

**UiO** : **Department of Physics**  
University of Oslo

**Parameters of the Capon beamformer in medical  
ultrasound imaging**

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# Abstract

In this thesis, the performance of the Capon beamformer is evaluated, using the Delay-And-Sum beamformer as a performance benchmark. The beamformers are utilized with simulated medical ultrasound data. The Capon beamformer is an adaptive high-resolution beamformer but it can be sensitive to errors in the assumed wavefield. This beamforming algorithm involves the estimation and inversion of a spatial covariance matrix. The robustness of this estimation and inversion can be increased by, e.g., applying spatial averaging when estimating the covariance matrix, as well as by adding a diagonal load to the estimated matrix.

The performance of the Capon beamformer is measured in terms of resolution and contrast when the subarray length and diagonal load are systematically varied. The condition number and the eigenvalues of the covariance matrix are utilized when assessing the stability with respect to the inversion of the covariance matrix. The resolution and contrast have been analysed using simulated scenes containing a set of scatterers, both with and without speckle, as well as a scene with a cyst in speckle, respectively. A variant of the Capon algorithm is the power Capon method, which is also assessed in terms of the separability of scatterers in a simulated scene.

In the resolution analysis, it is deemed that a subarray size between  $L_{\min} = \frac{1}{3}M$  and  $L_{\max} = \frac{2}{3}M$ , where  $M$  is the full length of the array, is robust. For the diagonal load, it is found that  $DL = 0.01 \frac{\text{tr}(R)}{\text{dim}(R)}$  is robust, dependent on the scene. Furthermore, the condition number of the covariance matrix is found to be a useful measure of stability in the covariance matrix inversion through analysis and can be applied as an aid when setting the subarray length and diagonal load.

An analysis of contrast metrics showed how the subarray length and diagonal load affect the measured contrast, combined with an analysis of how different regions of interest affect the contrast metrics. The measured contrast metrics are the Contrast Ratio, the Contrast-to-Noise Ratio, and the generalised Contrast-to-Noise Ratio. The contrast analysis shows that using the results obtained from analyzing the subarray length and diagonal load resolution leads to optimal outcomes in terms of contrast metrics.

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# Abbreviations

**CNR** Contrast-to-Noise Ratio. 25, 66–69, 73, 76

**CR** Contrast Ratio. 25, 66, 68–70, 73, 76

**DAS** Delay-And-Sum. xiii, 2, 15–18, 27, 32–34, 39–44, 47–50, 54, 57–59, 61, 62, 64–69, 71–73, 75

**DSB** Digital Signal Processing. 20, 34, 35, 37

**FWHM** Full Width at Half Maximum. 44, 63

**gCNR** generalised Contrast-to-Noise Ratio. 25, 26, 66, 68–70, 73, 76

**MLA** Multi-Line Acquisition. 19, 30, 31, 38–40, 48

**PDF** Probability Density Function. 25, 26

**ROI** Region Of Interest. 25, 66–70, 73, 76

**UiO** University of Oslo. 2, 34–36

**USTB** Ultrasound Toolbox. 32–37





# Chapter 1

## Introduction

### 1.1 Motivation

When creating medical ultrasound images, a probe transmits ultrasound waves into the body and then receives backscattered waves, where the amplitude depends on the physical properties of the tissue. The probe gathers these waves which are converted into electrical time signals. Then, an ultrasound signal processing system reconstructs these signals into images, before a final image sequence is displayed. By applying beamforming algorithms, the received probe element signals from the echoes are combined, by means of applying delays and weights to the individual sensor signals before they are stacked. Beamforming can be seen as a way to control how the signals are transmitted, or received by the probe, or both, and processed by the system, to provide a high-quality ultrasound image.

Beamformers can be divided into two categories; conventional beamformers and adaptive beamformers (Hoel Rindal, Austeng, and Rodriguez-Molares, 2020). Conventional beamformers are beamformers that combine the signals of a beam, by applying a predetermined set of weights and time delays. The delays are designed to increase the amplitude of the waves received from a certain direction. Adaptive beamformers, which are studied in more detail in this thesis, use additional information from the received data to adapt the element weights. This adaptation is hence based on the properties and statistics of the signal (Grythe, 2015).

The main conventional beamformer is the Delay-And-Sum beamformer, where the delays applied to the received signal at each sensor then follow a pre-determined model that does not depend directly on the received data. The sensor signals can also be weighted based on pre-determined window functions, such as Hamming, and Hanning (Agarwal, Tomar, and Kumar, 2021), before they are summed to form the beamformer output signal. A general drawback of the DAS beamformer is related to the width of the main lobe, which affects, e.g., how well-resolved targets in the examined scene are.

By applying adaptive beamforming, it is possible to achieve higher image resolution, due to the fact that the beamformer adapts to the actual scene and received data. There exist several adaptive beamformers, one being the Capon beamformer. The Capon beamformer is a minimum-variance beamformer which increases image resolution. The algorithm involves estimating and inverting the spatial covariance matrix which describes the relation between the data received on each probe element. The estimation and inversion of this matrix are not necessarily robust, as there might be only few time samples available for the estimation related to reconstructing a single image pixel. The limited amount of time samples and the associated uncertainty in a temporally averaged covariance matrix can result in a poor adaptive beamformer output, further resulting in decreased image quality and undesirable artefacts.

To avoid misleading information in the ultrasound image, a robust estimate and inversion of the covariance matrix should be performed for each pixel reconstruction. The estimate can be made more robust using several methods. The Digital Signal Processing group at the Department of Informatics, University of Oslo (UiO), has in previous studies applied several methods, such as averaging in time, averaging in space through subarray averaging, and adding a diagonal load. These approaches are described in depth in Synnevåg, Austeng, and Holm (2009a) and Synnevåg, Austeng, and Holm (2007a).

The methods of averaging in time and space enhance the estimate of the covariance matrix by increasing the accuracy of the signal estimate. It is generally important not to apply excessive averaging in time beyond the length of the ultrasound pulse, as this will lead to decreased image resolution in the range direction, and

## 1.2 Aims of thesis

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therefore to a blurred image.

As part of the Capon beamformer algorithm, the covariance matrix has to be inverted. The estimated covariance matrix can be non-invertible, or close to non-invertible, which results in an inaccurate result. By applying a diagonal load to the matrix, the invertibility of the matrix increases. It is desirable to avoid adding a too large amount of diagonal loading, as this may alter the statistics of the data described by the estimated covariance matrix. An excessive diagonal load results in a beamformer output that approaches the Delay-And-Sum beamformer output (Synnevåg, Austeng, and Holm, 2007b). The necessary amount of diagonal loading depends on the invertibility of the covariance matrix estimate, as well as robustness to errors. The latter is not part of the scope of this thesis. By looking at the condition number of the covariance matrix estimate, it is possible to estimate the amount of diagonal loading needed to attain a stable inversion of the covariance matrix, as the condition number of a matrix is a measure of matrix inversion stability.

## 1.2 Aims of thesis

Earlier research has provided insight and some rules of thumb with respect to the amount of subarray averaging and diagonal loading to apply. Still, there is no consensus within the ultrasound research community on the appropriate choices of subarray size and diagonal load. The overall objective of this thesis is to investigate the effects of these robustification methods and to provide additional guidelines on the choice of parameters.

An aim of this thesis is to examine the robustness of the Capon beamformer in terms of the covariance matrix estimation and inversion. This is done by analysing how the matrix concepts of eigenvalues and condition numbers can be used as a measurement of the stability of the estimate and inversion. This is done by measuring the resolution of the beamformer by analysing the separability of two scatterers in a simulated scene with varying distances between the scatterers. The resolution is measured when varying the subarray size and the applied diagonal load.

Another aim is to further analyze how robustification methods, such as subarray averaging and diagonal load, applied to the covariance matrix estimate affect a set of image contrast metrics. This is done using data from a simulated scene containing an anechoic cyst.

The following research questions are addressed in this thesis:

- Can the amount of diagonal load be determined based on the condition number of the covariance matrix estimate?
- Can the eigenvalues of the covariance matrix provide additional information on the resolution and separability of scatterers in a scene?
- To what extent are the eigenvalues of the estimated covariance matrix affected by applied subarray averaging and speckle?
- How is the separability of scatterers in a scene when applying the Capon beamformer affected by the amount of subarray averaging applied?
- How does applying speckle to the scene affect the scatter separability when varying the subarray size?
- How do diagonal loading and subarray averaging affect the contrast metrics in an anechoic cyst scene?
- How do the diagonal load and subarray averaging affect the contrast metrics when considering different regions of interest?

### 1.3 Thesis outline

In Chapter 2, fundamental concepts of wave physics, signal processing, and ultrasound imaging are introduced to provide the necessary background for understanding the work presented in this thesis. Some key aspects of applied linear algebra are also introduced.

Chapter 3 presents and illustrates the datasets utilized in this thesis, and how these datasets have been simulated. This chapter also presents the processing chain and the Ultrasound Toolbox, which is utilized in this thesis.

### 1.3 Thesis outline

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Chapter 4 presents and discusses the results obtained in this thesis. A comparison of two Capon beamformer algorithms is also provided, and a brief analysis of the general pixel resolution is done by examining the necessary amount of Multi-Line Acquisition. An analysis of the separability between two scatterers in a scene with no speckle is performed, before results are generated for scenes with speckle added to the data. Before these results are presented, a brief analysis of the speckle amplitude level is provided. Lastly, an analysis of contrast metrics for a simulated dataset of a cyst is done. Throughout this chapter, the results are discussed alongside their descriptions and display.

Finally, in Chapter 5, conclusions from the work presented in Chapter 4 are provided, answering the research questions listed above, and ideas for further work are discussed.



# Chapter 2

## Background

### 2.1 Sound waves and their properties

When researching medical ultrasound image reconstruction algorithms, it is important to understand the fundamentals of sound waves, how the waves propagate in mediums, and the properties of the waves. Some important physical properties and phenomena are the speed of sound in a medium, as well as the refraction, dispersion, absorption, and attenuation of a wave.

Sound waves which are audible for humans have a frequency between 20 Hz and 20 kHz, while ultrasound waves are inaudible and have a frequency above 20 kHz. Sound waves are caused by for instance vibrations at the source, and propagate through mediums as such, whether the medium is human tissue, air, or water. Audible sound waves and ultrasound waves used for imaging are pressure waves. Density and compressibility are the physical properties of the medium that decide at which *speed of sound* the pressure wave can propagate. For water, the speed of sound is typically 1500 m/s, and is mainly given by density, temperature, and salinity (Leroy, Robinson, and Goldsmith, 2008). The speed of sound in air is roughly 330 m/s (Pope, 1999).

For human tissue, the speed of sound lies between 1430 m/s and 1600 m/s depending on the tissue (Pope, 1999). A typical value applied as the speed

of sound in tissue in medical ultrasound imaging is 1540 m/s. The speed of sound often varies in human tissue within an examination area, depending on the composition of tissues (Duck, 2012). The speed of sound can also depend on the frequency. This phenomenon is denoted as *dispersion*.

When the wave encounters an area with a new acoustic impedance, e.g., moving from one kind of tissue to another, parts of the wave may be reflected at the interface. An example is shown in Figure 2.1. Acoustic impedance is the opposition a wave encounters in a medium, and is given by

$$Z = \rho c, \tag{2.1}$$

where  $Z$  is the acoustical impedance,  $\rho$  denotes the density of the medium, and  $c$  is the speed of sound within the medium. The reflection of an ultrasound wave traversing the interface between two media at normal incidence is given by the intensity reflection coefficient (Pope, 1999)

$$\alpha = \frac{(Z_2 - Z_1)^2}{(Z_2 + Z_1)^2}. \tag{2.2}$$

When the difference in acoustical impedance is large,  $\alpha$  will be large. A large  $\alpha$  means a large amount of the wave will be reflected at the interface.

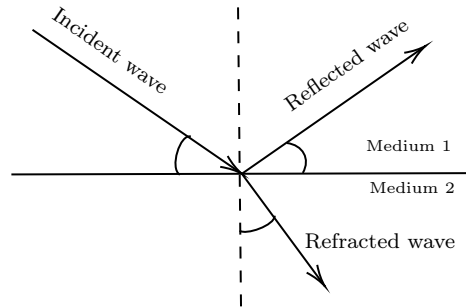
The different tissues may also *attenuate* the sound wave. Attenuation of a sound wave is a loss of wave amplitude when travelling through the tissue, as shown in Figure 2.2. Attenuation depends on properties such as density and speed of sound in the medium, and how much of the wave is refracted throughout the medium. When the ultrasound wave encounters a tissue with high stiffness, such as bone, the attenuation of the wave will be high (Pope, 1999). The attenuation of a wave when the wave travels through the medium is the main limitation of how far an ultrasound beam can travel.

Bone has a speed of sound of 4080 m/s (Pope, 1999) and a high density, which reflects most of the sound wave, which may result in a black shadow in an ultrasound image. This black shadow appears because most of the wave is

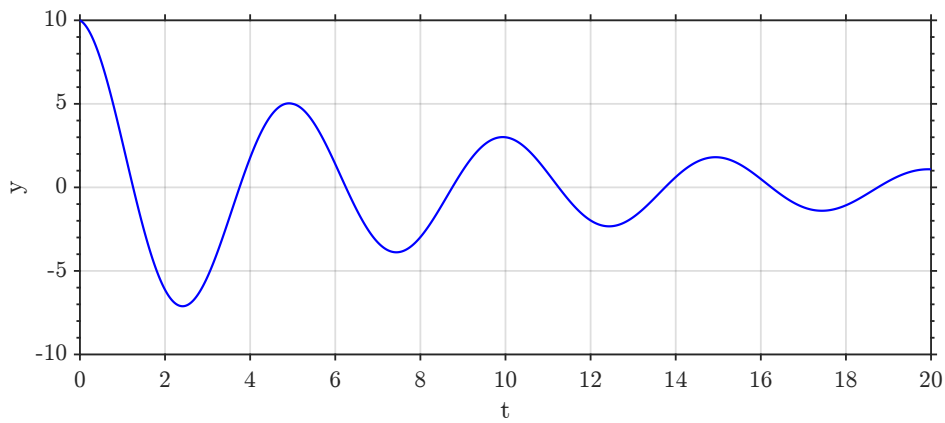


## 2.1 Sound waves and their properties

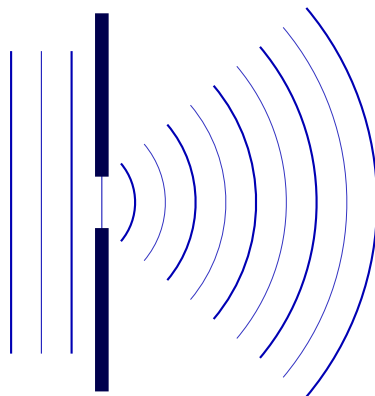
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**Figure 2.1:** An example of the refraction and reflection of a wave when coming upon the interface between two mediums.



**Figure 2.2:** An example of an attenuated signal.



**Figure 2.3:** An example of the diffraction of plane waves through a slit.

reflected, resulting in less energy being transmitted to the area behind the bone. Even though the wave may be fully reflected at the encountered surface of the bone, it is often possible to receive energy from the area behind it. The energy received depends on the size of the obstacle, in this example the bone, and is due to *diffraction* (Pope, 1999). Diffraction is the reason why a wave can “bend” around the corner of an object. An example of diffraction is given in Figure 2.3, which shows the diffraction of waves sent through a slit.

## 2.2 Digital sampling of physical signals

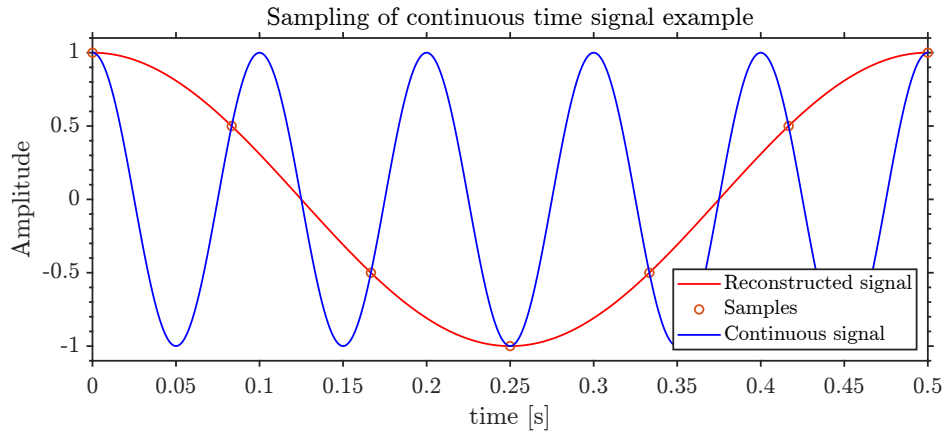
A physical signal, such as a sound wave, present in a medium can be recorded by an instrument, such as a hydrophone, a microphone or an ultrasound probe. The signal is then converted into a continuous electrical signal. This electrical signal is in turn converted to a digital signal through sampling. Sampling consists of collecting the value of a signal at points in time, i.e. converting a continuous-time signal to a discrete-time signal. This is typically done at a uniform sampling interval.

Conversion of a continuous-time signal to a discrete-time signal can be done without loss of information as long as the sampling of the continuous-time signal is sufficient. A sufficient sampling of a signal, according to the *Shannon-Nyquist sampling theorem* (Shannon, 1949), is to sample the signal using a sampling frequency at least twice as large as the highest frequency present in the signal. That is

$$f_{\text{sampling}} \geq 2 \cdot f_{\text{max}}. \quad (2.3)$$

This corresponds to sampling at least two samples per period of the highest-frequency signal component. This is done to prevent the recreation of the wrong signal.

One example of a badly sampled signal is shown in Figure 2.4 on the next page. The figure shows a well-sampled continuous-time signal, and the effect of sampling using a too small sampling frequency compared to the frequency in the signal. A physical recreation of the example in Figure 2.4 would result in a frequency of 2 Hz. This misrepresentation of a signal is called *aliasing* and is a limitation in sampling. Other limitations are noise in the sampled signal, e.g., noise in the form of quantification errors, or electrical noise from the probe. To be able to accurately reconstruct the original signal after sampling, it is important to satisfy the Shannon-Nyquist sampling theorem.



**Figure 2.4:** Example of poor sampling of a continuous-time signal. The blue line is a well-sampled discretisation of a continuous-time, periodic cosine signal with 10Hz. The red line is a possible reconstruction of the signal from the red samples when using a sampling frequency  $f_s = 1.2 \cdot f_{\text{high}} = 12$  Hz. The circles indicate where the samples are collected using a sampling frequency of  $f_s = 12$  Hz.

## 2.3 What is ultrasound imaging?

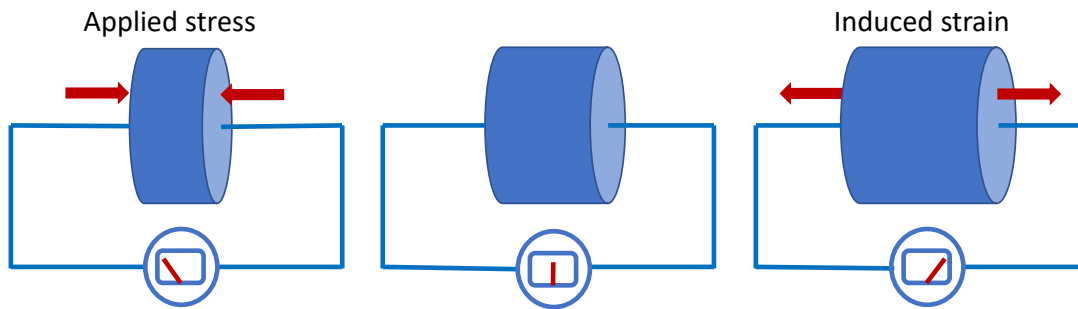
Ultrasound pulse-echo imaging is the process of transmitting and receiving ultrasound waves using a probe, before processing the received signal through various beamforming methods. Some beamforming methods will be presented in Section 2.4. The process of transmitting, receiving, and processing ultrasound waves is repeated quickly and continuously, which creates a live view of the imaged area.

An ultrasound probe is typically a set of elements made of piezoelectric materials and is used to transmit and receive ultrasound waves. Piezoelectric materials are ceramic crystals that vibrate when an alternating voltage is imposed on the material, as shown in Figure 2.5. This vibration generates pressure waves to be transmitted through a medium. Properties such as the frequency and strength of the alternating voltage imposed on the piezoelectric material decide the properties of the transmitted wave.

When the wave has been transmitted, it travels through the medium, for instance, the human body. The wave propagation changes when the wave encounters objects in the medium with different physical properties. A part of the wave is then

## 2.3 What is ultrasound imaging?

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**Figure 2.5:** An example of how piezoelectric material expands and compresses when an alternating voltage is applied.

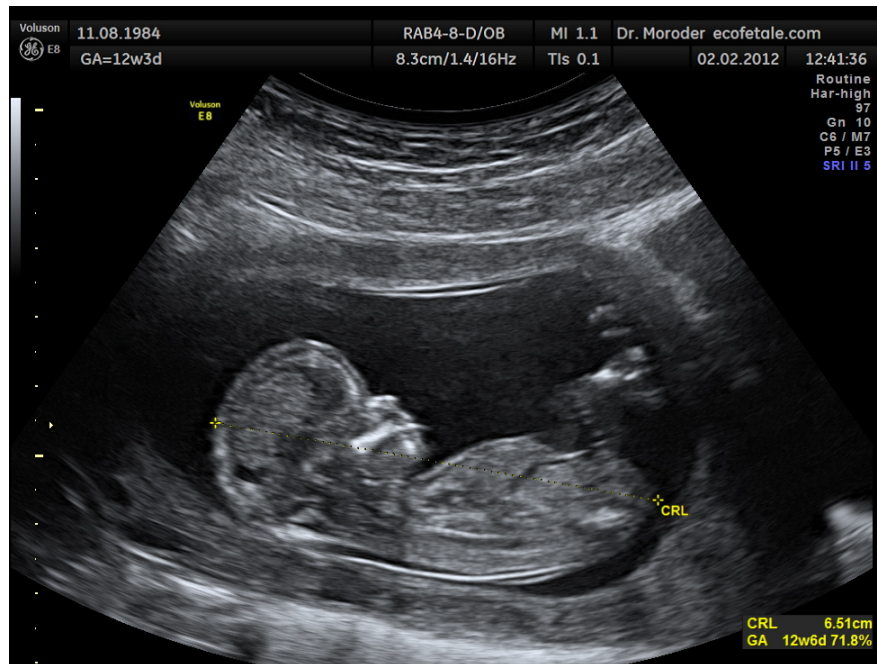
reflected.

