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X-ray Phase Contrast Allows Three Dimensional, Quantitative Imaging of Hydrogel Implants

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Abstract

Three dimensional imaging techniques are needed for the evaluation and assessment of biomaterials used for tissue engineering and drug delivery applications. Hydrogels are a particularly popular class of materials for medical applications but are difficult to image in tissue using most available imaging modalities. Imaging techniques based on X-ray Phase Contrast (XPC) have shown promise for tissue engineering applications due to their ability to provide image contrast based on multiple X-ray properties. In this manuscript, we investigate the use of XPC for imaging a model hydrogel and soft tissue structure. Porous fibrin loaded poly(ethylene glycol) hydrogels were synthesized and implanted in a rodent subcutaneous model. Samples were explanted and imaged with an analyzer-based XPC technique and processed and stained for histology for comparison. Both hydrogel and soft tissues structures could be identified in XPC images. Structure in skeletal muscle adjacent could be visualized and invading fibrovascular tissue could be quantified. There were no differences between invading tissue measurements from XPC and the gold-standard histology. These results provide evidence of the significant potential of techniques based on XPC for 3D imaging of hydrogel structure and local tissue response.

Key Terms

X-ray phase contrast; imaging; hydrogels; tissue engineering; micro-computed tomography

Introduction

There is a significant need for improved imaging methods for evaluating and monitoring biomaterials used in tissue engineering/regeneration, drug delivery, and cell therapies. Hydrogels are a particularly popular class of biomaterials, as they can be engineered with properties similar to soft tissues. Unfortunately, hydrogels present challenges when attempting to assess their properties in tissue. The material may be altered or lost due to

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disproportionate shrinkage during processing for histological analysis, and their high water content results in poor contrast in many non-invasive imaging modalities¹. Furthermore, histological methods are destructive and often only allow two dimensional assessment of the samples. Therefore, there is a need for the development of other methods for the assessment of hydrogel structure in tissue in order for these biomaterials to be employed for clinical therapies. The ideal method would allow for three dimensional (3D), quantitative, non-invasive evaluation of both the hydrogel structure itself and the soft tissue at the interface with the biomaterials in order to monitor both the performance of the material and repair of the damaged tissue.

A number of biomedical imaging modalities have been investigated to visualize and assess biomaterials for tissue engineering applications both *in vitro* and *in vivo*^{1, 17, 33}. These techniques have been utilized to study hydrogels with limited success. Autofluorescence exhibited by some natural^{6, 18, 42} and synthetic¹³ hydrogels allows for fluorescence imaging of structure and monitoring of degradation. However, this technique is primarily limited to *in vitro* applications. Hydrogels tagged with exogenous fluorescent makers^{5, 50} can be tracked both *in vitro* and *in vivo*, but poor tissue penetration means that these methods requires ectopic hydrogel implantation and the creation of a skin window chamber to mitigate scatter resulting from skin. Optical Coherence Tomography (OCT) has been implemented to image a variety of hydrogels for wound healing applications and has been successful for simultaneous monitoring of hydrogel degradation and skin formation^{27, 37, 41, 45}. However, the resolution of OCT decreases with increasing penetration depth limiting its broad application. Ultrasound (US) imaging has been used to quantify hydrogel volumes over several weeks *in vitro*^{16, 21} and to detect and quantify spatial difference in collagen gel microstructures³¹. US elastography can be employed to measure viscoelastic properties of degrading hydrogels *in vitro* and *in vivo*²⁵. While US methods have shown potential, they can be hindered by speckle noise and limited contrast of soft tissue formed in response to the hydrogel.

