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ABSTRACT

The use of intra-oral soft-tissue-engineered devices has demonstrated potential for oral mucosa regeneration. The aim of this study was to investigate the temporal expression of angiogenic biomarkers during wound healing of soft tissue reconstructive procedures comparing living cellular constructs (LCC) with autogenous free gingival grafts. Forty-four human participants bilaterally lacking sufficient zones of attached keratinized gingiva were randomly assigned to soft tissue surgery plus either LCC or autograft. Wound fluid samples were collected at baseline and weeks 1, 2, 3, and 4 post-operatively and analyzed for a panel of angiogenic biomarkers: angiogenin (ANG), angiostatin (ANT), PDGF-BB, VEGF, FGF-2, IL-8, TIMP-1, TIMP-2, GM-CSF, and IP-10. Results demonstrated a significant increase in expression of ANT, PDGF-BB, VEGF, FGF-2, and IL-8 for the LCC group over the autograft group at the early stages of wound repair. Although angiogenic biomarkers were modestly elevated for the LCC group, no clinical correlation with wound healing was found. This human investigation demonstrates that, during early wound-healing events, expression of angiogenic-related biomarkers is up-regulated in sites treated with LCC compared with autogenous free gingival grafts, which may provide a safe and effective alternative for regenerating intra-oral soft tissues (ClinicalTrials.gov number, NCT01134081).

KEY WORDS: tissue engineering, gingival recession, regenerative medicine, wound repair, regeneration, angiogenesis, clinical trial.

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Angiogenic Biomarkers and Healing of Living Cellular Constructs

INTRODUCTION

Gingival augmentation procedures, such as free gingival grafts, are performed routinely in an attempt to maintain an adequate zone of keratinized gingiva and improve gingival health in patients. However, there is morbidity associated with the harvesting of autogenous gingival grafts. Furthermore, the palate possesses limited tissue, providing a limited treatment area in some individuals. Tissue-engineered solutions such as a living cellular construct (LCC; CelTx[™], Organogenesis Inc., Canton, MA, USA) provide an unlimited source and can be a successful alternative to increasing the width of keratinized tissue without using autogenous palatal donor tissue. LCC is a living cellular and collagen construct, designed to regenerate intra-oral soft tissue comprised of type I bovine collagen and viable allogeneic human fibroblasts and keratinocytes isolated from human neonatal foreskin.

The living cellular construct evaluated in this study has been investigated for several different clinical indications, such as diabetic foot ulcers, venous leg ulcers, excision wounds, epidermolysis bullosa, and pressure ulcers (Falanga *et al.*, 1998; Falabella *et al.*, 1999; Veves *et al.*, 2001; Fivenson *et al.*, 2003). Recently, LCC was demonstrated to be safe and capable of generating keratinized tissue without the morbidity and potential clinical difficulties associated with donor site surgery (McGuire *et al.*, 2008).

The mechanism of action of LCC is still unknown. However, it is speculated to modulate healing by secondary intention of the surrounding soft tissues. Living cellular constructs act as a temporary wound covering that is eventually replaced by host cells. DNA of allogeneic fibroblasts and keratinocytes placed over wounds was no longer present after 6 wks of healing (Griffiths *et al.*, 2004). It is postulated that the population of live fibroblasts and keratinocytes improves the wound environment through growth factor interactions, matrix deposition and degradation, wound coverage, and a provision of responsive cells, leading to a clinically beneficial outcome (Sabolinski *et al.*, 1996). The purpose of this study was to investigate the expression of angiogenic biomarkers expressed during oral wound repair using living cellular constructs to regenerate soft tissue.



Figure 1. Patient distribution, time line, and living cellular construct transplantation. (A) Stratification of the study population was divided into 2 treatment groups at 3 clinical centers. (B) Study timeline. GCF, gingival crevicular fluid; WF, wound fluid; CWHS, clinical wound-healing score. (C) Sealed bioreactor with living cellular construct; (D) living cellular construct placed at the surgical site, 254 x 169 mm (300 x 300 DPI).

