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Fast spectral velocity estimation using adaptive techniques: \textit{In-vivo} results

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Abstract—Adaptive spectral estimation techniques are known to provide good spectral resolution and contrast even when the observation window (OW) is very short. In this paper two adaptive techniques are tested and compared to the averaged periodogram (Welch’s) for blood velocity estimation. The Blood Power spectral Capon (BPC) method is based on a standard minimum variance technique adapted to account for both averaging over slow-time and depth. The Blood Amplitude and Phase Estimation technique (BAPES) is based on finding a set of matched filters (one for each velocity component of interest) and filtering the blood process over slow-time and averaging over depth to find the power spectral density estimate. In this paper, the two adaptive methods are explained, and performance is assessed in controlled steady flow experiments and \textit{in-vivo} measurements.

The three methods were tested on a circulating flow rig with a blood mimicking fluid flowing in the tube. The scanning section is submerged in water to allow ultrasound data acquisition. Data was recorded using a BK8804 linear array transducer and the RASMUS ultrasound scanner. The controlled experiments showed that the OW could be significantly reduced when applying the adaptive methods without compromising spectral resolution or contrast.

The \textit{in-vivo} data was acquired using a BK8812 transducer. OWs of 128, 64, 32 and 16 slow-time samples were tested. Spectrograms with duration of 2 seconds were generated. Welch’s method required 128 samples to give a reasonable spectrogram, whereas the BPC only required 32 samples before the SNR became a limiting factor. The BAPES managed to display the spectrogram with sufficient quality at 16 slow-time samples.

I. INTRODUCTION

A typical B-mode ultrasound image consists of about 100 image lines, with each line being created by focusing the ultrasound transducer array on a single point on the current line. The backscattered waves are processed with dynamic receive focusing, and, ideally, only objects along the image line are interrogated. The velocity of moving blood can be estimated by imaging the same image line repeatedly. The slow-time signal (sampled with the pulse repetition frequency), for a specific depth, has a center frequency which is proportional to the axial velocity [1]. A common way of estimating the blood velocity is to estimate the power spectral density (PSD) of the slow-time signal. Displaying the PSD as a function of time (the so-called spectrogram) not only visualizes the blood velocity distribution, but also allows the operator to track the time-variations of the blood. The velocity distribution is important for detecting turbulent flow [2], [3] which can be used for diagnosing carotid artery stenosis [4].

In ultrasound imaging, the PSD is normally estimated using an averaged periodogram approach, also known as Welch’s method [5], [6]. However, as is well-known, the method suffers from either poor resolution or high leakage, or both, and to achieve sufficient spectral resolution, the duration of the observation window (OW) has to be long. This implies that a large number of transmissions (about 100) has to be carried out to obtain a sufficiently accurate PSD estimate. The following problems emerge:

- The temporal resolution of the spectrogram will be poor due to the long duration of the the averaging kernel used for computing the PSD estimate.
- Interleaved B-mode/Doppler sequences have to be used to update the B-mode image. The sequences can be interleaved so that every second transmission is a B-mode acquisition. This lowers the velocity range by a factor of two. Data can also be acquired in blocks, so that an entire B-mode image is acquired in one sweep. This leaves holes in the spectrogram.
- If the second solution is chosen, due to the lack of Doppler data during B-mode data acquisition, it is necessary to acquire B-mode data as seldom as possible, resulting in poor frame rate on the B-mode images.

A thorough explanation of the problems encountered in spectral Doppler ultrasound can be found in [7].

In this paper, we formulate a data-adaptive blood velocity spectral estimator based on the matched filterbank (MAFI) framework [6], [8]. The purpose is to investigate the possibility of decreasing the OW with maintained spectral resolution and contrast. With a decreased OW, more flexibility would be possible when designing the Doppler/B-mode sequences and all three of the aforementioned problems could potentially be solved.

II. METHODS

A. Data model

To estimate the blood velocity at a given location, a number of transmissions are carried out in the same direction. After focusing, the resulting image lines represent a time series over depth. Let $y_k(l)$ denote the available (stationary) data sample at slow-time (transmission-number) $l$ and fast-time (proportional to depth) $k$, of which the blood velocity spectrum is to be estimated. For a generic (axial) velocity $v_z$, $y_k(l)$ can
be expressed as [1] (assuming that In-phase and Quadrature
transducer, \( f \))

\[
y_k(l) = \alpha_v e^{j \left( \frac{2\pi}{f_{prf}} l + \frac{\pi}{2} \right)} + w_k(l),
\]

for \( k = 0, \ldots, K - 1 \) and \( l = 0, \ldots, L - 1 \), where \( \alpha_v \) denotes
the (complex-valued) amplitude of the sinusoidal signal at velocity \( v_z \),
assumed to be constant over the slow-time \( \omega \),
and \( w_k(l) \) is the residual (or noise) term containing the signal
components at all velocities different from \( v_z \). Furthermore,
\( \omega_v = 2\pi f_c \), with \( f_c \) denoting the center frequency of
the transducer, \( f_s \) the sampling frequency of the system,
\( f_{prf} \) the pulse repetition frequency, and \( c \) the speed of sound.
We note that the second term in the exponential of (1) is due to the
time shift occurring when the blood particles move between
transmissions. The problem of estimating the blood velocity
spectrum density can thus be expressed as forming an estimate of
\( |\alpha_v|^2 \), for each velocity of interest. For a given depth (fast-
time sample) \( k \), let

