficial region at 32°C and at 37°C, but not at 41°C. The collagen fibers in the transition and deep regions at 32°C seemed to be denser than those at 37°C. Scale bar = 10 μm; magnification, ×2000.

Figure 6: Glycosaminoglycan (GAG) and deoxyribonucleic acid (DNA) content.

(a) The GAG content per pellet at 37°C tended to be higher than that at 32°C, and that at 41°C was significantly lower than at the other two culture temperatures. (b) The DNA content per pellet at 41°C was significantly lower than at the other temperature environments. (c) The GAG content normalized by the DNA content (GAG/DNA) was significantly higher at 37°C than at 32°C on day 21. Values are presented as means ± 95% confidence intervals ($n = 6$ pellets/group; $**P < 0.01$).

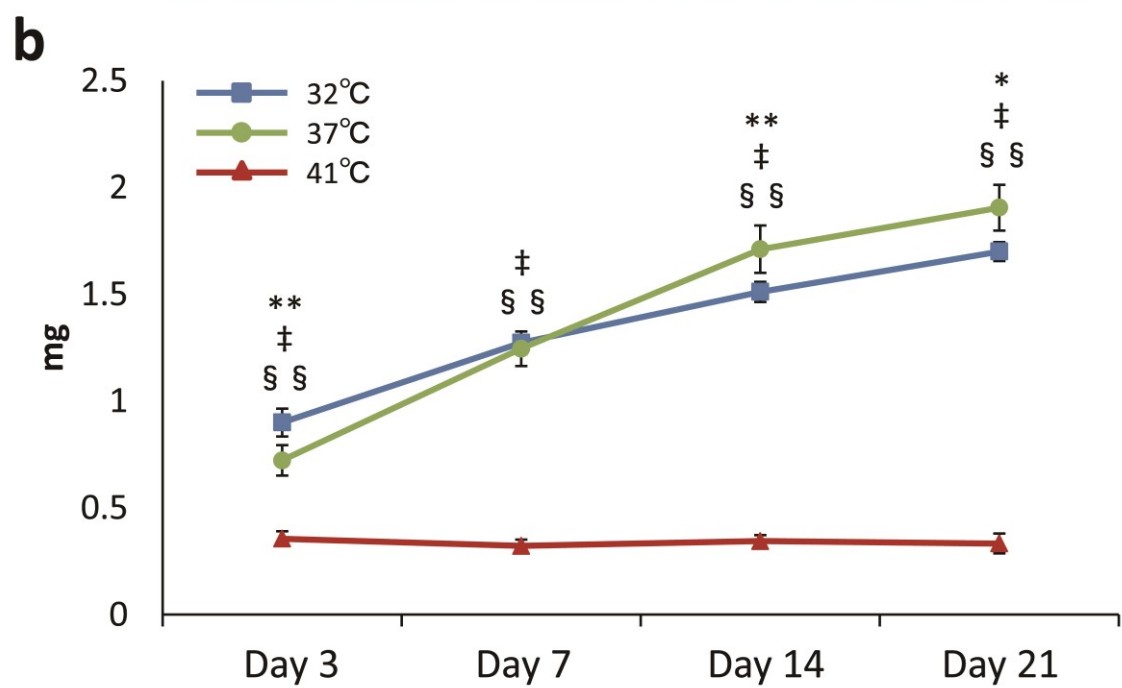
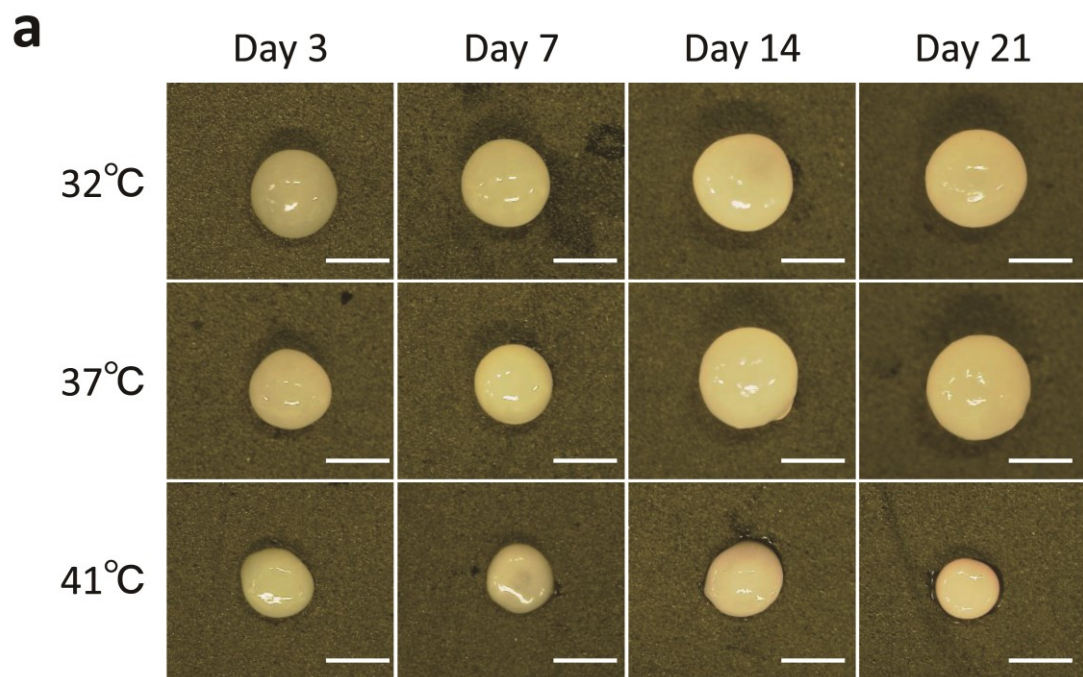


