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## **T<sub>2</sub>\* Relaxometry and Volume Predict Semi-Quantitative Histological Scoring of an ACL Bridge-enhanced Primary repair in a Porcine Model**

**Alison M. Biercevicz, BS<sup>1,4</sup>, Benedikt L. Proffen, MD<sup>2</sup>, Martha M. Murray, MD<sup>2</sup>, Edward G. Walsh, PhD<sup>3</sup>, and Braden C. Fleming, PhD<sup>1,4</sup>**

<sup>1</sup>Department of Orthopaedics, Warren Alpert Medical School, Brown University/Rhode Island Hospital, Providence RI

<sup>2</sup>Dept of Orthopaedic Surgery, Children's Hospital Boston, Boston MA

<sup>3</sup>Department of Neuroscience, Division of Biology and Medicine, Brown University, Providence, RI

<sup>4</sup>School of Engineering, Brown University, Providence, RI

### **Abstract**

Magnetic resonance imaging (MRI) variables, such as T<sub>2</sub>\* and volume, can predict the healing ligament structural properties. How these MR variables relate to semi-quantitative histology of the healing ACL is yet unknown. We hypothesized that T<sub>2</sub>\* and volume would predict the histological scoring of a healing ACL. Yucatan minipigs underwent ACL transection and received: bridge-enhanced ACL repair or no treatment. The surgical legs were harvested after 52 weeks and imaged using a high resolution 2-echo sequence. For each ligament the volume and median T<sub>2</sub>\* values were determined. The ACL specimens were then histologically analyzed using the advanced Ligament Maturity Index (LMI). The T<sub>2</sub>\* of the healing ligaments significantly predicted the Total LMI score as well as the Cell, Collagen and Vessel sub-scores; R<sup>2</sup>=0.78, 0.67, 0.65, and 0.60, respectively (p 0.001). The ligament volume also predicted the Total LMI score, Cell and Collagen sub-scores; R<sup>2</sup>=0.39, 0.33, 0.37, and 0.60, respectively (p 0.001). A lower ligament T<sub>2</sub>\* or a higher volume was associated with higher histological scores of the healing ligaments. This study provides a critical step in the development of a noninvasive method to evaluate ligament healing on a microscopic scale.

### **Keywords**

MRI; ACL; Histology; Ligament; Healing

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Corresponding Author: Braden C. Fleming, PhD, Bioengineering Labs, Coro West, Suite 404, Rhode Island Hospital, 1 Hoppin Street, Providence RI 02903; PH: 401-444-5444; FX: 401-444-4418; Braden\_Fleming@brown.edu.

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## Introduction

Current methods to evaluate ligament healing in animal models, such as histological evaluation or biomechanical testing, are crucial for tracking the healing response. However, these methods require biopsy or destructive testing and are not conducive for *in vivo* assessment. Using non-invasive MR imaging to evaluate graft maturation biomechanically, significant correlations were found between signal intensity and structural properties of the ACL reconstruction graft in an animal model.<sup>1-3</sup> The combination of both ligament/graft volume (amount of tissue) and signal intensity (tissue quality) significantly improved those predictions.<sup>1</sup> However, signal intensity is not ideal to document ligament healing as it is dependent on MR specifications, including manufacturer, acquisition parameters, and hardware effects that can vary between scanners.<sup>4</sup>

$T_2^*$  relaxation time, another MR variable, correlates with the level of tissue organization,<sup>5,6</sup> is well suited for imaging highly organized collagenous structures,<sup>5,7,8</sup> and is less sensitive to imaging parameters than signal intensity data.<sup>4</sup> Healing ligament volume, as defined by the distributions of  $T_2^*$  relaxation times within that ligament, can provide a predictive model of structural properties, where longer  $T_2^*$  times are associated with lower structural properties.<sup>9</sup> Longer  $T_2^*$  values in the meniscus were also found to be associated with worse biomechanical performance<sup>7</sup> and degeneration,<sup>8</sup> as defined by collagen content accessed via histological grading. While these findings between  $T_2^*$  relaxation time and histological outcomes in the meniscus are promising, the relationship of healing ACL  $T_2^*$  relaxation time with microscopic evidence remained unknown. More specifically, in both the research and clinical settings, it would be advantageous to understand what histological factors contribute to shorter  $T_2^*$  times in the healing ACL.

