

Acoustic Shadow Detection: Study and Statistics of B-Mode and Radiofrequency Data

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Abstract

An acoustic shadow is an ultrasound artifact occurring at boundaries between significantly different tissue impedances, resulting in signal loss and a dark appearance. Shadow detection is important as shadows can identify anatomical features or obscure regions of interest. A study was performed to scan human subjects (N=35) specifically to explore the statistical characteristics of various shadows from different anatomy and with different transducers. Differences in shadow statistics were observed and used for shadow detection algorithms with a fitted Nakagami distribution on radiofrequency speckle (RF) or cumulative entropy on brightness-mode (B-mode) data. The fitted Nakagami parameter and Entropy values in shadows were consistent across different transducers and anatomy. Both algorithms utilized adaptive thresholding, needing only the transducer pulse length as an input parameter for easy utilization by different operators or equipment. Mean Dice coefficients

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(\pm standard deviation) of 0.90 ± 0.07 and 0.87 ± 0.08 were obtained for the RF and B-mode algorithms, which is within the range of manual annotators. The high accuracy in different imaging scenarios indicate that the shadows can be detected with high versatility and without expert configuration. The understanding of shadow statistics can be used for more specialized techniques to be developed for specific applications in the future, including pre-processing for machine learning and automatic interpretation.

Keywords: Acoustic Shadow, Ultrasound, Speckle, Radiofrequency, Segmentation

1 Introduction

2 Ultrasound devices have become increasingly affordable and portable, en-
3 couraging applications such as point-of-care ultrasound (Bouhemad et al.,
4 2011), novice usage (Sippel et al., 2011), and analysis by machine learning
5 (Ghose et al., 2013). However, ultrasound is susceptible to unique artifacts
6 that increase the difficulty of interpretation and processing of images. One
7 artifact is an acoustic shadow, which occurs when an ultrasound wave crosses
8 a boundary of two materials with high impedance differences (Kremkau and
9 Taylor, 1986). The wave is almost completely reflected and depicted beyond
10 the boundary is a continuous dark region and a loss of anatomical features.
11 Shadows occur in air-tissue, tissue-bone, and tissue-lesion interfaces. Shad-
12 ows can aid interpretation, such as identifying gall stones (Good et al., 1979)
13 or spinal levels (Galiano et al., 2005). However, shadows, such as from poor
14 transducer contact, can lead to misinterpretation of anatomy, particularly by
15 novice users and automated processing algorithms. Thus, the identification
16 of shadows is an important preprocessing step in many applications.

17 Several methods have been used in literature to detect shadows and il-
18 lustrative examples are discussed. Geometric techniques model the path of
19 an ultrasound signal for an expected image along the scanline using a ran-
20 dom walk (Karamalis et al., 2012). Pixels are then flagged as a shadow
21 if it is below a heuristic confidence threshold of 0.25. However, geometric
22 techniques require knowledge of ultrasound transducer properties to param-
23 eterize random walk weights, such as the focal length, radius of curvature,
24 and thickness. The technique is therefore challenging to implement across
25 different ultrasound equipment. This also reduces applicability for machine

26 learning applications as accurate transducer parameter labels are required
27 for each image.

28 Pixel intensity methods ignore the transducer properties and analyze only
29 the graphical properties of an image (Hellier et al., 2010). Shadows have been
30 detected on brain images by analyzing the entropy along a scanline to flag pix-
31 els of sudden low entropy as a potential shadow. These techniques achieved a
32 comparable Dice similarity coefficient as geometric methods but require spe-
33 cific thresholding, window sizing, filtering, and image mask parameterization
34 for different anatomy and transducers. The drawback is again the need for
35 parameterization and tuning, which requires image processing expertise and
36 prior knowledge of specific applications.