The reflected waves, i.e. the echo, are received by the probe, and the wave is converted to digital signals, see Section 2.2. This conversion is the opposite of the transmission process, where the piezoelectric material vibrates due to the received wave, and is converted into an electric signal. This received electric signal is then digitally sampled. The digital signal is then processed by applying various beamforming algorithms, which will be presented in Section 2.4, generating a greyscale image, showing the final result of the ultrasound examination. A general example of an ultrasound image is shown in Figure 2.6 on the next page.

In general, when working with a phased array probe, an ultrasound image is created by transmitting and receiving a given number of beams within a sector. The received signals are then compounded into a polar image of  $t_x \times r_x$  pixels in cylindrical coordinates, i.e. the number of angular pixels is given by the amount of transmits  $t_x$ , and the number of range pixels is given by the number of recorded time samples  $r_x$  from each transmit. To view the image on a computer screen, the polar image is scan converted into a cartesian grid image that is being displayed.

### 2.3.1 Speckle

In most medical ultrasound images recorded in a physical medium, there will be noise-like signals called speckle. Speckle patterns in physical media are a result of natural inhomogeneity of the human tissue, i.e. reflections from structures that are



**Figure 2.6:** An example of an ultrasound image. The example shows an ultrasound image of a fetus in a sagittal scan at 30 weeks of pregnancy. The image is adapted from Moroder, 2012.

random and smaller than the size of the resolution cell (see Section 2.5.1). Speckle is then the sum of coherent and incoherent signals from at least 10 scatterers inside a resolution cell (Jensen and Nikolov, 2000). In ordinary ultrasound imaging, speckle is unavoidable due to the tissue inhomogeneity. A perfectly homogeneous medium will theoretically give a speckle-less result.

### 2.3.2 Near-field and far-field

All medical ultrasound imaging is applied in the near-field, where the transmitted wave maintains its spherical shape. In medical ultrasound imaging, a focused beam is applied. A focused beam will have a *focal zone*, and the shape of the beam transforms into an hourglass shape. The focal zone is the zone where the beam achieves the highest resolution.

## 2.4 Image reconstruction and beamforming

Beamforming can be seen as a way of controlling how the signals are transmitted or received or both. In the transmit process, beamforming is done by altering how the ultrasound wave is sent from each element. When the signal is received, the signal recorded on each element in the probe is transformed from a domain of position and time to  $x$  and  $y$  coordinates in an image. Beamforming is a form of spatial filtering.

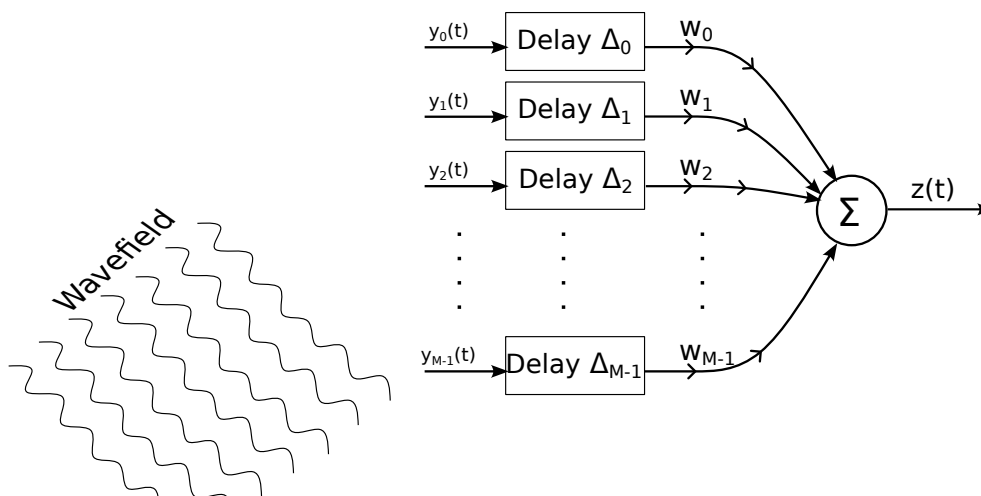
Beamformers are typically divided into two categories, namely conventional and adaptive beamformers (Rindal, 2019). The main beamformer applied in time is the DAS beamformer. DAS applies a set of delays and weights to the signal recorded at each element and sums these over all elements.

Adaptive beamformers are all beamformers where the signal alters how the beamforming is performed. This alteration is performed by continuously using information from the time signal to adapt, e.g., the element weights applied to each channel before summation. In addition, several beamformers exist that include a form of image weights, for instant coherence factor and phased coherence factor (Rindal, 2019). These act more like image-processing techniques. Such techniques will not be further discussed in this thesis.

### 2.4.1 Conventional beamforming – the Delay-And-Sum beamformer

Conventional beamforming is beamforming which applies a predetermined set of weights and delays at each element through both the transmit and receive beamforming process. As mentioned above, DAS is the main conventional beamformer. As the name implies, the delay is applied to the signal received at each element, and then all the signals are summed. A visual example of the DAS beamformer is shown in Figure 2.7.

When the reflections from interfaces in the scene are reflected back to the probe array, the transmitted signal arrives at different times over the array. This is due to the difference in travel time from the outer and middle array elements to a



**Figure 2.7:** A visualisation of the receiving part of the DAS algorithm, mathematically shown in Equation (2.4). The wave arrives at the elements at different times. The signal at each element is then delayed before being summed.

point in the tissue and back.

The difference in travel time can be compensated for by applying delays to the received signal from the probe elements. A set of optimal delays will result in the alignment of the signals at each element, so the summation of the received and delayed signals is optimized. This will result in the amplification of the signal received in the direction corresponding to the delays.

The DAS output is defined as

$$z_{\text{DAS}}(t) = \sum_{m=0}^{M-1} w_m y_m(t - \Delta_{m,t}), \quad (2.4)$$

where  $M$  denotes the number of elements in the array and  $m$  denotes the specific element number.  $w_m$  is the weight applied to the signal  $y_m$  from the element which has been delayed by the delay  $\Delta_{m,t}$ .

Equation (2.4) may also be written in a compact vector form;



## 2.4 Image reconstruction and beamforming

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$$z_{\text{DAS}} = \mathbf{w}Y. \quad (2.5)$$

$\mathbf{w}$  is then the weight vector containing the weights at each element, and  $Y$  contains the delayed signal recorded at each element.

### Expanding aperture

The lateral image resolution is dependent on aperture size and depth, and a general goal is to achieve a uniform resolution in the entire image. One aid to help achieve this is to utilize an expanding aperture. The resolution along the lateral axis is given by

$$x_{\text{res}} = z\Theta_{6\text{dB}} = z\frac{1.21\lambda}{D} = 1.21\lambda f\#. \quad (2.6)$$

The f-number, denoted  $f\#$ , is the ratio between the depth  $z$  and aperture size  $D$  (Rindal, 2019). Hence, in order to have a constant lateral resolution at all depths, the active aperture size is increased linearly as the imaging depth increases.

### 2.4.2 Adaptive beamforming – the Capon beamformer

An adaptive beamformer usually has a specific criteria to fulfil. One example of this criterion is to update the element weights by minimizing the expected output power of DAS, which is the minimum-variance beamformer, alternatively named the Capon beamformer (Capon, 1969). The Capon beamformer is typically a secondary step of DAS, however only the step of applying a set of delays to the received signals. The information from each element is then used to estimate the signal, and to update and adapt the beamformer continuously.

The beamformer estimates the weight vectors by estimating the signal, as mentioned above. The signal is estimated using what is called a *covariance matrix*. The aim of the minimum-variance Capon beamformer is to minimize noise and interference from other directions while maintaining the gain from the direction of interest, often denoted as the steering vector.

The DAS is applied as an initial step by applying a set of delays, but not summing the received signals. Then, the signal is received, and the covariance matrix  $R$ , which will be presented shortly, is estimated through the derivation of DAS from Equation (2.5):

$$E\{|z_{\text{DAS}}|^2\} = E\{\mathbf{w}^H Y\}^2 = E\{\mathbf{w}^H Y Y^H \mathbf{w}\}^2 = \mathbf{w}^H E\{Y Y^H\}^2 \mathbf{w} = \mathbf{w}^H \mathbf{R} \mathbf{w}. \quad (2.7)$$

The Capon beamformer is expressed as the solution to the following minimization problem:

$$\min_{\mathbf{w}} (\mathbf{w}^H \mathbf{R} \mathbf{w}), \quad \text{subject to} \quad \mathbf{w}^H \mathbf{a} = 1. \quad (2.8)$$

The analytical solution to the optimization problem above is (Synnevåg, Austeng, and Holm, 2009b)

$$\mathbf{w}_0 = \frac{\mathbf{R}^{-1} \mathbf{a}_0}{\mathbf{a}_0^H \mathbf{R}^{-1} \mathbf{a}_0}, \quad (2.9)$$

where  $\mathbf{a}_0$  is the steering vector. The steering vector gives information about the location parameters of the sources (Li, Stoica, and Wang, 2003). When the delays have been applied, the steering vector is simply a vector consisting of ones. The weight set from Equation (2.9) is then applied to a snapshot of time samples at all elements, resulting in the amplitude Capon beamformer output.

The power Capon beamforming method is derived by inserting Equation (2.9) as the set of weights applied in Equation (2.7), and leads to (Lukose and Mathurakani, 2016)

$$P_{\text{Capon}} = \mathbf{w}^H \mathbf{R} \mathbf{w} = \frac{1}{\mathbf{a}_0^H \mathbf{R}^{-1} \mathbf{a}_0}. \quad (2.10)$$

*The covariance matrix  $R$*  is a matrix containing a measure of the similarity between two samples recorded at two elements. By this, it is meant that the matrix element

## 2.4 Image reconstruction and beamforming

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$(m_1, m_2)$  in  $R$  represents the similarity between two samples at the probe elements  $m_1$  and  $m_2$ ;

$$R_{m_1, m_2} = \text{cov}(y_{m_1}(t), y_{m_2}(t)). \quad (2.11)$$

Here,  $x_m$  is the signal recorded at array element  $m$ , and  $t$  is the time sample in question.

The estimate of the covariance matrix, denoted as  $\hat{R}$ , is typically given as

$$\hat{R} = \frac{1}{N} \sum_{n=0}^N \mathbf{y}_m[n] \mathbf{y}_m[n]^H. \quad (2.12)$$

Here,  $N$  is the number of time samples,  $\mathbf{y}_m$  is the recorded signal at each sensor element, and  $\mathbf{y}_m^H$  denotes the complex conjugate transpose.

### 2.4.3 Multi-Line Acquisition

Multi-Line Acquisition (MLA) is a method frequently used in medical ultrasound imaging. The MLA method involves acquiring multiple received beams for each transmitted beam. There will normally be an equal amount of transmitted and received beams when no MLA is applied. When applying MLA to a set scenario, there are two main applications.

One implementation is to transmit fewer and wider non-overlapping beams while receiving an increased number of beams than transmitted beams. This implementation increases the frame rate while keeping the resulting amount of received beams the same. One drawback of this implementation is, however, the possible loss of resolution. When increasing the transmitted beamwidth, the axial resolution decreases. Targets in the scene may be missed due to the increased beamwidth, and even though the MLA method increases the axial resolution, it does not restore missed targets. Hence, MLA applied through this implementation increases the frame rate, but not necessarily the image quality.

The second main implementation is to transmit wider, overlapping beams while

receiving at an increased number of elements. This implementation does not increase the frame rate, but increases the image quality regarding the transmitted focus.

#### 2.4.4 Robustification

Adaptive beamformers, such as the Capon beamformer, are typically sensitive to errors on the probe array that effectively becomes an error in the applied steering or time delay. As mentioned in section 2.4.2, the Capon beamformer increases resolution by using an estimate of the covariance matrix. The estimate of the covariance matrix is based on the weighted time-signal, i.e. the result of applying a set of delays to the transmit dimension.

The covariance matrix estimate is based on rather short, non-stationary time-signals. One risk when working with such signals is that the signal model adapts poorly to the signal, as the signal model will not receive a sufficient amount of information about the signal. The estimation of the covariance matrix may then be challenging. However, by applying methods such as subarray averaging or time averaging, the estimation of the covariance matrix is robustified.

To avoid challenges with generally poor estimations of the covariance matrix, a robust estimate of the covariance matrix has to be found. As mentioned in the introduction, the Digital Signal Processing (DSB) group at the Institute of Informatics, University of Oslo, has applied some methods such as averaging in time and space, or adding a diagonal load to the covariance matrix. These have been presented by Synnevåg, Austeng, and Holm (2009a) and Synnevåg, Austeng, and Holm (2007a).

#### Time averaging

Ultrasound pulses are, as mentioned, non-stationary, meaning the covariance matrix estimate is based on only a short time snapshot. The time limitation may cause the covariance matrix estimate to be affected by speckle. This will further increase the variance of the covariance matrix estimate. By applying time averaging, the influence of speckle is reduced, causing the variance to be reduced,

## 2.4 Image reconstruction and beamforming

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ensuring a more stable pixel value.

As for all averaging methods, it is desired to average as much as possible, within reason. The maximum amount of averaging possible to apply is physically limited due to the non-stationarity of the pulse. However, it can be chosen based on the pulse length, as each part of the pulse may contain information about the imaged scene. The averaging is applied to the covariance matrix estimate, resulting in a stabilized estimate of each pixel value, as the pixel estimate will be less exposed to the variance of the speckle in the recorded scene.

### Subarray averaging

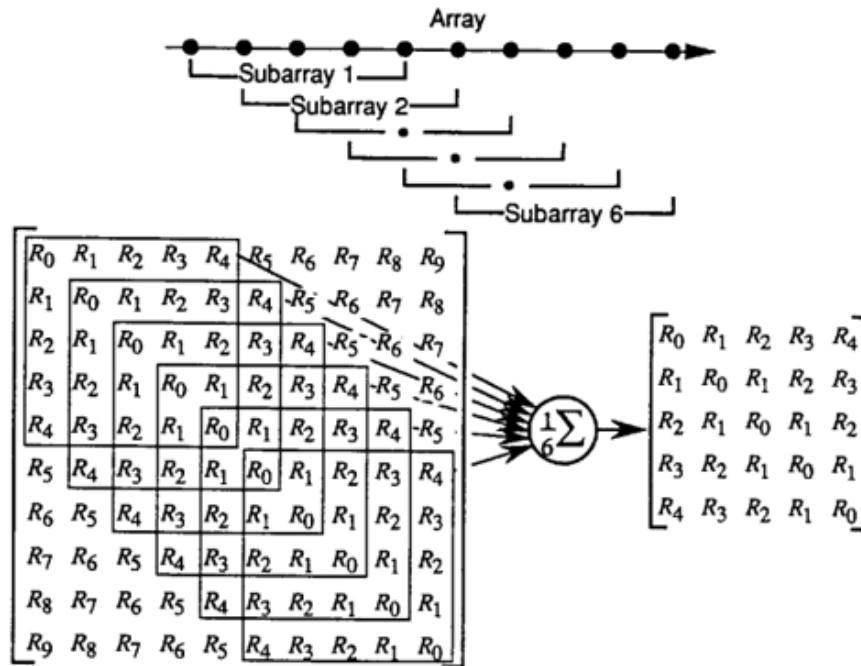
Subarray averaging is a space averaging method used to robustify the estimated covariance matrix, and is the main method to reduce the effect of coherent sources (Synnevåg, Austeng, and Holm, 2009b). Statistically, the information from all parts of the probe array is similar when delays have been applied. A spatially averaged covariance matrix will then, theoretically, contain the same information about the pulse and noise as the full covariance matrix.

For a full array consisting of  $M$  elements, one can divide the full array into  $P = (M - L + 1)$  subarrays of a length of  $L$  elements, and then apply adaptive beamforming to the subarrays. The covariance matrix estimation from Equation (2.12) is then averaged over all subarrays, and temporally averaged over  $2K + 1$  time samples, as such (Synnevåg, Austeng, and Holm, 2009b, Eq.8):

$$\hat{R} = \frac{1}{(2K + 1)(M - L + 1)} \sum_{k=-K}^K \sum_{l=0}^{M-L} \mathbf{y}_l[n - k] \mathbf{y}_l[n - k]^H. \quad (2.13)$$

An example of subarray averaging is shown in Figure 2.8, where an array with 10 elements is divided into 6 subarrays with a length of 5 elements. The covariance matrix estimate is then reduced from a  $10 \times 10$  matrix to a  $6 \times 6$  matrix.

When determining the subarray size  $L$ , there is a trade-off between the resolution and robustness of the Capon beamformer. A smaller subarray size robustifies the estimate of the covariance matrix, but it does reduce the resolution (Synnevåg,



**Figure 2.8:** A visualisation of the subarray averaging method. The original array has 10 elements, and the subarrays have a length of 5 elements. The covariance matrix is then reduced from a  $10 \times 10$  matrix, to a  $5 \times 5$  matrix through averaging. Figure borrowed from Johnson and Dudgeon, 1992, Fig.4.35.

Austeng, and Holm, 2007b).

### Diagonal loading

In the calculation of the weights (Equation (2.9)), the covariance matrix  $R$  is to be inverted. However, it is possible for the sample covariance matrix to be a non-invertible matrix. A non-invertible matrix does not have full rank, and will result in a matrix with infinitely positive and negative values in the matrix. A close-to-non-invertible matrix will give similar results. The result of the inversion of such matrices will result in an inaccurate inversion. The cause of non-inversion is the singularity of a matrix.

A *singular matrix* is a square matrix where the determinant is 0. In the case of a 2x2 matrix  $A = \begin{pmatrix} a & b \\ c & d \end{pmatrix}$ , the determinant is defined as (D. C. Lay, S. R. Lay, and McDonald, 2016)

## 2.5 Image metrics

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$$\det(A) = ad - bc. \quad (2.14)$$

Further, the inversion of the matrix  $A$  is defined as

$$A^{-1} = \frac{1}{\det(A)} \begin{pmatrix} a & b \\ c & d \end{pmatrix}. \quad (2.15)$$

The inversion is not defined if the determinant (Equation (2.14)) is zero. This is the case of singular matrices, as division by zero is not defined.

One way to make the inversion of the matrix is to add a diagonal load to the matrix, as such

$$\tilde{A} = A + DL * I. \quad (2.16)$$

Here,  $A$  denotes the matrix in question,  $\tilde{A}$  denotes the modified matrix, and  $I$  is the identity matrix, a matrix with ones at the diagonal and zeros elsewhere. The diagonal load is denoted  $DL$ , and is in this thesis defined as

$$DL = \text{regCoeff} \cdot \frac{\text{tr}(A)}{\text{dim}(A)}, \quad (2.17)$$

where  $\text{regCoeff}$  is a fraction representing the percentage of diagonal load factor applied,  $\text{dim}$  denotes the dimension of the matrix  $A$ , and  $\text{tr}$  is the trace of the matrix  $A$ , i.e. the sum of all diagonal elements of the matrix. The applied diagonal load reduces the singularity of the matrix, which in return increases the stability of the inversion.

## 2.5 Image metrics

The quality of an ultrasound image can be measured using different image metrics, such as resolution and contrast.

### 2.5.1 Resolution

One important image metric is the resolution, where a well-resolved image is desired to be able to distinguish all details from the imaged scene in the image. The depth resolution is given by

$$\delta r = \frac{c}{2B}, \quad (2.18)$$

and the lateral resolution is given by

$$\delta\beta = \frac{\lambda}{D}, \quad (2.19)$$

where  $c$  denotes the speed of sound in the medium,  $\lambda$  denotes the wavelength,  $B$  is the bandwidth of the signal, and  $D$  is the length of the array. Equation (2.18) and Equation (2.19) denotes the size of the resolution cell, i.e. the area in which two targets located within cannot be separated by the system.

To ensure a correctly sampled wave field, the Nyquist sampling theorem gives the following two-way ( $tr$ ) beam spacing constraint to be fulfilled (Bjåstad, 2009, Eq. 1.9)

$$\delta\theta_{\text{Nyquist}}^{tr} = \frac{\lambda}{D_{tx} + D_{rx}} [\text{rad}]. \quad (2.20)$$

Here,  $D_{tx}$  and  $D_{rx}$  denote the physical length of respectively the transmit and receive arrays.

One way to measure the resolution of an imaging system is to analyse a well designed scene, such as an image of a set of scatterers. By doing this, it is possible to measure the separability of scatterers in the scene, where the separability is described by a resolution criteria.

By analysing the intensity dip between two points in a scene, it is possible to describe whether or not the scatterers are separated by the beamformer. One criterion is the *-6 dB limit*, where the scatterers are considered separated if the



## 2.5 Image metrics

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amplitude dip between the two scatterers is 6 dB lower than the amplitude of the scatterers.

Another separability criterion is *the Sparrow limit*. Here, the scatterers are considered separated if the amplitude between the scatterers shows any dip.