The Ligament Maturity Index (LMI) has been used to assess histological sections of the early and late stages of healing.<sup>10,11</sup> Recently the LMI scoring system was adapted for later stages in “ligamentization”, where proliferation and remodeling of the tissue are more apparent. The adapted LMI assesses a number of complex factors critical to late stage ligament healing. These factors are evaluated with metrics, which form the Cell, Collagen and Vessel sub-scores and are then combined into the Total LMI score (Table 1).<sup>11</sup>

To date there have been no studies evaluating how  $T_2^*$  relaxation time and volume of a healing ligament can predict histological scoring. The objective of this study was to test a noninvasive method for predicting the histological outcomes of a bridge-enhanced ACL repair or a naturally healing ligament in a porcine model at 52 weeks of healing using  $T_2^*$  and volume. We hypothesized that MR derived measures of  $T_2^*$  and volume would be significant predictors of the histological scoring of a healing ACL after 52 weeks of healing.

## Methods

### Animal Model

This controlled laboratory experiment was approved by the Institutional Animal Care and Use Committee. Fifteen Yucatan minipigs (approximately 15 months of age) underwent unilateral ACL transection surgery as previously described.<sup>12</sup> Immediately following

transection, eight of the animals received bridge-enhanced ACL repair with an extracellular matrix-blood composite and seven were left untreated to heal naturally without bridge-enhanced repair (ACL transection).<sup>12</sup> It should be noted that these animals were part of another study evaluating the long-term effects of the bridge-enhanced ACL repair.<sup>9,11,12</sup> At 52 weeks, all surgical knees were harvested and immediately imaged.

### MR Imaging

All image data were acquired of the surgical knees using a 3T Siemens Tim Trio scanner (Erlangen, Germany) using a 20cm volume extremity coil.  $T_2^*$  relaxation time was estimated using high-resolution 3-D gradient echo (FLASH) data sets with parameters: TR=25ms, TE=7.36, 15.24ms (2 echoes), flip angle=12°, FOV=140mm, slice thickness=0.85mm (contiguous slices), reconstruction matrix size=512x512, 3 averages, bandwidth=130Hz/pixel, and scan time=19 minutes.<sup>9</sup> The bridge-enhanced repaired and naturally healing transected ligaments were manually segmented from the MR images in both the coronal and sagittal planes using commercially available software (Mimics 13.1, Materialise, Ann Arbor, Michigan). 3-D models of the healing ligaments were then created.<sup>1</sup> The total number of ligament voxels was used to determine the volume of the whole ligament (16.1 voxels equaled one mm<sup>3</sup>).

### MR Imaging $T_2^*$ Estimation

Using previously described methods,<sup>9</sup> Matlab (R2012b; MathWorks, Natick, MA) code was used to estimate  $T_2^*$  maps using the signal intensity (SI) relationship:

$T_2^* = \left( \frac{\ln SI_1 - \ln SI_2}{TE_2 - TE_1} \right)^{-1}$ ,<sup>9,13</sup> where  $SI_1$  and  $SI_2$  are the signal intensities corresponding to the echo times  $TE_1$  and  $TE_2$  where  $TE_2 > TE_1$  for each voxel. The ligament specific maps were produced by extracting the voxels corresponding to the ligament from the  $T_2^*$  maps using the 3-D models created from the segmented images. The median  $T_2^*$  value was determined for each ligament volume and then compared to the ligament scoring metrics as described below. The distributions of  $T_2^*$  for each ligament were found to be lognormal and positively skewed. Median  $T_2^*$  was used to represent the central tendency of each ligament as differences in the median value would be representative of the entire ligament distribution.