Figure 1

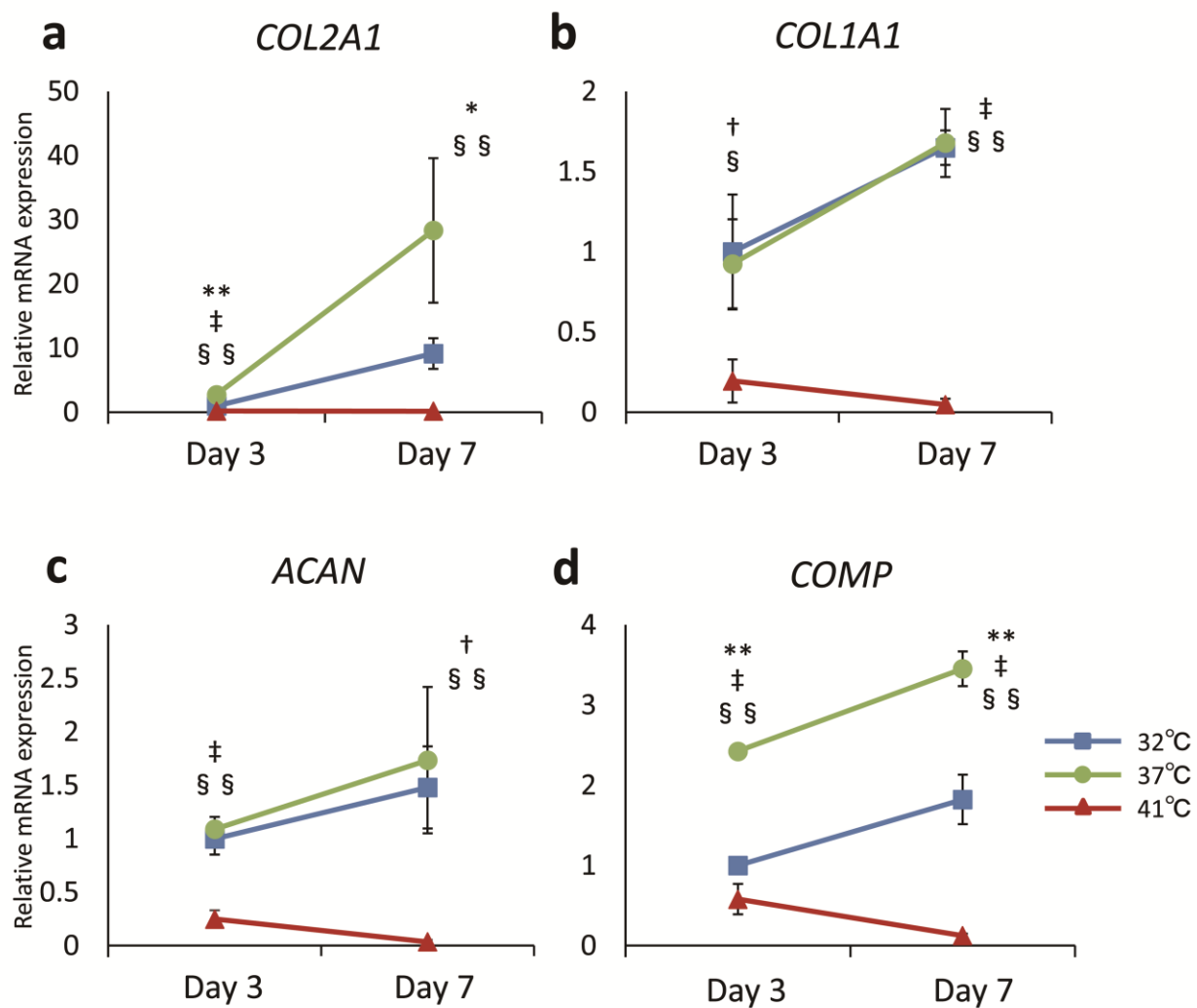


Figure 2