Magnetic Resonance Imaging (MRI) has also been employed to monitor hydrogels for tissue engineering and drug delivery applications. Implanted hydrogels were observed and their size quantified longitudinally with MRI^{36, 38}. MRI also allowed for the visualization of the hydrogels and changes in local tissue response to the implanted hydrogels^{11, 12, 39}. Further, MRI was used to reveal the integration of articular cartilage with a hydrogel by quantifying the width of transition and gradient between two materials in T2 weighted images³⁵. MRI has also been used to non-invasively track the degradation of hydrogels with the incorporation of contrast agents^{24, 26, 46}. New contrast mechanisms with MRI, such as chemical exchange saturation transfer MRI (CEST MRI), has enabled tracking degradation of hydrogels without the need for exogenous contrast agents²⁹. CEST MRI is a contrast mechanism for the detection of certain molecules with rapidly exchangeable protons that can be saturated with specific radiofrequency pulses⁴⁰. In Liang *et. al* the CEST MRI was limited to gelatin based hydrogels and may not correlate with structural changes to the gels but rather gelatin loss²⁹. While imaging depth is high with MRI making it better than US and optical methods, spatial resolution is on the order of hundreds of microns. Better

scanners are under development but, typically, spatial resolution is limited by signal to noise ratio, field of view, and acquisition time.

Hydrogels generate little or no contrast in absorption-based X-ray imaging techniques. However, imaging techniques based on X-ray phase contrast (XPC) have shown promise for imaging biomaterials and soft tissues with excellent depth penetration and high spatial resolution^{2, 9, 10, 22, 49}. Contrast in XPC imaging results from X-ray absorption, refraction and ultra small angle scatter (USAXS), allowing materials and tissues with low X-ray absorption properties to be visualized and discriminated without the need for exogenous contrast agents.

The advantages of XPC imaging stem from the fact that, at diagnostic X-ray energies, the refractive-index variations of tissue are generally orders of magnitude greater than variations in the X-ray attenuation coefficient²⁸. This is because refraction contrast decreases as the X-ray energy squared, whereas absorption contrast drops as the energy to the fourth power. Therefore, XPC imaging offers the potential for very-low-dose imaging⁴ by using X-ray energies that are traditionally considered too high to be useful for imaging soft tissue.

As opposed to conventional radiographic methods, XPC imaging requires the irradiating X-ray beam to possess a sufficient degree of coherence so that wave-like phenomenon such as refraction and interference can be observed. To understand the basic image formation principles of XPC imaging, consider an idealized scenario in which a monochromatic X-ray plane-wave $U_i(x,y,z)$, propagating along the z -axis, irradiates an object. The transmitted wavefield $U_t(x,y,z=0)$ on the contact plane behind the object is given by³⁴.

$$U_t(x,y;z=0)=U_i(x,y,z)\exp[j\phi(x,y) - \frac{\mu(x,y)}{2}] \equiv A(x,y)\exp[j\phi(x,y)],$$

where

$$\phi(x,y) = -\frac{2\pi}{\lambda} \int dz \delta(r) \quad \text{and} \quad \mu(x,y) = -\frac{4\pi}{\lambda} \int dz \beta(r),$$

λ is the wavelength, and $\delta(r)$ and $\beta(r)$ denote the real and imaginary components of the X-ray refractive index. The quantities $\phi(x,y)$ and $\mu(x,y)$ represent the projected phase and linear attenuation coefficient that characterize the object, respectively. Note that if the intensity $I(x,y;0)=|U_t(x,y;0)|^2$ on the contact plane were recorded, the resulting image would correspond to a conventional absorption-based radiography that is determined solely by the X-ray absorption properties.

An imaging system allows phase-contrast effects to be observed only when the transmitted X-ray wavefield $U_t(x,y;0)$ is modified in some way prior to recording the intensity of the modified wavefield. In this way, variations in $\phi(x,y)$ introduced by the object can result in variations in the recorded intensity. The variations in the recorded image that are due to

variations in $\varphi(x,y)$ are referred to as phase contrast (PC) effects. In general, the raw measured image will have a mixed contrast that arises from both PC and conventional absorption contrast. In some implementations, multiple images are acquired that correspond to different configurations of the imaging system, and then computational methods can be utilized to un-mix the contrast mechanisms and obtain separate images that contain only PC or absorption contrast. Additionally, for some implementations, a third image can be computed that contains information regarding the ultra-small-angle-scattering properties of the object.

