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Perfusion estimation using contrast enhanced three-dimensional subharmonic ultrasound imaging: an *in vivo* study

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Abstract

Objectives—The ability to estimate tissue perfusion (in mL/min/g) *in vivo* using contrast-enhanced three-dimensional (3D) harmonic and subharmonic ultrasound imaging was investigated.

Materials and Methods—A Logiq 9 scanner (GE Healthcare, Milwaukee, WI) equipped with a 4D10L probe was modified to perform 3D harmonic imaging (HI; $f_{\text{transmit}} = 5$ MHz and $f_{\text{receive}} = 10$ MHz) and subharmonic imaging (SHI; $f_{\text{transmit}} = 5.8$ MHz and $f_{\text{receive}} = 2.9$ MHz). *In vivo* imaging was performed in the lower pole of both kidneys in five open-abdomen canines after injection of the ultrasound contrast agent (UCA) Definity (Lantheus Medical Imaging, N Billerica, MA). The canines received a 5 $\mu\text{L}/\text{kg}$ bolus injection of Definity for HI and a 20 $\mu\text{L}/\text{kg}$ bolus for SHI in triplicate for each kidney. Ultrasound data acquisition was started just prior to injection of UCA (in order to capture the wash-in) and continued until washout. A microvascular staining technique based on stable (non-radioactive) isotope-labeled microspheres (Biophysics Assay Laboratory Inc, Worcester, MA) was used to quantify the degree of perfusion in each kidney (the reference standard). Ligating a surgically exposed branch of the renal arteries induced lower perfusion rates. This was followed by additional contrast-enhanced imaging and microsphere injections to measure post-ligation perfusion. Slice data were extracted from the 3D ultrasound volumes and used to generate time-intensity curves off-line in the regions corresponding to the tissue samples used for microvascular staining. The mid-line plane was also selected from the 3D volume (as a quasi-2D image) and compared to the 3D imaging modes. Perfusion was estimated from the initial slope of the fractional blood volume uptake (for both HI and SHI) and compared to the reference standard using linear regression analysis.

Results—Both 3D HI and SHI were able to provide visualization of flow and, thus, perfusion in the kidneys. However, SHI provided near complete tissue suppression and improved visualization of UCA flow. Microsphere perfusion data were available for 4 canines (1 was excluded because of

an error with the reference blood sample) and showed an average perfusion of 9.30 ± 6.60 and 5.15 ± 3.42 mL/min/g pre and post ligation, respectively. The reference standard showed significant correlation with overall 3D HI perfusion estimates ($r=0.38$; $P=0.007$), but correlated more strongly with 3D SHI ($r=0.62$; $P<0.001$). Additionally, these results showed an improvement over the quasi-2D HI and SHI perfusion estimates ($r=-0.05$ and $r=0.14$) and 2D SHI perfusion estimates previously reported by our group ($r=0.57$)

Conclusions—In this preliminary study, 3D contrast-enhanced nonlinear ultrasound was able to quantify perfusion *in vivo*. 3D SHI resulted in better overall agreement with the reference standard than 3D HI and was superior to previously reported 2D SHI results. 3D SHI outperforms the other methods for estimating blood perfusion, due to the improved visualization of the complete perfused vascular networks.

Keywords

Subharmonic imaging; harmonic imaging; 3D ultrasound; ultrasound contrast agents; perfusion estimation

Introduction

The ability to quantify blood flow in tissue, organs and tumors can provide information necessary to diagnose abnormalities or monitor treatment. It has been shown that by combining ultrasound imaging with microbubble based contrast agents it possible to increase the sensitivity to blood flow in both macro and microvasculature¹⁻³. Hence, contrast-enhanced ultrasound imaging has been used widely as a tool for blood perfusion imaging⁴⁻¹⁴. Renal quantitative perfusion imaging using both wash-in and destruction-reperfusion of ultrasound contrast agents (UCA) has been able to produce good correlation ($r>0.9$) with absolute perfusion^{4, 5}. Perfusion derived parameters such as time-to-peak, peak-intensity, wash-in slope and mean transit time also have shown promise in identifying altered hepatic hemodynamics⁷ as well as in microvascular applications monitoring antiangiogenic treatment of tumors^{9-11, 13}. Determination of thresholds for identification of critical changes in perfusion and development of parametric perfusion maps in cases of acute ischemic stroke using contrast-enhanced ultrasound have also been studied^{12, 15}. Parameters (such as arterial liver perfusion, portal-venous perfusion) derived from time-intensity curves using curve-fitting techniques for studying malignant liver tumors have shown moderate correlation with dynamic contrast-enhanced computed tomography (CT) in terms of quantifying perfusion⁸. Also, the use of UCA wash-out (presence and timing) in the differential diagnosis of hypervascular liver masses has proven to be feasible⁶. Most perfusion studies to date have been performed using conventional two-dimensional (2D) ultrasound (i.e., they are limited to a single scanning plane)⁴⁻¹⁴. By limiting the measurement to a single plane, it may be difficult to accurately gauge a volumetric parameter such as blood perfusion. Using a 3D ultrasound transducer, assessment of volumetric data is possible.