37 Machine learning methods have gained significant interest in medical
38 imaging analysis. To our knowledge, no machine learning method has demon-
39 strated the capability of general shadow detection from multiple types of
40 anatomy. Deep learning methods have identified features in a specific im-
41 age sets that contain shadows, such as neuroanatomical regions in cranial
42 scan (Milletari et al., 2017) or spinal levels in a posterior scan (Hethering-
43 ton et al., 2017). Although machine learning has the potential of providing
44 automated feature recognition in multiple applications, a large data set is re-
45 quired for an algorithm to recognize certain features. Ultrasound imaging is
46 highly variable due to unique artifacts, operator techniques, and equipment.
47 In addition, shadows are a common feature that occur in various imaging
48 scenarios. Previous techniques focused on a single anatomical region and
49 training data was from a consistent imaging scenario. However, it is difficult
50 to construct a training data set with the generality required to recognize

51 shadows in different scenarios usable for a variety of ultrasound applications.

52 There are two objectives to this paper. First, to address the need for
53 understanding general characteristics of shadows, a study was conducted to
54 scan multiple anatomy and transducers specifically to analyze the statistics of
55 different types of shadows. Second, to address existing needs for versatile de-
56 tection with minimal parameterization, previous methods were then extended
57 utilizing statistical thresholding of radiofrequency (RF) or brightness-mode
58 (B-mode) data to detect shadows from various imaging scenarios. The two
59 methods are illustrated in a flowchart in Fig. 1.

60 **Materials and Methods**

61 *Data Collection*

62 Ultrasound RF and B-mode data were acquired by scanning 37 adult
63 participants with informed written consent, approved by the University of
64 British Columbia Research Ethics Board (Study ID: H18-01199). The scans
65 included a forearm scan near the distal end of the pronator quadratus, an
66 elbow scan near the cubital fossa, and a rib scan on the anterior surface of
67 right ribs 11-12. Each scan was taken with both a curvilinear (Model C5-
68 2/60, Ultrasonix Medical Corporation, Richmond, BC, Canada) and linear
69 (Model L14-5/38, Ultrasonix Medical Corporation, Richmond, BC, Canada)
70 transducer. Different transducer settings were used for each anatomical re-
71 gion and transducer, summarized in Table 1. Shadows were expected to
72 occur due to superficial and deep bones and from an air gap created by the
73 lateral edges of the transducer not being in flush contact with the skin. The
74 experiment was designed to generate a dataset from various imaging scenar-

ios to explore general shadow characteristics and to validate the versatility
of the two simple shadow detection methods. A pulse length of 2mm was
used for both transducers.

Radiofrequency Speckle Analysis

To analyze shadows, windows of speckle were analyzed on the RF signal. Speckle occurs from interference of randomly distributed microscopic scatterers, resulting in a granular appearance on the image. To produce B-mode images, manufacturers often use image enhancement algorithms, such as logarithmic compression, nonlinearly alter speckle patterns. B-mode image formation can also be manipulated by an operator to visually enhance an image, such as adjusting time-gain compensation or dynamic range. Thus, the underlying speckle analysis in RF signals can provide shadow detection usable across different machines and operators. However, the original speckle pattern contains information of the acoustic interactions in tissue. (Burckhardt, 1978). By analyzing the RF signal distribution, we can statistically characterize the distributions in tissue compared to shadow regions. We expect tissue to resemble speckle modeled by known distributions and expect shadow to resemble different distributions, which may be a mixture of lessened speckle due to the signal loss and background electronic noise. Previous studies have attempted despeckling methods on images containing shadows (Aysal and Barner, 2007) by using filters based on the Rayleigh-like distributions. As such, even if shadow regions do not exactly resemble known speckle distributions, they may still be characterized to a sufficient extent with known distributions for a maximum likelihood fit. The fitted parameters can then be used to differentiate between shadow and non-shadow regions.

100 One of the first models for speckle is the one parameter Rayleigh distribu-
 101 tion to model the probability density of a random walk (Burckhardt, 1978).
 102 The Rayleigh distribution is capable of modeling fully developed speckle,
 103 which does not occur in limited scattering (Tuthill et al., 1988). More gen-
 104 eralized models have been applied such as the Rician, Homodyned-K, and
 105 Nakagami distributions to characterize speckle (Destrepes and Cloutier,
 106 2010). The utility of speckle has been demonstrated in the literature to
 107 classify tumorigenicity of breast lesions (Byra et al., 2016) or levels of liver
 108 fibrosis (Ho et al., 2012) by categorizing image regions based on the speckle
 109 pattern. Shadow characterization presents a simpler problem as shadow and
 110 non-shadow regions contain significantly different speckle patterns. Thus,
 111 the Nakagami distribution expressed in Eq. 1 was chosen to model speckle.
 112 The Nakagami distribution provides greater generality than the Rayleigh
 113 distribution while being more computationally efficient than the Rician or
 114 Homodyned K distributions (Destrepes and Cloutier, 2010):