\[
y_k(l) \triangleq \left[ y_k(l) \ldots y_k(l + N - 1) \right]^T,
\]

for \( \ell = 0, \ldots, L - 1 \), with \( L = \tilde{L} - N + 1 \) representing the total
number of slow-time vectors in the time-series. By introducing
\( \phi = \omega / f_s \) and \( \psi = -2\omega v_z / c f_{prf} \), (2) can be written as

\[
y_k(\ell) = \alpha_v e^{j \phi + j \psi} + w_k(\ell),
\]

where \( w_k(\ell) \) is formed similar to \( y_k(\ell) \), and

\[
a_\psi \triangleq \left[ 1 \ e^{j \psi} \ldots e^{j \psi(N-1)} \right]^T.
\]

B. The averaged periodogram (Welch)

The averaged periodogram is a well known and well used
method for estimating the PSD and is given by

\[
P_{\text{welch}}(\psi) = a_\psi^H \hat{R} a_\psi,
\]

where

\[
\hat{R} = \frac{1}{K T} \sum_{k=0}^{K-1} \sum_{l=0}^{L-1} y_k(l)y_k^H(l),
\]

is the estimate of the auto-correlation matrix of the data.

C. Blood Spectral Power Capon (BPC)

The Blood Spectral Power Capon estimate of the PSD is
formed by the following optimization problem

\[
h_\psi = \arg \min_{h_\psi} h_\psi^H \hat{R} h_\psi \text{ subject to } h_\psi^H a_\psi = 1.
\]

This can be interpreted as finding a filter \( h_\psi \), for each
frequency of interest, which minimizes the filtered power of
the signal, under the constraint that the amplitude of the frequency
component of interest, passes through the filter unaffected.

An analytical solution to (7) can be found [6],

\[
\hat{P}_{\text{BPC}}(\psi) = \frac{1}{a_\psi^H \hat{R}^{-1} a_\psi}.
\]

D. Blood Amplitude and Phase Estimation (BAPES)

Here, a matched filter bank is used to estimate the PSD
of the blood process. The matched filterbank is found by
designing a filter (for each frequency of interest) based on the
following optimization problem

\[
h_\psi = \arg \min_{h_\psi} h_\psi^H Q_w h_\psi \text{ subject to } h_\psi^H a_\psi = 1,
\]

where \( Q_w \) is the covariance matrix of the noise and interference
term \( w_k(l) \). It should be stressed that the matrix \( Q_w \),
changes as different frequency components \( a_\psi \) are probed.

Solving the optimization problem in (9) is not as straightforward as solving (7). This is because no direct estimate of \( Q_w \) is available. The suggested solution is to use a 'two stage approach', where an initial amplitude estimate of the frequency of interest is found, and used to estimate the matrix \( Q_w \). Thereafter, the solution to (9) can be found. The details
of this process, however, are out of scope for this paper and the
interested reader is referred to [9].

III. RESULTS

In this section, we will evaluate the performance of the
discussed estimators using both steady state flow measured
in a circulating flow rig, and in-vivo measurements on the
common carotid artery.

A. Flow rig measurements

The three methods were tested on a circulating flow rig
with a blood mimicking fluid flowing in the tube. The scanning
section is submerged in water to allow ultrasound data
acquisition. Data was recorded using a BK8804 linear array
transducer and the RASMUS system [10]. The \( f_{prf} \) was 1.89
kHz and the maximum velocity in the tube was 0.05 m/s.
The velocity of the fluid was measured in the tube center.
The transmitting aperture consisted of the 16 central elements
focused with an F-number of 3.3. The data was processed with
dynamic receiving focusing using a Hanning apodization over all
128 receiving elements.

To obtain a reliable reference measurement, a weighted
averaged periodogram was used. The OW was chosen to be
\( N = 256 \) samples to ensure sufficient spectral resolution.
The number of samples used in the averaging was \( \tilde{L} = 400 \).
The number of samples used for averaging in the fast-time
direction was \( K = 40 \). Prior to PSD estimation, the data
was weighted using a Hamming window. 144 independent PSD
estimates were created and the mean of these are represented
as the light gray plot in Figs. 2 and 1. The reader should be
aware that the ultimate goal is to reduce the total number of
observations of the blood process \( \tilde{L} \). For the results in this
section, this is, however, not possible. This is due to the need
for a reference spectrogram, which is based on a weighted
averaged periodogram, which needs a long averaging kernel to
perform adequately. In the section describing the in-vivo results,
however, the averaging kernel will be chosen to depend on the
OW, and the benefits from having a shorter OW can be fully
exploited.