**Histological Scoring**—Following imaging, the specimens were frozen and stored at -20 degrees Celsius until mechanical testing. The knees were then prepared and mechanically tested to determine structural properties.<sup>11,12</sup> Following biomechanical testing, knees were cut sagittally through the center of the ACL. Bone-ligament-bone sections were formalin fixed, decalcified (DELTA-Cal, Delta Products Group, Aurora, IL), and paraffin embedded. 7  $\mu$ m thick sections were cut with a microtome, placed onto custom glass slides (Corning 75x50 mm Plain Microscope Slides, Corning Incorporated, Corning, NY), and stored at 4° C until staining with hematoxylin and eosin (H&E) or  $\alpha$ -smooth muscle actin antibodies (SMA).<sup>11</sup> Cell density, morphology, as well as collagen formation (evaluated under polarized light with a 137nm wavelength filter) were assessed on the H&E slides, while SMA stained sections were used to determine vascularity.

Using the advanced LMI (Table 1), the sagittal sections from the central portions of the ligaments were scored in five regions; 1mm into the ligament from either tibial or femoral insertion site, and three regions in between. When choosing the five regions for histological analysis, the examiner selected regions that were free of synovium and not visibly deformed by biomechanical testing.<sup>11</sup> The five regions from each ligament were separately scored according to the cell, collagen and vessel criteria. The cell, collagen and vessel sub-scores for each ligament were then determined by averaging the scores for each of the five regions using the sub-scores respective criteria. The resulting cell, collagen and vessel sub-scores for each ligament were then summed to determine the total LMI score, representing the cumulative effects of healing (Table 1). All investigators were blinded to the treatment group for all histological and MRI evaluations. Additionally, the intraclass correlation coefficient between examiners using the Ligament Maturity Index has been shown to be greater than .90.<sup>14</sup>

**Data Analysis**—The repaired and naturally healing transected ligaments were grouped together and analyzed as a single data set. First, single linear regression models were used to separately test the relationships between: 1) ligament median  $T_2^*$  and histological scores and, 2) MR volume and histological scores (Total LMI, Cell sub-score, Collagen sub-score, and Vessel sub-score).

Subsequently, both ligament median  $T_2^*$  and ACL volume were included in a multiple linear regression model to predict the histological scores. For all regression equations, the R-square values and p-values for the models were reported as indicators of the strength of the relationships. Individual p-values of the independent variables in the multiple regression equation were used to assess the relative contribution of ligament median  $T_2^*$  and volume to the prediction. 95% confidence intervals (Figure 1) and standard error (SE) (Tables 2 and 3) of the regression equations were also determined as a means to assess the accuracy of the prediction equations.

## Results

The median ligament  $T_2^*$  of bridge-enhanced repaired and naturally healing transected ligaments significantly predicted Total LMI score as well as Cell, Collagen and Vessel sub-scores;  $R^2=0.78, 0.67, 0.65,$  and  $0.60,$  respectively (all  $p < 0.001$ ) (Figure 1, Table 2). The MR ligament volume was also a predictor of Total LMI score as well as Cell, and Collagen sub-scores;  $R^2=0.39, 0.33,$  and  $0.37,$  respectively ( $p=0.012, p=0.025,$  and  $p=0.016$ ) (Figure 1, Table 2). Ligament volume only approached significance for predicting the Vessel sub-score;  $R^2=0.25,$   $p\text{-value}=0.059$  (Figure 1, Table 2). In general a lower median ligament  $T_2^*$  or a higher ligament volume was associated with indications of healing or higher histological scores of the repaired and transected ligaments (Figures 1, 2, 3).

Using ligament median  $T_2^*$  in conjunction with volume, the multiple linear regression model predictions for Total LMI score as well as Cell, Collagen and Vessel sub-scores were marginally improved;  $R^2=0.82, 0.70, 0.71,$  and  $0.61,$  respectively (all  $p < 0.001$ ) (Table 3).