$$\Phi(x, m, \omega) = 2\left(\frac{m}{\omega}\right)^m \frac{1}{\Gamma(m)} x^{(2m-1)} e^{\frac{-m}{\omega} x^2} \quad (1)$$

115 where x is RF intensity, m is the shape parameter or Nakagami m parameter,
 116 ω is a scale parameter and $\Gamma(m)$ is the gamma distribution.

117 To characterize shadows, the raw RF data was first processed by com-
 118 puting the echo envelope of each scanline with a Hilbert transform. This was
 119 performed on an averaged RF signal from three image frames. This creates
 120 a pre-scan converted image, visually similar to B-mode but without filtering
 121 to alter speckle. Next, the RF image was divided into overlapped windows
 122 with a width of a single RF scanline and a length of three times the pulse

length. We expect the width of a single RF scanline to be on the order of magnitude of a resolution cell, which is on the same order of magnitude as the correlation length (Wagner and Insana, 1988). The window length was demonstrated in literature to be sufficiently large to capture multiple wavelengths and scattering events while being small enough to be useful in differentiating different regions on the millimeter scale (Byra et al., 2016). Next, each window was fit to a Nakagami distribution using a maximum likelihood estimate to compute a map of Nakagami parameters m and ω , as shown in Fig. 1.

Then, for each ultrasound image, Otsus method was applied to its Nakagami ω map to automatically compute a ω threshold for each individual image as we expect separate distributions for shadow and non-shadow regions. This was sufficient as the ω parameter is significantly different for shadow regions with abundant speckle and non-shadow regions with minimal speckle. Then, for each scanline, the axially deepest data point that is above the threshold is labeled as the shadow boundary and all data points below are labeled as a shadow.

The Nakagami shape parameter, m , was also investigated, though there was not sufficient delineation between parameter values in shadow and non-shadow regions for this parameter to be effective in thresholding. The distributions of the two parameters are displayed for shadow and non-shadow regions in Fig. 4.

B-mode Scanline Analysis

Many ultrasound machines do not provide access to RF data for speckle analysis. Thus, a previous pixel-intensity shadow detection method on B-

mode images was modified and extended. Scanline entropy was investigated on B-mode images to characterize different types of shadows, but with the addition of adaptive thresholding of entropy to address the need for usability with minimum configuration. B-mode analysis was performed on an averaged image from three image frames, similar to RF analysis. First, the cumulative scanline entropy is computed for each pixel, similar to the “Rupture Criterion” (Hellier et al., 2010), with the window size fixed as three times the pulse length, η , as defined in Eq. 2. This is the same window size as the RF analysis.

$$S_{i,j} = \sum_{k=1}^{3\eta} I(i-k,j) \log_2 \frac{I(i-k,j)}{I(i+k,j)} + I(i+k,j) \log_2 \frac{I(i+k,j)}{I(i-k,j)} \quad (2)$$

where $S_{i,j}$ is the cumulative entropy at pixel i on scanline j , η is the pulse length, and $I(i)$ is the gray level, or intensity, of pixel (i,j) . For the case of curvilinear images, radial scanlines were linearly interpolated between the two symmetric lateral edges of the image.

Next, Otsu’s method is applied onto the entropy map of each image to automatically compute a threshold entropy value, similar to RF analysis. The intuition of the threshold is different than in RF analysis. In RF analysis, the threshold separates patches of intense and minimal speckle. In B-mode analysis, the threshold separates pixels of a shadow boundary, which has high entropy, and pixels away from shadow boundary, which include shadow and non-shadow regions. Thus, shadows can be identified by finding the last pixel on a scanline with an entropy higher than the threshold, representing a bright shadow boundary.