### 2.5.2 Contrast

#### Contrast Ratio

Contrast Ratio (CR) is a ratio between the mean background value and the mean value in a Region Of Interest (ROI), and is defined as (Smith, Lopez, and Bodine, 1985)

$$\text{CR} = \frac{\mu_{\text{ROI}}}{\mu_{\text{background}}}. \quad (2.21)$$

Here,  $\mu$  denotes the expected value of the power of the beamformed signal, i.e.  $\mu = E\{|b|^2\}$  of the ROI and the background region.

#### Contrast-to-Noise Ratio

The Contrast-to-Noise Ratio (CNR) was first introduced by Patterson and Foster (1983) as *contrast-to-speckle ratio*, later denoted as contrast-to-noise ratio. The CNR is defined as

$$\text{CNR} = \frac{|\mu_{\text{ROI}} - \mu_{\text{background}}|}{\sqrt{\sigma_{\text{ROI}}^2 + \sigma_{\text{background}}^2}}, \quad (2.22)$$

where  $\sigma = \sqrt{E\{|b|^2 - \mu^2\}}$  is the standard deviation of the power of the beamformed signal in a set region.

#### Generalized Contrast-to-Noise Ratio

The generalised Contrast-to-Noise Ratio (gCNR) is related to the total overlap area of the Probability Density Function (PDF) derived from the two regions ROI

and background region (Rodriguez-Molares et al., 2020). The overlap OVL is given by

$$\text{OVL} = \int_{-\infty}^{\epsilon_0} p_{\text{ROI}}(x)dx + \int_{\epsilon_0}^{\infty} p_{\text{background}}(x)dx, \quad (2.23)$$

where  $\epsilon_0$  is the optimal threshold between the two PDFs to avoid misdetection between the two regions, and  $p$  denotes the PDF value. The relationship between the gCNR and OVL is (Rodriguez-Molares et al., 2020)

$$\text{gCNR} = 1 - \text{OVL}. \quad (2.24)$$

## 2.6 Linear algebra aspects

### 2.6.1 Eigenvalues of a matrix

If one has a probe with  $M$  elements which receive a set of waves, the received signal can be viewed in two ways. One way is to consider the signal recorded at each element as a function of time, while another way is to consider one time sample and a snapshot from each sensor. In the case of considering a snapshot of all elements, these samples can be used in the estimation of the covariance matrix  $R$ . From  $R$ , it is possible to calculate the eigenvalues, which will return  $m$  values from an  $m \times m$  matrix. When considering an analysis as such, the number of significant eigenvalues will represent the number of significant waves received at the probe elements. The vector space spanned by the eigenvectors having a corresponding significant eigenvalue represents the space spanned by the significant waves received.

A wave arriving from the far-field arrives as a plane wave. This can be described with one eigenvector and the corresponding eigenvalue. An analysis of the eigenvalues will then show a significant eigenvalue, representing the plane wave. When speckle is present in the data, the eigenvalues will show an increased level of small eigenvalues, as there is more signal present in the data.

## 2.6 Linear algebra aspects

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A wave arriving from the near-field will be a spherical wave and will arrive at the probe slightly curved. Depending on the curvature, this may result in a few or several significant eigenvalues with corresponding eigenvectors.

The size of the array will affect the size of the covariance matrix, which in turn affects the number of eigenvalues, as the size of the array coincides with the number of eigenvalues. A near-field wave received by the probe has a spherical shape until the delay step of DAS has been applied. If the applied delays return a perfect plane wave, there will be one set of eigenvalues representing the wave. However, if the estimate of delays is not perfectly accurate, or the wave arrives from a direction slightly off from the direction of interest, the received wave may still have a slightly curved wavefront. A full array is then more affected by this curved wavefront than a smaller fraction of the full array, which in turn may increase the number of significant eigenvalues.

The magnitude of the eigenvalues also increases when the subarray size increases, as a larger array receives more energy from the wave.

### 2.6.2 Condition number of a matrix

The condition number of a matrix is a measure of how sensitive the result is to variations in the input data. A matrix can be defined as well- or ill-conditioned. A well-conditioned matrix will return a more stable result of the matrix inversion (Equation (2.15)) (Cheney and Kincaid, 2007, p. 321) (D. C. Lay, S. R. Lay, and McDonald, 2016, p. 132).

A 2-norm condition number of a square matrix  $A$  is defined as (Cline, Moler, Stewart, and Wilkinson, 1979, Eq.1.6)

$$k(A) = \|A\| \|A^{-1}\|. \quad (2.25)$$

This can be written as

$$k(A) = \frac{\lambda_{\max}}{\lambda_{\min}}. \quad (2.26)$$

If the eigenvalues are particularly large or small, round-off errors in the calculations are inevitable. Round-off errors occur if the value is getting infinitely big or small, which is the situation when dividing a value by a respectively very small or large value.

When the covariance matrix is singular, the condition number will be infinitely high. When working with sample covariance matrices in the case of beamforming, this can for instance happen with simulated data sets with no speckle, in pixels with a value of 0, or if the sources are coherent. Then, a slight change in the covariance matrix element will have a big effect on the stability of the matrix inversion. Matrix inversion and non-invertible matrices have been presented in Section 2.4.4.

The condition number represents the stability of the matrix inversion, where  $\log_{10}(k(A))$  is an estimation of the loss of precision. This is an upper boundary, representing the number of digits that may be miscalculated. For example,  $k(A) = 10^5$  gives an upper boundary of 5 digits. What is considered a too large condition number depends on the precision of the calculation tool applied. *MATLAB* has a precision of 16 digits, meaning a matrix with a condition number above  $10^{16}$  is an ill-conditioned matrix (Cheney and Kincaid, 2007).

One way to avoid an infinitely high condition number is to apply a robustification method, the most common method being adding a diagonal load to the matrix. The diagonal loading method was presented in Section 2.4.4.

# Chapter 3

## Methods

### 3.1 Datasets

The datasets used in this thesis are all simulated using *Field II Ultrasound Simulation Program* (2023), version 3.30. The probe is identical between all datasets. The simulated datasets have been generated both with and without speckle. A simulated speckle-less scene has initially been used when examining resolution, before speckle is added to repeat the resolution examinations.

#### 3.1.1 Setup of the probe, the scene, and beamforming parameters

All data has been simulated using the same probe, the *P4* phased array transducer with 64 elements (Vega, 2015). The transducer array properties are shown in Table 3.1, and the simulation parameters are listed in Table 3.2 on the following page. The element height of the probe is reduced from the original P4 phased array transducer, to counteract the elevation focus of the original probe. This is done to avoid introducing effects that could influence the results if not properly addressed.

In some cases, some beamforming parameters have been set. Table 3.3 on the next page contains the parameters set when beamforming. These parameters have

| Probe properties       | Value      |
|------------------------|------------|
| Element width          | 0.25 mm    |
| Element height         | 5 mm       |
| Speed of sound $c$     | 1540 mm    |
| Centre frequency $f_c$ | 2.56 MHz   |
| Pulse length           | 2.5 cycles |
| Number of elements     | 64         |

Table 3.1: Probe properties based on the P4 phased array transducer, applied when simulating scenes.

| Simulation parameter name  | Value                     |
|----------------------------|---------------------------|
| Number of transmits, $t_x$ | 128                       |
| Simulated depth            | 0 to 60 mm                |
| Simulated width            | $-30^\circ$ to $30^\circ$ |
| Depth of scatterers        | 30 mm                     |
| Depth of cyst              | 30 mm                     |
| Cyst radius                | 5 mm                      |

Table 3.2: The applied variables when simulating scenes.

| Parameter name                          | Value                     |
|---|---------------------------|
| Subarray size, $L$                      | $\frac{1}{3}M \approx 21$ |
| Diagonal load                           | 1 %                       |
| $f\#$                                   | 1.7                       |
| Expanding aperture window               | Boxcar                    |
| $MLA_{\text{resolution work}}$ factor   | $MLA = 3$                 |
| $MLA_{\text{contrast work}}$ factor     | $MLA = 1$                 |
| Lateral distance $d$ between scatterers | 2 mm                      |

Table 3.3: The default beamforming and dataset parameters when a value needs to be set for analysis.

### 3.1 Datasets

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been set based on earlier research and current standards.

Due to the expanding aperture (Section 2.4.1), the subarray size is adapted to keep the number of subarrays the same at each depth. At a depth of 30 mm, the adapted subarray size is  $L_{\text{new}} = 19$  elements, keeping the corresponding fraction of  $L = \frac{1}{3}M$ . The MLA factors listed are the factor of which the MLA increases the number of axial samples.

#### Choice of transmits, $t_x$

From Equation (2.20), the Nyquist sampling theorem states that the beam spacing requirement is

$$\delta\theta_{\text{Nyquist}}^{\text{tr}} = \frac{\lambda}{D_{tx} + D_{rx}} = \frac{0.60156 \cdot 10^{-3}}{2 \cdot 64 \cdot 0.25 \cdot 10^{-3}} = 0.0188[\text{rad}] = 1.0771^\circ. \quad (3.1)$$

The full array size is then  $D_{tx} = D_{rx} = N_{\text{elements}} \cdot \text{element width}$ . Further, the relationship between the number of transmits,  $t_{x,\text{Nyquist}}$ , within a full scan for the Nyquist sampling theorem is

$$\theta_{\text{tot}} = \delta\theta_{\text{Nyquist}}^{\text{tr}} \cdot t_{x,\text{Nyquist}}, \quad (3.2)$$

which results in the following amount of necessary beams according to the Nyquist criterion

$$t_{x,\text{Nyquist}} = \frac{\theta_{\text{tot}}}{\delta\theta_{\text{Nyquist}}^{\text{tr}}} = \frac{60^\circ}{1.0771^\circ} = 55.7 \text{ transmits}. \quad (3.3)$$

By applying  $t_x = 128$  transmits, the constraint given by the Nyquist sampling theorem is met, and the pulse is correctly sampled. However,  $t_x = 128$  results in an oversampled signal in the lateral direction, which results in a visually pleasing beamformer output.

### 3.1.2 Simulations of the scenes containing scatterers

The simulated datasets have been simulated with points at a 30 mm depth and different lateral locations in the scene. One example of a beamformed scene without speckle is shown in Figure 3.1, a scene where there are two points located with a lateral distance of 2 mm. This specific simulated scene has been used when a set distance between the scatterers is necessary, e.g., when examining the effect of subarray averaging. When the lateral distance between the scatterers is altered for examination of scatter separability, the applied distances are 0, 0.5, 1, 1.5, 2, and 5 mm.

### 3.1.3 Simulation of speckle scene

Speckle is added to the simulated scenes to replicate a more realistic scene. The speckle simulation is done separately from the scatter simulation.

The speckle scene has been simulated by generating  $10^6$  randomly located scatterers in the full scene. This is to ensure statistically correct speckle, which is when there are 10 or more random scatter points within a resolution cell, as mentioned in Section 2.3.1. The simulated speckle dataset beamformed with the DAS beamformer in USTB is shown in Figure 3.2.

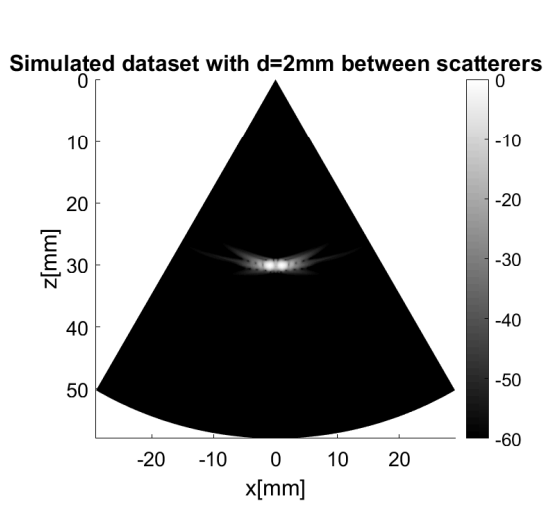
The result of combining the datasets visualised in Figure 3.1 and Figure 3.2 is shown in Figure 3.3. The speckle level is  $-30$  dB.

### 3.1.4 Simulated cyst target

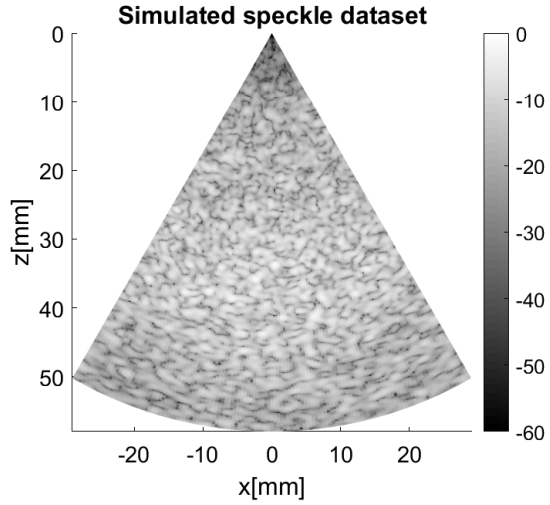
When measuring contrast, a simulated dataset containing an anechoic cyst is used. This is done using the same method as presented in Section 3.1.3. The cyst, visualised in Figure 3.4, is then placed by removing all random scatterers within a circle located at a depth of 30 mm, with a radius of 5 mm.



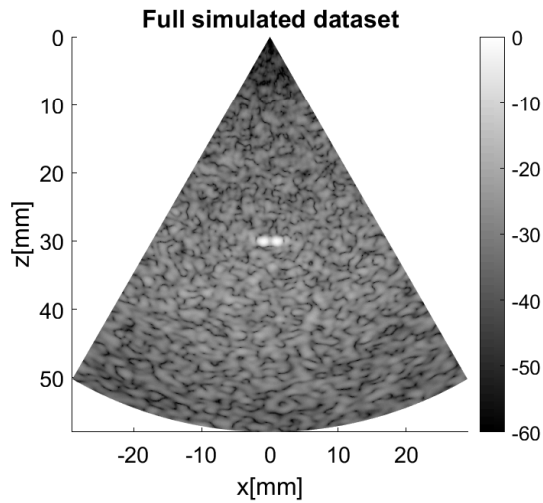
### 3.1 Datasets



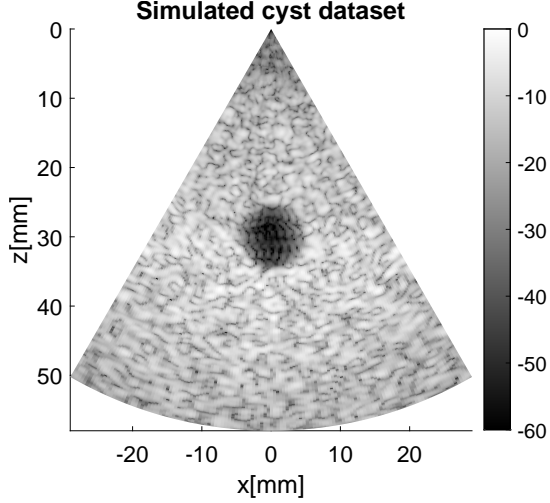
**Figure 3.1:** A simulated dataset of a speckle-less scene with two scatterers located at a depth of 30mm, with a distance of 2mm between them. The data has been beamformed using the implementation of the DAS beamformer in the Ultrasound Toolbox (USTB).



**Figure 3.2:** A simulated speckle dataset, beamformed using the DAS beamformer.



**Figure 3.3:** The beamformed result of the combined simulated dataset with two scatterers and simulated speckle. The two scatterers are located at a depth of 30mm, with a lateral distance of 2 mm. The speckle is at a level of  $-30$  dB. The full dataset has been beamformed using the DAS beamformer.



**Figure 3.4:** A simulated dataset containing a cyst located at a depth 30 mm, with a 5 mm radius. The data has been beamformed using the DAS beamformer.

## 3.2 The Ultrasound Toolbox

USTB is a MATLAB toolbox for processing ultrasonic signals. Its aim is to facilitate the comparison of imaging techniques and the dissemination of research results. USTB covers processing techniques for tissue and flow visualisation, as well as other image reconstruction techniques. (*Ultrasound Toolbox* n.d.).

USTB was utilized for beamforming and visualisation. The toolbox features well-tested beamformer implementations, such as the DAS beamformer.

### 3.2.1 Implementation of *getCapon* into the Ultrasound Toolbox

One of the first steps in this thesis was to implement a stand-alone Capon beamformer algorithm, named *getCapon*, into USTB as a postprocess. The *getCapon*-algorithm dates back to 2009 at the DSB group at UiO.

First, the code was updated slightly regarding modernisation in the MATLAB syntax, before some new functionalities were added. The main functionality implemented in the algorithm is the possibility of applying expanding aperture (section 2.4.1) to the data, followed by an update of the subarray averaging method (section 2.4.4) to adapt to the expanding aperture.

After this, *getCapon* was implemented as a post-process in the USTB, to utilize the full functionality of the USTB. The applied functionality mentioned above is kept, as well as adapting the algorithm to fit the USTB syntax.

## 3.3 Code availability

The source code for this thesis is available on Github through the following link:

<https://github.com/helenewold/MasterThesis>.

Simulated datasets can be reproduced using the scripts located in the folder *Source Code/Dataset generation scripts*

# Chapter 4

## Results and discussion

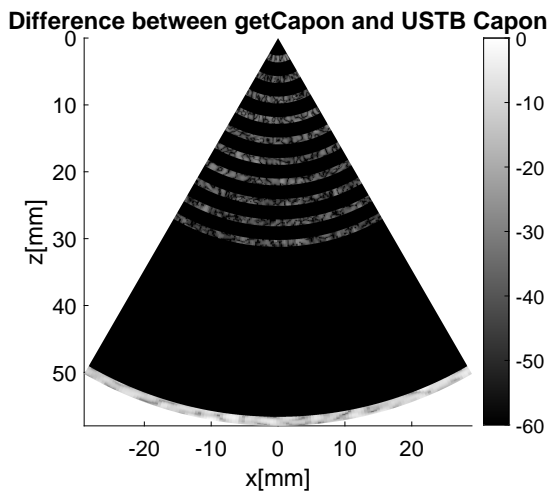
### 4.1 Implementation of the Capon beamformer into the Ultrasound Toolbox

One of the first tasks in this master project was to implement an already existing Capon beamforming algorithm developed by the DSB group at UiO as a post-process operation in the Ultrasound Toolbox (USTB) framework.

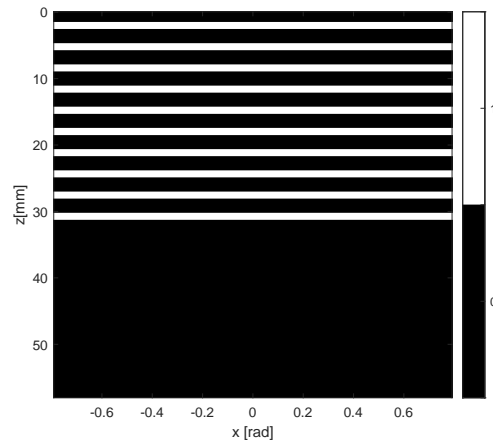
However, there already exists an implementation of the Capon beamformer in the USTB. Both implementations calculate the weight vector as shown in Equation (2.9) on page 18, which theoretically should result in identical numerical output from the two algorithms. It is then useful to compare the processing results from the two implementations, to examine the result regarding the output similarities.

A natural way to examine the similarity between the implementations is by calculating and visualising the numerical difference between their outputs when given the same input. The numerical difference between all pixel values in the image is calculated by

$$\Delta = \left| \frac{|z_A|}{\max(|z_A|)} - \frac{|z_B|}{\max(|z_B|)} \right|, \quad (4.1)$$



**Figure 4.1:** The numerical difference between the outputs from the UiO *getCapon* algorithm and the USTB Capon minimum-variance algorithm. The input was the simulated speckle dataset.



**Figure 4.2:** The numerical difference between the subarray length on the beamformed outputs from the UiO *getCapon* algorithm and the USTB Capon minimum-variance algorithm. The white represents where there are differences between the two methods.

where the complex output from the UiO *getCapon* implementation was applied as  $z_A$ , and the complex output from the USTB Capon algorithm as  $z_B$ . The beamformer input was the recorded dataset presented in Section 3.1.3, visualised in Figure 3.2.

The result of the calculation is shown in Figure 4.1, where a striped pattern is seen along the depth axis. The explanation for this are in the numerical differences which occur when applying both expanding aperture and the subarray averaging method, see respectively Section 2.4.1 and Section 2.4.4. The subarray size changes according to the expanding aperture approach, as the ratio between the subarray length and the full array length is kept constant.

Figure 4.2 visualises the numerical differences in subarray length  $L$  applied in the two Capon beamforming algorithms. As is noticeable in the figure, the pattern coincides with the striped pattern visible in Figure 4.1. The differences from Figure 4.1 are barely visible to the naked eye when visually analysing the full output of both algorithms. The numerical difference is below  $-20$  dB, which,

## 4.2 Applied Multi-Line Acquisition

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when visually comparing the two outputs, are close to non-distinguishable.

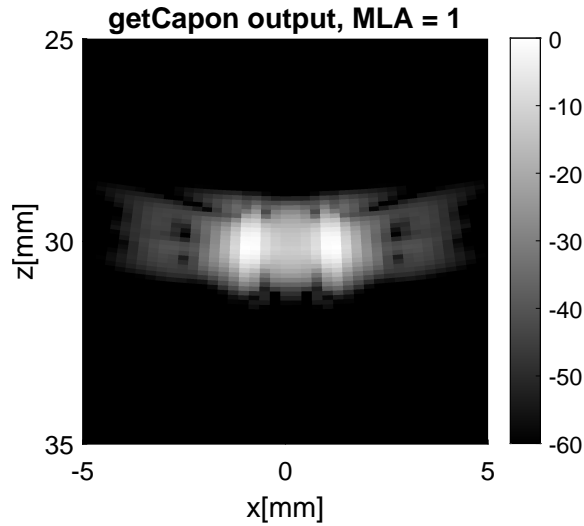
The reason for the difference in output and applied subarray lengths has not been located. One possible reason for how this difference occurs might be the way the two implementations apply the time averaging with respect to the first and last samples, which then may affect the application of the expanding aperture. The number of averaged time samples and the number of depth samples where there are differences between the two implementations, are the same, supporting the presented possible reasoning.