## Discussion

A quantitative tool for assessing the healing of an ACL post-operatively could be valuable to both researchers and clinicians. Ligament  $T_2^*$  and volume has been used to assess structural properties of a healing ligament,<sup>9</sup> but has not yet been used to quantitatively assess ligament healing via histology. In this study, the MR derived terms of median ligament  $T_2^*$  and volume significantly predicted the semi-quantitative histological scores in healing bridge-enhanced repaired and natural healing transected ligaments. In general, a shorter median  $T_2^*$  time and larger volume were associated with better histological outcome scores (i.e., improved healing) (Figure 2). As expected, the MR ligament volume was not as strongly associated with histological outcomes as median  $T_2^*$  as indicated by respective p-values and  $R^2$  values of the single regressions (Table 2). This difference in prediction strength between median  $T_2^*$  and volume is likely due to the microscopic nature of the histological assessments. Few of the histological evaluation criteria account for gross ligament size, while the majority of the criteria relate to elements of tissue organization and remodeling and it is these factors that have been found to effect  $T_2^*$  times.<sup>5,6,8</sup> Furthermore, we found that by combining the median ligament  $T_2^*$  and MR volume in a multiple regression equation (Table 3), the ability to predict the histological outcome scores was only marginally improved as observed by the small  $R^2$  increases between the single and the multiple linear regression models. This marginal improvement in the multiple regression model is due to the  $T_2^*$  term dominating the regression prediction and the volume data having a weaker association with the histological outcome criteria, which can be observed by the higher individual p-values for volume from the multiple regression analysis (Table 3).

Using ligament median  $T_2^*$  to predict the Total LMI score ( $R^2=0.78$ , p-value<0.001) was stronger than the  $T_2^*$  prediction of individual sub-scores (Cell  $R^2=0.67$ , p-value<0.001; Collagen  $R^2=0.65$ , p-value<0.001; and Vessel  $R^2=0.60$ , p-value<0.001)(Table 2). This difference in prediction strength is a result of the cumulative nature of the Total LMI score, which accounts for the complexity of the cellular, collagen and vascular aspects of the healing process for the whole ligament. Furthermore, the Total LMI assesses healing throughout five different regions of the ligament, and as such, we would expect it to relate to the median ligament  $T_2^*$ , a measure representing the whole ligament.

Our findings align with a prior MR investigation looking at signal to noise quotient (a measure of SI) in healing ACL grafts using an ovine model.<sup>3</sup> Qualitative histological assessment of graft healing was found to be present with grafts displaying a lower SI and higher structural properties.<sup>3</sup> The qualitative nature of the histological assessments in this previous study did not allow for a direct prediction analysis. Building on these results, we found that by using  $T_2^*$  relaxation time we could predict a semi-quantitative histological score (LMI) for assessing ligament healing and maturation. Additionally, the relationship between our MR variable  $T_2^*$  and histological scores tells a similar narrative to the qualitative findings of the previous ovine study. We found low  $T_2^*$  values were associated with greater cell density and collagen organization where the previous study found similar histological results for grafts with low SI.

MR imaging has also been used to find a correlation between  $T_2^*$  and level of collagen organization assessed with semi-quantitative histology in the meniscus. Short  $T_2^*$  times were identified with more densely packed collagen fibers and longer  $T_2^*$  values were associated with less densely packed collagen and meniscal damage.<sup>8</sup> While our study used a different means to acquire  $T_2^*$  times, linking collagen structure with shorter  $T_2^*$  values aligns with the findings presented here-in. This further suggests that  $T_2^*$  can serve as a proxy for tissue organization and that remodeling in healing ligaments can be identified with these MR methods.