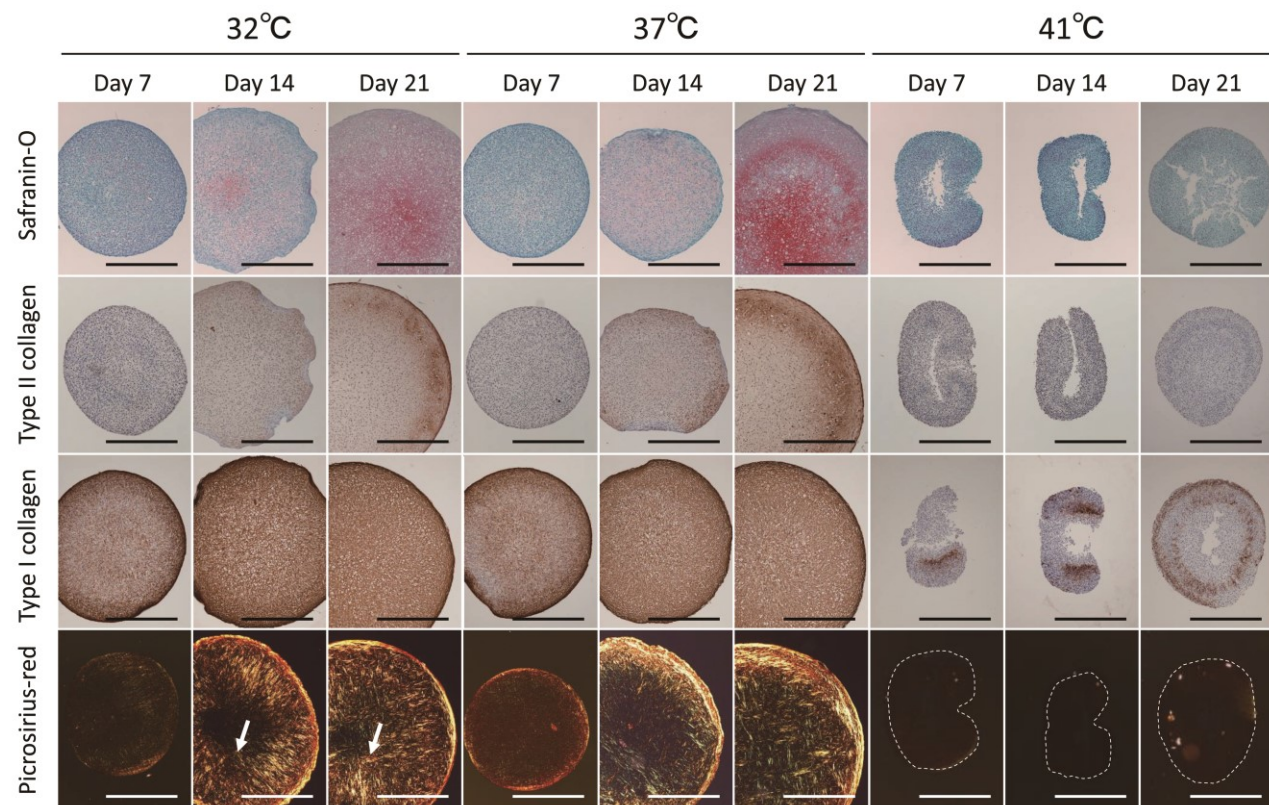


Figure 3

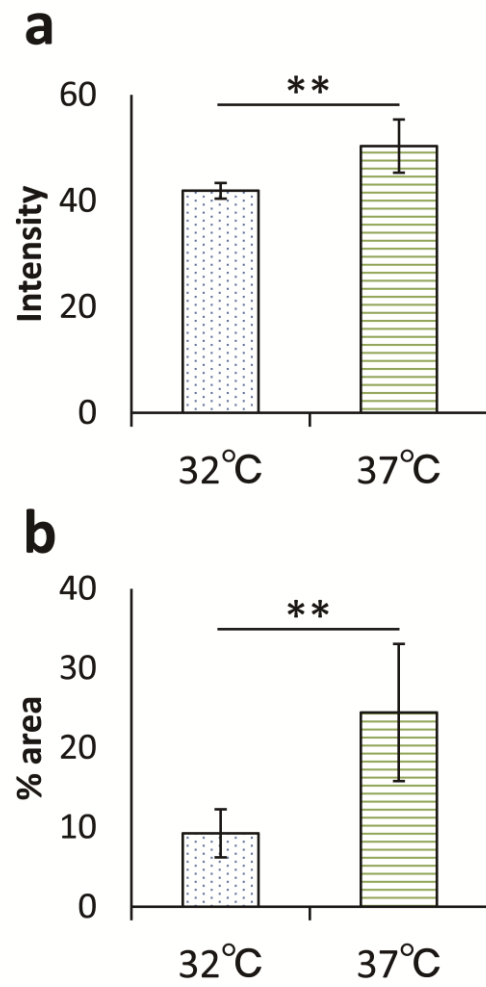