Under sufficient acoustic excitation and correct transmit frequency, UCAs are able to act as nonlinear scatters, producing a wide range of frequency components in the received spectra¹⁶. These frequency components range from the subharmonic ($f_0/2$) to higher harmonics ($n \cdot f_0$) of the transmitted frequency (f_0). Selectively receiving at a specific harmonic allows for improved contrast detection by substantially reducing linear echoes produced by tissue^{14, 17-23}. One such contrast specific imaging mode that has now been implemented on most commercial ultrasound scanners is the second harmonic or harmonic imaging (HI) mode. However, HI suffers from low contrast to tissue ratio due to second harmonic generation in the surrounding tissue. Generation of the subharmonic frequency component is UCA specific. By receiving at the subharmonic frequency ($f_0/2$) it is possible

to achieve near complete tissue suppression²⁴. The feasibility of subharmonic imaging (SHI) has been shown by both our group^{24–27} and others^{28–31}. In a previous study by our group, a modified Logiq 9 scanner with a 4D10L probe and capability to perform 3D SHI was validated *in vitro* and *in vivo*³². In this study, we hypothesized that a combination of 3D imaging and SHI would be able to estimate perfusion (in mL/min/g) *in vivo* and perform better than 3D HI as well as previously reported 2D perfusion estimates.

Materials and Methods

Animal Preparation

All animal studies were performed in accordance with National Institute of Health guidelines for use of laboratory animals and approved by our institution's Animal Use and Care Committee. *In vivo* studies were performed on five canines (mean weight, 21.4±1.1 kg) sedated with an intravenous injection of propofol (Abbot Laboratories, Chicago, IL, USA; dose 7 mL/kg). The canines were placed on a warming blanket to maintain body temperature. A facemask with 4 to 5% isoflurane (Isothesia; Abbot Laboratories, North Chicago, IL) was used for introduction of anesthesia, which was maintained with 0.5 to 2% isoflurane during the entire procedure. A 4F intravascular catheter (for administration of microspheres) was introduced into the left ventricle of the heart through the carotid artery and another 5F intravascular catheter (for collecting the reference blood sample) was inserted into the aorta (above the level of the renal artery) through a femoral artery. The intravascular catheters were positioned under ultrasound guidance. Finally, an 18-gauge angiocatheter (for administration of UCA) was placed in left forelimb peripheral vein. A mid-line abdominal incision was made to provide direct placement of the probe on the kidney.

Ultrasound Imaging Setup

A Logiq 9 US scanner (GE Healthcare, Milwaukee, WI) equipped with a 4D10L probe (Footprint, 50 X 58 mm; bandwidth, 3.5–11.9 MHz; field of view, 37.4 mm) was modified to perform 3D HI ($f_{\text{transmit}} = 5$ MHz and $f_{\text{receive}} = 10$ MHz; two cycle transmit pulses) and 3D SHI ($f_{\text{transmit}} = 5.8$ MHz and $f_{\text{receive}} = 2.9$ MHz; four cycle transmit pulses) based on results from a prior study³². Pulse inversion was implemented for both imaging modes to suppress linear signal components from the received signals. The acoustic output of the system was previously measured for these configurations and found to have a peak mechanical index of 0.33 and 0.36 at the focus for HI and SHI (at maximum transmit settings) respectively³².

In Vivo Harmonic and Subharmonic Imaging

In vivo imaging was performed in the lower pole of each kidney. A sterile transducer cover (Civco Medical Instruments, Kalona, IA) was used to cover the ultrasound probe while imaging. Initially, a baseline image was acquired in conventional B-mode to determine position and orientation of transducer. An acoustic output of 12% and gain of 20 dB were selected as optimum settings based on earlier qualitative observations of trial injections in both HI and SHI modes. These settings allowed minimization of tissue signals without suppressing contrast agent visualization. All system settings were maintained constant for both HI and SHI to allow direct comparison of their performance. In order to acquire 3D US scan volumes, a region-of-interest (ROI) was selected in B-mode. The size and position of the ROI was the same for HI and SHI. The size and volume angle of the ROI determined the volume acquisition rate (V_R ; 1.6–3.0 Hz). After the ROI was selected, 3D HI and SHI were performed.

The canines received a weight-based bolus injection of Definity (Lantheus Medical Imaging, North Billerica, MA) followed by a 5 mL saline flush through the 18-gauge catheter placed in the forelimb peripheral vein. Bolus injection dosage of 5 $\mu\text{L}/\text{kg}$ and 20 $\mu\text{L}/\text{kg}$ were used for HI and SHI respectively. These ultrasound contrast agent concentrations have been identified as optimal based on our previous experiences with each of the imaging modes^{26, 32, 33}. Wilson et al³⁴ also used similar concentrations for their study comparing diagnosis of focal liver masses between contrast enhanced and unenhanced ultrasound. The differences in ultrasound contrast agent concentrations used for each imaging mode are attributed to the ability of HI to more easily detect ultrasound contrast agent within the microvasculature due to higher resolution at increased frequencies and weaker signals associated with subharmonic emissions. Image acquisition was started prior to injection of contrast agent and recorded until UCA washout. Harmonic imaging and SHI were performed in triplicate for each kidney with an interval of 15 minutes or more between injections to avoid any cumulative effect of the UCA and allow for a return to baseline conditions. After the first set of 12 injections (3 per imaging mode per kidney), a microvascular staining technique^{35, 36} based on stable (nonradioactive), isotope labeled microspheres (Biophysics Assay Laboratory [BioPAL] Inc, Worcester, MA) was used as a reference standard to quantify the degree of perfusion. The microspheres are 15 μm in diameter and are suspended in normal saline containing Tween 80 and 0.01% Thimerosal. For this study approximately 10 million microspheres (gold was used for pre-ligation; concentration, 2.5 million spheres/mL) were injected through the 4F catheter in to the left ventricular chamber. Immediately after microspheres injection, a reference blood sample was drawn from the 5F catheter into a 20 mL syringe using an automatic withdrawal pump (Syringe Pump 11 Plus; Harvard Apparatus, Holliston, MA). The withdrawal rate was set to 10 mL/min, and blood was collected for a 1-minute period, resulting in a 10 mL reference blood sample.

