FUZZY ENTROPY BASED NONNEGATIVE MATRIX FACTORIZATION FOR MUSCLE SYNERGY EXTRACTION

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ABSTRACT

The concept of muscle synergies has proven to be an effective method for representing patterns of muscle activation. The number of degrees of freedom to be controlled are reduced while also providing a flexible platform for producing detailed movements using synergies as building blocks. It has previously been shown that small components of movement are crucial to producing precise and coordinated movement. Methods which focus on the variance of the data make it possible to overlook these small components in the synergy extraction process. However, algorithms which address the inherent complexity in the neuromuscular system are lacking. To that end we propose a new nonnegative matrix factorization algorithm which employs a cross fuzzy entropy similarity measure, thus, extracting muscle synergies which preserve the complexity of the recorded muscular data. The performance of the proposed algorithm is illustrated on representative EMG data.

Index Terms— Matrix Factorization, NMF, Fuzzy Entropy, Muscle Synergies, EMG

1. INTRODUCTION

The human body is capable of great dexterity and coordination, giving it the capability to produce a wide array of precise detailed movements but, as such, it is a highly complicated and also redundant system [1]. With such large numbers of degrees of freedom, how the central nervous system implements a control strategy is of interest in numerous different fields, from robotics to rehabilitation [2]. The concept of synergies, that is, common patterns of muscle activations which can be reused and serve as building blocks to produce detailed movements, has been gaining traction in recent years [3, 4, 5]. Synergies provide an approach which reduces the number of degrees of freedom to be controlled individually whilst providing the flexibility and adaptability to produce a wide range of motions.

Effective assessment of muscle synergies requires an accurate method for extracting the synergies. The standard approach to muscle synergy extraction is to use a nonnegative matrix factorization (NMF). That is, given a data matrix $X \in \mathbb{R}^{N \times N}$ and positive integer $K < \min \{X, N\}$ find nonnegative matrices $W \in \mathbb{R}^{X \times K}$ and $H \in \mathbb{R}^{N \times N}$

$$\min_{W,H} D(Y \mid WH) \quad \text{subject to} \quad W \geq 0, \quad H \geq 0,$$

where $D(Y \mid WH)$ is an appropriately selected measure of goodness of fit [6], with the most common NMF algorithms being the alternating least squares (ALS) and the multiplicative Lee-Seung algorithms [7]. The consideration of the choice of algorithm used to extract muscle synergies has been limited to only a relatively small number of studies. For instance, the authors in [3] have compared different factorization algorithms including NMF, principal component analysis (PCA) and independent component analysis (ICA), and NMF algorithms based on signal dependent noise have been developed in [8] and [9].

While many of the approaches to identifying muscle synergies perform well in terms of producing a general factorization of the data, they do not always consider the full problem [10]. The success of the factorization is often based on assessing the variance accounted for (VAF) of the extracted synergies. However, success should be based not only on finding a representation which can, to a large extent, fit the data but it is also crucial to consider the task being performed. The reconstructed data should be capable of producing movement which could complete the required task [2, 11].

Of particular interest is the ability to produce not just movement in terms of the broad strokes but also the fine, precise movement which produces detailed coordinated actions. Our previous work has shown that improvements in coordination of finger movement are represented in the small components of the synergistic movement [12]. Therefore, if we want to successfully extract muscle activity which represents not only to the majority of the movement but also the subtle differences which account for better coordination we need to consider alternative measures as VAF may not be sufficient.

When it comes to the classification and assessment of changes in muscle activity recorded via surface electromyography (EMG), entropy is a commonly used tool [13, 14, 15], with measures based on approximate entropy (ApEn) [16] being a popular choice. ApEn provides a family of statistics to allow estimation of the rate of information generation in a time series, with sample entropy ( SampEn) [17] being a modification of ApEn aimed at reducing the bias inherent in the calculation of ApEn. However, both ApEn and SampEn rely on an absolute decision boundary to determine whether two vectors are similar or not. In many systems, particularly biological systems which display a wide range of variability, such an absolute classification is not always realistic and the algorithm becomes sensitive to the choice of this decision boundary. In contrast fuzzy entropy (FuzzyEn) [18] employs a continuous fuzzy decision boundary to give a relative degree of similarity, resulting in a more consistent measure of entropy which is less sensitive to parameter selection [19]. Furthermore, FuzzyEn has been shown to be an ef-
fective measure of EMG data [18, 20].