Using only two echo times adds uncertainty to the determination of  $T_2^*$  by overestimating the relaxation time value.<sup>4,15</sup> However, a two echo estimation has been shown to correlate with a six echo determination of  $T_2^*$ , indicating the  $T_2^*$  estimation used here captures the inter-specimen relative differences.<sup>16</sup> Additionally, this approach served to limit scan time (19 min) and post-processing time, important factors to consider for eventual clinical translation.<sup>13,16,17</sup> Histological scoring was performed after a freeze-thaw cycle, to minimize potential differential effects of freezing on the histological findings of the ligaments all knees were subject to the same freeze thaw protocol. Preceding histological analysis, mechanical testing was performed.<sup>18</sup> However, a quasi-static protocol was used to test the ligaments (ramp speed 20mm/min) allowing the testing to be halted once a drop in load occurred to avoid complete disruption of the ligament and to minimize irreversible changes during the mechanical testing. All ligaments in this study failed within the midsubstance allowing histological assessment of the areas out side the area of local disruption. It is also reasonable to assume that the failure testing did not alter cellular and vascular content. For this study, we assumed that median graft  $T_2^*$  and volume were the only MR variables that would account for LMI histologic scores. It is possible that other MR analyses, such as diffusion weighted imaging to quantify fiber alignment, could improve correlations to histology. Lastly, the MR images collected in this study were collected postmortem eliminating problems with motion artifact. Future work will evaluate the efficacy of using a  $T_2^*$  imaging method to predict structural properties *in vivo* for longitudinal studies.

In this study, median ligament  $T_2^*$  and MR volume were found to be predictive of semi-quantitative histological scores assessing healing of the ACL in a porcine model at 52 weeks post operatively. This study provides a critical step in the development of a non-invasive method to predict healing on a microscopic level for bridge-enhanced repair. This technique may prove beneficial as a surrogate outcome measure for healing and may have clinical implications for tracking post-operative changes with ligament healing in patient cohort studies and randomized controlled trials.