Next, ischemic areas of the renal tissue were created by ligating a surgically exposed branch of the renal artery feeding the lower pole of the kidney. These low perfusion areas were imaged for 6 contrast injections (3 HI and 3 SHI). This was followed by an injection of a different isotope-labeled microsphere (either lutetium or samarium) and withdrawal of a 10 mL reference blood sample. This process was repeated for the other kidney. No attempts were made to standardize the ligations, because the intent was to produce a range of different perfusion stages. After completion of the experiments, the dogs were euthanized by an intravenous injection of Beuthanasia (0.25 mg/kg; Schering-Plough Animal Health Corporation, Summit, NJ). After euthanasia, the kidneys were harvested and 1 mL of methylene blue dye was injected through the main renal artery to identify areas of infarction/ ischemia (see Fig 1). The kidney was then cut in half, corresponding to the imaging plane, and 4 small sections (approximately $1 \times 1 \times 1 \text{ cm}^3$; arterial-lateral, arterial-medial; posterial-lateral and posterial-medial) of tissue from the lower pole (scanning region) were extracted. All tissue samples were weighed (average tissue sample weight: $0.90 \pm 0.20 \text{ g}$) and stored in pre-weighed sample vials. The tissue and blood samples were stored in an oven (Imperial III incubator, Model No 305, Lab-Line Instruments Inc, Melrose Park, Illinois) at 50° C for about 12 hours and then sent to BioPAL's facilities for activation and analysis. Results of the assay were reported as the microsphere concentration of each tissue sample normalized to the microsphere concentration in the reference blood sample, which provides a measure of absolute perfusion (Q_P , in mL/min/g) according to

$$Q_P = \frac{\sum \text{spheres in tissue} / \sum \text{spheres in reference}}{\text{weight of tissue}} \times Q_R \quad (1)$$

where, Q_R is the reference withdrawal rate (in mL/min)²⁵.

Image Processing and Analysis

The 3D digital clips (stored as DICOM files) obtained from each injection were transferred from the scanner to a desktop computer (Dell Optiplex 960, Dell Inc, Austin, Texas) for offline analysis. Volume information was extracted from the DICOM files into MATLAB (Version 2012a, Mathworks Inc, Natick, MA) using custom-made GE (GE Global Research, Niskayuna, NY) extraction software. Acquired volumes contained about 45–55 image slices/volume over a 19° volume angle. All image processing was performed using the Image Processing Toolbox in MATLAB. To standardize image processing, all HI and SHI volumes were color scaled to the same level.

For each injection, the average image intensity per volume (which is the sum of the image intensity across all slices per volume divided by the number of slices in the volume) was calculated across the entire time sequence. These averaged intensities were used to generate a time-intensity curve. Figure 2 shows a time-intensity curve (normalized to the maximum intensity) generated for a single injection pre-ligation. Based on the time-intensity curve, the time point of peak contrast wash-in (T_{pk}) was determined. The individual slice intensities of the volume corresponding to T_{pk} were plotted as a distribution across the volume with reference to the midline plane. Figure 3 shows the slice intensity distribution from the lateral to the medial sections of the volume corresponding to T_{pk} identified in figure 2. To correlate with the lateral and medial tissue samples previously obtained, the peak slice (± 3 slices) in both the lateral and medial parts of the slice intensity distribution were selected. The selected slice groups were isolated from the rest of the volume and used for all further image processing. The image intensity of the isolated volumes was averaged across the slices and a cine loop over the entire time sequence was generated. Next, a radiologist was asked to select ROIs (5×5 pixels) corresponding to the tissue samples (arterial-lateral, arterial-medial, posterial-lateral and posterial-medial) from the respective (medial or lateral) cine loops. The average ROI image intensity was used to generate a time-intensity curve. A 5th order moving average filter was applied to the time-intensity curve to reduce noise induced by changes in speckle pattern arising from slight variations in the scan plane with respiratory and cardiac motion (see Figure 4). Baseline intensity was determined by averaging the first ten time-points on the curve.

Assuming that the total harmonic or subharmonic backscattering cross section is the sum of the subharmonic cross sections of all individual bubbles in the volume of interest at low bubble concentrations³⁷, a quantifiable parameter termed the fractional blood volume (FBV) similar to the definition of the fractional moving blood volume^{38–40} can be defined. The FBV in an ROI can be estimated as the contrast signal intensities from the blood normalized by the intensity of 100% blood (i.e., a normal blood vessel) obtained at the same depth and with overlying tissues^{38, 39}. Perfusion was estimated from the initial slope of the FBV uptake (rFBV) as

$$rFBV = \frac{dFBV}{dt} \quad (2)$$

where the initial slope was defined as the slope from 10% to 30% above baseline (scaling relative to the difference between baseline and peak enhancement). Perfusion estimation from the initial slope of the time-intensity curve was used based on experience from a previous 2D perfusion study²⁵. The perfusion estimates were averaged over 3 injections to obtain the final result. The variability of the measurements was assessed as the standard deviation divided by the mean.