Figure 4

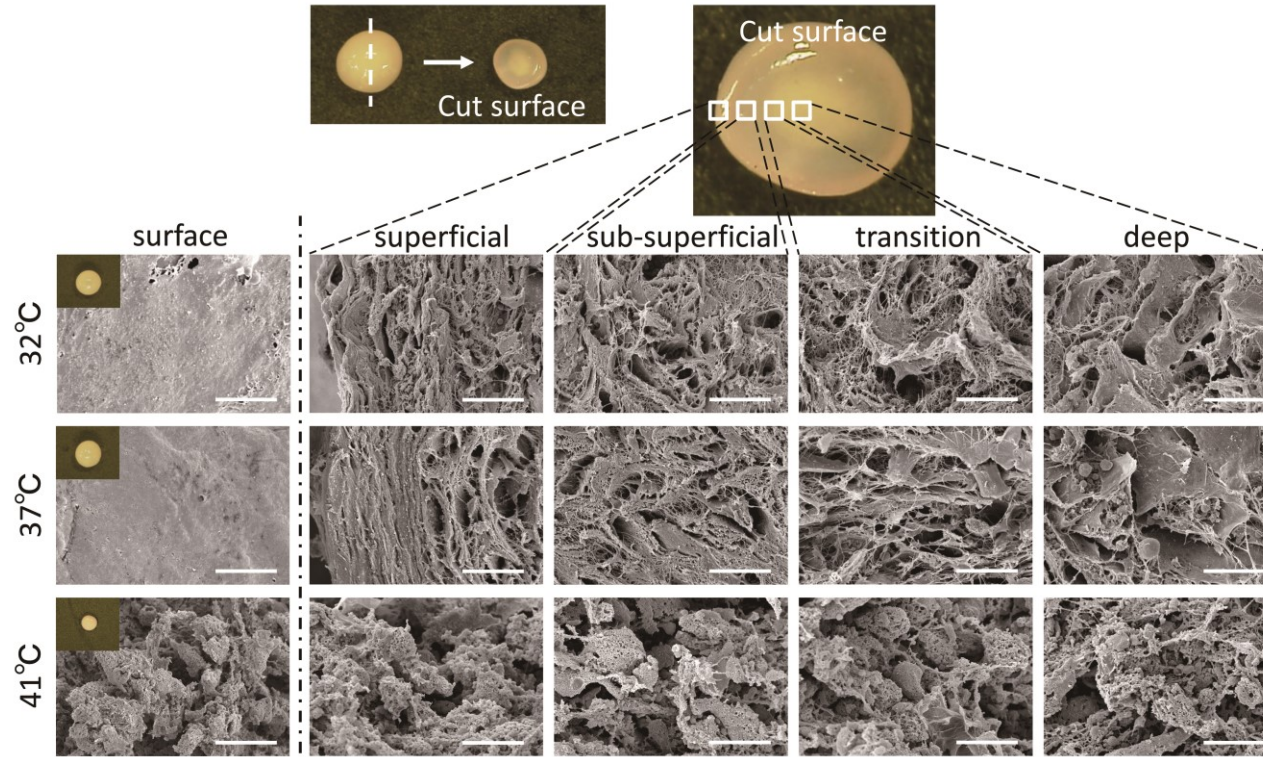


Figure 5