170 *Validation*

171 A trained annotator (RH) manually outlined the boundary of the shadow
172 regions on B-mode images. The manual regions were used as a gold standard,
173 as manual identification is common in clinical practice and has been used in
174 previous literature for comparison (Hellier et al., 2010). A Dice coefficient
175 was computed to compare similarity of manual and automated shadow de-
176 tection. The manual outline was used to define four regions for classification
177 of statistical parameters: a non-shadow region above the boundary, a shadow
178 region below the boundary, a “transition region”, which is a window defined
179 as three pulse lengths long axially below the boundary, and a “deep shadow
180 region”, which is the data below the transition region. The validation was
181 repeated with the RF and entropy window increased and decreased by 50%.
182 The Ljung-Box Q-test was used to measure residual autocorrelation of the
183 Dice coefficients. A Wilcoxon rank sum test has been performed between
184 Nakagami parameter values in shadow and non-shadow regions and between
185 entropy values in shadow and non-shadow regions.

186 **Results**

187 Examples of detected shadows from both methods are highlighted in gray
188 in Fig. 2 in different shadow detection scenarios. The Dice coefficients for
189 both methods for different anatomy and transducers are shown in Table
190 2. The mean Dice coefficients (\pm standard deviation) were 0.90 ± 0.07 and
191 0.87 ± 0.08 for RF and B-mode methods. Manual annotation was repeated
192 five times with a mean Dice coefficient of 0.92 ± 0.02 for all images and trans-
193 ducers. The Dice coefficient did not change by more than 0.03 when the

194 window size was varied by 50%.

195 With the benefit of a varied dataset, general statistics of shadows can
196 be analyzed, as summarized in Table 3 and Table 4. The distributions of
197 Nakagami parameters and entropy for the different regions are visualized
198 in Fig. 4. For shadow detection, the parameters differentiating a shadow
199 and non-shadow are of particular interest. Shadows were observed to have a
200 mean Nakagami ω parameter of 4.14 ± 0.40 and a mean entropy of $1.03 \pm$
201 0.29 whereas non-shadows were observed to have a mean ω of 6.24 ± 0.92
202 and 2.20 ± 0.81 . Wilcoxon rank sum p values were less than 0.002 between
203 Nakagami parameter distributions in shadow and non-shadow regions and
204 less than 0.001 between entropy distributions in shadow and non-shadow
205 regions, indicating that shadow and non-shadow regions have statistically
206 different distributions for ω and entropy. The values of entropy and Nakagami
207 ω are consistent across different transducers and anatomical regions. The
208 variance of entropy and Nakagami ω in one imaging region and transducer
209 setting is less than the variance across different regions and transducers for
210 shadows and non-shadows.

211 Discussion

212 The RF and B-mode shadow detection developed achieved a comparable
213 Dice similarity coefficient to manual detection for all anatomy and transducer
214 types ($p < 0.025$). The previous studies using B-mode entropy reported a
215 mean Dice coefficient of 0.91 ± 0.07 between manual annotators (Hellier et al.,
216 2010). An important feature of shadow detection is being able to differentiate
217 between a shadow and simply high attenuation of the signal. Both scenarios

218 result in an eventual loss of signal. Shadow detection, however, has a char-
219 acteristic high intensity shadow boundary before a significant loss in signal,
220 compared to gradual signal losses in attenuation. The high Dice similarity
221 coefficient indicates that both methods were capable of this distinction. This
222 is also visualized in Fig. 2, where regions of low intensity without a bright
223 shadow boundary were correctly labeled as non-shadow. The high accuracy
224 supports the versatility of the detection method as both methods are able
225 to identify shadows across different anatomy and transducers with minimum
226 configuration.

227 For a general observation for shadows, the computed Nakagami ω param-
228 eters of all manually outlined shadows indicate that there is a statistically
229 significant difference between shadow and non-shadow regions, regardless of
230 anatomy and transducer and even with the error in the transition regions
231 considered. The speckle and its statistics from shadows is thus distinct from
232 the speckle created by tissue, muscle, or fat. This observation can be utilized
233 in the future for further analysis of shadows.