As entropy measures have proven effective in characterizing EMG data, and any reconstructed data from a factorization of such EMG data must produce muscle activations capable of replicating the desired movement. It is therefore a natural extension to include an entropy measure into the muscle synergy extraction. Entropy rates have previously been used in blind source separation [21, 22], and for NMF corentropy [23] has been shown to provide an accurate similarity measure [24, 25, 26]. We therefore propose to include a FuzzyEn based measure into the update of the NMF algorithm.

The rest of this paper is organised as follows, in Section 2 we first introduce the cross FuzzyEn (C-FuzzyEn) [27] measure and then derive the update of a gradient descent NMF algorithm using a C-FuzzyEn similarity measure. In Section 3 we then test the algorithm for muscle synergy extraction on EMG data and compare the algorithm performance with that of the standard alternating least squares algorithm. Finally, Section 4 concludes the paper.

2. FUZZY ENTROPY NMF

One of the keys to successful NMF is the selection of the distance measure between the data matrix, Y, and the decomposed matrices, W and H. Standard measures such as the L2 norm distance or the Kullback-Leibler divergence are linear similarity measures and may not be appropriate in situations where the data contains nonlinearities or are extremely noisy. With that in mind, and in order to preserve the complexity of the data, our algorithm employs the C-FuzzyEn between Y and W \cdot H as a similarity measure between the data and the decomposition. The measure is first applied row-wise between Y and W \cdot H in order to update W, followed by column-wise for the update of H. Such an approach ensures both the individual muscle activity and the interaction between the muscles is represented as accurately as possible by the decomposition.

2.1. Cross Fuzzy Entropy

FuzzyEn was introduced to overcome limitations of ApEn and SampEn due to their reliance on the use of the Heaviside function. As discussed previously both algorithms employ an absolute boundary in their determination of whether two vectors are similar or not. In contrast FuzzyEn employs an exponential function to give a continuous degree of similarity between vectors based on their closeness [18]. As a natural extension of FuzzyEn, C-FuzzyEn has been introduced in order to assess the degree of similarity between two different signals using fuzzy entropy [27, 28]. The C-FuzzyEn of two time series of N points,

\[ u(i) : 1 \leq i \leq N \]
\[ v(j) : 1 \leq j \leq N , \]

is estimated from the N − m vectors of length m,

\[ x_m(i) = \{ u(i+k) : 0 \leq k \leq m-1 \} - \overline{u}(i) \]
\[ y_m(j) = \{ v(j+k) : 0 \leq k \leq m-1 \} - \overline{v}(j) , \]

where the mean values of the vectors, \( \overline{u}(i) \) and \( \overline{v}(j) \), are removed to eliminate any baseline and allow comparison of their similarity based only on their shape rather than absolute position. The distances between any pair of vectors is given in terms of the infinity norm, that is the maximum absolute difference between their scalar components

\[ d_{i,j}^m = \| x_m(i) - y_m(j) \|_\infty , \]

Their degree of similarity can then be defined in terms of a fuzzy function of this distance. In this case the exponential function \( \exp(-\lambda p/r) \), where \( p \) defines the gradient of the boundary and \( r \) the width of the boundary, such that

\[ D_{i,j}^m(p,r) = \exp \left( -\frac{(d_{i,j}^m)^p}{r} \right) . \]

The overall similarity function for vectors of length \( m \), \( \phi^m \), then becomes

\[ \phi^m(p,r) = \frac{1}{N-m} \sum_{i=1}^{N-m} \left(\frac{1}{N-m} \sum_{j=1}^{N-m} D_{i,j}^m(p,r) \right) , \]

and in the same vein the similarity function of vectors of length \( m+1 \) is given by

\[ \phi^{m+1}(p,r) = \frac{1}{N-m} \sum_{i=1}^{N-m} \left(\frac{1}{N-m} \sum_{j=1}^{N-m} D_{i,j}^{m+1}(p,r) \right) . \]

The C-FuzzyEn of the two time series can be expressed in terms of the ratio of the negative log of the conditional probability \( \phi^{m+1}/\phi^m \) which for a finite time series is equivalent to

\[ \text{C-FuzzyEn}(m, p, r, N) = \ln \phi^m(p,r) - \ln \phi^{m+1}(p,r) . \]

For simplicity the terms C-FuzzyEn(m, p, r, N), \( \phi^m(p,r) \) and \( \phi^{m+1}(p,r) \) will be referred to as C-FuzzyEn, \( \phi^m \) and \( \phi^{m+1} \) in the following derivation.