Next, the three methods described in Section II were tested
as a function of OW. Again, 144 PSD estimates were created
of the flow process. The mean of the PSD estimates for $N = 128$ can be seen in Fig. 1. $L$ was still chosen to be 400 and $K = 40$. All three methods are able to resolve the spectrum of the blood signal. Welch’s method displays somewhat more spectral smoothing compared to BPC and BAPES.

![Power density spectra for N = 128](image1)

**Fig. 1.** The mean of the PSD estimates for $N = 128$, $L = 400$ and $K = 40$. The reference PSD is given in light gray. All three methods are able to resolve the spectrum of the blood signal. Welch’s method displays more spectral smoothing compared to BPC and BAPES.

It is well known that for long OWs $^1$ Welch’s methods has good spectral resolution, so the interesting thing is to investigate the methods performance when $N$ is small. In Fig. 2, the mean of 144 PSD estimates for $N = 4$, $L = 400$ and $K = 40$ can be seen. It is obvious that Welch’s method fails miserably in finding the PSD. Also the BPC has trouble in finding the PSD and displays one sharp peak instead of the true distribution. Only the BAPES can be considered to resolve the PSD as well as the reference method.

![Power density spectra for N = 4](image2)

**Fig. 2.** The mean of the PSD estimates for $N = 4$, $L = 400$ and $K = 40$. The reference PSD is given in light gray. Welch’s method fails miserably in finding the PSD. Also the BPC displays one sharp peak instead of the true distribution. Only the BAPES can be considered to resolve the PSD as well as the reference.

The mean relative bias compared to the maximum of the reference PSD is given by

$$
\text{MRB} = 10 \log_{10} \left\{ \frac{1}{P_{\text{ref}}} \sum_{p=0}^{P-1} \frac{\tilde{P}(\psi_p) - P_{\text{ref}}(\psi_p)}{P_{\text{ref}}(\psi_p)} \right\}, \quad (10)
$$

where $\tilde{P}(\psi_p)$ is the mean of the PSD estimate in question, $P_{\text{ref}}(\psi_p)$, is the reference PSD and $P$ is the number of frequency values investigated. The MRB is given in Fig. 3 for the three methods. Welch’s method displays worst performance when $N$ is small. Welch’s method and the BPC, both improve their MRB as $N$ grows whereas the BAPES has a relatively constant MRB as a function of $N$ showing superior performance at small $N$.

![Mean relative bias](image3)

**Fig. 3.** The MRB for the three methods. BAPES shows the best performance and is not improved as $N$ grows. The BPC is better than Welch’s method and improves in terms of MRB as $N$ grows.

### B. Measurements on the common carotid artery

*In-vivo* measurements were also carried out. The RASMUS system was used when collecting the data using a BK8812 linear array transducer. The transmitted pulse was a single cycle sinusoid at 5 MHz and the $f_{prf}$ was 12 kHz. To make the data narrowband, the received echoes were filtered using an eight cycle sinusoid at 5 MHz before further processing. The common carotid artery of a healthy 36 year old male was scanned by one of the authors. The spectra were plotted as a function of time, so that the full temporal velocity distribution could be visualized. The OW was varied and the spectrograms for $N = 128$ and $N = 16$ can be seen in Fig. 4 and Fig. 5, respectively.

The number of segments in slow-time used when estimating the covariance matrix was chosen to be $L = N + 1$. This was done to achieve the improved time-resolution possible when decreasing the OW. The number of fast-time samples was still $K = 20$ and approximately 1.7 seconds of data were generated. The first experiment involved $N = 128$ and the result can be viewed in Fig. 4 at a dynamic range of 40 dB. When using a long OW, all three methods are able to show the spectrogram properly. The sidelobes visible in the Welch spectrogram are a result of the rectangular weighting of the data. These could easily be removed using another weighting at the expense of spectral resolution.

Next, a shorter OW was chosen. Here $N = 16$ and the result can be seen in Fig. 5 with a dynamic range of 40 dB. The spectral resolution of Welch’s methods is much to poor to reproduce the spectrogram correctly. The BPC has sufficient spectral resolution and so has BAPES. BAPES displays slightly better signal to noise ratio than the BPC.
Fig. 4. The spectrogram for the three methods when \( N = 128 \) with a dynamic range of 40 dB. All three methods are able to properly show the spectrogram. The sidelobes visible in the Welch spectrogram are a result of the rectangular weighting of the data. These could easily be removed using another weighting at the expense of spectral resolution.

Observe the finer temporal detail for the BPC and BAPES in the spectrograms in Fig. 5 compared to Fig. 4. This results from the shorter temporal averaging when estimating the spectrogram.

IV. CONCLUSION

In this paper, we have tested two data-adaptive blood velocity spectral estimators (BPC and BAPES). The two methods were compared to the averaged periodogram (Welch’s method) via flow measurements in a circulating flow rig and in-vivo measurements on the carotid artery. It has been shown that, by using the adaptive methods the OW used for estimating the spectrogram can be significantly decreased with maintained spectral resolution. This offers more flexibility when acquiring spectral Doppler data and B-mode data simultaneously. Therefore, the frame rate of the B-mode images could potentially be increased. Furthermore, the time resolution of the spectrogram will increase, offering finer temporal details to be studied.

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