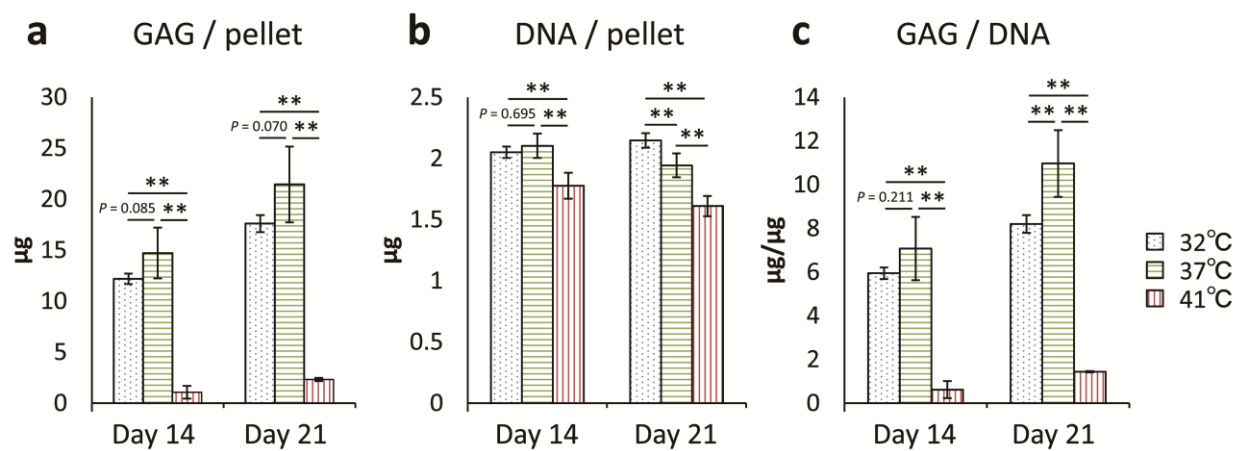
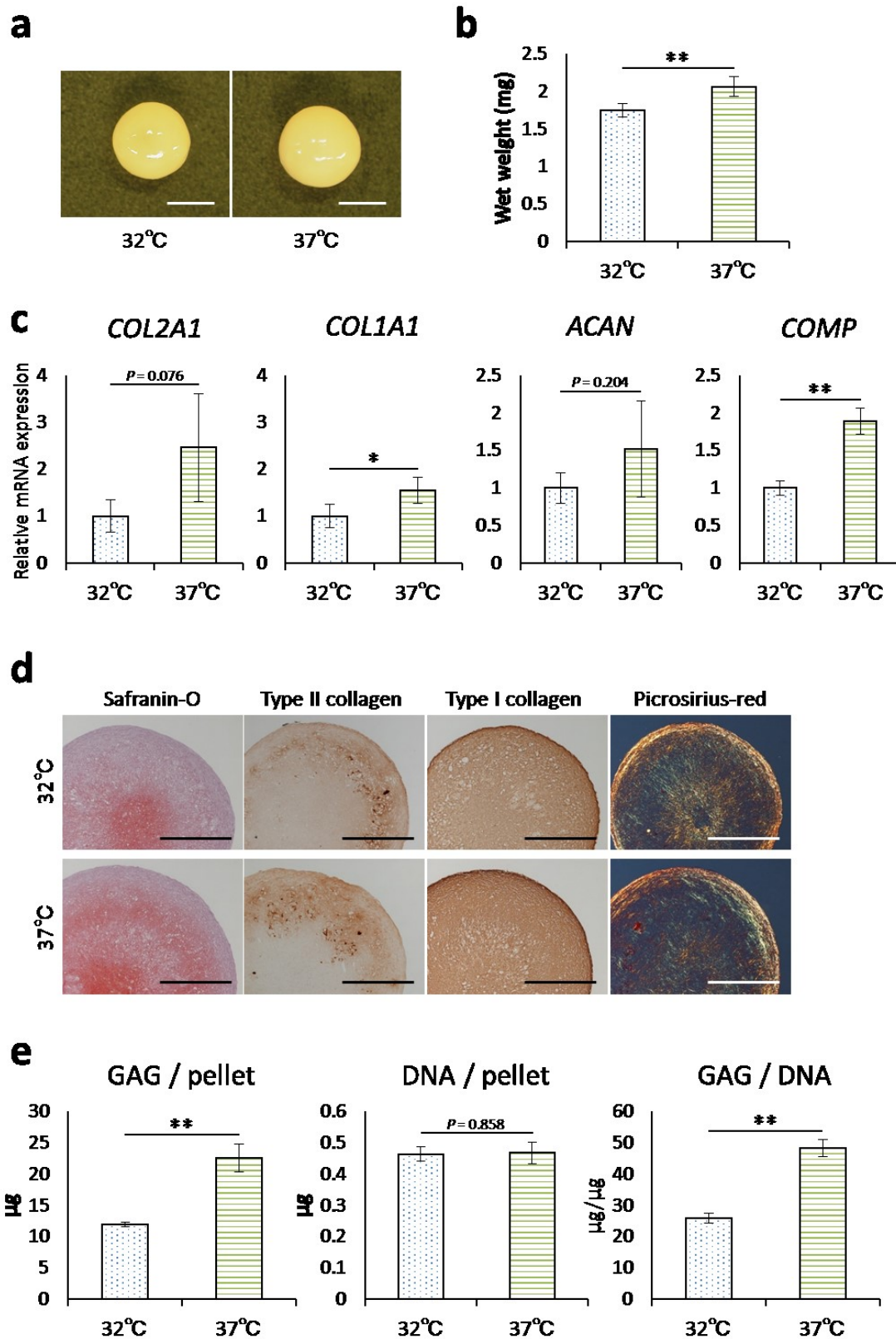