2.2. Proposed NMF Update

First we define the measure of goodness of fit \( D(Y||WH) \) in the NMF algorithm (1) in terms of the C-FuzzyEn (10) of the elements of \( Y \) and \( W \cdot H \). Then using a gradient descent rule, where the gradient of the cost function is taken row-wise with respect to the individual elements of \( W \), and column-wise with respect to the elements of \( H \), we have the updates

\[ [W]_{x,k} \leftarrow [W]_{x,k} - \eta^W \nabla_{[W]_{x,k}} D(Y||WH) , \]
\[ [H]_{k,n} \leftarrow [H]_{k,n} - \eta^H \nabla_{[H]_{k,n}} D(Y||WH) , \]

where \( \eta^W \) and \( \eta^H \) are the respective learning rates. For simplicity, we will consider the cost function in terms of its constituent parts, \( \ln \phi^m \) and \( \ln \phi^{m+1} \), and take the gradient of each separately.

For the row-wise measure of C-FuzzyEn we substitute

\[ y^m_x(i) = \{ Y_{x,i}, \ldots , Y_{x,i+m-1} \} - [Y^m]_{x,i} , \]
\[ w^m_h(j) = \{ [WH]_{x,j}, \ldots , [WH]_{x,j+m-1} \} - [WH]^m]_{x,j} , \]

into the distance measure (6). Then taking the gradient of \( \ln \phi^m \) with respect to the components of \( W \) gives

\[ \nabla_{[W]_{x,k}} \ln \phi^m = \frac{1}{\tilde{\phi}^m} \sum_{i=1}^{N-m} \sum_{j=1}^{N-m} \left( \frac{p}{r} D_{i,j}^m (d_{i,j}^m)^{p-1} \left( [H]_{k,n} - [WH]^m_{k,n} \right) \right) \cdot \left( w^m_h(j)(c^m) - w^m_h(i)(c^m) \right) \]

\[ \Phi^m_{x,k} \]

where \( c^m \in [0, m-1] \) such that the \( c^m \)th elements of \( y^m_x(i) \) and \( w^m_h(j) \), that is \( y^m_x(i)(c^m) \) and \( w^m_h(j)(c^m) \) respectively, are the
elements which correspond to the maximum absolute difference between the vectors
\[
\|y^m_x(i)(c^m) - wh^m_x(j)(c^m)\| = \|y^m_x(i) - wh^m_x(j)\|_\infty. \tag{16}
\]

In the same manner if we consider the column-wise measure of C-FuzzyEn, calculating the distance measure between the vectors
\[
y^m_x(i) = \{y_{i,n}, \ldots, y_{i+m-1,n}\} = \{y^m_x(i)\}_{k,n}, \tag{17}
\]
and
\[
wh^m_x(j) = \{WH_{j,n}, \ldots, WH_{j+m-1,n}\} - \{WH^m_x(j)\}_{k,n}, \tag{18}
\]
taking the gradient of \(\ln \phi^m\) with respect to the components of \(H\) results in
\[
\nabla_{[H]_{k,n}} \ln \phi^m_x = \frac{1}{\phi^m_x} \sum_{i=1}^{X-m} \sum_{j=1}^{X-m} \left( \frac{p}{r} D_{i,j}^{m} \left( d_{i,j}^{m} \right)^{p-1} \right. \nonumber \\
\cdot \left. \left( [W]_{j+c,m,k} - [WH]_{j+k,c} \right) \right) \cdot \left( y^m_x(i)(c^m) - wh^m_x(i)(c^m) \right) \right) - \frac{1}{\phi^m_x X^{p+1}} \cdot \nonumber \\
\sum_{i=1}^{N-m} \sum_{j=1}^{N-m} \left( \frac{p}{r} D_{i,j}^{m+1} \left( d_{i,j}^{m+1} \right)^{p-1} \left( [H]_{k,j+c,m+1} - [WH]_{k,j+k,c} \right) \right) \cdot \left( y^{m+1}_x(i)(c^{m+1}) - wh^{m+1}_x(i)(c^{m+1}) \right). \tag{19}
\]