There are four implementations of X-ray PC imaging that are under development including Propagation-Based PC Imaging, X-ray Interferometry, Differential PC Imaging using X-ray Gratings, and Analyzer Based Imaging using Crystals. In analyzer methods (Figure 1B), an X-ray beam is prepared by use of a crystal monochromator, which results in a collimated and monochromated incident wavefield. This wavefield irradiates an object and is subsequently diffracted by an analyzer crystal. Based on properties of the analyzer crystal, X-rays travelling at or near the crystal's Bragg angle are selected and ultimately passed on to the intensity detector. The analyzer crystal is rotated to different angles (on a microradian scale) and only the X-rays that satisfy Bragg's condition will reflect and be detected. Measurements are taken at several different angles of the analyzer crystal in order to generate an angular intensity curve known as a rocking curve (Figure 1C). When an object is present in the pathway, changes in this rocking curve can be detected and used to reconstruct images that represent the absorption, refraction, and USAXS properties of the object.

The main objective of this study was to investigate an analyzer-based XPC imaging technique for 3D imaging of a model hydrogel system and soft tissue structure in a model tissue engineering system. We have previously shown that this technique allows visualization of hydrogels in culture and tissue sections⁹. XPC computed tomography (CT) is expected to enable imaging of both biomaterial and tissue structures in 3D and allow for quantification of tissue features. The tissues are imaged without the introduction of exogenous contrast agents resulting in contrast based on material and tissue properties alone.

Materials and Methods

Materials

Polyethylene glycol (PEG, Mn 8000), fibrinogen from human plasma (Fg, 50–70% protein), thrombin from human plasma (Tb), 2-hydroxy-2-methylpropiophenone (Irgacure 1173), acryloyl chloride (98%), and triethylamine (TEA, 99.5%) were obtained from Sigma-Aldrich (St. Louis, MO). Sodium chloride (99.5%), dichloromethane (99.9%), and ethyl ether (anhydrous) were obtained from Fisher Scientific (Hampton, NH).

Preparation of fibrin loaded porous PEG hydrogels

The samples imaged were part of a larger study that was previously published²³. A salt leaching technique that has been previously described^{13, 14} was adapted to make porous PEG scaffolds containing Tb. Briefly, PEG-diacrylate (PEG-DA) was synthesized by reaction of PEG (Mn 8000) with acryloyl chloride and TEA. The product was purified with

dichloromethane dissolution and ethyl ether precipitation. The purified PEG-DA (250mg/ml) was then dissolved in saturated salt water, with Tb (100U/ml) and Irgacure 1173 (0.5% w/v, photoinitiator) added to form the hydrogel precursor. Two hundred microliters of the precursor was mixed with 200mg of sieved salt crystals (100–150 μ m in diameter) and polymerized under UV light (365nm) for 5 minutes. The salt crystals were then leached out in 50ml of DI water with agitation for 3 hours, replacing the DI water every hour.

Excessive water on the hydrogels surface was mostly removed by blotting with sterile gauze. The hydrogels were air-dried for 1 hour to allow evaporation of water in the pores. Three hundred microliter of Fg solution (10mg/ml, 20mg/ml or 40mg/ml) was dropped onto the porous hydrogel surface, and the solution absorbed into the pores by gravity and capillary force. The hydrogels were incubated at room temperature for 30 minutes to allow interaction of Fg with Tb to form a fibrin gel within the pores and then rinsed with DI water.

Animal model

All animal experiments were carried out at Edward Hines, Jr. VA Hospital, and the procedures were approved by the institutional animal care and use committee (IACUC). Fibrin loaded PEG hydrogels with 0, 10, 20 or 40mg/ml Fg were synthesized and prepared under sterile conditions. Polypropylene shells were fabricated in the shape of a top hat, with a height of 4 mm, a diameter of 10.5 mm, and an integral sewing ring of 3.55 mm extending from the open end of the hat.