Figure 6

Supplementary Table

Supplementary Table 1. Primer sequences for qRT-PCR

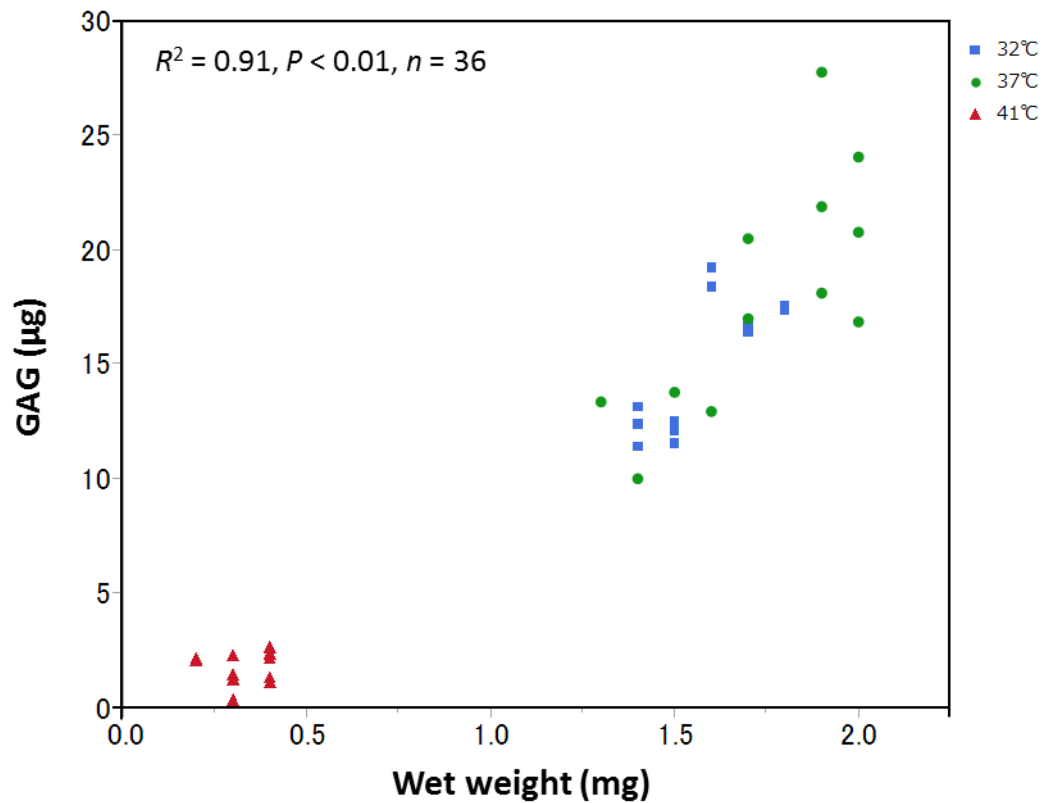
	Sense (5'–3')	Antisense (5'–3')	Length (bp)
<i>COL2A1</i>	GCTATGGAGATGACAACCTGGCTC	CACTTACCGGTGTGTTTCGTGCAG	256
<i>COL1A1</i>	CAGAACGGCCTCAGGTACCA	CAGATCACGTCATCGCACAAAC	101
<i>ACAN</i>	GAATTCCTGGCGTGAGAAC	GGGGATGTTGCGTAAAAGAC	107
<i>COMP</i>	AACAGTGCCCAGGAGGAC	TTGTCTACCACCTTGTCTGC	191
<i>RPL13a</i>	AAGTACCAGGCAGTGACAG	CCTGTTTCCGTAGCCTCATG	100
<i>YWHAZ</i>	TGCTTGCATCCCACAGACTA	AGGCAGACAATGACAGACCA	126



Supplementary Figure 1: Results from donor B.