234 In RF detection, both false positive and false negative errors most fre-
235 quently occurred immediately below a shadow boundary as opposed to B-
236 mode detection where errors were in various regions. To study the frequent
237 areas of error further, the “transition region” immediately below a man-
238 ually annotated shadow boundary and a “deep shadow region” below the
239 transition region was investigated. The Nakagami ω parameter of transition
240 regions of all anatomy and transducers were within a standard deviation of
241 both shadow and non-shadow regions. The deeper shadow regions were ob-
242 served to have a lower Nakagami ω parameter than shadow regions and with

243 a lower standard deviation as summarized in Table 3. The spread of the
 244 speckle also significantly decreases after the transition region. This indicates
 245 that the transition region cannot be fully distinguished from either a shadow
 246 or non-shadow and presents as it is statistically similar to the two. This
 247 is likely the cause of the errors, as the speckle distribution is much more
 248 consistent in the deep shadow regions compared to any other region. Phys-
 249 ically, speckle interactions appear to gradually lessen after a brightest point
 250 on a scanline, possibly due to incomplete total reflection at a boundary. The
 251 boundary is thus is not an instantaneous division between non-shadow and
 252 shadow, rather, there is a transition region with statistics between a shadow
 253 and non-shadow before the speckle fully resembles a shadow.

254 In the transition region of B-mode images, the entropy values were similar
 255 but consistently higher than non-shadow values. This is expected as entropy
 256 is the highest when there is the greatest change in pixel intensity, which oc-
 257 curs at a shadow boundary, even with the a non-instantaneous non-shadow
 258 to shadow transition. However, the averaged entropy of all non-shadow re-
 259 gions have a greater spread than the Nakagami parameters, likely due to
 260 the differing operator settings used. Thus, B-mode detection may not be as
 261 consistent as RF detection.

262 As both RF and B-mode images search for a threshold for the start of a
 263 shadow, it is possible to misinterpret a beginning of a shadow as a reverber-
 264 ation artifact. Reverberation at a shadow boundary would cause a similar
 265 bright region followed by a dark region, which visually appears like a shadow
 266 boundary despite being an artifact in a shadow region. This is addressed by
 267 considering directionality when searching for the start of a shadow boundary

268 such that the first shadow boundary when traversing down a scanline is in-
269 terpreted as a beginning of a shadow and any further shadow boundaries are
270 interpreted as reverberation artifacts. Fig. 2 shows shadow detection with a
271 reverberation artifact underneath a shadow caused by the radial joint.

272 There is a limitation with analysis using the Nakagami distribution in
273 that the fitted Nakagami distribution to model scatterers change depending
274 on transducer frequency. Previous literature observed that in the 36-58MHz
275 frequency range, the Nakagami m parameter decreased near the theoretical
276 lower limit compared to a higher Nakagami m parameter value at 10MHz
277 signal (Cloutier et al., 2004). This was reported to be due to the spatial
278 organization of the cells being "on the order of a fraction of the wavelength"
279 and a Nakagami distribution cannot model the scatterers of red blood cells at
280 this frequency. Due to this and from limitations of the equipment used in our
281 study, we cannot conclude that shadow detection with Nakagami analysis will
282 be accurate in higher frequencies beyond the values tested. Future studies are
283 required to analysis the performace of shadow detection in higher frequencies.
284 Diagnostic ultrasound commonly uses a frequency range of 2-15MHz (Jensen,
285 2007) and the shadow detection method is expected to not be applicable
286 in most use cases without issues from the high frequency behaviour of the
287 Nakagami distribution.

288 There is a limitation for diagnostic usage of the proposed shadow method
289 in cases where acoustic shadowing does not exhibit the characteristic bright
290 boundary followed by a dark region. In cases where there is partial or incom-
291 plete shadowing, such as small calcifications in the placenta (Abramowicz
292 and Sheiner, 2008). In these cases, there is a resemblance of a shadow, where

293 the calcification is brighter and the region below is noticeably darker, but
294 not with a brightness difference as extreme as shadowing from the ulna and
295 the regions below retain speckle similar to tissue. Although calcifications
296 are pathologically important to recognize, the proposed shadow detection
297 method would likely be unable to detect the partial shadowing from these
298 calcifications. The proposed method would be applicable only in cases of
299 more complete shadowing, which would still be practical for significant gall
300 and kidney stones, for instance.