Male Lewis rats (300g~400g) were anesthetized initially with 5% isoflurane via a nose cone and maintained at 2–4% isoflurane/35% oxygen mixture during the procedure. Their backs were shaved and skin scrubbed with isopropyl alcohol followed by a povidone-iodine antiseptic solution. A longitudinal incision was made along the spine and the skin separated using blunt dissection. Six polypropylene shells containing the hydrogels were implanted subcutaneously and secured with four evenly spaced sutures through the sewing ring for each animal. The skin incision was closed using 4-0 nylon suture after implantation. At 2 and 3 weeks after implantation the animals were euthanized with Nembutal (pentobarbital 100 mg/kg) and perfusion fixed with 4% paraformaldehyde and the implanted samples harvested. Samples were then stored in formalin prior to imaging, which has a minimal effect on tissue contrast evaluated by analyzer-based systems³².

XPC imaging - Multiple Image Radiography (MIR)

Implanted hydrogels submerged in deionized water (Figure 1A) were imaged using a MIR imaging system (Figure 1B) at the National Synchrotron Light Source at Brookhaven National Laboratory (Beamline X15A)⁴⁷. The X-ray beam produced by the synchrotron passes through a monochromator consisting of two parallel Bragg crystals to create a collimated monochromatic incident beam⁴⁷. A monochromated 20 keV beam was utilized for X-ray imaging. The detector employed was an X-ray Imager VHR 1:1, CCD (Photonic Science Limited, UK) sensor (400 \times 4008 pixels) with a detector pixel size of 9 μ m. Using a perfect flat-zone silicon [333] analyzer crystal reflection, the measurement data were acquired at 11 angular positions ranging from -4 to $+4$ μ rad to generate a rocking curve for

each pixel in the detector^{3, 8, 15}. Five hundred tomographic intensity measurements were acquired over a 180° angular range for each analyzer-crystal orientation. The acquisition time at each crystal orientation was one second making total acquisition time 12.5 hours. At each tomographic view angle, three parametric MIR images that represent projected absorption, refraction and USAXS properties of the sample were computed⁴³. From knowledge of the three MIR images computed at each tomographic view angle, a filtered back projection reconstruction method was employed to reconstruct volumetric images of the three MIR properties. Three different samples, one for each condition, representing different fibrinogen concentrations and harvest times were imaged.

Histology

Following imaging, the samples were dehydrated in a graded ethanol series (Leica TP1020) and paraffin embedded using routine histological methods. Serial sections (5 µm) were cut using a Leica RM2200 microtome and stained for hematoxylin and eosin (H&E) using standard protocols. Stained sections were imaged using an Axiovert 200 inverted microscope with 5× microscope objective (Carl Zeiss MicroImaging, Inc. Gottingen, Germany) equipped with an AxioCam MRC5 color digital camera (Carl Zeiss). The camera and computer controlled X-Y-Z stage allowed tiling of multiple images to build high resolution (1.08 µm/pixel) images of entire tissue cross sections. Axiovision 4.2 image analysis software (Carl Zeiss) allowed automated control of all aspects of acquisition and processing.

Quantification and Analysis

The depth of tissue invasion into the hydrogels was measured in Axiovision 4.2 image analysis software for all samples in both XPC refraction images and H&E images. Invasion from the underlying tissue was measured manually at ten different locations in six sections for each sample to get an average depth of tissue invasion. All statistical data are expressed as means ± standard deviations. Statistical significance in tissue invasion between X-ray and histological measurements was determined using student's t-test, with a *p* value of less than 0.05 considered significant.

Results

XPC Imaging

The PEG hydrogels were harvested from the animals with the underlying tissue and the entire sample imaged by use of the MIR method to produce three separate two dimensional projection images representing the X-ray absorption, refraction, and ultra small angle X-ray scatter (USAXS) properties of the sample (Figure 2). Upon close examination, the hydrogel and soft tissue could be identified in all three images. However, the underlying muscle could be more clearly identified in the refraction and USAXS images (Figure 2B & C). The hydrogel was difficult to identify in the absorption radiograph.