In order to compare the performance of the 3D ultrasound modes to 2D, the midline plane of each volume was isolated to generate a quasi-2D image sequence for each injection. The ROIs previously selected for the 3D volumes were applied to the 2D image sequence. However, since it is not possible to differentiate between medial and lateral sections (unless scanned separately) in 2D, individual anterior and posterior ROI were selected and compared to the average of the medial and lateral perfusion values obtained from the tissue samples for their respective anterior and posterior regions.

Statistical Analysis

The 3D HI and SHI perfusion estimates were compared to the microsphere perfusion results and to each other using linear regression. The Pearson's correlation coefficients along with the RMSE were calculated for the entire data set for both HI and SHI. Perfusion data were further analyzed in subsets of location (anterior and posterior) and perfusion state (pre and post ligation). All four combinations of location and state were also compared to the reference standard. A two-way analysis of variance using the mixed effects model was used to evaluate the effect of biological variability (dog, location within the kidneys and perfusion state) on the perfusion estimates. All statistical analyses were performed using SPSS 20 (IBM Corporation, Armonk, New York) with $P < 0.05$ considered as statistically significant.

Results

Both 3D HI and 3D SHI were able to provide excellent visualization of vascularity within the lower pole of the kidney after administration of UCA. However, it was evident that the ability to delineate the vessel from surrounding tissue was better in 3D SHI due to almost complete tissue suppression. The 3D renderings were able to show vascularity in fine detail and the ability to freely rotate on any of the three-axes of imaging allowed for visualization of vasculature, which would otherwise not be appreciated in 2D (as shown in Figure 5). In Figure 5 there is noticeable reduction in contrast enhancement post-ligation. A Doppler overlay of the scanning region confirms the reduction of blood flow and hence reduction in contrast perfusion (Figure 6).

A total of 108 UCA injections were administered throughout the study (in one canine scanning was performed only on a single kidney). The data from one canine was discarded due to a problem with the reference blood sample acquired for microvascular staining. The remaining data yielded 336 time-intensity curves (for 3D HI and SHI) based on ROIs selected in the anterior, posterior, medial and lateral regions of the kidney. After averaging, a total of 56 perfusion estimates were available for each mode i.e., HI and SHI. However, 4 perfusion estimates were removed from HI due lack of information owing to shadowing artifacts in the posterior region of the kidney and 2 perfusion results from the microvascular staining technique yielded no value. In order to allow for direct comparison of performance, all 6 data points were removed from the correlations. The variability of the perfusion estimates was on average 0.456 (range of 0.364 – 0.657) for 3D HI and 0.488 (range of 0.306 – 0.681) for 3D SHI.

Overall (i.e., pre and post-ligation for anterior and posterior), both 3D HI ($r=0.38$; $P=0.007$) and 3D SHI ($r=0.62$; $P<0.001$) showed significant correlation with the microsphere reference standard and with each other ($r=0.73$; $P<0.001$). However, when analyzed in subsets of perfusion state and location, 3D SHI had better correlation with the reference standard compared to 3D HI in all cases except one (post-ligation posterior) as shown in Table 1. Harmonic imaging did not show significant correlation with the reference standard in the subsets of pre-ligation, pre-ligation anterior, pre-ligation posterior and post-ligation posterior data. Subharmonic imaging also showed no significant correlation in the subsets of

post-ligation posterior and pre-ligation anterior, but correlated significantly in the remaining subsets. When present in sufficient concentrations shadowing artifacts from overlying contrast agent usually impacts posterior regions of scanning. It is therefore not surprising that ligation induced low perfusion states combined with the posterior scanning plane produced greater variability in the results. However, it is surprising that the pre-ligation anterior subset does not show significant correlation in both the 3D SHI and 3D HI perfusion estimates. The best 3D SHI perfusion estimates occurred in the subsets of posterior and pre-ligation posterior ($r=0.79$; $P<0.001$). The overall perfusion data set and the best linear fit of data for 3D HI (regression line: $Q_p = 0.20 rFBV + 4.25$) and 3D SHI (regression line: $Q_p = 0.36 rFBV + 3.82$) and their corresponding RMSE are shown in Figure 7. Analysis of variance was performed only on the overall perfusion data set. Results showed that there was no significant effect of canine (HI: $P=0.60$ and SHI: $P=0.94$), and that there were no significant interactions between location and perfusion state (HI: $P=0.19$ and SHI: $P=0.11$). Significant interaction was observed between location, perfusion state and canine for 3D SHI ($P=0.007$) but not for 3D HI ($P=0.10$).

Overall quasi-2D perfusion estimates did not show any significant correlations (HI: $r = -0.05$, $P=0.824$ and SHI: $r=0.14$, $P=0.5$) with the microsphere perfusion results. Correlation analysis for the subset of location and perfusion states did not show any significant correlation with the reference standard (see Table 2) except for SHI post-ligation in the posterior location ($r=0.90$, $P=0.04$) albeit based on just 5 data points.