When applying temporal averaging, the covariance matrix of a sample  $y_m(n)$  is averaged over the covariance matrices of the surrounding  $t$  samples, i.e. the samples between  $y_m(n - \frac{t}{2})$  and  $y_m(n + \frac{t}{2})$ . For the first  $\frac{t}{2} - 1$  samples, there are not  $\frac{t}{2}$  prior time samples to include in the averaging of the covariance matrix. Then, the lack of sufficient prior time samples has to be accounted for. This can be done by either averaging the covariance matrix over all available prior time samples, or by not including the first  $\frac{t}{2} - 1$  time samples. The same applies to the last  $\frac{t}{2} - 1$  time samples. The USTB implementation includes a method which averages using the covariance matrices of all available prior samples if there are less than  $\frac{t}{2}$  samples available before or after sample  $y_m(n)$ .

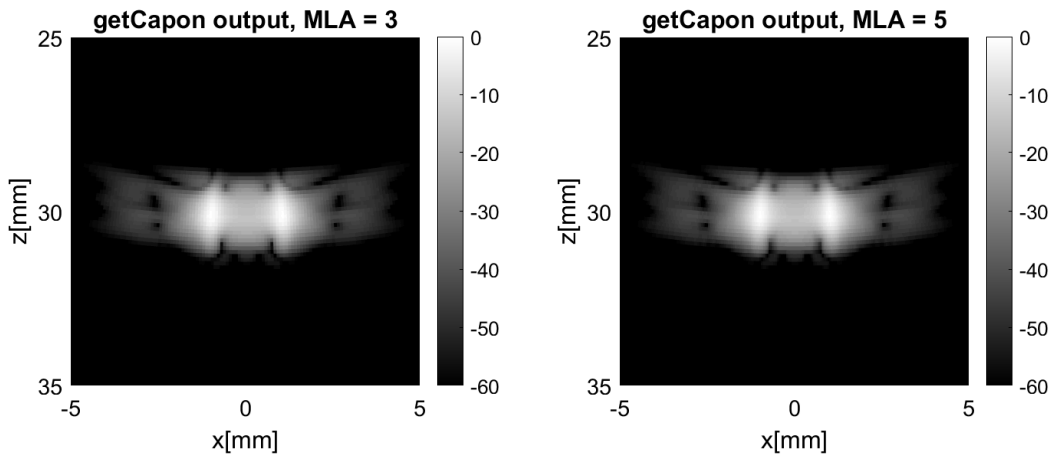
Both implementations have been analysed with respect to Equation (2.9) and the subarray averaging method (section 2.4.4 on page 21), and are then considered mathematically correct. The *getCapon* implementation exhibited a reduced computational time, and since there is no located reason for the numerical differences, the *getCapon* algorithm from the DSB group is utilized. As the *getCapon* algorithm is separate from the USTB, other than the inclusion as a post-process, it is possible to modify the algorithm without impacting the USTB core structure.

## 4.2 Applied Multi-Line Acquisition

It is important to have a well-sampled image, as a real-time visualisation can be misrepresented if the image is poorly sampled in space. In the case of a scene containing scatterers somewhere in the scene, a poor spatial sampling may cause



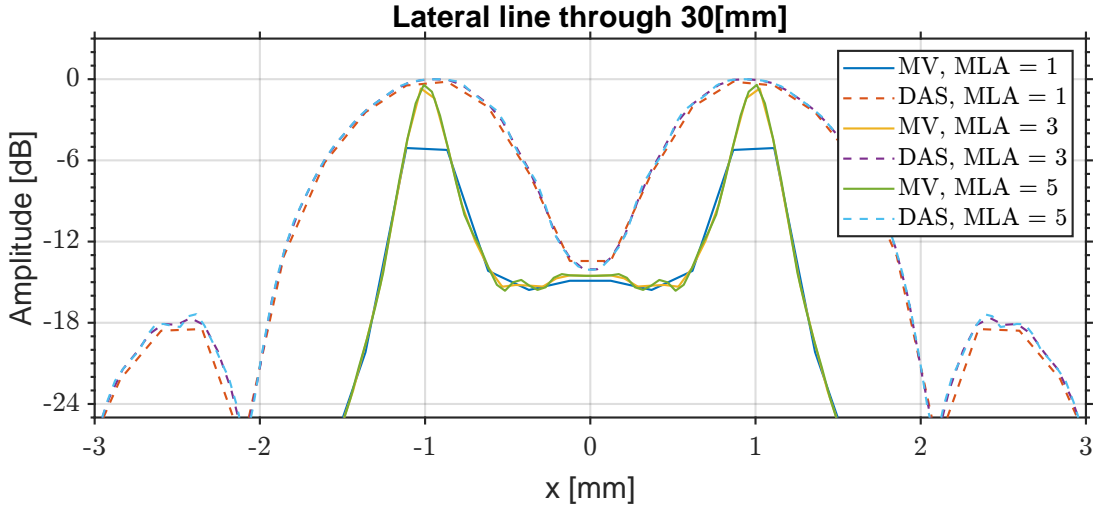
(a) Result from the Capon Beamformer when no MLA is applied.



(b) Result from the Capon Beamformer when an MLA factor of  $MLA = 3$  is applied. (c) Result from the Capon Beamformer when an MLA factor of  $MLA = 5$  is applied.

**Figure 4.3:** The result from the Capon Beamformer when applying two different MLA factors of 3 and 5, as well as no MLA, to a dataset with two scatterers at 30 mm depth, and 2 mm distance between. The figure visualises how the pixel size decreases when the MLA factor increases.

## 4.2 Applied Multi-Line Acquisition



**Figure 4.4:** Resolution through the lateral axis at a depth of 30 mm, showing the separability between two scatterers at a distance of 2 mm using different MLA factors for both the DAS and Capon beamformer. The DAS beamformer results are visualised using a dashed line, and the Capon beamformer results are visualised using solid lines.

a missed peak at the scatter locations.

Increasing the lateral resolution demands a denser lateral sampling. One way of increasing the lateral resolution is to apply Multi-Line Acquisition (MLA), see Section 2.4.3. The amount of MLA is adjustable, and a choice regarding this amount must be made. To keep the results comparable throughout this thesis, a consistent use of MLA is beneficial. Therefore, it is useful to analyse how MLA affects the separability and spatial resolution. This analysis was done with the Capon beamformer, where the input is the dataset containing two scatterers located 2 mm apart at a depth of 30 mm. The dataset was presented in Section 3.1.2 and visualised in Figure 3.1.

The notation  $\text{MLA} = \text{value}$  is applied throughout this discussion, to represent the factor by which the received beams have been increased. No applied MLA is denoted as  $\text{MLA} = 1$ . The analysis was done with  $\text{MLA} = 1, 3, \text{ and } 5$ . The dataset has been simulated using  $t_x = 128$  transmit beams, as noted in Table 3.2.

Figure 4.3 shows the image reconstructed using the Capon beamformer. The subfigures of Figure 4.3 visualise the processing with mMLA when using various

factors of  $MLA = 1, 3$ , and  $5$ .

When comparing Figure 4.3a with Figure 4.3c, namely the results from the Capon Beamformer using an MLA of respectively no MLA and  $MLA = 5$ , it is clear that the lateral resolution in Figure 4.3c is improved from Figure 4.3a. In Figure 4.3a, the pixels are more visible along the lateral axis compared to Figure 4.3b and Figure 4.3c. In Figure 4.3, it is shown that a higher MLA gives a noticeably enhanced resolution using the Capon beamformer, with more precise locations of the scatters with respect to the width of the scatterers.

As seen in Figure 4.4, the DAS beamformer represented by the dashed lines performs similarly for the different MLA factors. It is possible to see how the DAS beamformer already is over-sampled with 128 transmit beams when considering the case of  $MLA = 1$  at 30 mm depth. An increased MLA-factor will result in an even more over-sampled beamformer output for the DAS beamformer. The main difference between the DAS cases shown in Figure 4.4, is the difference in the amplitude level at  $x = 0$  mm. Here, the amplitude of the  $MLA = 1$  case is approximately 1 dB higher than for the  $MLA = 3$  and  $MLA = 5$  cases.

For the Capon beamformer, the case of  $MLA = 1$  is clearly undersampled, as indicated by the chopped scatter peak. By chopped, it is meant that the amplitude peak at the scatterers is flat. This chopped peak is not visible in Figure 4.3. For this undersampled case, the beamformer misses the scatter location peaks. The amplitude is approximately 5 dB lower for the  $MLA = 1$  case than for the two cases of  $MLA = 3$  and  $MLA = 5$ . The two cases with applied MLA do result in a well-sampled signal when examining the resolution at 30 mm depth. Regarding the amplitude around 0 mm, the three MLA cases perform equivalently. The higher MLA-factors do, however, cause a slightly increased amount of perturbations.

Regarding the two MLA factors 3 and 5, the resolution analysis results shown in Figure 4.4 using both the DAS and the Capon beamformer are very similar. Throughout this thesis, the beamforming is performed with an applied MLA factor of 3 to achieve a well-sampled signal along the lateral axis at 30 mm depth. An MLA-factor of 3 increases the lateral pixel count from 128 to 384 pixels.



### 4.3 Resolution

#### 4.3.1 Scatterers in a speckle-less scene

One of the main image metrics is the resolution of objects in the recorded scene. The following section will examine how the Capon beamformer performs with respect to the separability of scatterers in the scene, applying the DAS beamformer as a performance benchmark. The impact of the applied subarray averaging on the separability will be studied. The separability analysis is initially done to define a limit of lateral distance between the two scatterers where both the DAS and Capon beamformer results in separated scatter location with the parameter set discussed in Section 3.1. The eigenvalues of the estimated covariance matrix are also examined.

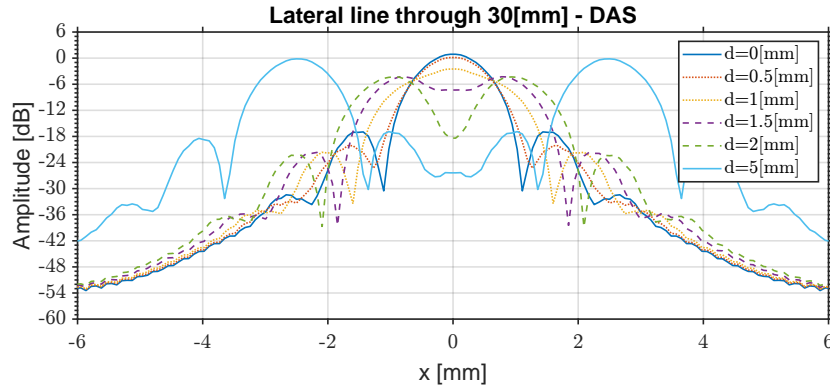
##### **Increasing distance between scatterers**

First, the separability of scatterers in a scene is analysed when the distance between the scatterers in the scene is varied. The scatterers are always simulated symmetrically around the probe centre axis at a 30 mm depth.

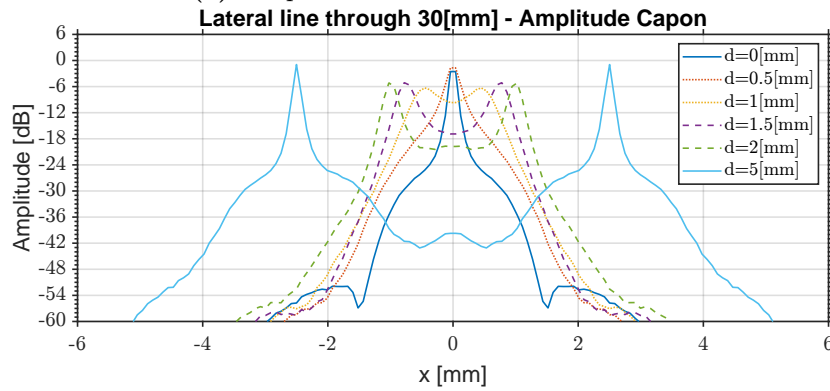
##### **Separability**

Figure 4.5 shows the amplitude along a lateral line at 30 mm for different beamformed outputs, where the beamformed scenes contain two scatterers at 30 mm depth. The difference between the scenes is the distance between the two scatterers. Figure 3.1 visualises an example of the beamformed output of such a scene. The inter-scatter distances applied are 0, 0.5, 1, 1.5, 2, and 5 mm. Each subfigure visualises the amplitude of the different scenes beamformed using the DAS beamformer, the amplitude Capon beamformer, and the power Capon beamformer.

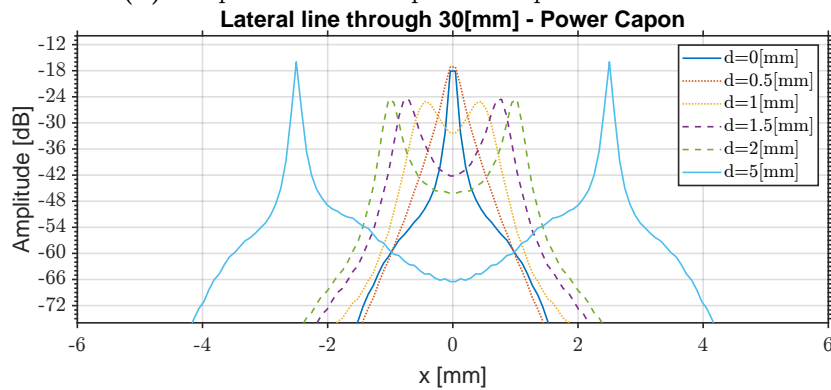
The results from Figure 4.5a and Figure 4.5b have all been normalized by the maximum value in the case of DAS applied to the 5 mm case. This has been done to ensure a representable comparison between the results. It will also be possible to see if one scene is more prone to signal cancellation when the amplitude is compared with another scene, which is not as apparent when the images are



(a) Output from the DAS beamformer.



(b) Output from the amplitude Capon beamformer.



(c) Output from the power Capon beamformer.

**Figure 4.5:** The resolution from the three methods of DAS, amplitude Capon and power Capon, for different scenes with varying distances between two scatterers. The distances between the scatterers are respectively 0, 0.5, 1, 1.5, 2, and 5 mm, and the scatterers are located at 30 mm depth.

### 4.3 Resolution

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individually normalized. The result from Figure 4.5c is, however, not normalized, as the power Capon method in the `getCapon` algorithm has an unlocated error. Correcting the power Capon implementation has not been prioritized in this thesis, as the separability can be measured between scatterers, and the effect of subarray averaging is still visible.

The power Capon result does however visualise how the amplitudes of the scatterers are about 12 dB lower than the corresponding results from the amplitude Capon. The difference between the amplitude of the scatterers when the inter-scatter distances are 0 mm and 1 mm is about 6 dB, which is approximately 2 dB bigger difference than from corresponding results from the amplitude Capon method. This difference may, however, be reduced when proper normalization and the proper implementation of the power Capon method have been implemented.

From all subfigures in Figure 4.5, it is noticeable how the peak amplitude of the scatterers from the dataset using two scatterers at 0 mm and 0.5 mm apart is enhanced compared to the other cases, as well as the peak amplitude from the dataset with two scatterers at 1 mm when using DAS in Figure 4.5a. The max amplitude then decreases when the scatterers are located 1.5 and 2 mm apart, before increasing for a 5 mm distance. This is due to the beampattern of the array and beamformer, which represent how the intensity of a beam varies dependent on direction and distance from the source. If the scatterers are positioned so that one scatterer is located within the sidelobe of another scatterer, it results in a reduction in the amplitude of the scatterers.

For the case of two scatterers located with 0 mm distance between, i.e. located at the same place in the scene, the scatter amplitude is chopped. The missed peak is a result of undersampling. The scatterers are located between two receive lines, and as the effective point spread function is narrow, the peak is missed, and the apparent amplitude of the peak is lowered. For the Capon beamformer, the amplitude increases slightly when the two scatterers are moved 0.5 mm from each other, the received lines are closer to the two scatterers.

For two scatterers with 1.5 mm when using the DAS beamformer, a dip between the scatterers is noticeable. According to the Sparrow Limit, presented in Section 2.5.1,

the two scatterers are then separable. This is not the case if one is to consider the  $-6$  dB limit. The two scatterers located 1 mm apart when using the amplitude Capon beamformer are also separated when considering the Sparrow Limit.

A noticeable difference between the images reconstructed using the DAS and both the Capon beamforming implementations is in the width of the located scatterers for all scenarios, regarding, e.g., the Full Width at Half Maximum (FWHM). FWHM is the width of the scatterers when the amplitude is reduced by half from the maximum amplitude. The DAS beamformer generally results in a larger estimated width of the scatterers.

### Eigenvalues

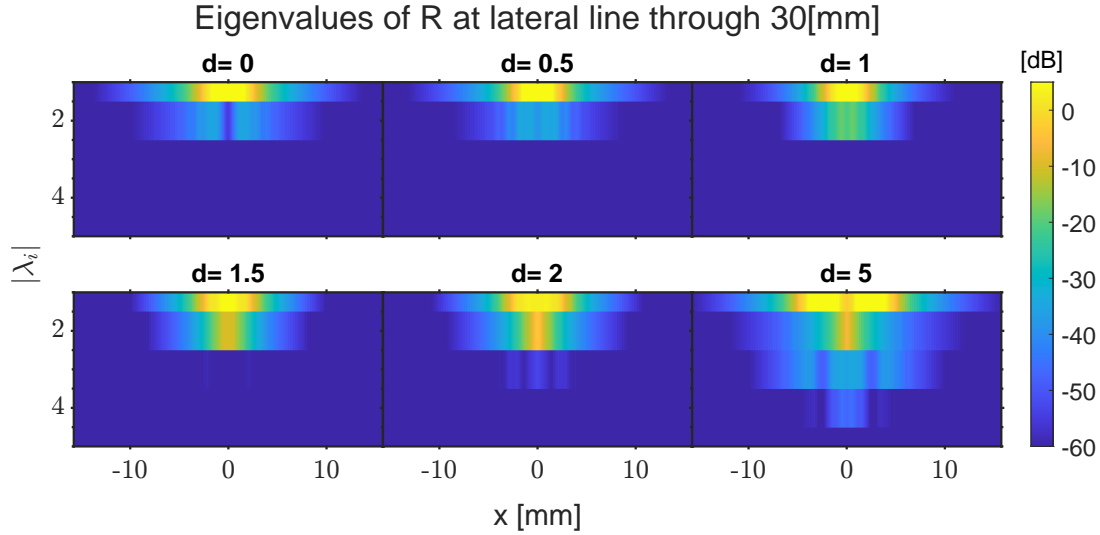
Figure 4.6 shows the eigenvalues of the covariance matrix through a lateral line at 30 mm, and Figure 4.7 the axis along  $0^\circ$ . The eigenvalues are calculated using the *eig()* function in MATLAB. The eigenvalues have been sorted, locating the highest values as  $\lambda_1$ .

The eigenvalues are visualised in decibel. As the eigenvalues are quite small, perturbations in the eigenvalues then appear more significant. This will help derive more information about how, e.g., the increasing distance between the scatterers affects the eigenvalues.

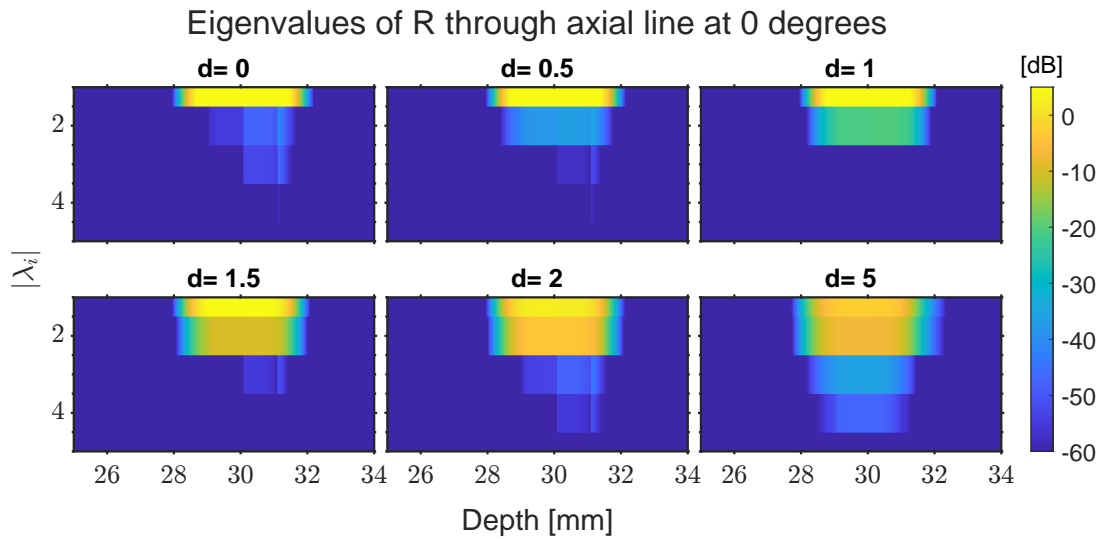
None of the eigenvalues has been normalized, other than a limitation to the colour bar, to exclude eigenvalues close to zero. This is done to achieve a good representation of the eigenvalues for the different scenes when compared to each other.