In order to further examine the tissue and hydrogel structure, the two-dimensional refraction images obtained at each tomographic view angle were used to reconstruct CT slices of the samples (Figure 3). In the volumetric refraction CT image, the hydrogel, invading

fibrovascular tissue and underlying skeletal muscle could all be clearly identified (Figure 3). Reconstructed absorption and USAXS CT images did not contain significant contrast and detail in hydrogel and soft tissue structure (data not shown). The hydrogel consisted of a speckle pattern likely resulting from the internal porous structure. Skeletal muscle structure could be easily identified in the images, with distinct edges identified at what is likely divisions between individual fascicles. The fibrovascular tissue at the interface between the hydrogel and muscle had a much more heterogeneous structure.

The Refraction CT Slices were rendered to generate 3D volumes of the samples (Figure 4 and Supplemental Video 1). The renderings enable 3D visualization of the entire sample (Figure 4A). The hydrogel could be identified due to its textured appearance. Further, the invading tissue front could be clearly seen (black arrows).

Comparison with Histology

Following XPC imaging, the samples were processed for histology and H&E stained as the gold standard method of analysis. The stained sections were matched to the CT slices for identification and confirmation of the tissue structures observed (Figure 5). XPC imaging allowed identification of similar tissue features seen in the histological images. Fascicles in the underlying muscle could be seen as striations in the XPC refraction image and the bulk structure could be matched between images. The location and bulk shape of the hydrogel was apparent in both images and the pores seen in the histology likely contributed to the texture seen in the hydrogels in the XPC images⁹.

The front edge of tissue invading into the hydrogel structure could be identified in both XPC and histology images (Figure 5). The volume of tissue regenerated is a critical issue in tissue engineering and the depth of invaded tissue represents a critical measurement directly related to the volume of tissue formed. The invasion depth was quantified from histology and XPC images (Figure 6). Both techniques resulted in data showing a general trend of increasing invasion depth with time. In addition, depth did not vary with the concentration of fibrinogen within the hydrogels regardless of analysis methods. Overall, there were no significant difference in the average invasion depths measured using XPC or the gold standard method (quantitative histomorphometry).

Discussion

There is a need for techniques that allow three-dimensional, quantitative analysis of hydrogel and soft tissue structure in tissue engineering applications. XPC imaging techniques have been shown to enable visualization of hydrogels based on differences in the X-ray refractive index values of hydrogels and tissue⁹. In this study, we have expanded on these results, showing that XPC CT imaging allows 3D visualization and quantification of hydrated soft tissues and hydrogels. The XPC images show clear distinction between underlying muscle, invading fibrovascular tissue and the hydrogel without the need for exogenous contrast agents (Figures 4 and 5). Furthermore, when matched to histology, XPC images demonstrated comparable details in both hydrogel and soft tissue structure.

Hydrogels were implanted with varying concentrations of fibrinogen in the pores and harvested at two different time points in order to investigate the role of composition and time on tissue response. The depth of tissue invasion is an important characteristic that relates to the overall volume of engineered tissue generated, which is a critical aspect related to the potential success of a strategy. Depth could be identified and measured in all the samples from XPC refraction data and the results were not different from values measured in corresponding histology. The average depth of tissue invasion measured from the XPC images did not depend on fibrinogen concentration or time for these samples. However, in this study, only a subset of explanted samples from the full larger study were imaged as a proof of concept showing the potential of XPC as a tool for studying hydrogels in tissue engineering applications. Regardless, these findings suggest that the XPC technique would have led to the same conclusions in regards to tissue invasion as previously presented in Jiang et al.²³. However, more samples would be needed to provide sufficient power for comparing the different groups as performed with histology in the previous work. Regardless, our results show that XPC can allow analysis comparable to histology but with the added benefit of 3D imaging capabilities and without sectioning and staining the samples.