Discussion

Perfusion estimates based on the initial slope of the UCA time-intensity curve from volumetric data obtained using HI and SHI have been evaluated *in vivo*. Overall perfusion estimates showed significant correlation (3D HI: $r=0.38$; $P=0.007$ and 3D SHI: $r=0.62$; $P<0.001$) with absolute perfusion measurements obtained using the reference standard (isotope labeled microspheres) and with each other ($r=0.73$; $P<0.001$). However, 3D SHI resulted in better correlation with reference standard as compared to 3D HI in all subsets of location and perfusion state except for one case (post-ligation posterior). Analysis of perfusion estimates by subsets of location and perfusion state showed the best correlations with the reference standard in post-ligation for 3D HI ($r=0.66$, $P=0.001$), posterior and pre-ligation posterior for 3D SHI ($r=0.79$, $P<0.001$). This is an improvement over the 2D SHI perfusion estimation study, which only favored high perfusion rates in the anterior of the kidneys ($r=0.73$)²⁵.

Quasi-2D estimates were obtained by selecting the midline plane of the volumetric data and these did not exhibit agreement with the reference standard (HI: $r = -0.05$, $P=0.82$ and SHI: $r=0.14$, $P=0.50$). The 2D data was extracted from a single plane of an entire 3D volume and correlated to an approximation of the perfusion (average of medial and lateral perfusion used for correlation). Ideally, imaging performed using a 2D probe along the medial and lateral sections of the kidney would provide better perfusion estimates to compare with the 3D estimates. A previous study by our group to estimate perfusion *in vivo* using 2D SHI (in the medial and lateral plane) produced a good overall correlation ($r=0.57$) with the reference standard (also microsphere based perfusion values) and maximum correlation ($r=0.73$) in the pre-ligation anterior subset²⁵.

The importance of volumetric imaging for quantitative perfusion measurements was highlighted in a study that compared the errors in 2D perfusion estimates with changes in transducer positioning⁴¹. The repeatability of 2D and 3D perfusion estimates was evaluated and showed that 3D imaging was able to produce a standard deviation of less than 3% of the mean as compared to 10% (as high as 22%) for 2D. In a similar study, millimeter-sized

deviations in transducer position produced errors between 6.4 to 40.3% among perfusion estimators used for generating quantitative maps of tumor vascularity⁴².

A study by Paltiel et al⁴³, evaluated the ability to quantify testicular parenchymal perfusion in an *in vivo* model of unilateral testicular torsion using volumetric contrast-enhanced ultrasound. Their perfusion estimates showed excellent correlation (up to $r=0.90$) with absolute perfusion rates (microspheres used for reference standard)⁴³. However, for their study a mechanical arm was used to hold the probe in position during contrast agent infusion (as compared to a bolus injection for this study). Furthermore, the entire testis was selected as an ROI (an unfeasible approach in larger organs such as the kidney) for volumetric analysis reducing the error caused by subjective selection of a smaller ROI. Other groups have attempted to quantify blood flow in mL/min rather than in mL/min/g^{44, 45}. To the best of our knowledge, this is the first attempt to enable quantification of absolute perfusion via perfusion estimates determined using contrast-enhanced 3D nonlinear ultrasound imaging.

Although perfusion estimates have shown significant correlations with the reference standard, it is important to acknowledge the limitations of our study. There were problems associated with acoustic shadowing artifacts in the posterior region of the kidney. No motion compensation (other than a basic smoothing filter) was applied to reduce breathing induced motion artifacts. Finally, the tissue samples extracted for determining perfusion were about $1 \times 1 \times 1 \text{ cm}^3$ in size, making it difficult to accurately select a corresponding ROI in the volumetric data (which induces subjective error) to produce perfusion estimates. Future studies will involve applying motion compensation, developing methods to automate selection of ROI and possible human studies.

Conclusions

Quantitative perfusion estimation using contrast-enhanced 3D nonlinear ultrasound imaging is shown to be feasible *in vivo*. Specifically, 3D SHI perfusion estimates exhibit good correlation with the reference standard ($r=0.62$) as compared to HI ($r=0.38$). This is an improvement over a previous 2D SHI perfusion estimation study ($r=0.57$) by our group and highlights the feasibility of 3D SHI and importance of volumetric imaging for making accurate perfusion measurements.

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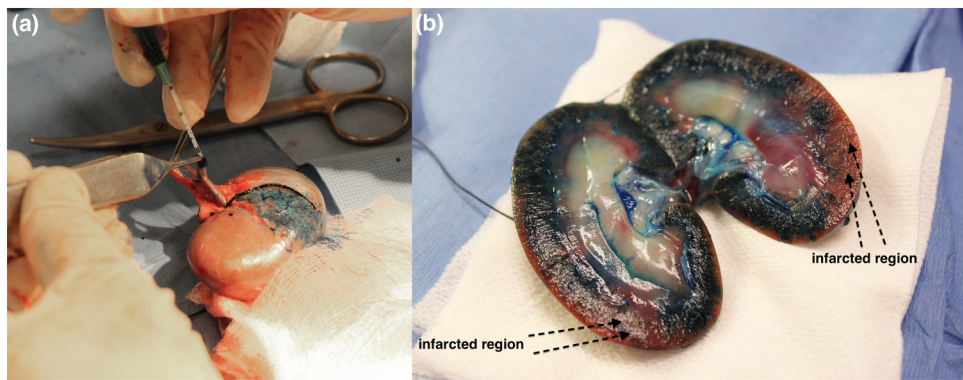


Figure 1. (a) Injection of methylene blue dye via renal artery. (b) Normal and infarcted regions visualized after making a mid-line cut.