MATERIALS & METHODS

This investigation enrolled 44 participants from 3 clinical centers (Fig. 1A). This sample size was selected for feasibility rather than to power specific hypotheses, given that this is a first-time-in-humans investigation. Patients involved in the study possessed bilateral mucogingival defects with $\leq 1 \text{ mm of}$ attached gingiva located in contralateral quadrants requiring mucogingival surgery to augment the zone of keratinized and attached gingiva. The patients in this study were a cohort of 96 research subjects participating in a pivotal clinical trial designed to evaluate living cell constructs as an alternative to tissue from the palate to enhance oral soft-tissue regeneration and wound healing who agreed and consented to participate in this investigation (Clinical Trial Registration No. NCT00587834) (McGuire et al., 2010). The participants provided periodontal wound fluid (WF) after mucogingival surgeries with either LCC (Fig. 1C) or autogenous free gingival grafts (autograft). The 2 surgical sites of each patient were randomly selected to receive LCC as a donor material in one site (Fig. 1D) or a conventional autograft using keratinized tissue from the palate as the donor material at the contralateral site. A computer-generated randomization scheme provided allocations for both treatment assignments, and order of administration was contained in sealed envelopes. Informed consent was obtained at the initial visit prior to any research-related treatment or procedure. The study protocol conformed to the ethical guidelines as reflected in obtainment of approval by the University of Michigan's Human Subjects Research Review Committee and the Western Institutional Review Board (WIRB). This human trial investigation was registered at the NIH Clinical Trials Registry (Clinical Trial Registration No. NCT01134081).

Inclusion/Exclusion Criteria

Individuals were enrolled in the study if they were between 18 and 70 yrs of age and possessed bilateral recession-type defects manifested by an insufficient zone of attached gingiva ($\leq 1 \text{ mm}$), not requiring root tooth coverage. Patients were excluded: if they displayed a shallow vestibule, severe gingival recession, and tooth mobility (Grade ≥ 2); if they had any systemic conditions that could compromise wound healing and preclude periodontal surgery (i.e., diabetes mellitus, cancer, HIV, bone metabolic diseases); if they had received, within 2 mos prior to study entry, systemic corticosteroids, immunosuppressive agents, intravenous bisphosphonates, radiation therapy, and/or chemotherapy that could compromise wound healing and preclude periodontal surgery; if they had used any tobacco product within 3 mos; if they had only molar teeth suitable for soft-tissue grafting; if they had known hypersensitivity to bovine collagen; and if they had received an investigational drug or biological/ bioactive treatment within 30 days prior to study enrollment.

Wound-healing Score

During the first four-week follow-up visits, the healing condition of the soft tissues at the surgical site was examined by visual inspection. A secondary endpoint, clinical woundhealing score (CWHS), was recorded to reflect the extent of healing at the surgical site and to further evaluate the significance of angiogenic biomarkers on clinical wound healing. A wound-healing scale was created uniquely to classify the possible post-surgical healing characteristics: 0 = Mature wound healing; 1 = Erythema; 2 = Bleeding; 3 = Graft mobility; 4 = Suppuration; and 5 = Necrosis.



Figure 2. Wound fluid collection at the borders of the wound during the 4 wks post-operatively for both autograft and living cellular construct. 156 x 179 mm (300 x 300 DPI).

Periodontal Wound Fluid Collection

Prior to the surgical procedure, gingival crevicular fluid (GCF) was collected from the 3 study teeth in each participant. One tooth received the autograft, and the other received the LCC. The GCF samples were taken from the mesio-buccal site of each tooth. The area around each sample site was air-dried, and the supra-gingival plaque biofilm was removed. A methylcellulose paper strip (Periopaper[®], ProFlow Inc., Amityville, NY, USA) was gently inserted into the gingival sulcus until slight resistance was felt. The fluid sample was then collected for 30 sec, and the strips were placed in Eppendorf tubes and kept on ice.

Periodontal wound fluid was collected from the test and control graft sites at the 4 corners of the surgical site (Fig. 2). The surgical site was gently dried with gauze to remove excess saliva. Methylcellulose strips were gently inserted at the corners of the grafts. WF was collected as with GCF. Both GCF and WF samples were subsequently kept on dry ice and stored at -80°C until needed for analysis. The collection of WF occurred at wks 1, 2, 3, and 4 after surgery (Fig. 1B).

Angiogenic Biomarker Analysis

Prior to biomarker analysis, GCF and WF were thawed at RT. In addition, the 4 WF samples from each site were pooled, and proteins were eluted as previously described (Giannobile et al., 1995). Angiogenic biomarker expression was quantified with the use of a human angiogenesis custom array kit for the presence of 10 different biomarkers simultaneously according to the manufacturer's protocol (Quantibody®, Ray-Biotech, Inc., Norcross, GA, USA), which included recombinant protein standards for standard curve generation. The biomarkers selected for the array were: TIMP-1, TIMP-2, VEGF, IP-10, PDGF- BB, GM-CSF, IL-8, FGF-2, angiogenin (ANG), and angiostatin (ANT). Slides were scanned and measured for fluorescent signal intensity, and data were imported to RayBio® Antibody Array Analysis software for assessment (including data sorting, averaging signal intensities, background subtraction, normalization, and obtaining protein levels that were then exported for statistical "Statistical analysis; see below, Analysis").