As the upper left figure in both Figure 4.6 and Figure 4.7 on the facing page represents the scene with two scatterers located in the same position, it is not possible to measure any separability. As noticeable in the upper left figure in Figure 4.6, there is a slight decrease in the magnitude of the second largest eigenvalues in the area surrounding 0 mm. The decrease in the eigenvalue magnitude is caused by the axial line being gathered nearly directly through the point. The transmitted pulse is then directed straight at the scatterers, and the received, delayed signal is then a plane wave which hits all elements simultaneously. The covariance is

### 4.3 Resolution



**Figure 4.6:** Eigenvalues of the covariance matrix along a lateral line going through a depth of 30 mm. The scatterers are located at 30 mm depth, with a distance of 0, 0.5, 1, 1.5, 2, and 5 mm between the scatterers.



**Figure 4.7:** Eigenvalues of the covariance matrix through the azimuth axis at an angle of  $0^\circ$ . The scatterers are located at 30 mm depth, with a distance of 0, 0.5, 1, 1.5, 2, and 5 mm between the scatterers.

then high between all elements from this pixel position, resulting in a covariance matrix where all matrix entries are approximately the same value. Such a matrix is called a all-ones matrix.

When considering scenes with a larger distance between the scatterers, the received signal contains a more distributed contribution of the waves reflected by the two scatterers. The covariance between all elements then decreases slightly, causing the covariance matrix to lose the all-ones characteristic. This reduction also increases the number of significant eigenvalues, hence reducing the eigenvalue magnitudes.

In general, it is found that the eigenvalues gathered along the axial line through 30 mm presented in Figure 4.7 are skewed at greater depths. This skewness is a result of the direction of arrival of the signals, as the reflected wave arrives at the probe array dependent on the distance between the element and the localization of the point.

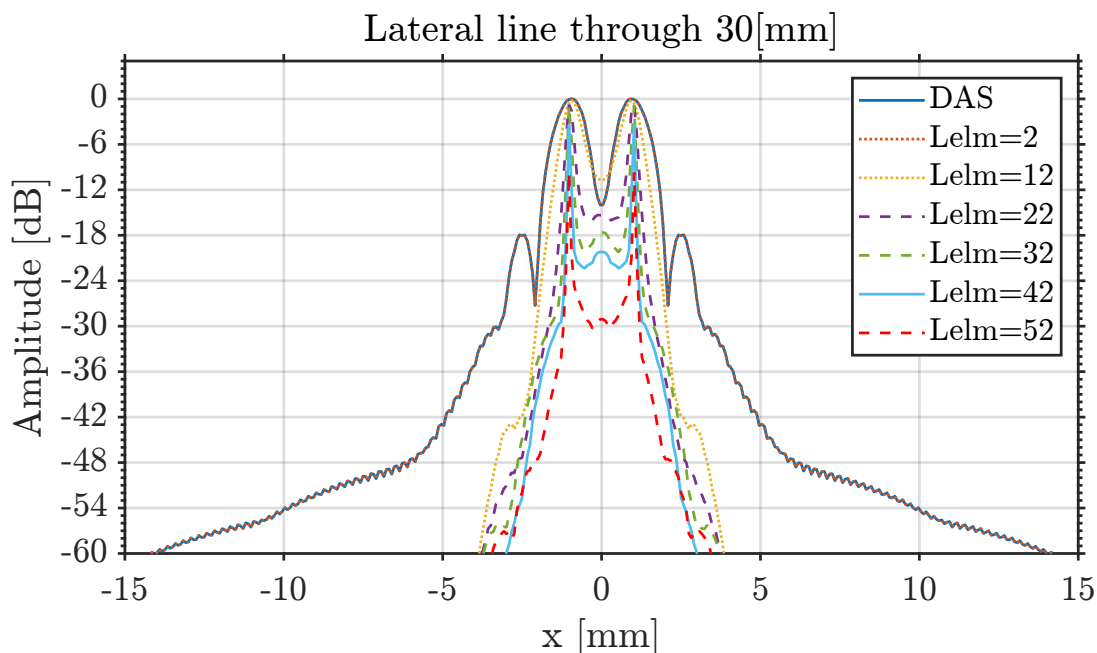
Now, are the eigenvalues of the covariance matrix applicable for defining the separability of two scatterers? It is possible, as the magnitude of the largest eigenvalues is experiencing a decrease between two well-separated scatterers. However, this method fails to prove new insights compared to a simple analysis of the amplitude.

The eigenvalues do, however, provide some insight into the invertibility of the covariance matrix. If a covariance matrix only has minuscule eigenvalues, the inversion of the matrix may suffer from large numerical errors. This will be covered shortly in Section 4.3.1 when the condition number of the matrix is to be examined.

Generally, a reduction in eigenvalue magnitude between the scatterers becomes increasingly notable when the distance between the scatterers increases. When speckle is added to the scene at a later stage in this thesis (Section 4.3.3), there is expected an increase in the magnitude of the more significant eigenvalues relative to the most significant eigenvalue. Speckle decreases the correlation between the signals recorded at each element, which then causes the covariance matrix to have a speckle component that tapers off away from the diagonal. This increases the orthogonality of the covariance matrix columns, which further increases the spread

### 4.3 Resolution

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**Figure 4.8:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers when the subarray length is 2, 12, 22, 32, 42, and 52 elements.

of the eigenvalues.

#### Analysis of different subarray lengths

Now, the second analysis regarding resolution in this thesis is an analysis of how the subarray length affects the separability of two closely located scatterers.

The analysis below is performed with a scene containing two scatterers located at a depth of 30 mm, 2 mm apart. This scene was chosen as the scatterers are separable when considering the  $-6$  dB limit for both the DAS and Capon beamformer, as shown in Section 4.3.1. The two scatterers will then always be separable when increasing the subarray length, as they are separable when applying DAS. Compared to the DAS result, the width and amplitude of the scatterers can be affected by an increased subarray length.

The notation of subarray length throughout the discussion is  $L = no. array elements$  of the full array.

## Separability

Figure 4.8 shows the result when the subarray size is varied in the minimum-variance Capon beamformer pipeline. It visualises the extent to which the subarray size affects the separability of two scatterers. When looking at the result from DAS and a subarray size  $L = 2$ , the results are close to identical. The two lines plotted for the two results overlap each other, indicating a similar result. When applying spatial averaging using  $L = 1$ , the result of the Capon beamformer is theoretically the DAS beamformer with an applied uniform weight, as this corresponds to summing the signal in both receive and transmit directions after applying a delay.

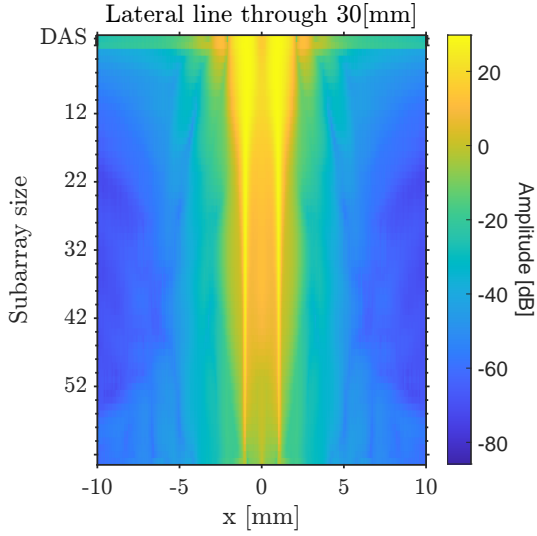
The difference between two subarray sizes applied is expected to be quite small when the subarray sizes differ by one element, as the amount of averaging only slightly increases. However, when increasing the subarray size in increments of 10, a more noticeable difference is measurable. As shown in Figure 4.8, the amplitudes of the two scatterers are decreasing between each 10-element increment of the subarray size when passing a certain  $L$ . With  $L = 2, 12, 22$ , and  $32$ , the amplitude at the scatterers is close to the DAS result, only decreasing with 1 dB at most. When applying  $L = 42$ , the amplitude at the peaks is reduced with 3 dB. The amplitude when  $L = 52$  is reduced by about 10 dB.

The decrease in signal amplitude is a possibly consequence of two effects. The first effect is instability in the covariance matrix estimate, as the averaging applied decreases when  $L$  increases. Subarray averaging reduces the effect of coherent signals (Synnevåg, 2009), and too little averaging causes signal cancellation. The second effect is the resolution of the image possibly being too coarse due to a low value of the MLA, such that the Capon beamformer does not properly capture the peak of the scatter, causing the amplitude to be reduced.

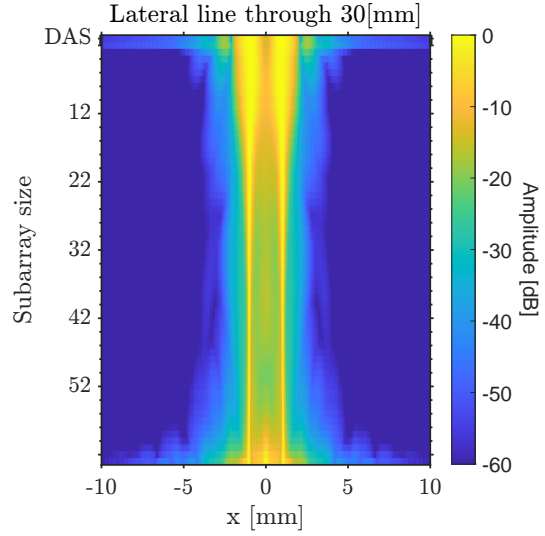
In addition to the calculated amplitude of the two scatterers, the general separability between the two is useful to examine. The decibel level of the dip between the two scatterers implies how well the two scatterers are separated. One expected result is the DAS and the  $L = 2$  having a nearly identical separation, which is the case as shown in Figure 4.8. The separation between the two scatterers when



### 4.3 Resolution



**Figure 4.9:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers when the subarray length  $L$  increases from 2 to 63 elements. The top line is the result from DAS. All results have been normalized in regard to the DAS output.



**Figure 4.10:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers when the subarray length  $L$  increases from 2 to 63 elements. The top line is the result from DAS. All results have been normalized according to the maximum of each beamformed output.

considering  $L = 12$  is however worse compared to the result of the two smaller subarray sizes, i.e. DAS and  $L = 2$ . Here, the scatterers have a separation of about 15 dB, while the result from  $L = 12$  has a separation level of 10 dB.

From a subarray size of  $L = 22$  elements and up, the dip will become slightly larger in the middle of the two scatterers.  $L = 22$  has a magnitude reduction of approximately 16 dB, and  $L = 32$  of approximately 18 dB. When moving on to the results from  $L = 42$  and 52, the general signal amplitude decreases with respectively 3-4 dB and 10 dB.

An additional aspect to notice in Figure 4.8 is the width of the two scatterers. For DAS and subarray of 2 elements, the width at the  $-6$  dB resolution limit is about 1 mm, while the increase to  $L = 12$  elements gives a decreased width of about 0.75 mm. Both  $L = 22$  and  $L = 32$  return a decreased width of about 0.2 mm at  $-6$  dB. When further increasing the subarray size, the apparent width

of the scatterers decreases. This results in a width of about 0.1 mm for both scatterers. The increase in subarray size is, however, at a cost of signal intensity. From this, it is possible to see that the separation as well as the width of the scatterers increases when the number of elements in the subarray increases, up to a certain subarray size. With a subarray size of about 50 % of the full array size, there is a limitation of the resolution according to the  $-6$  dB limit when normalized by the result from DAS.

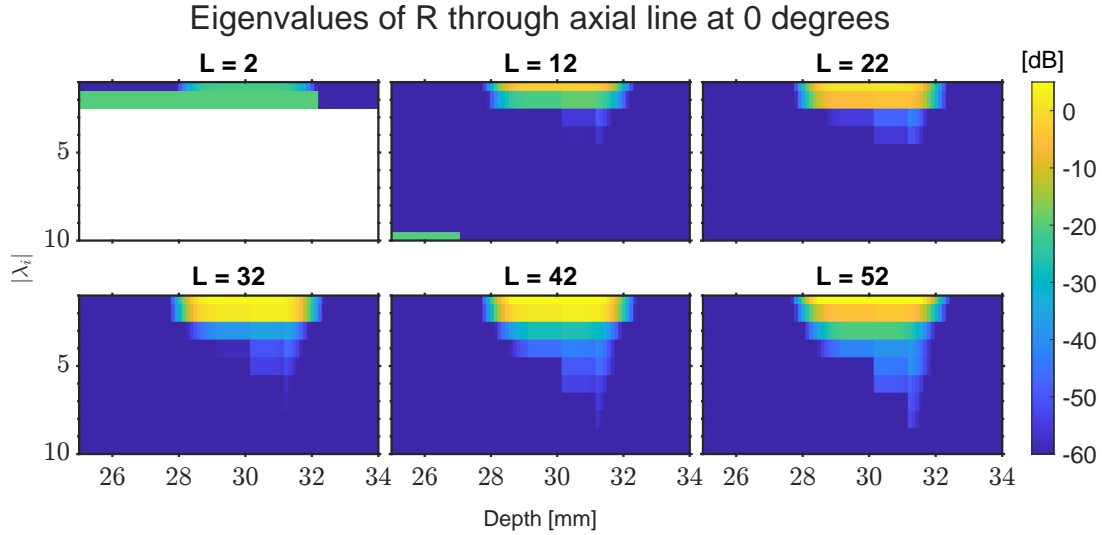
As shown in Figure 4.9, the intensity loss follows the increase of subarray size. The apparent width of the scatterers is about 2 mm, and narrows when the subarray size passes about 16 elements. This equals  $\frac{1}{4}$  of the full array size. When passing  $L \approx 42$  elements in the subarray, i.e.  $\frac{2}{3}$  of the full array, the results will approach signal cancellation according to the  $-6$ dB limit when compared with the DAS beamformer.

The results visualised in Figure 4.10 is corresponding to the result in Figure 4.9, however with a different normalization. Figure 4.9 is normalized in regard to the DAS result, while Figure 4.10 is normalized in regard to the maximum value from each output. Figure 4.10 then shows how the scatter width from  $L = 40$  elements to  $L = 60$  elements is comparable. This was mentioned in regard to Figure 4.8, i.e. how the signal mainly loses intensity when the subarray length is increased past a certain point. When the subarray length increases, a slight increase in amplitude appears around  $x = 0$  mm, which is between the two scatterers. The small amplitude increase is visible for smaller subarray sizes, and is visible in Figure 4.8. The increased magnitude visible in Figure 4.10 is caused by the scatter peak being decreased relative to the amplitude around  $x = 0$  mm.

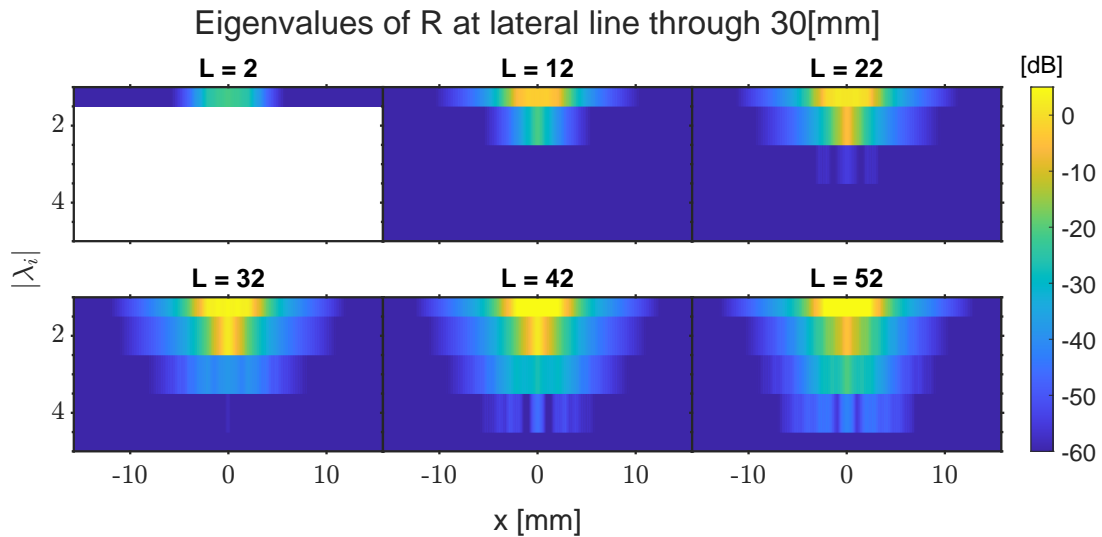
### **Eigenvalues of the subarray analysis**

Figure 4.11 shows the eigenvalues of the covariance matrices gathered along the azimuth axis along  $0^\circ$ . It is noticeable that the largest eigenvalues increase when the subarray size increases. This is also the case with the number of significant eigenvalues for the covariance matrices, and is expected, as introduced by Section 2.6.1. Hence, there is no new, significant information provided when analysing the eigenvalues at this stage.

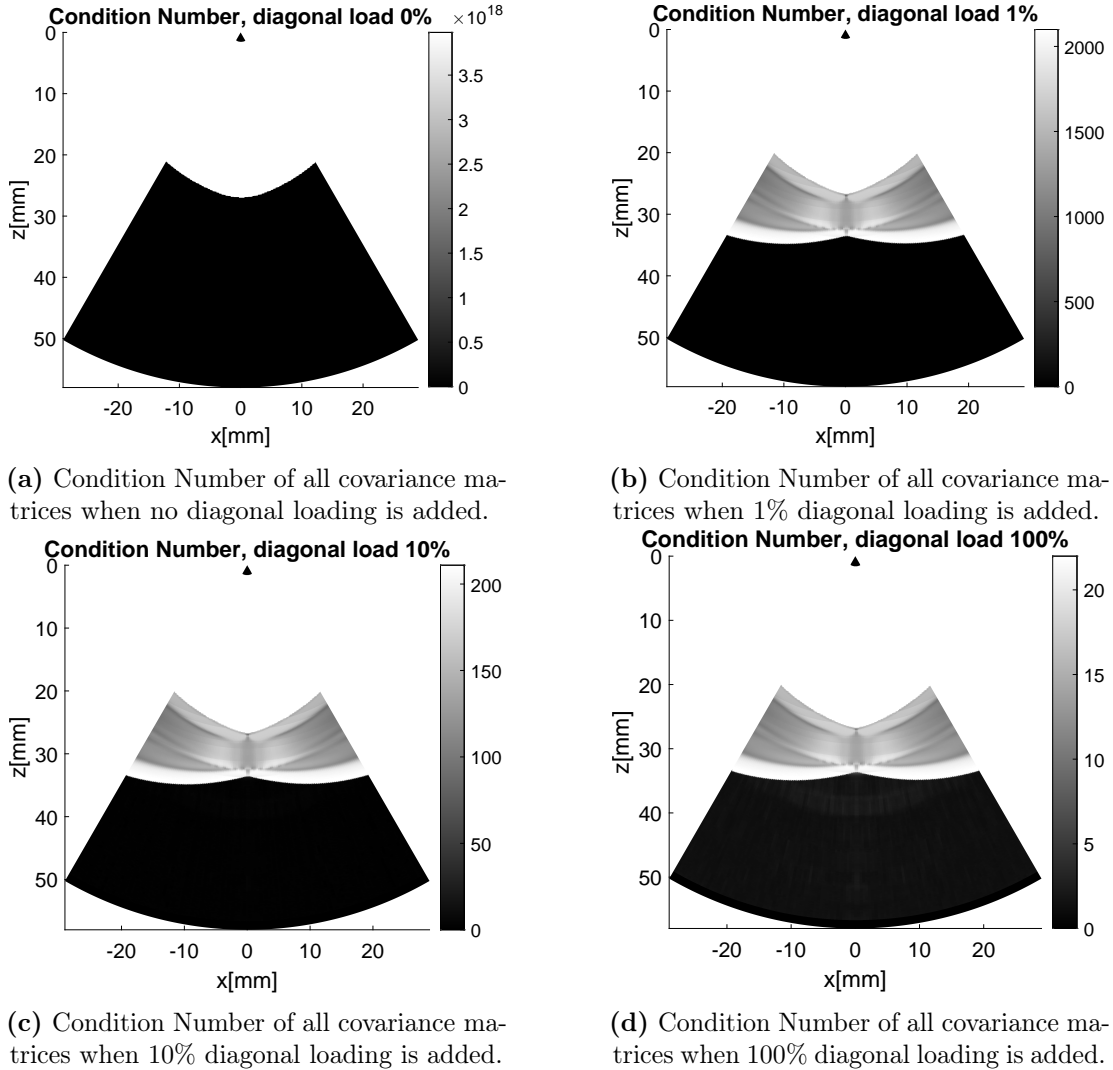
### 4.3 Resolution



**Figure 4.11:** Eigenvalues of the covariance matrix through the azimuth axis at an angle  $0^\circ$ . The scatterers are located at 30 mm depth, with a distance of 2 mm between them. The result has been calculated using the subarray lengths of 2, 12, 22, 32, 42, and 52 elements.



**Figure 4.12:** Eigenvalues of the covariance matrix through the azimuth axis at a depth of 30 mm. The scatterers are located at 30 mm depth, with a distance of 2 mm between them. The result has been calculated using the subarray lengths of 2, 12, 22, 32, 42, and 52 elements.



**Figure 4.13:** A visual representation of the condition number of the covariance matrices when the diagonal load is 0, 1, 10, and 100 %. The white field above 25 mm is due to the covariance matrix being a non-invertible matrix, leading to *NaN* result. The dataset is a speckle-less dataset with two scatterers at 30 mm depth with 2 mm between the scatterers.

## 4.3 Resolution

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### Condition number of the covariance matrix

As mentioned in Section 2.6.2, the condition number is in linear algebra a measure of the stability of a matrix inversion, and applying a diagonal load to the matrix before inversion increases the inversion stability, see Section 2.4.4. The effect of diagonal loading on the stability of matrix inversion will be examined next by analysing the condition number of the matrix. The calculations have been done using the dataset presented in Section 3.1.2 and visualised in Figure 3.1, as input to the Capon beamformer. How much the subarray averaging applied affects the condition number will also be considered shortly.