Following the same approach to calculate the gradients with respect to both \(W\) and \(H\) for \(\ln \phi^{m+1}\) gives the gradient of the C-FuzzyEn cost function with respect to the components of \(W\) as
\[

\nabla_{[W]_{x,k}} = \nabla_{W_{x,k}} \ln \phi^m_x \cdot \nabla_{W_{x,k}} \ln \phi^m_x + 1 \cdot \frac{1}{\phi^m_x} \sum_{i=1}^{X-m} \sum_{j=1}^{X-m} \left( \frac{p}{r} D_{i,j}^{m} \left( d_{i,j}^{m} \right)^{p-1} \right. \nonumber \\
\cdot \left. \left( [H]_{k,j+c,m} - [WH]_{k,j+k,c} \right) \right) \cdot \left( y^m_x(i)(c^m) - wh^m_x(i)(c^m) \right) \right) \right) \cdot \left( y^{m+1}_x(i)(c^{m+1}) - wh^{m+1}_x(i)(c^{m+1}) \right). \tag{20}
\]

and with respect to the components of \(H\) as
\[

\nabla_{[H]_{k,n}} = \nabla_{H_{k,n}} \ln \phi^m_x \cdot \nabla_{H_{x,k}} \ln \phi^m_x + 1 \cdot \frac{1}{\phi^m_x} \sum_{i=1}^{X-m} \sum_{j=1}^{X-m} \left( \frac{p}{r} D_{i,j}^{m+1} \left( d_{i,j}^{m+1} \right)^{p-1} \right. \nonumber \\
\cdot \left. \left( [W]_{j+c,m+1} - [WH]_{j+k,c} \right) \right) \cdot \left( y^m_x(i)(c^m) - wh^m_x(i)(c^m) \right) \right) \right) \cdot \left( y^{m+1}_x(i)(c^{m+1}) - wh^{m+1}_x(i)(c^{m+1}) \right). \tag{21}
\]

Substituting (20) and (21) into (11) and (12) respectively gives us the NMF update with a C-FuzzyEn based cost function distance measure.

### Table 1. Muscles recorded on each channel and location over muscle for muscles with multiple recordings - upper indicating closer to the body (origin) and lower closer to the hand (insertion).

<table>
<thead>
<tr>
<th>Channel</th>
<th>Muscle</th>
<th>Location</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Extensor digitorum</td>
<td>upper</td>
</tr>
<tr>
<td>2</td>
<td>Anconeus</td>
<td></td>
</tr>
<tr>
<td>3</td>
<td>Flexor carpi ulnaris</td>
<td>lower</td>
</tr>
<tr>
<td>4</td>
<td>Pronator teres</td>
<td></td>
</tr>
<tr>
<td>5</td>
<td>Flexor carpi radialis</td>
<td>lower</td>
</tr>
<tr>
<td>6</td>
<td>Flexor carpi radialis</td>
<td>lower</td>
</tr>
<tr>
<td>7</td>
<td>Palmaris longus</td>
<td></td>
</tr>
<tr>
<td>8</td>
<td>Pronator teres</td>
<td></td>
</tr>
<tr>
<td>9</td>
<td>Extensor carpi ulnaris</td>
<td>lower</td>
</tr>
<tr>
<td>10</td>
<td>Extensor digitorum</td>
<td></td>
</tr>
<tr>
<td>11</td>
<td>Extensor carpi radialis brevis</td>
<td></td>
</tr>
<tr>
<td>12</td>
<td>Extensor carpi radialis longus</td>
<td></td>
</tr>
<tr>
<td>13</td>
<td>Abductor pollicis brevis</td>
<td></td>
</tr>
<tr>
<td>14</td>
<td>Abductor digitii minimi</td>
<td></td>
</tr>
<tr>
<td>15</td>
<td>Biceps brachii</td>
<td>upper</td>
</tr>
<tr>
<td>16</td>
<td>Biceps brachii</td>
<td>lower</td>
</tr>
</tbody>
</table>