Statistical Analysis

For each angiogenic biomarker, the primary outcome was the within-subject difference between the concentrations

at the autograft and LCC surgical areas. These differences were measured in each participant prior to surgery, as well as 1, 2, 3, and 4 wks after surgery, leading to a series of 5 longitudinal, within-mouth differences for each individual. For each marker, the differences at all 5 time-points were analyzed simultaneously by a linear mixed model (LMM), in which the actual marker levels were modeled as a function of time (categorical), treatment group, and the interaction of time and treatment, while accounting for the repeated measures on each individual with a random subject effect. Given that both time and treatment are categorical, our approach was a generalized version of repeatedmeasures ANOVA, since our LMM allowed for differing variability in marker levels at each time-point. Within each LMM, we also tested for a difference between the 2 treatment groups with respect to the overall time pattern in marker levels. A Bonferroni adjustment for multiple comparisons was applied to the p-values presented in Fig. 3. Statistical significance was defined as a *p*-value < 0.05.



Figure 3. Expression of angiogenic biomarkers of living cellular construct (LCC) and autograft from baseline to week 4. (**A**) Mean expression of VEGF. Inter-group analysis (*) demonstrated a statistically significant expression of VEGF for living cellular constructs at week 1 (p < 0.05). (**B**) Mean expression of IL-8. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**D**) Mean expression of FGF-2. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**D**) Mean expression of FGF-2. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**F**) Mean expression of PDGF-BB. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**F**) Mean expression of ANT. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**F**) Mean expression of ANT. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**F**) Mean expression of ANT. Inter-group analysis (*) demonstrated a statistically significant expression for living cellular constructs at week 1 (p < 0.05). (**F**) Mean expression of ANT. Inter-group analysis (*) demonstrated a statistically significant expression of TIMP-1. (**H**) Mean expression of TIMP-2. Bars indicate standard error measurements (SEM). 319 x 374 mm (300 x 300 DPI).

Characteristics	Autograft, $N = 44$	Living Cellular Construct, N = 44	P-value
Age (yrs)			
Mean (SD)		47.1 (13.1)	ND
Gender			
Male	25 (57%)		ND
Female	19 (43%)		ND
Race			
Caucasian		40	ND
African-American	1		ND
Hispanic	2		ND
Smoking status			
Non-smokers	27 (60%)		ND
Former smokers	17 (40%		ND
Mean clinical measurements (mm)			
Keratinized tissue width	1.43 ± 0.69	1.41 ± 0.72	NS
Attached gingiva width	0.08 ± 0.79	0.02 ± 0.76	NS

 Table.
 Patient Baseline Demographics

ND, not determined; NS, not significant.

RESULTS

In total, 44 individuals were evaluated in the investigation (Table). Wound-healing scores measured after surgical procedure decreased during the 4 time-points (Appendix Table). No differences were found between groups regarding wound-healing scores (Appendix Fig. 1). Hence, the expression of the angiogenic biomarkers tested did not significantly correlate with the wound-healing score (Appendix Fig. 2).

VEGF concentrations over time for the LCC and autograft groups are shown in Fig. 3A. Both groups demonstrated an increased expression of VEGF at week 1 compared with baseline, with a progressive reduction from weeks 2-4. By week 1, LCC showed a significant increase in VEGF levels compared with autograft (p < 0.05). In addition, over time, LCC demonstrated significantly higher VEGF levels compared with autograft (p < 0.05). Endogenous ANG released over time for both LCC and autograft groups is shown in Fig. 3B. ANG expression levels reached their peak expression at week 1 and progressively decreased for both the LCC and autograft groups. No statistically significant difference was noted between groups.

IL-8 expression over time for all groups is demonstrated in Fig. 3C. Both groups showed an increase in IL-8 expression by 1 wk. Further, LCC demonstrated significantly higher IL-8 expression levels compared with autograft (p < 0.05), which was also confirmed in the overall time analysis. FGF-2 WF levels are depicted in Fig. 3D. LCC demonstrated a higher expression of FGF-2 compared with autograft only at week 1 (p < 0.05). PDGF-BB released over time for all groups is demonstrated in Fig. 3E. At week 1, LCC demonstrated a significantly higher expression compared with autograft (p < 0.05), which was also noted at the overall time analysis (p < 0.05).