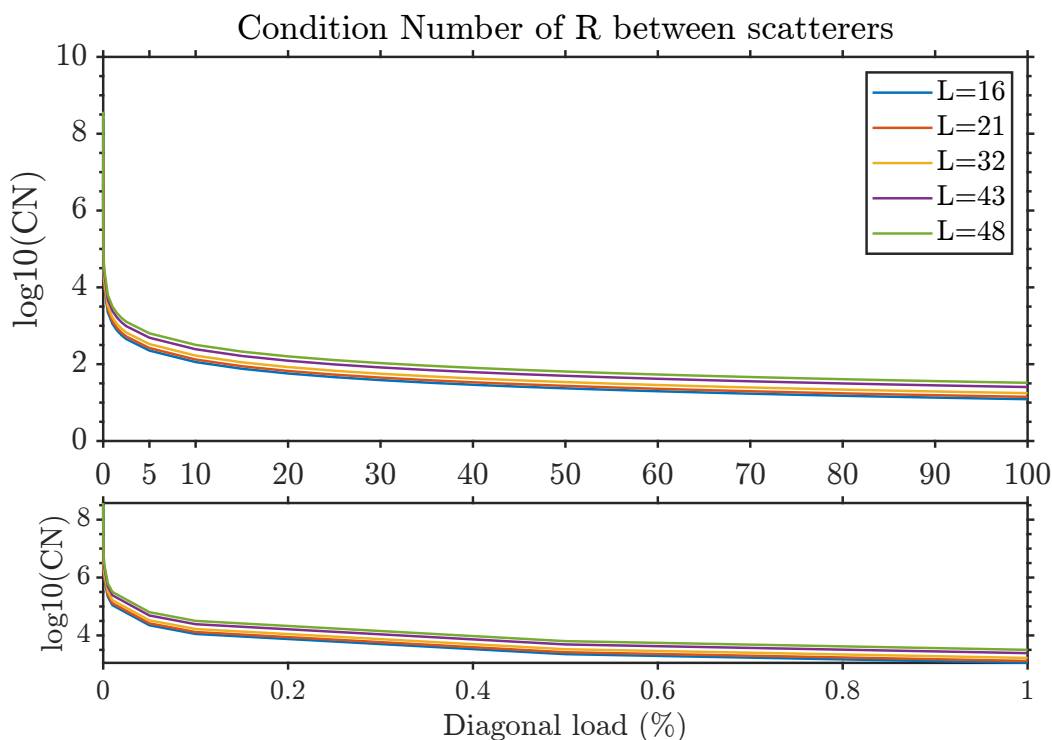
The logarithmic value of the condition number,  $\log_{10}(CN)$ , denotes the maximum amount of digits that are in danger of being miscalculated by the matrix inversion (Section 2.6.2). What then defines a well-conditioned or ill-conditioned matrix, i.e. large or small condition number, is the precision of the calculation tool. MATLAB has a precision of 16 digits. For example, if one were to think a sufficient limit of a large condition number is 50 % of the calculation precision, a condition number below  $\log_{10}(CN) = \log_{10}(10^8) = 8$  digits is acceptable.

### Added diagonal loading

As mentioned in Section 2.4.4, diagonal loading is a method to make the covariance matrix inversion more robust. The amount of diagonal load necessary to apply depends on, e.g., the input data of the beamformer, the precision of the applied calculation tool, and the extent of error present in the beamformer.

From the results of the condition number calculations in Figure 4.13, where the condition number is represented by the colour bar, it is possible to see how the condition number is reduced with a factor of 10 when the diagonal load increases with a factor of 10. The condition number results are visually similar between all subfigures in Figure 4.13. The condition number decreases drastically when a small diagonal load of 1% is applied, as seen from the magnitude of the colour bar, causing the stability of the matrix inversion to increase.

Even though it is important to have a good precision of the matrix inversion, it is also important to avoid altering the covariance matrix estimate significantly



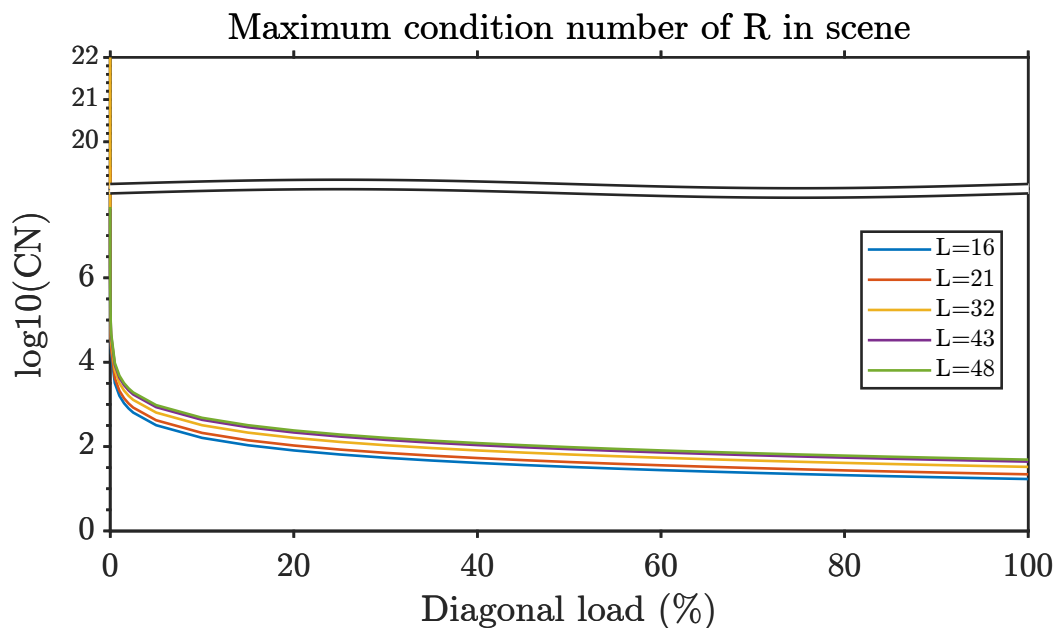
**Figure 4.14:** Logarithmic value of the condition number with increasing diagonal load, when the covariance matrix between the two scatterers is analysed. Each line represents a subarray average size. The upper plot shows the full range of diagonal loading considered. The lower plot shows the limited first percentage of the applied diagonal load.

by adding an excessive amount of diagonal load. As previously indicated in the introduction, an excessive diagonal load applied to the covariance matrix will result in a beamformer output that approaches the DAS beamformer output (Synnevåg, Austeng, and Holm, 2007b). The covariance matrix estimate is based on physical time signals, and major alterations will cause the covariance matrix estimate to deviate excessively from the original covariance matrix. Hence, the amount of diagonal loading applied must be kept within boundaries.

Figure 4.14 shows the logarithmic value of the condition number calculated for the covariance matrix in a point in the middle of the two scatterers in the scene. Figure 4.14 has been calculated using a set of subarray lengths, namely  $L = \frac{1}{4}, \frac{1}{3}, \frac{1}{2}, \frac{2}{3},$  and  $\frac{3}{4}$  of the full array of  $M = 64$  elements. Here it is noticeable how much a small increase in the diagonal load robustifies the matrix inversion,

### 4.3 Resolution

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**Figure 4.15:** Logarithmic value of the maximum condition number from the calculations of the covariance matrix from the Capon minimum-variance beamformer from each tested diagonal load. Each line represents a subarray size.

rather independent of the subarray length.

Diagonal loading is an aid to reduce numerical errors in the covariance matrix estimate inversion. However, a stage is reached where the applied diagonal load does not affect the stability of the matrix inversion as much. At this stage, the applied diagonal load is no longer as beneficial. This can be seen in Figure 4.14 and Figure 4.15, where the stability of the covariance matrix inversion is represented by the logarithmic value of the condition number. Between 10 % and 100 % diagonal loading, the calculation precision represented by the logarithmic condition number only decreases by approximately 1 digit when considering Figure 4.14. Hence, a diagonal load above 10 % alters the covariance matrix excessively when considering how limited the increase of numerical stability of the matrix inversion is.

The differences in the condition numbers between the various subarray lengths are small, but worth noticing. A larger subarray is in need of more diagonal load if a certain condition number is sought. As seen from Figure 4.14,  $L = 16$  calls

for a diagonal load of 10 % to achieve  $\log_{10}(CN) \approx 2$ , while the corresponding logarithmic condition number for  $L = 48$  calls for a diagonal load between 40 % and 50%.

When calculating the condition number of the covariance matrix estimate from the point located between the two scatterers, the condition number is consistently below a value of approximately  $10^9$ . This indicates a matrix inversion with a numerical accuracy of at least 45 % when utilizing MATLAB. This is not the case when considering the maximum finite condition number of the entire scene, as shown in Figure 4.15. When comparing Figure 4.14 and Figure 4.15, the same dependency of the diagonal load is visible for all subarray sizes, and a similar trajectory. The maximum finite condition number is, however, greater than the condition number from the middle point between the scatterers.

Again, a reduced subarray size results in a lowered condition number when considering the same diagonal load. There is therefore a need for less spatial averaging when applying a set diagonal load to achieve a certain matrix inversion stability measured by the condition number.

If a set matrix inversion stability, represented by the condition number, is desired for a certain subarray length, the diagonal load necessary for this condition number can be calculated through a similar examination as shown in Figure 4.14.

Several positions in the scene were considered when analysing the condition number of the covariance matrix, but not all are presented here. The trend when increasing diagonal load is however consistently following a trajectory close to the function  $f(x) = \frac{1}{x}$ . In general, the condition number is applicable to define the stability of the matrix inversion, which corresponds to an increased stability of the Capon beamformer output. By increased stability of the Capon beamformer output, it is in this context meant that the inversion is not the cause of large errors in the beamformer output.

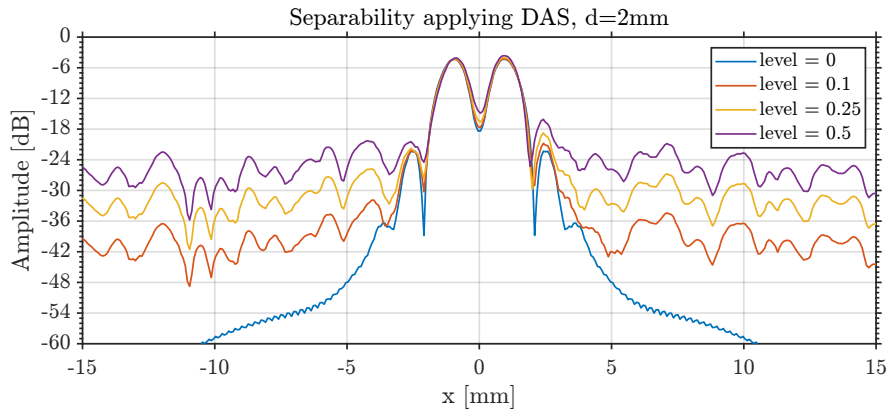
### 4.3.2 Applying speckle and defining a speckle level

In this analysis, an appropriate speckle level for simulations is examined. There will always be speckle in datasets recorded in a physical medium, as mentioned

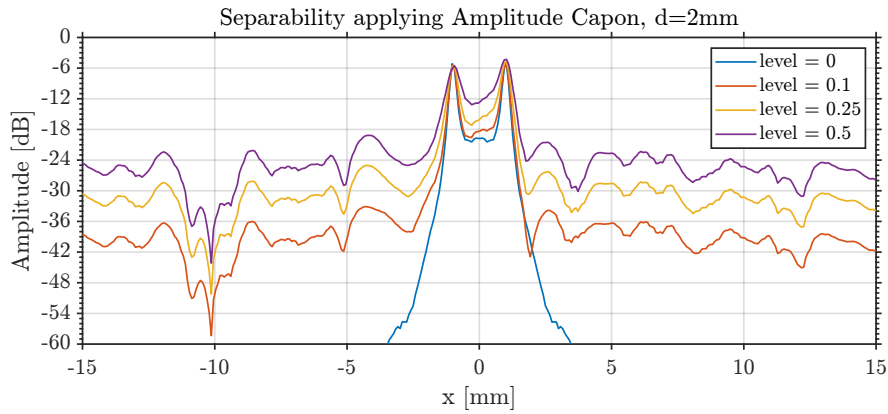


### 4.3 Resolution

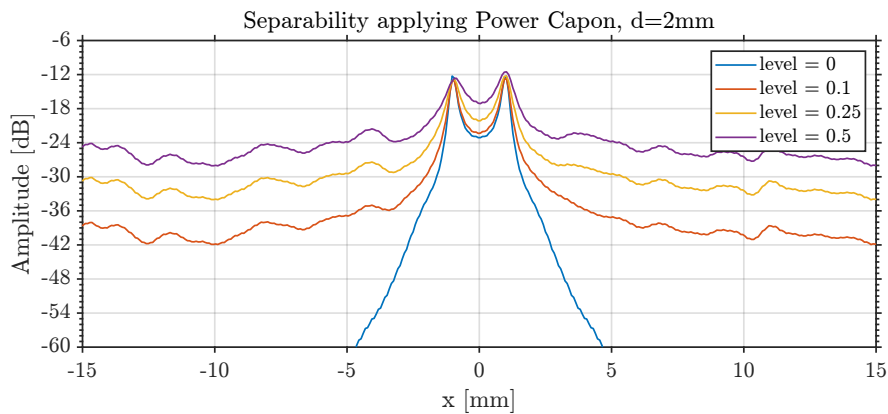
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**Figure 4.16:** The effect of increasing speckle level compared to the scatter amplitudes, using the DAS beamformer.



**Figure 4.17:** The effect of increasing speckle level compared to the scatter amplitudes, using the amplitude Capon beamformer.



**Figure 4.18:** The effect of increasing speckle level compared to the scatter amplitudes, using the power Capon beamformer.

in Section 2.3.1. Therefore, all examinations done thus far should be repeated, applying a scene also containing speckle. The speckle and scatter datasets have both been simulated using the same maximum amplitude. In order to identify the scatterers in the speckle scene reliably, the amplitude of the speckle data has to be reduced.

The reduction of speckle amplitude is done by simply multiplying the amplitude of the simulated speckle with a fraction. For this analysis, these fractions are  $\frac{1}{10}$ ,  $\frac{1}{4}$ , and  $\frac{1}{2}$ . The effect is visualised by measuring how the separability between the scatter scatterers is affected by the speckle level in the scene.

### **The effect on separability when varying the speckle level**

From Figure 4.5a and Figure 4.5b on page 42, it was possible to see how two scatterers in a speckle-less scene were separable according to the  $-6$  dB limit for both the DAS and Capon beamformers when the scatterers had a distance larger than 2 mm between them. A significant change in separability is not expected for this scene, other than a possible increase in amplitude between the scatterers.

The amplitude of the external area of the two scatterers was reduced significantly compared to the amplitude of the scatterers. When speckle is added, it is natural that the amplitude outside of the two scatterers will increase relative to the amplitude of the scatterers, as the speckle creates a consistent amplitude level in the image. It was also noticeable how the amplitude in the area between the two scatterers decreased when the distance between the scatterers increased.

As seen in Figure 4.16 through Figure 4.18, the amplitude is reduced by approximately 12 dB from the scatter peak when considering the amplitude at  $-4$  mm for the 0.25 speckle level in all three figures. For 6 mm, the amplitude is reduced by approximately 18 dB. These reductions are slightly less for the power Capon method, however, mostly for the amplitude at 6 mm, which is only reduced by approximately 12 dB.

For the power Capon method shown in Figure 4.18, it is noticeable that the speckle amplitude level is comparable to the amplitude Capon method and DAS, and the scatter amplitude has increased with approximately 10 dB compared

### 4.3 Resolution

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to the corresponding result from Figure 4.5c. The increase in amplitude is only present for the power Capon result, however, the power Capon method is not normalized, as in Section 4.3.1.

It is noticeable when comparing Figure 4.17 and Figure 4.18 that the speckle is drastically smoothed when applying the power Capon method. At  $x = -10$  mm, there is an arbitrary amplitude dip of approximately 12 dB visible when applying the amplitude Capon method. However, the dip is smoothed when utilizing the power Capon method.

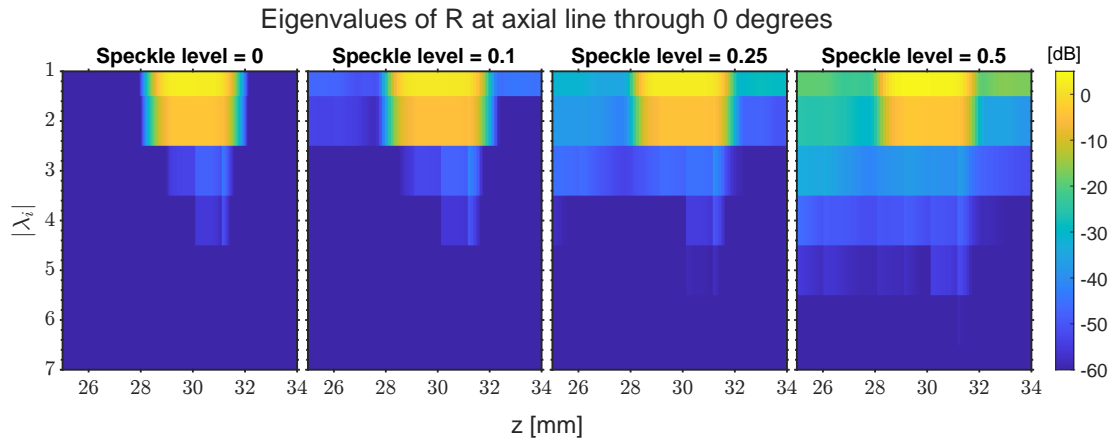
The result in Figure 4.16, Figure 4.17 and Figure 4.18 supports the expectations mentioned, as they visualise how the speckle amplitude level increases, only slightly affecting the amplitude in the area between the two scatterers. The latter change is larger for the amplitude and power Capon beamformer than for the DAS. The width of the scatterers increases slightly when using the amplitude Capon and power Capon beamformers, due to the amplitude increase between the two scatterers.

#### **Eigenvalues of the covariance matrix when increasing the speckle level**

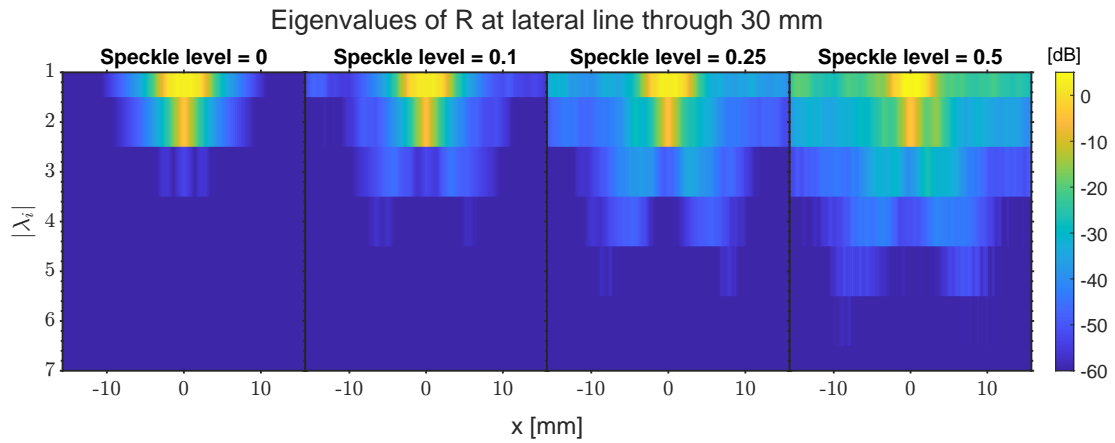
From Section 4.3.1 and Section 4.3.1, we saw how a speckle-less scene would result in mainly two significant eigenvalues. Due to the decrease in covariance between the recorded element signals when speckle is added, it is expected that the amount of noticeable eigenvalues increases in both the axial and lateral directions.

A way to either confirm or deny this is to analyse the eigenvalues when the speckle level is varied. The result is shown in Figure 4.19 and Figure 4.20, for the axial direction through  $0^\circ$  and the lateral direction through 30 mm, respectively. From these figures, it is seen that there is a noticeable increase in the general eigenvalue magnitudes when the speckle level increases, which confirm the expected increase in amount of noticeable eigenvalues.

The speckle level chosen for the following analysis is  $\frac{1}{4}$ , i.e. a relative amplitude of  $-30$  dB. This is due to the speckle not affecting the visibility of the scatterers in the scene, while still maintaining a reasonable level to challenge the beamformer.



**Figure 4.19:** The eigenvalues from the covariance matrix from an axial line going through  $0^\circ$ . The speckle level increases from no speckle in the leftmost plot to  $\frac{1}{10}$ ,  $\frac{1}{4}$ , and  $\frac{1}{5}$  from left to right. The scene used is two scatterers located at 30 mm depth with 2 mm distance between the scatterers.



**Figure 4.20:** The eigenvalues from the covariance matrix from a lateral line through 30 mm. The speckle level increases from no speckle in the leftmost plot to  $\frac{1}{10}$ ,  $\frac{1}{4}$ , and  $\frac{1}{5}$  from left to right. The scene used is two scatterers located at 30 mm depth with 2 mm distance between the scatterers.