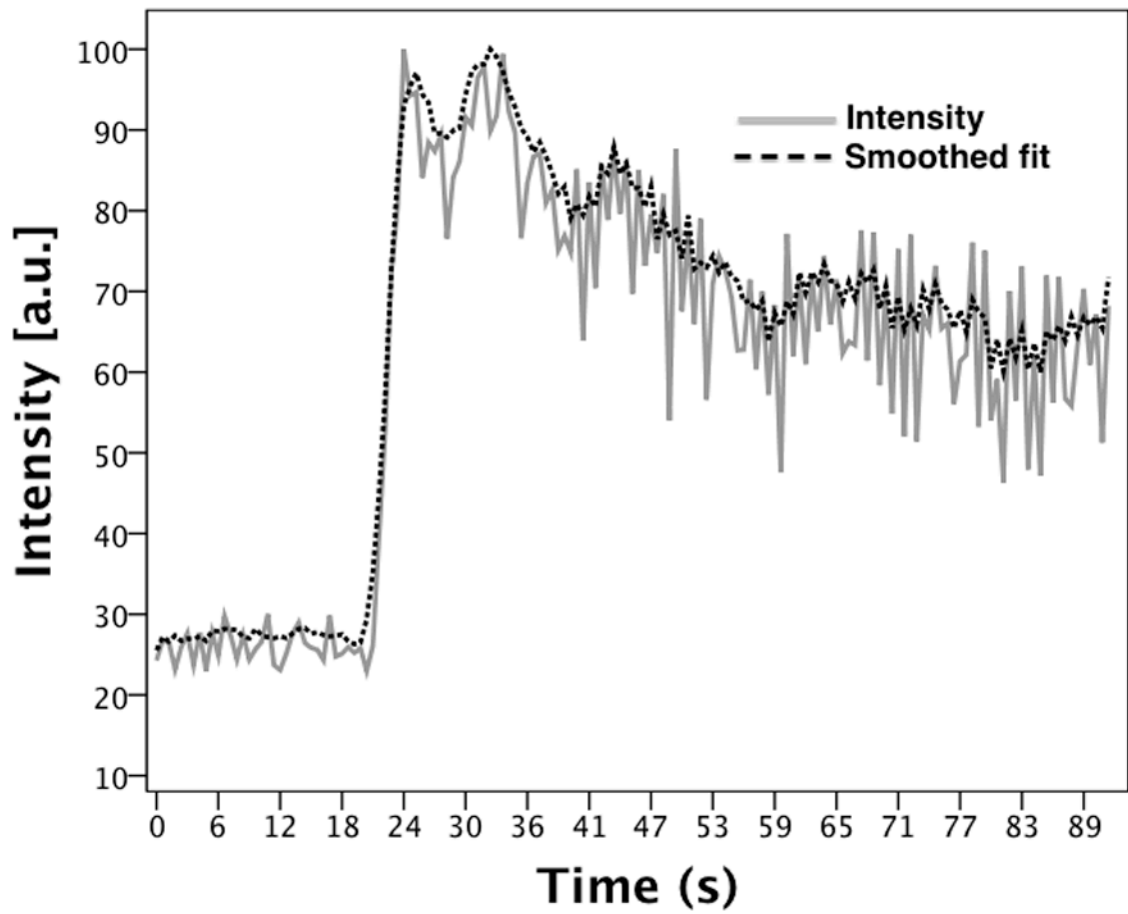


Figure 2. Subharmonic time-intensity curve generated from the averaged image intensities for a single injection (pre-ligation). The point of peak contrast wash-in is labeled.