301 In previous literature, shadows were defined qualitatively (Kremkau and
302 Taylor, 1986) as a sudden loss of signal and brightness. The observed transi-
303 tion region in this study suggests that the qualitative definition of a shadow
304 may be insufficient for accurate detection. One algorithm may detect the
305 shadow starting immediately after the brightest location, or another may
306 use a convention such as a full width at half maximum to define where the
307 signal has sufficiently low intensity to resemble the start of a shadow. There
308 is a decision point required for a clear definition for where a shadow begins
309 to improve shadow detection accuracy, both from a signaling perspective for
310 image processing and a visual perspective for manual inspection.

311 The findings in this study result in several implications. First, the statis-
312 tics of acoustic shadows have been investigated on a dataset with shadows
313 occurring from multiple scenarios as opposed to specific cases where shadows
314 are observed. This provided a more generalizable observation that shad-
315 ows can be characterized by distinctive speckle distributions in different of
316 anatomy and equipment and that there exists a transition region before the
317 loss of speckle in a shadow. Second, the shadow detection methods demon-

318 strated high accuracy, indicating that the same shadow detection method
319 can be used with different transducer or imaging location. In future stud-
320 ies, the speckle statistics observed can be used to develop further models for
321 anatomical features containing shadows. In machine learning algorithms, an
322 initial network could be used with the shadow detection methods presented.
323 Future studies would also have to take into consideration the most frequent
324 source of error of shadow detection as the shadow boundary.

325 **Conclusions**

326 Acoustic shadows from different imaging scenarios were investigated. RF
327 and B-mode methods were developed for acoustic shadow detection requiring
328 only the transducer pulse length as the input parameter. When comparing to
329 manual detection, the methods achieved a Dice similarity coefficient within
330 range of manual observers. The work focused on applying shadow detection
331 and statistical analysis to a varied dataset of three different anatomical loca-
332 tions and two different transducer to provide a representative understanding
333 of general acoustic shadows. The statistics of acoustic shadow indicate that
334 shadows contain a distinct speckle distribution compared to non-shadows and
335 the speckle characteristics transition at the shadow boundary. The statistical
336 findings of shadows can aid interpretation of ultrasound images in the future
337 using speckle analysis. The versatility of the shadow detection method has
338 the potential to improve the interpretation of ultrasound images with shadow
339 artifacts or to serve as a pre-processing step for machine learning methods.

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404 **Figure Captions**

405 **Figure 1:** Processing steps for Radiofrequency (RF) and B-mode shadow
406 detection. RF processing is used if RF data is available and involves
407 fitting the Nakagami distribution onto the echo envelope of each RF
408 scanline before adaptive thresholding with Otsus method. In many
409 cases, there may only be access to B-mode image data, for which an
410 entropy map is computed and similar adaptive thresholding is used to
411 detect shadows.

412 **Figure 2:** A visualization of the B-mode and RF parameter maps. The b)
413 Entropy Map was computed from processing of the a) original B-mode
414 image and the d) Nakagami ω map was computed from the c) echo
415 envelope. Note that the echo envelope contains noticeable speckle,
416 which has been used to fit a Nakagami distribution to characterize
417 shadow. The region at depth 2.50 cm and scanlines 32-40 is attenuation
418 and not a shadow. This is an important distinction in shadow detection
419 and both maps show the region as below a threshold to flag a shadow
420 boundary.

421 **Figure 3:** A comparison of the original B-mode images, the detected shad-
422 ows manual detection, RF detection, and B-mode detection. Both
423 detection methods perform similarly to manual detection. Both meth-
424 ods perform slightly less accurately on curvilinear images, likely due
425 to the reduced resolution from interpolating the scanlines. Most errors
426 of RF detection occur near the shadow boundary, likely due to the
427 transitioning speckle from non-shadow to shadow.

Figure 4: Histograms of Nakagami parameters and entropy values in shadow and non-shadowing regions. The Nakagami ω has a more noticeable delineation between shadowing and non-shadowing distributions compared to the Nakagami m parameter and was used as the only parameter to threshold shadow boundaries. The entropy distributions for shadow and non-shadow differ as entropy is very minimal in continuous dark shadow regions. Although entropy varies in non-shadow regions, thresholding can be used to detect a shadow boundary where at some point along a scanline, the entropy increases above a threshold and remains low afterward to resemble the shadow distribution.