While the imaging was performed on *ex vivo* samples at a synchrotron facility, efforts are underway for developing in-lab small animal XPC imagers that could lead to longitudinal *in vivo* XPC imaging^{7, 30, 44, 48}. Our findings, combined with studies showing *in vitro* imaging of hydrogel samples² and full live mice⁷ with an in-lab XPC system indicate the significant potential for imaging of hydrogels in animals with micro-focus X-ray tube. Depending on the implementation of the in-lab XPC system, the polychromatic nature of the micro-focus X-ray tube can be overcome. With the in-lab analyzer based system a monochromator is utilized to filter the beam⁴⁸ while for a grating-based system, as source grating can be used to create the necessary degree of coherence⁷. Additionally, a number of other technical issues will need to be overcome. Specifically, optimized data-acquisition strategies need to be coupled with advanced image reconstruction algorithms in order to overcome the long data-acquisition times and relatively high radiation doses currently required for small animal imaging. Recent results in these areas provide enthusiasm for this possibility^{19, 20}.

PEG hydrogels are being explored for a broad range of biomedical applications. The ability to evaluate the structure of the hydrogels in tissue is critical to evaluating and optimizing their performance. While the PEG hydrogels used in this study were not degradable, in most applications they are made degradable by the introduction of sequences susceptible to hydrolysis or enzymatic cleavage within network crosslinks. Based on their hydrogel contrast within the refraction images observed in this study, it may be possible to track the degradation of degradable PEG hydrogels longitudinally by quantifying overall volume or based on changes in the internal structure.

XPC images also revealed important tissue detail in the underlying skeletal muscle. Detail in the hierarchical structure could be clearly identified, including boundaries between fascicles. This structure is critical to the function of skeletal muscle and may be lost in traumatic or degenerative disease. XPC could be used to identify damaged muscle tissue as well as to evaluate muscle tissue engineering strategies.

The results presented here establish that XPC imaging can reveal features of PEG hydrogels and soft tissues. XPC CT allowed for 3D visualization of hydrogels and quantification of soft tissue invasion. These results suggests that XPC could further benefit tissue engineering by allowing imaging of hydrogels in engineered tissues and monitoring of the local, and integrated, soft tissue response.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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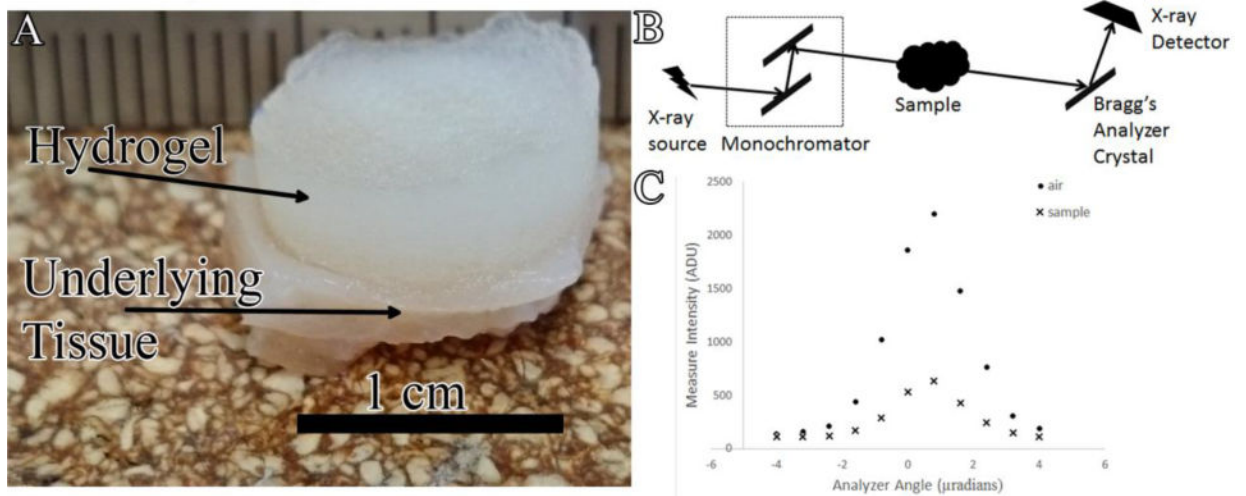


Figure 1.