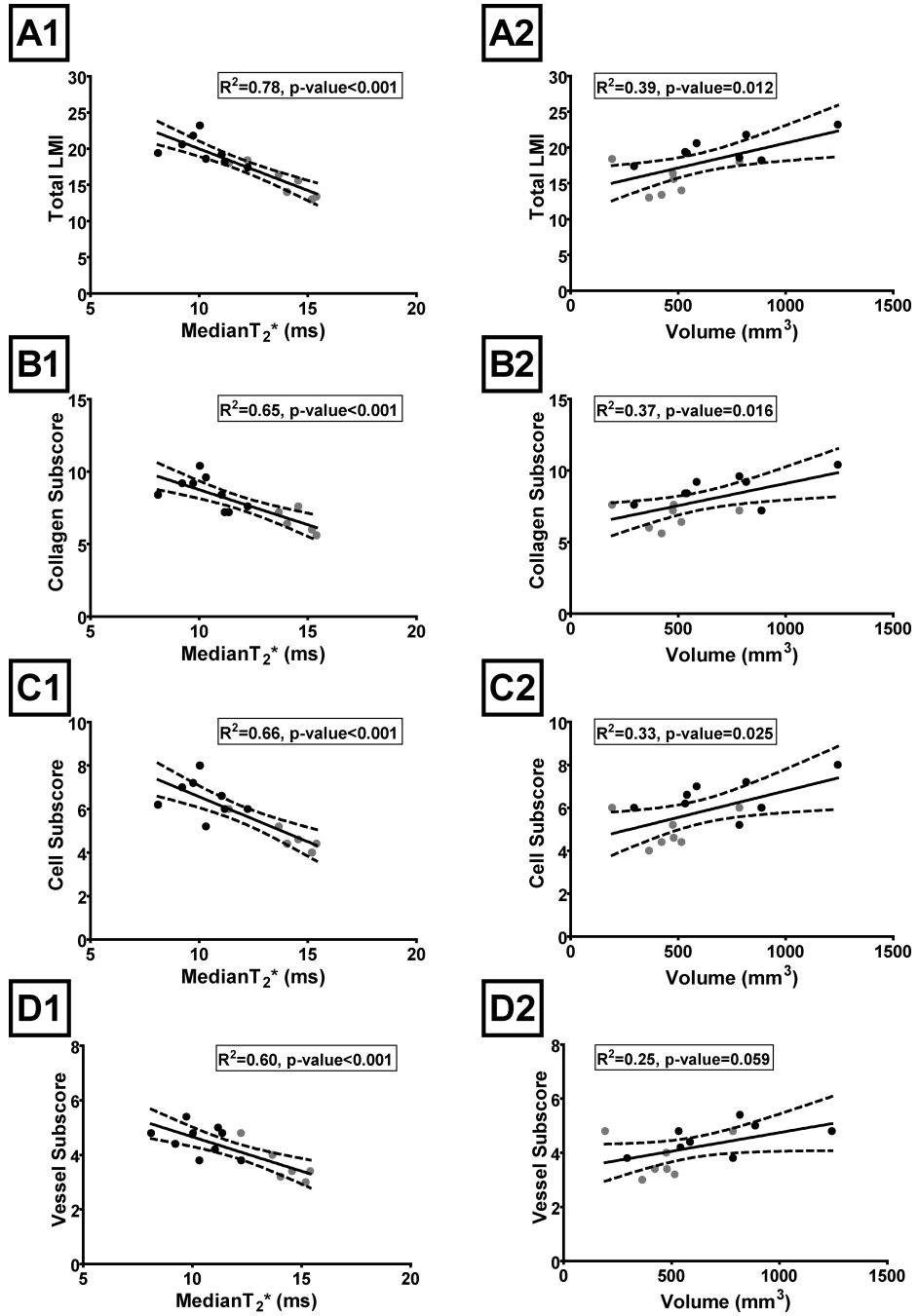
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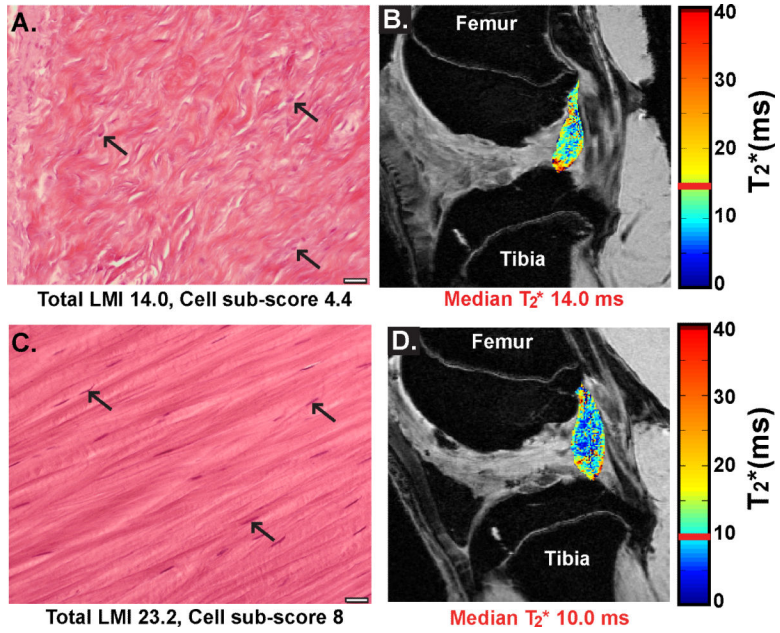
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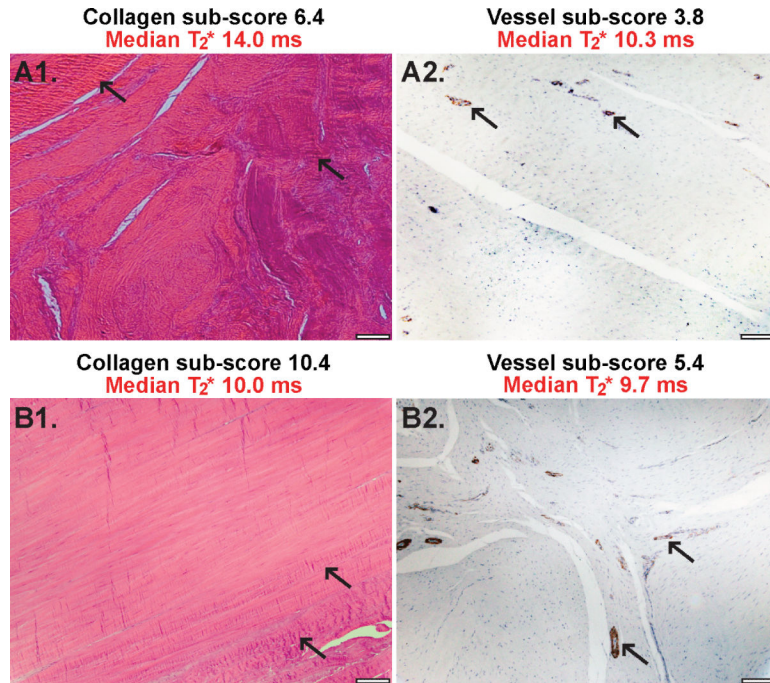