ANT levels over time for both groups are shown in Fig. 3F. At week 1, LCC showed an increased expression in ANT expression compared with autograft (p < 0.05). The overall time analysis also demonstrated a significantly higher expression in

favor of the LCC group (p < 0.05). TIMP-1 expression over time for all groups is shown in Fig. 3G. LCC demonstrated a modestly increased concentration of TIMP-1 at weeks 1 and 2 compared with autograft, but did not show significant difference between groups. The expression of TIMP-2 over time for both groups is illustrated in Fig. 3H. At week 1, the LCC group showed an increase in TIMP-2 levels compared with autograft, but, again, did not show significant difference between groups.

Due to minimal-no expression of GM-CSF and IP-10, no statistical analysis was performed involving these proteins (data not shown).

DISCUSSION

The oral wound-healing field has reported few studies investigating soft-tissue-engineered biomaterials for regeneration of mucosal tissue (Izumi *et al.*, 2003; McGuire and Nunn, 2005; McGuire *et al.*, 2008). However, to our knowledge, this study was the first investigation to evaluate the angiogenic biomarker profiles of autologous gingival grafts and soft-tissue-engineered constructs. The rationale for analyzing angiogenic biomarkers is that angiogenesis is an essential feature of normal wound repair, which is partly responsible for the formation of fibrovascular tissue containing fibroblasts, collagen, and blood vessels that are the hallmark of an established healing response.

During the study, VEGF was highly expressed during the first week after the surgery, and its levels progressively decreased to the fourth week of healing. Many studies have described the presence of VEGF collected directly from the surgical wound fluid.

The presence of VEGF was analyzed in surgical WF, and it was demonstrated that VEGF production and VEGF-mediated angiogenic activity would rise in the early hypoxic wound and then fall when neovascularization was complete and wound perfusion was restored (Nissen *et al.*, 1998). Additionally,

VEGF together with other angiogenic factors have also been found in the WF of oral wounds (Cooke *et al.*, 2006; Sakai *et al.*, 2006).

IL-8 is a chemokine produced by macrophages, epithelial cells, and endothelial cells. Koch and co-workers have demonstrated that IL-8 is involved in angiogenesis-dependent disorders such as rheumatoid arthritis, tumor growth, and wound repair (Koch *et al.*, 1992). IL-8 has also been reported to enhance endothelial cell proliferation, survival, and regulation of angiogenesis (Li *et al.*, 2003). In the current study, both living cellular constructs and autografts displayed higher expression of IL-8 during week 1, with higher expression for living cellular constructs over time.

Tissue inhibitors of metalloproteinases (TIMPs) have been found to have potential roles as growth factors, survival factors, growth inhibitors, and inhibitors of angiogenesis (Stetler-Stevenson and Seo, 2005; Chirco *et al.*, 2006; Stetler-Stevenson, 2008). The current investigation was able to detect the expression of both TIMP-1 and TIMP-2 during all evaluation periods, which peaked at week 1. Living cellular constructs showed an overall higher expression of both TIMP-1 and TIMP-2 compared with autograft. Osborne and Schmid evaluated the production of TIMPs by living cellular constructs and demonstrated that the induced expression of TIMP-1 and TIMP-2 may have the potential to counteract the imbalance between matrix production and degradation and thus may support wound reepithelialization (Osborne and Schmid, 2002).

It is important to note that while several angiogenic biomarkers appeared to be elevated at early time-points, this study was not able to translate this directly to angiogenesis actually occurring *in vivo*. A proxy measure of this was the erythema component of the CWHS, which was notably present during weeks 1 and 2. However, studies evaluating the early wound healing of mucosal and skin wounds demonstrated that the presence of angiogenic biomarkers was positively related to increased blood vessel formation (Szpaderska *et al.*, 2005; Kumar *et al.*, 2009).

Although the present study compared the differences in angiogenic biomarker expression between living cellular constructs and autografts, it is important to understand that both living cellular constructs and autografts heal by two different healing processes. Autografts heal by primary intention repair. Conversely, a living cellular construct is not integrated into the surrounding tissues, but rather, it modulates the healing activity of the underlying and adjacent tissue. It is postulated that the supply of live fibroblasts and keratinocytes improves the wound environment through growth factor interactions, matrix deposition and degradation, wound coverage, and a provision of responsive cells, leading to therapeutic functions (Sabolinski *et al.*, 1996).

In conclusion, the results of this study demonstrated that, during early wound-healing events, expression of angiogenicrelated biomarkers is up-regulated in sites treated with LCC compared with autogenous free gingival grafts. The use of LCC may affect the regeneration of intra-oral soft tissues or the treatment of chronic soft-tissue lesions.

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