A) Photograph of explanted hydrogel-tissue sample B) Schematic of MIR imaging system at the NSLS X-ray beam produced by synchrotron is filtered to a single energy through monochromator and passed through the sample X-rays travelling at or near the Bragg's angle of the analyzer crystal are reflected up to the X-ray detector C) Example of rocking curves acquired in the absence (●) and presence (x) of hydrogel sample.

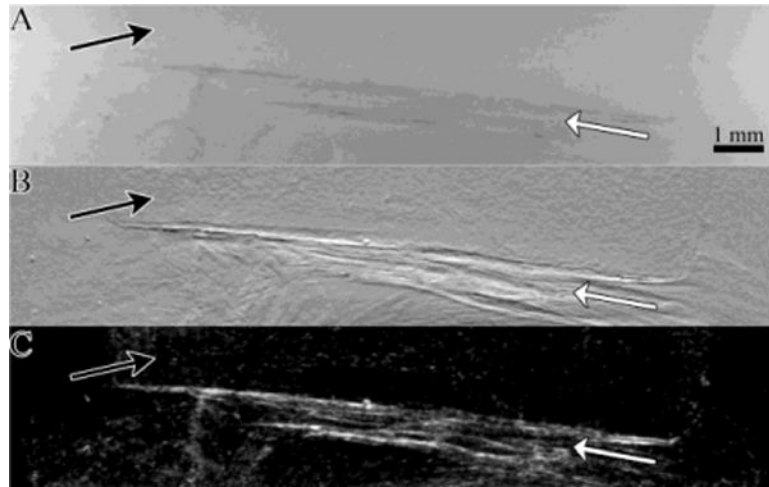


Figure 2.

A) Absorption b) Refraction and c) USAXS images of a 1 week explanted hydrogel-tissue sample generated using MIR. The underlying tissue (white arrows) produces contrast in all three X-ray property images. The hydrogel (black arrows) produces the greatest contrast in the refraction image.

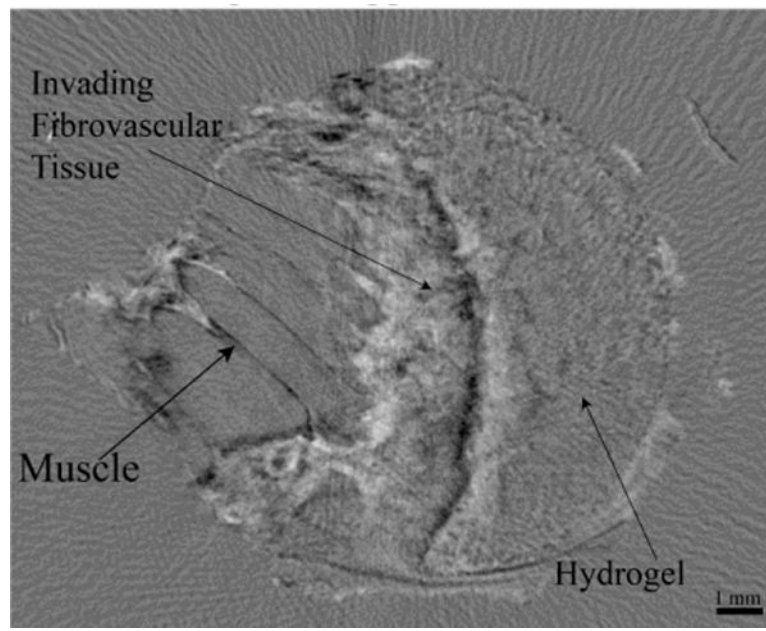


Figure 3. Refraction XPC CT slice of a hydrogel-tissue sample. The hydrogel, fibrovascular tissue, and underlying skeletal muscle can all be clearly identified.