**Figure 1.** The healing ligament histology scores (A- Total LMI, B- Collagen Sub-score, C- Cell Sub-score, and D- Vessel Sub-score) as a function of ligament median T<sub>2</sub>\* value (A1, B1, C1, and D1) and volume (A2, B2, C2, and D2) in the linear regression models. Dashed lines represent 95% confidence interval. The ligaments that received bridge-enhanced ACL repair are depicted with black circles and ligaments that were transected and left to heal naturally are depicted with gray circles.





**Figure 2.**  
A) Example ligament histology image with a low total ligament score and cell sub- score (Total LMI 14.0, Cell sub-score 4.4). Arrows indicate cell nuclei not clearly aligned with longitudinal axis of collagen fibers. Also note the collagen fibers lack a distinct longitudinal axis. B) The associated T<sub>2</sub>\* ligament map for the low total LMI histology image overlaid on the original DICOM image. C) Example ligament histology image with a high total ligament score and cell sub-score (Total LMI 23.2, Cell sub-score 8). Arrows indicate cell nuclei aligned with longitudinal axis of collagen fibers. D) The associated T<sub>2</sub>\* ligament map for the high total LMI histology image overlaid on the original DICOM image. Histology images are H&E stained at 40X magnification, scale bar indicates 20µm. The color bars in the T<sub>2</sub>\* maps represent the range of T<sub>2</sub>\* values in the ligament with the median T<sub>2</sub>\* value for the ligament highlighted in red. The MR images are a sagittal view of the femoral notch with the femur at the top of the image and the tibia at the bottom. For the MR images shown TE = 7.36 ms.



**Figure 3.**

A1) Example H&E stained polarized image with a collagen sub-score of 6.4 and median ligament  $T_2^*$  of 14 ms. Arrows indicate areas with collagen crimp not distinctly aligned with fiber longitudinal axis. A2) Example SMA stained image with a vessel sub-score of 3.8 and median ligament  $T_2^*$  of 10.3 ms. Arrows indicate smooth muscle like actin rich muscularis layer around arterioles visible in the interfascicular regions. B1) Example H&E stained polarized image with a collagen sub-score of 10.4 and median ligament  $T_2^*$  of 10 ms. Arrows indicate collagen crimp aligned with fiber longitudinal axis. B2) Example SMA stained image with a vessel sub-score of 5.4 and a median ligament  $T_2^*$  of 9.7 ms. Arrows indicate smooth muscle like actin rich muscularis layer around arterioles visible in the interfascicular regions. All histological images at 10X magnification, scale bars indicate 100 $\mu$ m.

**Table 1**

Criteria used to determine the advanced Ligament Maturity Index (LMI). The five regions from each ligament were separately scored according to the cell, collagen and vessel criteria. The cell, collagen and vessel sub-scores for each ligament were then determined by averaging the scores for each of the five regions using the sub-scores respective criteria. The resulting cell, collagen and vessel sub-scores for each ligament were then summed to determine the total LMI score representing the cumulative indications of healing.