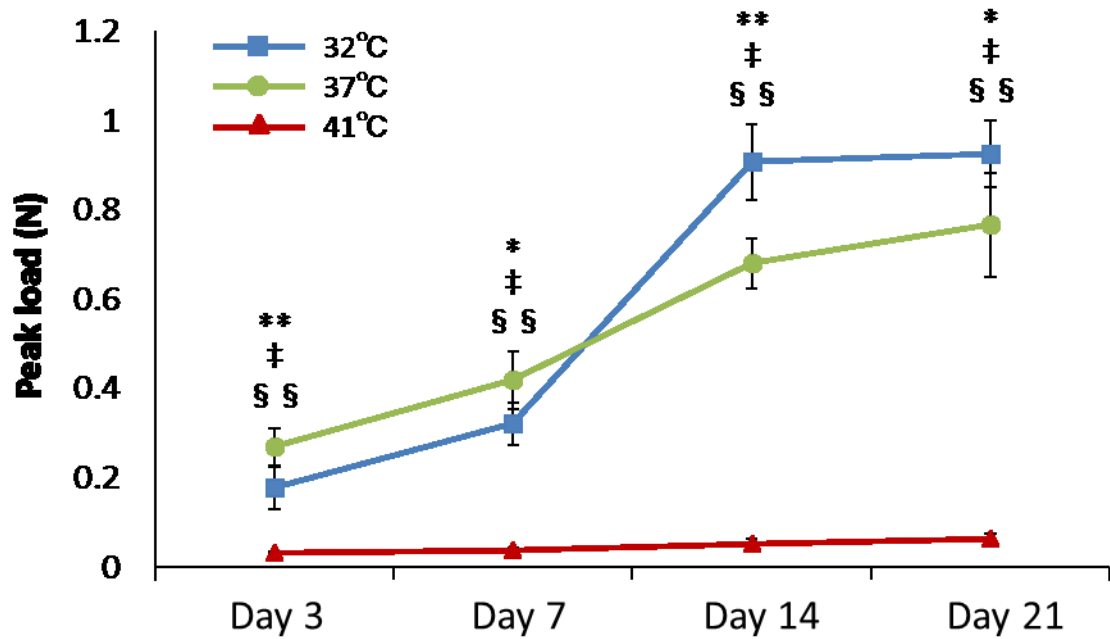
The pellets from donor B cultured at 32°C and 37°C were harvested at 3 and 21 days to confirm the reproducibility of the findings from donor A. (a) Representative images of the generated pellets at 21 days. Scale bar = 1 mm. (b) Wet weight of the pellets at 21 days. The wet weight was heavier at 37°C than at 32°C. Values are presented as means $\pm$ 95% confidence intervals ($n = 6$ pellets/group; $**P < 0.01$). (c) Relative mRNA expressions of *COL2A1*, *COL1A1*, *ACAN*, and *COMP* at 3 days. These genes showed similar trends to those observed for donor A (Fig. 2), except for *COL1A1*. *COL1A1* was slightly, but significantly higher at 37°C than at 32°C. Values are presented as means $\pm$ 95% confidence intervals ($n = 3$ pellets/group; $*P < 0.05$, $**P < 0.01$). (d) Representative images of the histological and immunohistochemical findings at 21 days. Similar results as for donor A (Fig. 3) were obtained. Scale bar = 500 μ m; magnification, $\times 100$. (e) GAG and DNA content per pellet and the GAG/DNA ratio at 21 days. Similar results as for donor A (Fig. 6) were obtained. Values are presented as means $\pm$ 95% confidence intervals ($n = 6$ pellets/group; $**P < 0.01$).

Abbreviations: mRNA, messenger ribonucleic acid; *COL2A1*, collagen type II A1; *COL1A1*, collagen type I A1; *ACAN*, aggrecan; *COMP*, cartilage oligomeric matrix protein; GAG, glycosaminoglycan; DNA, deoxyribonucleic acid.



Supplementary Figure 2: The correlation between the glycosaminoglycan (GAG) content and the wet weight.

The correlation between the GAG content and the wet weight examined using the Pearson's correlation coefficient has a strong correlation ($R^2 = 0.91$, $P < 0.01$, $n = 36$; blue square, 32°C; green circle, 37°C; red triangle, 41°C).



Supplementary Figure 3: Biomechanical analysis.

To analyze biomechanical property of the generated pellets cultured at 32°C, 37°C, and 41°C, an unconfined compression test was performed using a mechanical testing instrument with the use of a 3-mm diameter indenter (Autograph AG-X; Shimadzu, Kyoto, Japan). Each pellet on days 3, 7, 14, and 21 was compressed uniaxially in a testing chamber filled with PBS at room temperature. A pre-load of 0.01 N was applied and allowed to equilibrate for 60 sec. Then the loading was applied at a strain rate of 0.005 mm/s (up to a 50% strain), and peak load (N) was obtained.

The peak load was higher at 37°C until day 7. However, after day 14, it was inverted, and the pellets generated at 32°C showed a significantly higher peak load. The peak load at 41°C was significantly lower than others at all days. Values are represented as means $\pm$ 95% confidence intervals ($n = 6$ pellets/group; $*P < 0.05$, 32°C vs. 37°C; $**P < 0.01$, 32°C vs. 37°C; $\ddagger P < 0.01$, 32°C vs. 41°C; $\S\S P < 0.01$, 37°C vs. 41°C).