## 4.3 Resolution

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### 4.3.3 Scatterers in a speckle-scene

#### Separability of scatterers when increasing the inter-scatter distance

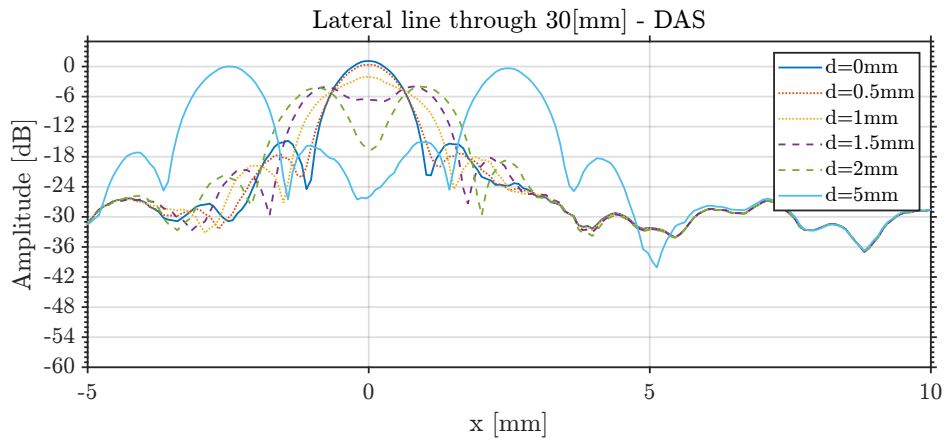
As shown in Figure 4.21, the applied speckle only affects the separability of the two scatterers slightly when the distance between the scatterers increases. The DAS beamformer output shown in Figure 4.21a is not greatly affected by the applied speckle to the scene, other than generally increasing the amplitude level in the ultrasound image.

The amplitude and power Capon beamformer results, shown in respectively Figure 4.21b and Figure 4.21c, maintain the width of the scatterers when speckle is applied. The increased speckle level does, however, reduce the separability of the scatterers located 1 mm apart, as this scene no longer is separable, regardless of resolution criteria.

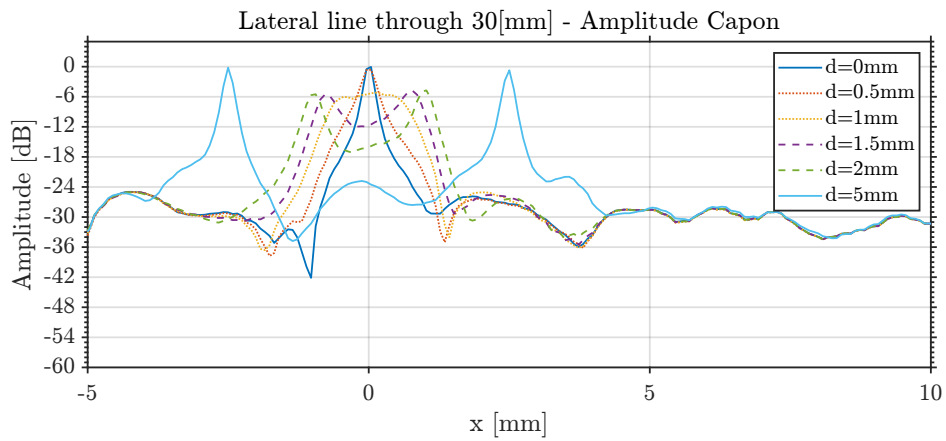
A general matter to note is how the amplitude level between the scatterers around  $x = 0$  mm is affected. When working with a speckle-less scene, this level was rather low,  $-40$  dB for the amplitude Capon beamformer with two scatterers located 5 mm apart. After speckle is applied, this level is raised to approximately  $-25$  dB. The DAS beamformer performs rather similarly whether the speckle is applied or not, the output is not symmetric due to the speckle.

For the power Capon beamformer, the amplitude of the scatterers is increased when speckle is applied. For the speckle-less result, shown in Figure 4.5c on page 42, the amplitude of the scatterers are between  $-16$  and  $-26$  dB for the different cases. When speckle is applied, the amplitude is raised approximately 10 dB, resulting in a scatter amplitude between  $-8$  and  $-14$  dB. Some of this increase in amplitude is probably due to the results not being normalized.

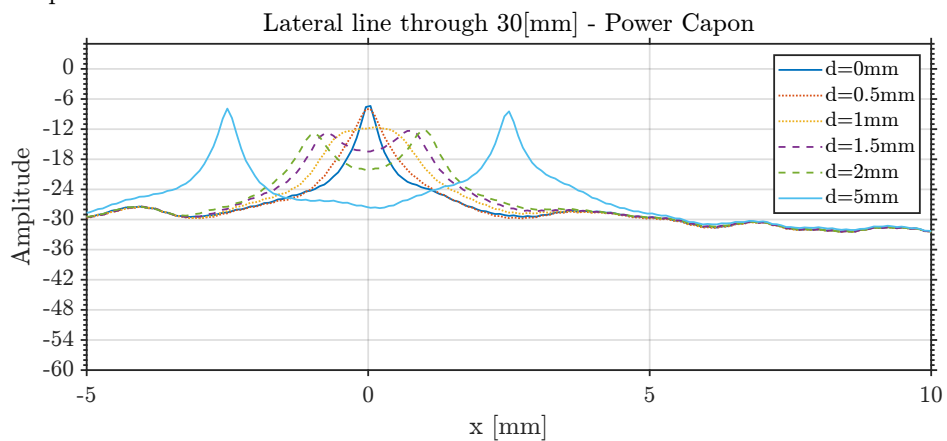
Further, the amplitude between the two scatterers has increased for all cases, causing the amplitude dip to be reduced. This affects the separability of the  $d = 1$  mm case, as this case is not separable according to any resolution criteria after speckle is applied. This shows that even though the power Capon method reduces the variance of the speckle, the separability of scatterers in a speckle scene may be greatly affected compared to the amplitude Capon method.



(a) Resolution for different distances between two scatterers, using the DAS beamformer.



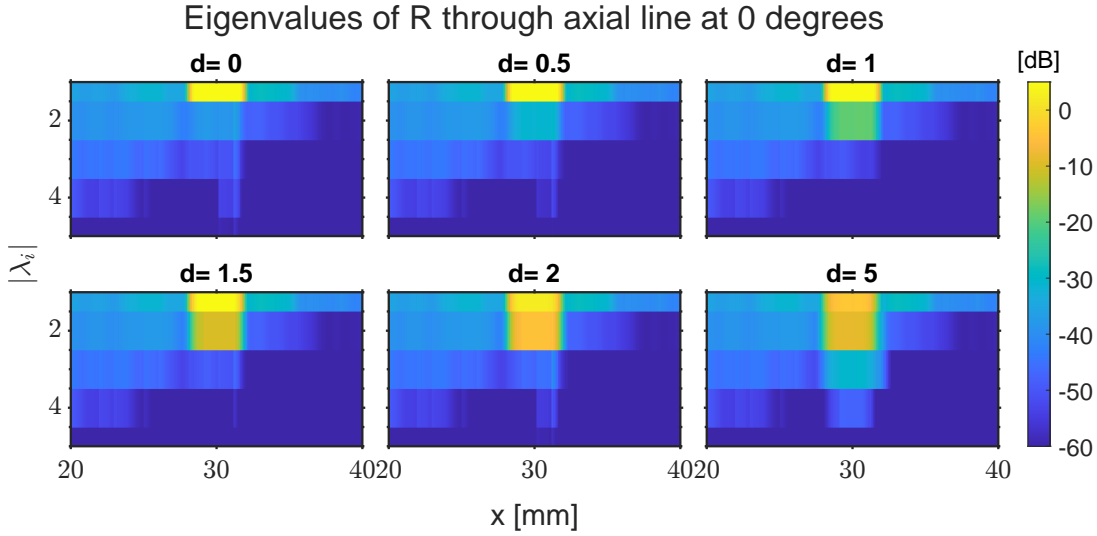
(b) Resolution for different distances between two scatterers, using the amplitude Capon beamformer.



(c) Resolution for different distances between two scatterers, using the power Capon beamformer

**Figure 4.21:** The resolution from the three beamforming methods DAS, amplitude Capon, and power Capon, for different scenes containing speckle and two scatterers located at varying lateral distances. The distances between the scatterers are respectively 0, 0.5, 1, 1.5, 2, and 5 mm, and the scatterers are located at 30 mm depth.

### 4.3 Resolution



**Figure 4.22:** Eigenvalues of the covariance matrix through the azimuth axis at an angle  $0^\circ$ . The scatterers are located at 30 mm depth, with varying inter-scatterer distances. The distances are 0, 0.5, 1, 1.5, 2, and 5 mm.

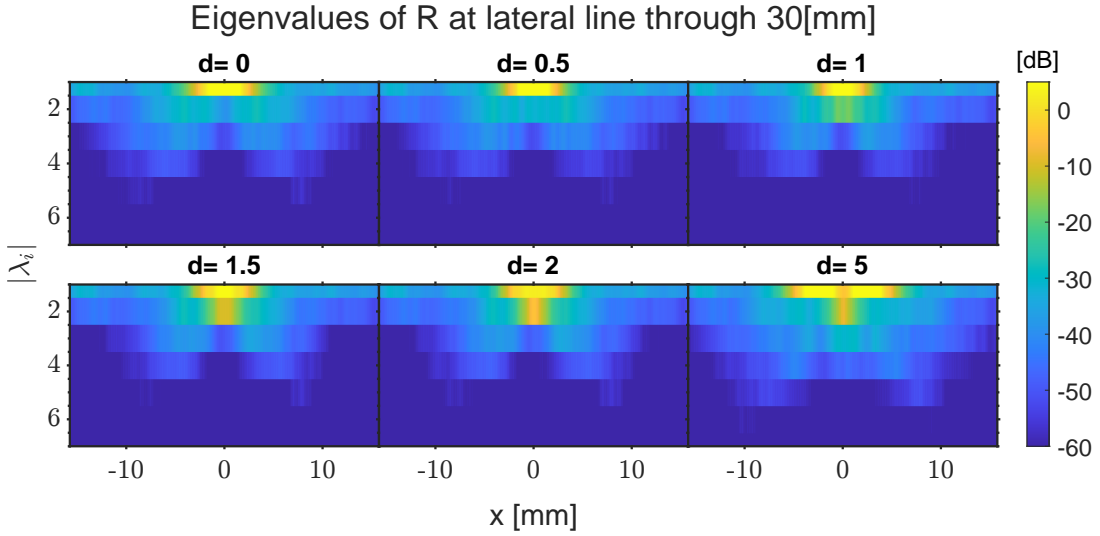
#### Corresponding eigenvalues when increasing inter-scatterer distances

Moving on to the eigenvalues of the covariance matrix in the scene with applied speckle, one can see how the amount of eigenvalues increase from when the speckle-less scene was examined. There are still two more significant eigenvalues around the scatterers, but the general scenes have a higher amount of noticeable eigenvalues, as shown in Figure 4.22 and Figure 4.23. This increase is due to the decrease in covariance between the probe elements, and is expected as discussed in Section 4.3.2.

#### Analysis of different subarray lengths

Here, the effect of subarray averaging is examined. As in Section 4.3.1, the scene examined contains two scatterers at 30 mm depth located 2 mm apart, however, now with applied speckle.

The main expectation regarding the separability of the scatterers is that the FWHM is increased when applying speckle to the scene, when comparing the same subarray length for the datasets with and without speckle. Other than this,



**Figure 4.23:** Eigenvalues of the covariance matrix through the lateral axis at a depth of 30 mm. The scatterers are located at 30 mm depth, with varying inter-scatterer distances. The distances are 0, 0.5, 1, 1.5, 2, and 5 mm.

it is expected that the average amplitude background level increases, as seen in Section 4.3.2, due to the applied speckle.

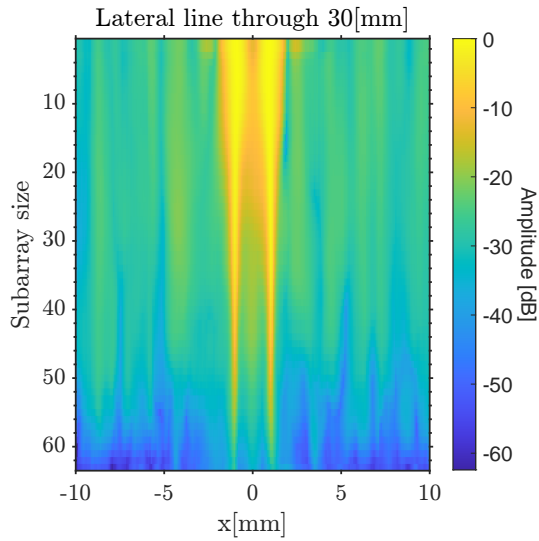
### Resolution

Figure 4.24 shows the resolution from the Capon amplitude beamformer of all subarray sizes when normalized by the maximum DAS result. The scatter width is increased compared to the speckle-less result (Figure 4.9), and the width is maintained until  $L \approx \frac{1}{3}M$ . This is at a later stage than the speckle-less result shown in Figure 4.9, where the width decreased rapidly after about  $L = 16$  elements, i.e.  $L = \frac{1}{4}M$ . When considering where the scatter peak is reduced below  $-6$  dB compared to DAS, the approximate upper limit of  $L = \frac{2}{3}M$  is similar between the cases with and without speckle.

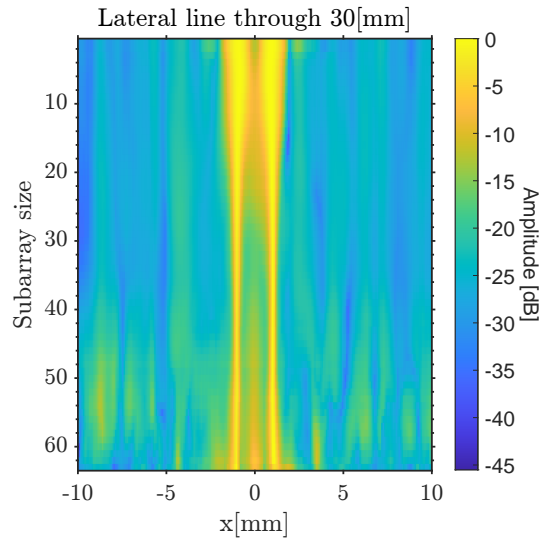
As expected, the apparent width of the scatterers has increased when speckle is applied. The separation regarding the general magnitude reduction between the two scatterers is also reduced, as the magnitude level between the scatterers is increased, as seen in Figure 4.26. Hence, the performance of the Capon beamformer suffers slightly due to the speckle.



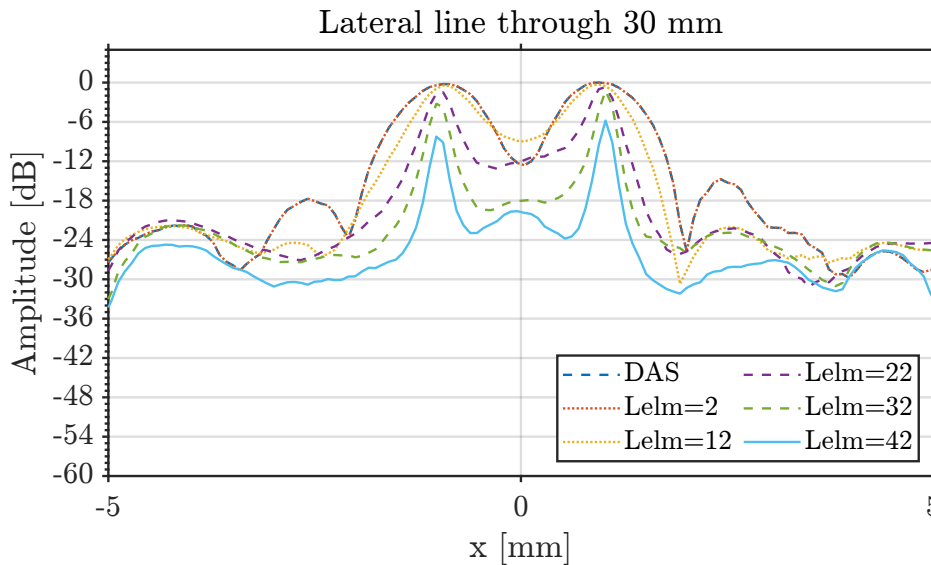
### 4.3 Resolution



**Figure 4.24:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers in a speckle scene when the subarray length  $L$  increases from 2 to 63 elements. All results have been normalized in regard to the DAS output.



**Figure 4.25:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers in a speckle scene when the subarray length  $L$  increases from 2 to 63 elements. All results have been normalized according to the maximum of each beamformed output.

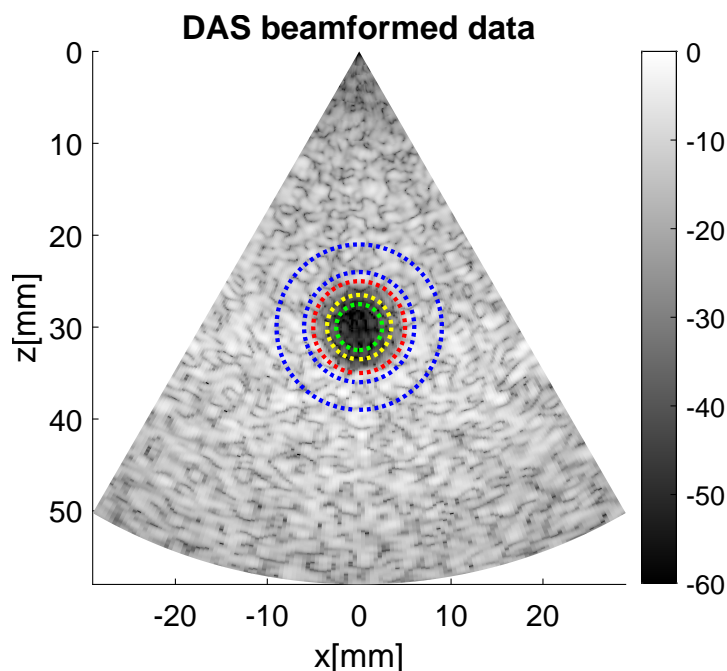


**Figure 4.26:** The amplitude of the signal through the lateral axis at a 30 mm depth. The figure shows the separability between two scatterers in a speckle scene when the subarray length is 2, 12, 22, 32, and 42 elements.

## 4.4 Contrast

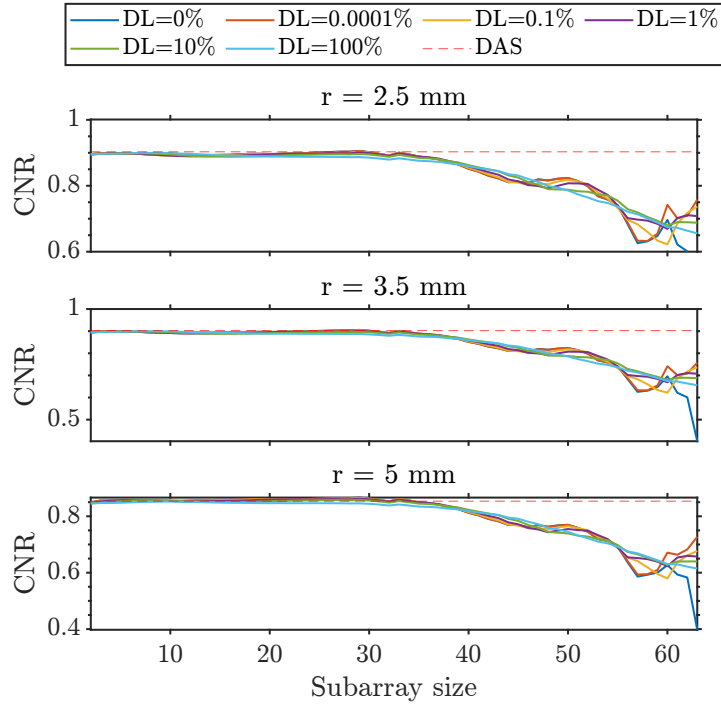
Further, the images are analysed using a set of contrast metrics, to study how subarray averaging and the amount of diagonal loading affect the image contrast when utilizing the amplitude Capon beamformer. The contrast metrics analysed are the CR, CNR, and gCNR, presented in Section 2.5.2. The contrast metrics are calculated using three different ROIs, while keeping the same background region.

The three ROIs are implemented to study how the different contrast metrics are affected by the chosen ROI. The cyst has a correct, physical radius of 5 mm. In an ideal case, the cyst is perfectly reconstructed by the system. This is not the case, however, due to both physical limitations, and how the applied beamformer operates. The DAS beamformer, e.g., has a wide mainlobe, which smoothes the theoretical edge of the cyst. When applying the Capon beamformer, the cyst



**Figure 4.27:** The beamformed output of a scene with a cyst, beamformed by DAS, containing a set of circles visualising the regions applied when working with contrast metrics. The blue circles mark the inner and outer limits of the background region. The green circle marks a border with a radius of 2.5 mm,  $ROI_1$ . The yellow circle marks a border with a radius of 3.5 mm,  $ROI_2$ . The red circle marks the theoretical border of the cyst,  $ROI_3$ .