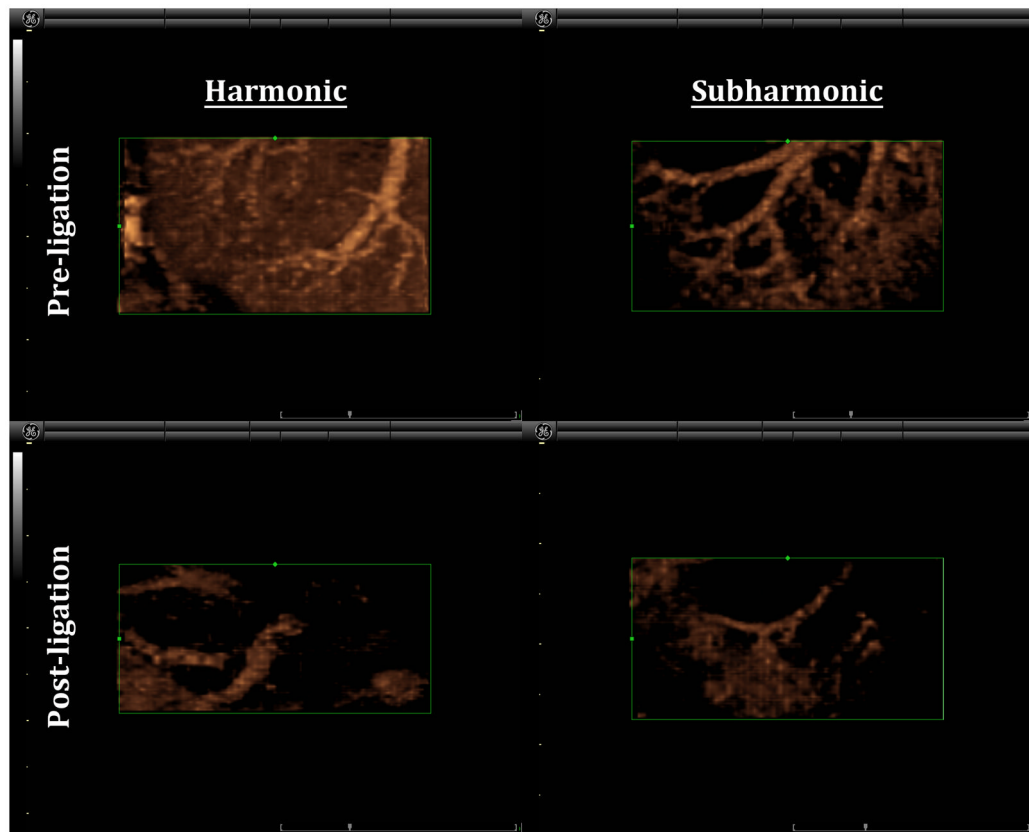


Figure 3. Slice (image intensity) distribution across the volume (from lateral to medial sections). The dashed line is a representation of the mid-line plane. The slices highlighted in black were isolated from the entire volume for further image processing (as lateral and medial volumes).

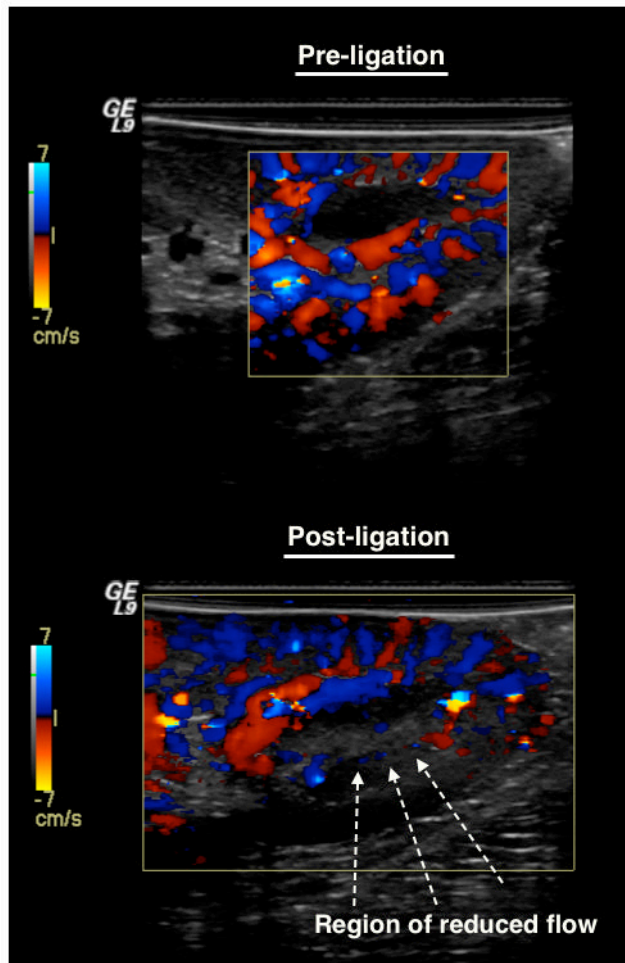


Figure 4. Subharmonic time-intensity curve generated from an ROI selected in the lateral anterior section before and after smoothing (solid and dashed line respectively).

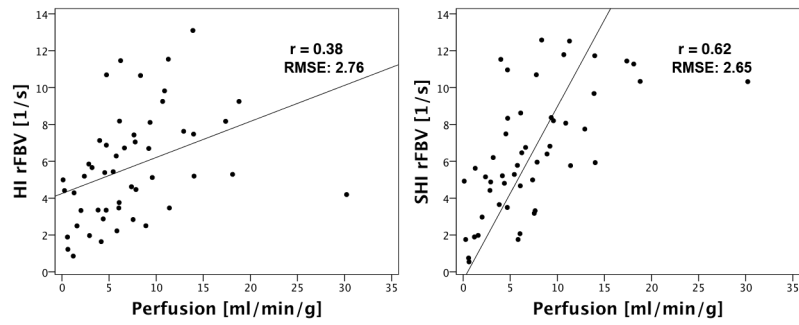


Figure 5. Comparison of 3D HI and SHI pre and post ligation (same canine and same kidney). Clear suppression of tissue is seen in the 3D SHI rendering as compared to HI. Reduced flow is visualized (post-ligation) in both the modes.