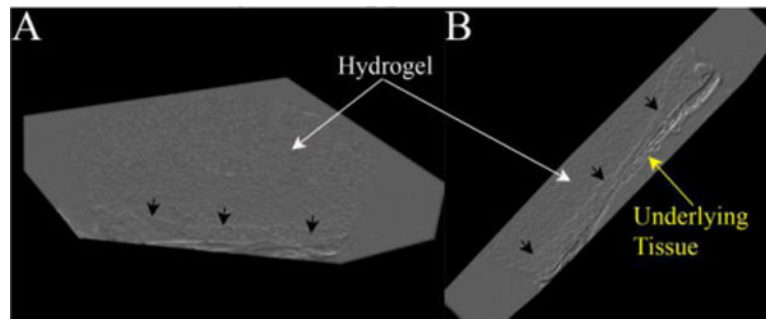


Figure 4. Volume renderings of a hydrogel-tissue sample displaying the 3D structure of the hydrogel (white arrows), invading tissue front (black arrows) and underlying muscle (yellow arrow).

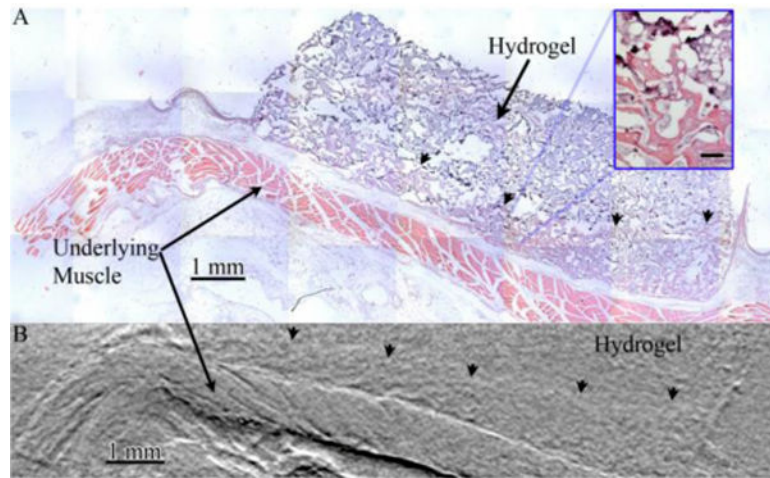


Figure 5. Images of a hydrogel-tissue sample acquired via A) sectioning of the sample followed by H&E staining and widefield optical microscopy and B) digital slices of an XPC refraction CT of whole sample volumes. Structure in the underlying muscle is visible in both images as is the front of invading fibrovascular tissue (arrowheads). Inset displayed magnified view of hydrogel (dark purple) and invading fibrovascular tissue. Scalebar in inset is 100 μm.

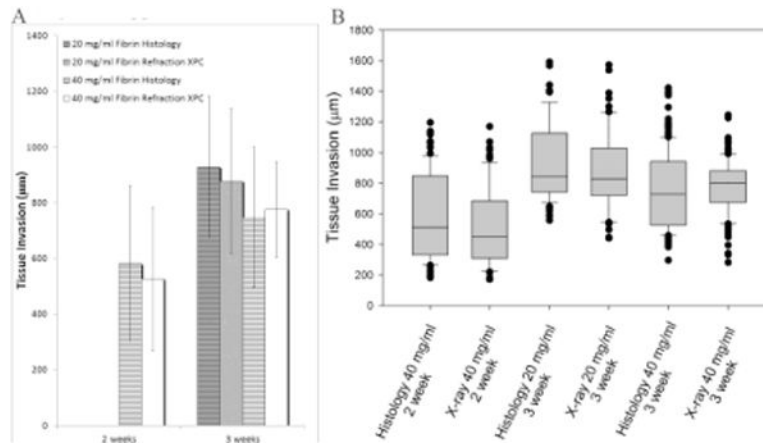


Figure 6.

A) Bar graph displaying the mean and standard deviation of tissue invasion depth measured from both histology (striped) and XPC (solid) images B) Box plot displaying median tissue invasion depth along with 10th, 25th, 75th, and 90th percentiles as vertical boxes with error bars for both histology and XPC images XPC provides similar results to the gold standard histological techniques with more precise measurement and less variations.