#### 4.4 Contrast

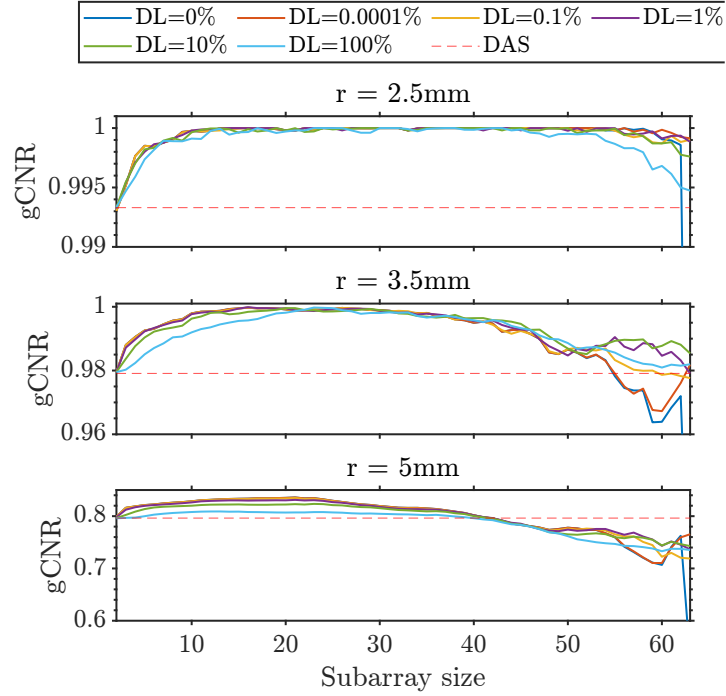


**Figure 4.28:** The measured CNR when subarray size increases, with an applied diagonal load of 0%, 0.0001%, 0.1%, 1%, 10%, and 100%. The CNR from the DAS output is used as a benchmark, shown as a red, dashed line. Each subplot visualises the corresponding CNR from three different applied ROIs. The upmost subplot is  $ROI_1 = 2.5$  mm, the middle subplot is  $ROI_2 = 3.5$  mm, and the bottom subplot is  $ROI_3 = 5$  mm.

edge is sharpened due to the beamformer being a high-resolution method, but elements such as subarray averaging may affect this. This is assessed in this section.

The limits of the different regions are shown in Figure 4.27. The three ROIs are circles around the cyst centre, with radiuses of 2.5 mm, 3.5 mm, and 5 mm, denoted respectively as  $ROI_1$ ,  $ROI_2$ , and  $ROI_3$ .  $ROI_3$  has a radius of 5 mm, as the cyst has a theoretical radius of 5 mm. The other ROI radiuses are chosen to have a set of reasonable ROI for the final, beamformed image.  $ROI_1$  will mainly only contain the data from the cyst, while  $ROI_2$  may contain some speckle, dependent on, e.g., subarray averaging.

The beamformed output from a set of subarray lengths is shown in Figure 4.31



**Figure 4.29:** The measured gCNR when subarray size increases, with an applied diagonal load of 0%, 0.0001%, 0.1%, 1%, 10%, and 100%. The gCNR from the DAS output is used as a benchmark, shown as a red, dashed line. Each subplot visualises the corresponding gCNR from three different applied ROIs. The utmost subplot is  $ROI_1 = 2.5$  mm, the middle subplot is  $ROI_2 = 3.5$  mm, and the bottom subplot is  $ROI_3 = 5$  mm.

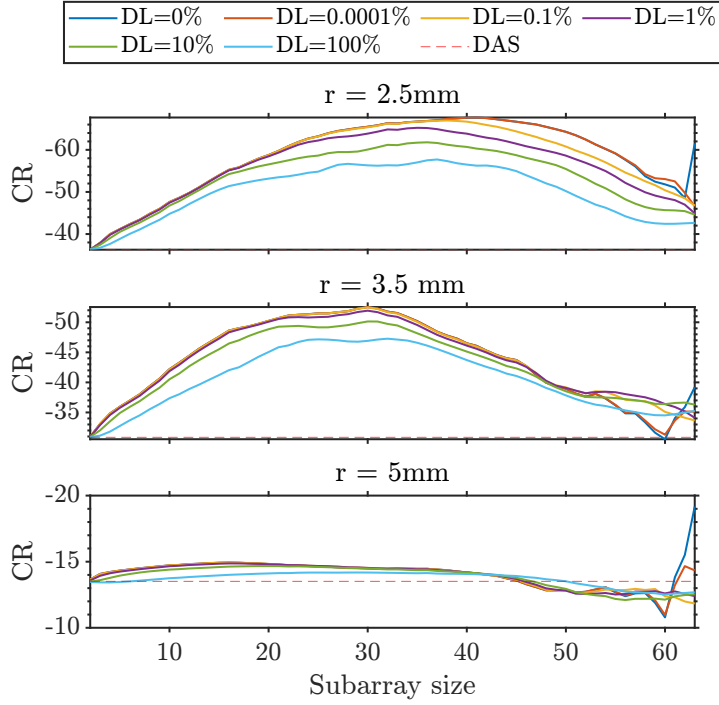
on page 71 and Figure 4.32 on page 72. The figures have been beamformed by applying  $L = 10, 20, 32, 40,$  and 60 elements, shown in the figures from Figure 4.31b to Figure 4.32c.

Figure 4.28, Figure 4.29 and Figure 4.30 show how the contrast metrics, CNR, gCNR, and CR, respectively, are affected by the subarray averaging, for a set of applied diagonal loads.

As shown in Figure 4.28, the CNR of the Capon beamformer is comparable with the CNR of the DAS beamformer for all subarray sizes up to approximately half the of the full array, for all applied diagonal loads. After the subarray size passes  $L \approx \frac{1}{2}M = 32$ , the CNR suffers. This goes for all ROIs accounted for.

The gCNR does however increase from the DAS gCNR when the subarray length

#### 4.4 Contrast



**Figure 4.30:** The measured CR when subarray size increases, with an applied diagonal load of 0%, 0.0001%, 0.1%, 1%, 10%, and 100%. The CR from the DAS output is used as a benchmark, shown as a red, dashed line. Each subplot visualises the corresponding CR from three different applied ROIs. The upmost subplot is  $ROI_1 = 2.5\text{ mm}$ , the middle subplot is  $ROI_2 = 3.5\text{ mm}$ , and the bottom subplot is  $ROI_3 = 5\text{ mm}$ .

increases. How much depends on the considered ROI. As seen in Figure 4.32c, which is the beamformed output when  $L = 60$ , the speckle has changed, causing the  $g\text{CNR}$  to increase. The  $g\text{CNR}$  of  $ROI_1$  and  $ROI_2$  approaches a value of 1, while the real cyst ROI peaks at  $g\text{CNR} \approx 0.84$ .

The CR shown in Figure 4.30 shows how the Capon beamformer has a better CR compared to the DAS CR for all subarray lengths when considering  $ROI_1$  and  $ROI_2$ . The best CR is achieved when applying  $ROI_1$ , and is achieved with  $L \approx 42$ , and 0% or 0.0001% applied diagonal loading.

From all figures showing the contrast metrics with a varying subarray length, namely Figure 4.28, Figure 4.29, and Figure 4.30, it is noticeable how the applied diagonal load does not greatly affect the CNR and  $g\text{CNR}$  other than some

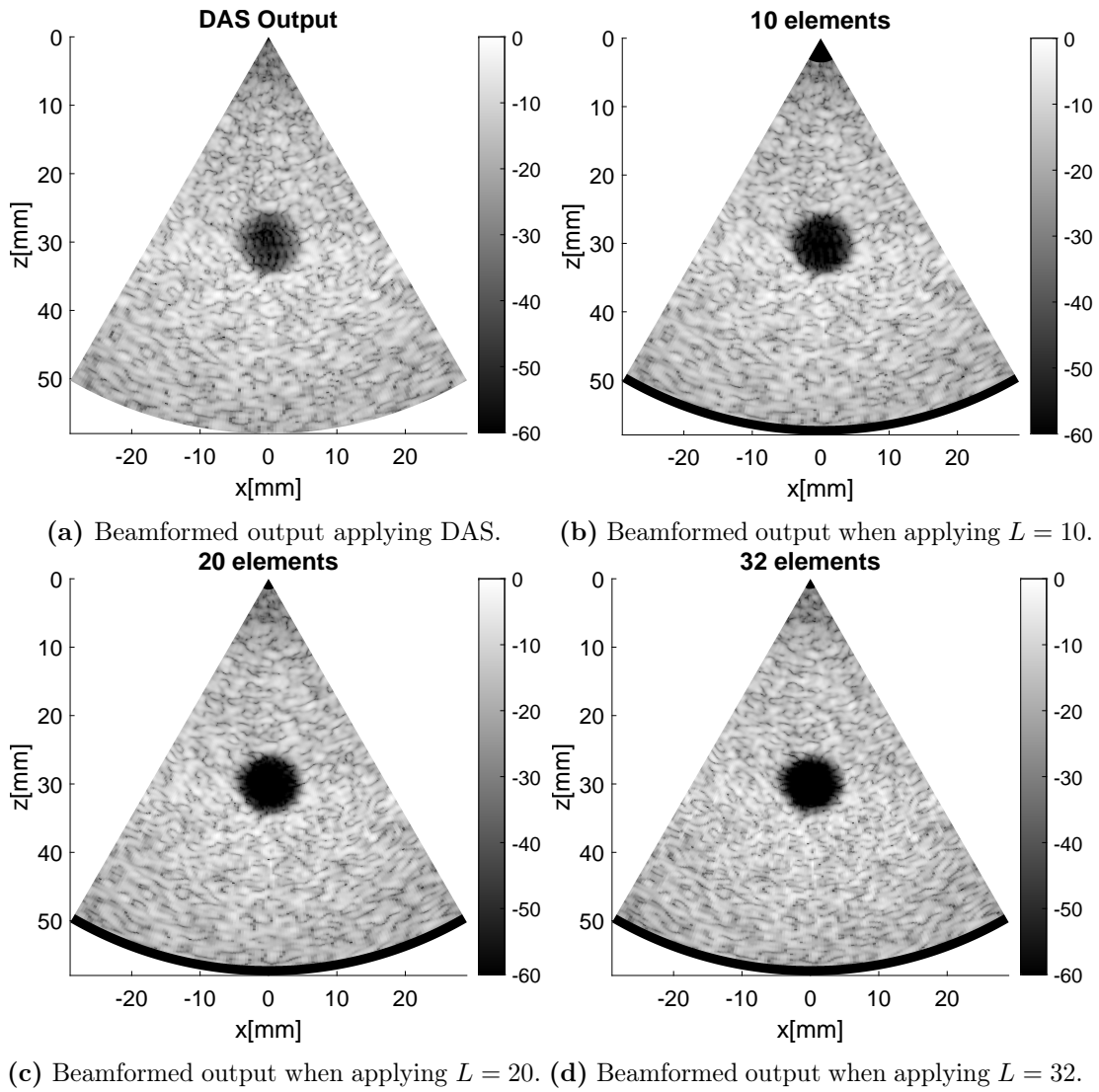
deviations. The general trend for the CR is that the more applied diagonal load, the lower CR, as shown in Figure 4.33a on page 73. The contrast metrics are hence affected slightly by the applied diagonal load, and are differently affected by the amount of subarray averaging.

Based on the result shown in Figure 4.28, passing a subarray size of  $L \approx 32$  reduces the gCNR. From the gCNR shown in Figure 4.29, a subarray size above  $L \approx 10$  results in an increased gCNR. Even though all metrics depend on the ROI chosen, the CR is more influenced by the ROI when considering the subarray size.  $ROI_2$  does however show that a subarray size between  $L \approx 16$  and  $L \approx 32$  results in the best CR, which states that the visual contrast toward the actual cyst edge with a subarray size in that interval.

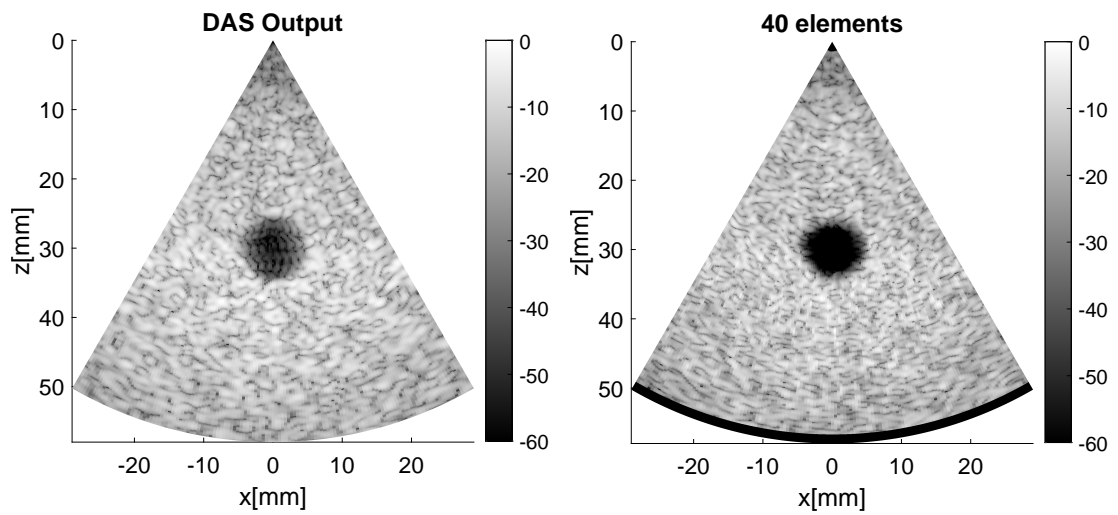
The effect of diagonal loading for a set of subarray sizes is shown in Figure 4.33 on page 73. However, the results from Figure 4.28, Figure 4.29, and Figure 4.30 visualise how the diagonal loading and subarray size are connected regarding the contrast metrics.

## 4.4 Contrast

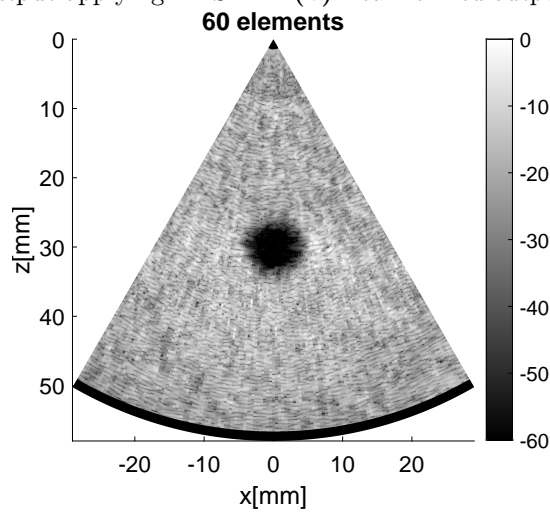
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**Figure 4.31:** The beamformed output applying DAS and the Capon beamformer, applying the subarray sizes  $L = 10, 20$ , and  $32$  elements.



(a) Beamformed output applying DAS. (b) Beamformed output when applying  $L = 40$ .

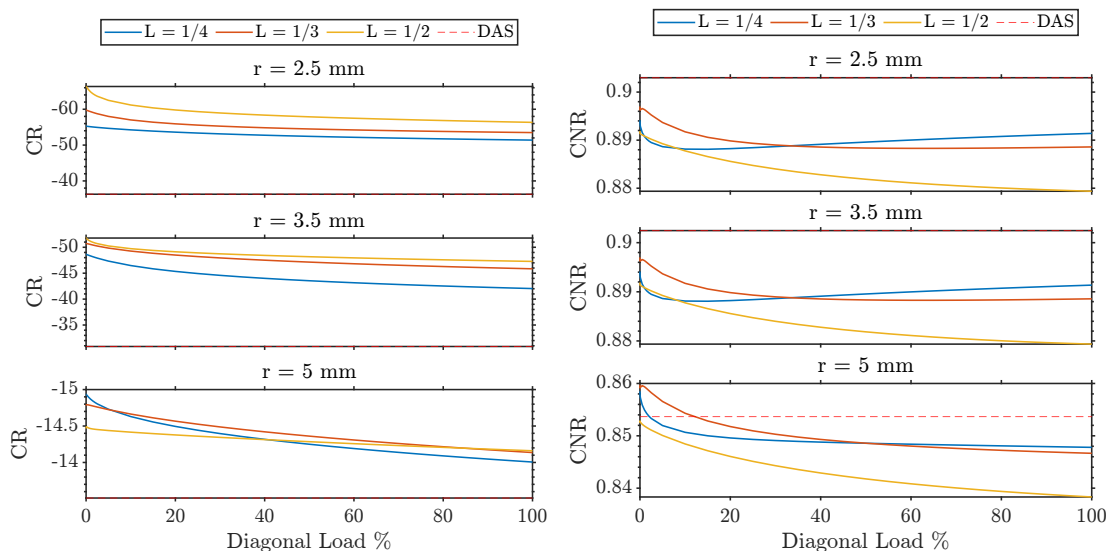


(c) Beamformed output when applying  $L = 60$ .

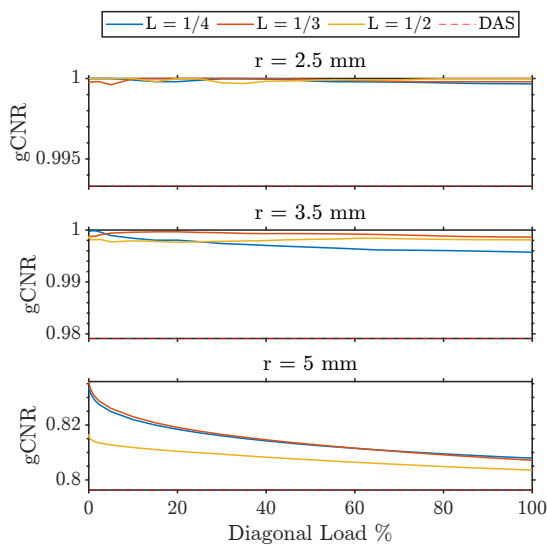
**Figure 4.32:** The beamformed output applying DAS and the Capon beamformer, applying the subarray sizes  $L = 40$ , and  $60$  elements.



## 4.4 Contrast



(a) The measured CR when applied diagonal load increases. Each subplot visualises the corresponding CR from three different applied ROIs. (b) The measured CNR when the applied diagonal load increases. Each subplot visualises the corresponding CNR from three different applied ROIs.



(c) The measured gCNR when the applied diagonal load increases. Each subplot visualises the corresponding gCNR from three different applied ROIs.

**Figure 4.33:** The contrast metrics CR, CNR, and gCNR measured for three different ROIs when increasing the diagonal load, for set subarray sizes of  $L = \frac{1}{4}M$ ,  $\frac{1}{3}M$ , and  $\frac{1}{2}M$ . The DAS output is used as a benchmark, shown as a red, dashed line. Each upmost subplots are  $ROI_1 = 2.5$  mm, each middle subplots are  $ROI_2 = 3.5$  mm, and each bottom subplots are  $ROI_3 = 5$  mm.



# Chapter 5

## Conclusion and further work

In this thesis, an analysis of the performance of the Capon beamformers was done to evaluate resolution and contrast, using the DAS beamformer performance as a benchmark. The analysis was applied to simulated medical ultrasound data, and was based on parameter studies to assess how the amount of subarray averaging and diagonal load affect the ultrasound image resolution and contrast. The study is done by examining the separability between two scatterers in a scene. The contrast metrics are examined from a scene containing a cyst, as well as the stability of the covariance matrix inversion measured by the condition number. The eigenvalues of the covariance matrix were examined to assess the potential use to understand beamformer performance.

When analysing the resolution as a function of a varying subarray size, it was found that two scatterers located at a lateral distance of  $d = 2$  mm apart at a depth of 30 mm are well resolved for a subarray size between  $L_{\min} = \frac{1}{4}M$  and  $L_{\max} = \frac{2}{3}M$ , where  $M$  is the full array length. Through analysis of scenes with two scatterers in a speckle background, the lower boundary  $L_{\min}$  is increased to  $L_{\min} = \frac{1}{3}M$ , while the upper boundary  $L_{\max}$  is maintained.

The condition number of the covariance matrix estimate provides insight into the stability of the matrix inversion, and was shown to be an applicable measure of matrix inversion stability with respect to the applied diagonal load. From the

perspective of image quality, it was shown that a small amount of diagonal loading, i.e.  $\approx 1\%$ , resulted in a significant increase in stability of the covariance matrix inversion. The increased stability results in a more accurate weight vector.

The eigenvalues did not provide any additional information regarding the resolution and separability of scatterers in a scene, as a general analysis of the amplitude provided this information. By applying subarray averaging, the eigenvalues was shown to have an increased amount of significant eigenvalues, and the eigenvalues had an increased magnitude. An increased speckle level also increased the amount of significant eigenvalues. However, these observations did not provide any additional information regarding the resolution of the two scatterers either.

By analysing the contrast metrics CR, CNR, and gCNR when varying both the applied diagonal load and subarray size, as well as considering three different ROIs, it was shown that a subarray size between  $\frac{1}{3}M$  and  $\frac{1}{2}M$  overall results in better Capon beamformer image contrast. For the diagonal load, an applied load of  $\approx 1\%$  is deemed sufficient to increase the stability of the matrix inversion. The applied diagonal load affects neither the resolution nor the contrast greatly. There were no significant deviations from these findings when considering different ROIs, other than the ROI at the theoretical edge of the cyst. This ROI was affected differently by the varied subarray averaging and diagonal loading, as this ROI contained larger amounts of the surrounding speckle dependent on the applied subarray averaging and diagonal loading. Hence, the results from the contrast analysis and the resolution analysis coincide.

Primarily, the conducted research within the current thesis has been carried out exclusively by applying simulated data. Consequently, it is crucial to replicate the study using recorded in-vivo data to confirm the discovered findings within the current thesis.

The conclusion made in this thesis shows what an appropriate choice of subarray averaging and diagonal load are to get optimal resolution and contrast in a simulated scene. The scene of which the analysis was done in the current thesis is however quite homogeneous regarding the location of the scatterers.

One further approach would be to consider an increased variability of scatter

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locations regarding both lateral and axial positions. This could help when building more generalised conclusions on the subarray averaging and diagonal loading parameters.

Setting the diagonal load based on a condition number criterion is a proposed approach. By doing this, one can examine how the Capon beamformer depends on diagonal load to maintain a given matrix inversion stability. Thereby, a possible method of it could be to adaptively decide the diagonal load for each pixel. This might also include adapting the subarray size  $L$  to increase the matrix inversion stability.

Further, a study of the statistics of the resolution and contrast metrics can be done over multiple speckle scenes. In the current thesis, only one speckle scene was considered. A study of how these statistics change when altering the subarray averaging and diagonal loading for different scenes could help build the foundation of when to choose what parameters regarding the applied subarray averaging and diagonal loading.



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