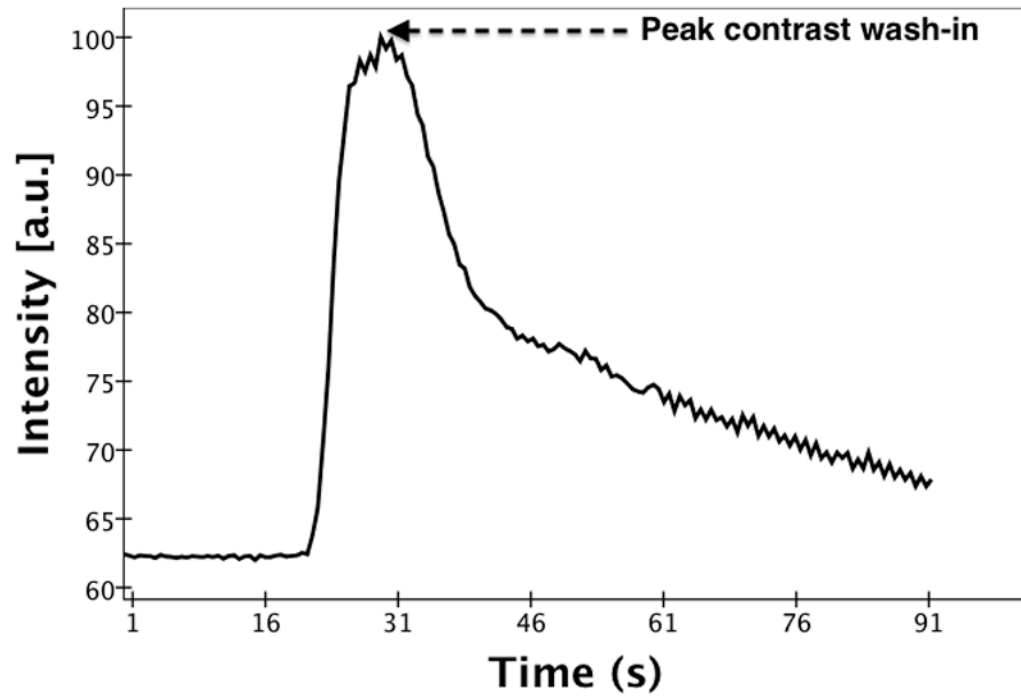


Figure 6.
Color Doppler visualization of blood flow before and after ligation

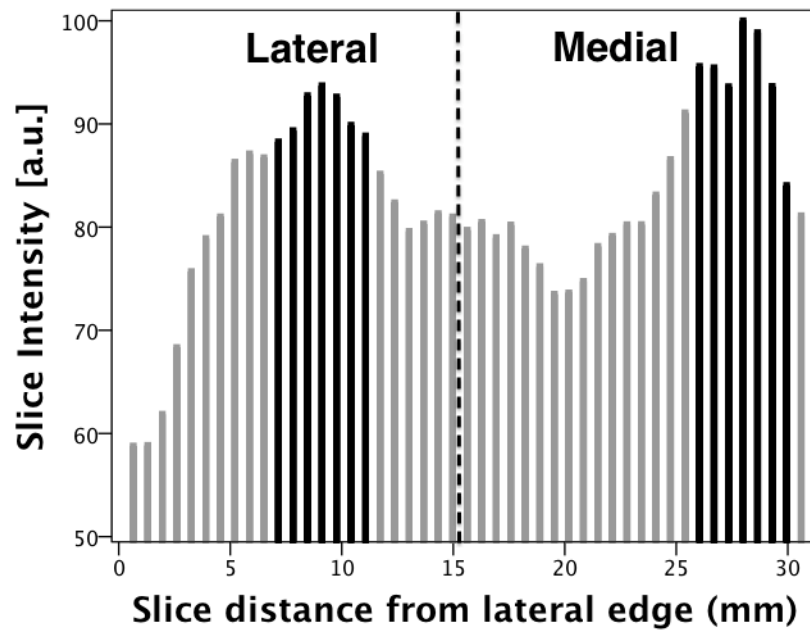


Figure 7. Overall (a) harmonic (b) subharmonic perfusion estimates versus microsphere perfusion (in mL/min/g)

TABLE 1

Regression Results From the Overall and Regional Subsets of the 3D HI and 3D SHI Perfusion Estimates Correlations With the Reference Standard.

Subset	HI		SHI	
	<i>r</i>	<i>P</i>	<i>r</i>	<i>P</i>
Overall (n = 50)	0.38	0.007	0.62	<0.001
Anterior (n=28)	0.41	0.033	0.49	0.008
Posterior (n=22)	0.45	0.034	0.79	<0.001
Preligation (n = 27)	0.21	0.285	0.66	<0.001
Postligation (n=23)	0.63	0.001	0.70	<0.001
Preligation anterior (n=14)	0.27	0.344	0.44	0.12
Preligation posterior (n=13)	0.24	0.437	0.79	<0.001
Postligation anterior (n= 14)	0.58	0.03	0.71	0.004
Postligation posterior (n=9)	0.58	0.101	0.47	0.197

3D indicates 3-dimensional; HI, harmonic imaging; SHI, subharmonic imaging.

TABLE 2

Regression Results From the Overall and Regional Subsets of the Quasi-2D HI and Quasi-2D SHI Perfusion Estimates Correlations With the Reference Standard.

Subset	HI		SHI	
	<i>r</i>	<i>P</i>	<i>r</i>	<i>P</i>
Overall (n=24)	-0.05	0.824	0.14	0.5
Anterior (n=13)	-0.12	0.709	-0.22	0.459
Posterior (n = 11)	-0.15	0.616	0.61	0.035
Preligation (n=13)	-0.02	0.951	0.01	0.963
Postligation(n=11)	-0.15	0.633	0.22	0.484
Preligation anterior (n=7)	-0.24	0.603	-0.22	0.634
Preligation posterior (n=6)	0.02	0.965	-0.24	0.602
Postligation anterior (n=6)	0.26	0.617	0.31	0.498
Postligation posterior (n=5)	-0.82	0.023	0.90	0.042

2D indicates 2-dimensional; HI, harmonic imaging; SHI, subharmonic imaging.