Cell Sub-Score (total = 8 pts)		Collagen Sub-Score (total = 12 pts)		Vessel Sub-Score (total = 6 pts)	
Criteria	pts	Criteria	pts	Criteria	pts
<b>Presence of inflammatory cells</b>		<b>Width of bundles</b>		<b>Density of blood vessels</b>	
Necrosis	0	No bundles	0	None present	-1
Polymorphonuclear cells	1	Width less than 50 $\mu$ m	2	more than 200% present	0
No inflammatory cells	2	Width greater than 50 $\mu$ m	4	150%-200% present	1
				Less than 150% present	2
<b>Nuclear aspect ratio (NAR) of fibroblasts</b>		<b>Bundle orientation with long axis of ligament</b>		<b>Vessel orientation with long axis of ligament</b>	
No cells	-1	No bundles	-2	No vessels oriented	-2
Average NAR less than 2 (round)	0	less than 50% oriented	0	Less than 30% oriented	-1
Average NAR 2-4	1	50%-75% oriented	2	less than 50% oriented	0
Average NAR greater than 4 (elongated)	2	75%-100% oriented	4	50%-75% oriented	1
				75%-100% oriented	2
<b>Nucleus of fibroblast aligned with fascicles and long axis of ligament</b>		<b>Collagen Crimp</b>		<b>Vessel maturity</b>	
No cells	-2	No crimp	-2	No vessels seen	0
Less than 30% of cells oriented	-1	less than 25% crimp	0	Capillaries only present	1
30%-50% oriented	0	25%-75% crimp	2	Arterioles present	2
50%-75% oriented	1	Crimp with normal length present	4		
75%-100% oriented	2				
<b>Number of fibroblasts</b>					
None	-1				
Greater than 2 $\times$ normal	0				
Between 1.5 $\times$ and 2 $\times$ normal	1				
Less than 1.5 $\times$ normal	2				

**Table 2**

Summary of the healing ligament histology score single linear regression prediction equations as a function of ligament median  $T_2^*$  value or volume.

Histology Scoring Dependant Variable	Model	MR Independent Variable	Coefficient	p-value	SE of the estimate	R <sup>2</sup>
<i>Total LMI</i>	Single Regression	<i>Median T2*</i>	-1.15	<0.001	1.4	0.78
		<i>Volume</i>	0.0069	0.012	2.3	0.39
<i>Cell Subscore</i>	Single Regression	<i>Median T2*</i>	-0.415	<0.001	0.7	0.67
		<i>Volume</i>	0.002	0.025	1	0.33
<i>Collagen Subscore</i>	Single Regression	<i>Median T2*</i>	-0.49	<0.001	0.6	0.65
		<i>Volume</i>	0.003	0.016	1.1	0.37
<i>Vessel Subscore</i>	Single Regression	<i>Median T2*</i>	-0.25	<0.001	0.6	0.60
		<i>Volume</i>	0.001	0.059	0.7	0.25

**Table 3**

Summary of the healing ligament histology score multiple linear regression prediction equations as a function of ligament median T<sub>2</sub>\* value and volume.

Histology Scoring Dependant Variable	Model	MR Independent Variable	Coefficient	Individual p-value	Regression p-value	SE of the estimate	Regression R <sup>2</sup>
<i>Total LMI</i>	<i>Multi Regress</i>	<i>Median T2*</i>	-0.99	<0.001	<0.001	0.8	0.82
		<i>Volume</i>	0.0027	0.106			
<i>Cell Subscore</i>	<i>Multi Regress</i>	<i>Median T2*</i>	-0.36	0.002	<0.001	0.7	0.70
		<i>Volume</i>	0.0009	0.254			
<i>Collagen Subscore</i>	<i>Multi Regress</i>	<i>Median T2*</i>	-0.40	0.003	<0.001	0.8	0.71
		<i>Volume</i>	0.0014	0.159			
<i>Vessel Subscore</i>	<i>Multi Regress</i>	<i>Median T2*</i>	-0.23	0.006	<0.001	0.5	0.61
		<i>Volume</i>	0.0004	0.500			