438 **Tables**

439 **Table 1:** Transducer properties for different imaging scenarios.

440

	Anatomy	Frequency	Depth	Gain
Linear Transducer (L14-5/38)	Forearm	11.0MHz	5.0cm	50%
	Elbow	11.0MHz	5.0cm	40%
	Ribcage	5.0MHz	10.0cm	30%
Curvilinear Transducer (C5-2/60)	Forearm	4.0MHz	5.0cm	50%
	Elbow	4.0MHz	5.0cm	40%
	Ribcage	3.3MHz	10.0cm	30%

441 **Table 2:** Mean Dice coefficients for different imaging scenarios \pm standard
442 deviation.

		RF	B-Mode
Linear (L14-5/38)	Forearm	0.91 \pm 0.05	0.89 \pm 0.06
	Elbow	0.94 \pm 0.06	0.90 \pm 0.07
	Ribcage	0.87 \pm 0.09	0.84 \pm 0.06
Curvilinear (C5-2/60)	Forearm	0.89 \pm 0.05	0.86 \pm 0.08
	Elbow	0.93 \pm 0.04	0.90 \pm 0.09
	Ribcage	0.83 \pm 0.08	0.83 \pm 0.10
Mean	All Anatomy	0.90\pm0.07	0.87\pm0.08

443 **Table 3 :** The mean Nakagami ω and Entropy values of different anatomy,

transducer, and shadowing region \pm standard deviation. Values are consistent among different transducers and anatomical regions. The variance of entropy and Nakagami ω in one imaging region and transducer setting is less than the variance across different regions and transducers for shadows and non-shadows.

	Linear (L14-5/38)			Curvilinear (C5-2/60)		
	Forearm	Elbow	Ribcage	Forearm	Elbow	Ribcage
Nakagami ω (Log Scale)						
Shadow	4.15 \pm 0.45	4.18 \pm 0.45	4.04 \pm 0.42	4.22 \pm 0.32	4.19 \pm 0.40	4.08 \pm 0.37
Non-Shadow	6.19 \pm 0.96	6.49 \pm 0.97	6.29 \pm 0.95	6.54 \pm 0.88	6.29 \pm 1.04	5.64 \pm 0.71
Transition	4.94 \pm 0.62	5.36 \pm 0.62	4.96 \pm 0.38	5.26 \pm 1.02	5.37 \pm 0.99	4.59 \pm 0.92
Deep Shadow	4.13 \pm 0.43	4.16 \pm 0.43	4.03 \pm 0.41	3.93 \pm 0.20	4.09 \pm 0.30	4.03 \pm 0.26
Entropy (Log Scale)						
Shadow	0.92 \pm 0.22	1.10 \pm 0.36	1.04 \pm 0.27	1.06 \pm 0.28	0.96 \pm 0.21	1.10 \pm 0.37
Non-Shadow	2.34 \pm 0.96	2.34 \pm 0.80	2.14 \pm 0.82	1.67 \pm 0.82	1.75 \pm 1.14	1.88 \pm 0.42
Transition	2.45 \pm 0.62	2.56 \pm 0.53	2.15 \pm 0.51	2.18 \pm 1.21	1.93 \pm 1.10	1.99 \pm 1.10
Deep Shadow	0.71 \pm 0.43	0.89 \pm 0.26	0.92 \pm 0.40	0.98 \pm 0.21	0.82 \pm 0.19	1.04 \pm 0.26

Table 4 : The mean Nakagami ω and Entropy values of all anatomy and transducers for different shadowing regions \pm standard deviation.

	Mean Nakagami ω (Log Scale)	Mean Entropy (Log Scale)
Shadow	4.14 ± 0.40	1.03 ± 0.29
Non-Shadow	6.24 ± 0.92	2.02 ± 0.81
Transition	5.08 ± 0.77	2.21 ± 0.84
Deep Shadow	4.06 ± 0.34	0.89 ± 0.27