### 3. ALGORITHM PERFORMANCE

#### 3.1. EMG Data

The EMG data has previously been described in [29, 30], in brief the data was recorded from 5 healthy right-handed subjects, all of whom gave informed consent prior to the experiment. The EMG data was acquired using a g.USBamp (Guger Technologies, Graz, Austria) amplifier. The data was sampled at 4.8kHz and before analysis was bandpass filtered between 10-200Hz, rectified and down-sampled by a factor of 10. Surface electrodes were placed over 12 different muscles located in the upper arm, forearm and hand of the subjects, with 4 of the larger muscles having electrodes placed over 2 different locations giving a total of 16 channels (listed in Table 1) with a reference electrode placed on the subjects earlobe. The experimental procedure involved the subject performing a series of simple hand and wrist movements. For the purposes of illustrating the performance of our algorithm the trials representing movement related to grasping a cylindrical object were selected. For each subject the grasping movement was repeated 3 times which were manually divided into separate trials giving a total of 15 trials analysed.

#### 3.2. Algorithm Performance

The C-FuzzyEn has three parameters which need to be set. The parameter, \(m\), is the length of the vectors to be compared, a larger value of \(m\) may result in a more detailed representation of the dynamics but also requires either large quantities of data - which can be difficult to obtain for biological datasets, or a wider decision boundary which will consequently lead to a loss in information. An appropriate choice of vector length for short biological datasets is a value of \(m = 2\). The parameters \(p\) and \(r\) relate to the gradient and width, respectively, of the exponential function determining the fuzzy boundary. Too large and we risk information loss, and too small and the influence of noise is increased. Typically for \(p\) small integer values greater than 1 are used [19], in this case \(p = 2\). For experimental data it can be beneficial to set the value of \(r\) as a function of the standard deviation of the data, thus, allowing comparison of data with different amplitudes [17]. In this case as we consider the updates of \(W\) and \(H\) in terms of the rows and columns of \(Y\), for the updates
of $W$, $r = 0.2 \times \sigma_{Y,x}$, and for the updates of $H$, $r = 0.2 \times \sigma_{Y,n}$. The learning rates of the NMF were determined empirically with $\eta^W = 0.8$ and $\eta^H = 0.1$.

The performance of the proposed algorithm was compared against a standard alternating least squares (ALS) NMF algorithm. To compare how well the decomposition from each algorithm fit the data, the Akaike Information Criteria (AIC) [31] was used

$$\text{AIC}(\theta) = -2L + 2n_p,$$

where $L$ is the maximum log-likelihood and $n_p$ is the number of independent parameters in the model. As the true number of synergies is unknown we compared both algorithms across a range of different numbers of bases, $k$. The AIC was determined for the lowest residual error between the extraction and the original data for each value of $k$ for both algorithms. The average of the AIC across all trials for both algorithms, with the number of synergies ranging from 3 to 15, are shown in Table 2. These results indicate that, across all synergies, the proposed algorithm outperforms the standard ALS algorithm giving a better representation of the data in terms of the AIC.

In relation to muscle synergy extraction, the basis vectors contained within each column of the matrix $W$ represent the synergy coefficients illustrating how much each muscle contributes to the synergy. The matrix $H$ represents the time-varying coefficients of each basis across all the samples. To further compare the performance of the two algorithms we consider differences/similarities in the synergy coefficients contained within the $W$ matrices extracted via each algorithm. Figure 1 provides a representative illustration of the synergies extracted for $k=3$ for a single trial. Importantly, although there was only a small improvement in the AIC of the proposed algorithm over the ALS, as can be seen, there was a noticeable difference in the synergies extracted. The two different algorithms provide a significantly different representation of how the muscles contributed to the synergies. With the ALS NMF having a much more even spread of the contributions across all of the muscles. Whereas, the FuzzyEn NMF considered three muscles in particular

![Fig. 1. Sample extracted synergies for k=3, blue bars - FuzzyEn NMF, yellow bars - ALS NMF.](image)

Table 2. Comparison of the average AIC for the proposed FuzzyEn NMF and the ALS NMF across a range of number of synergies.

<table>
<thead>
<tr>
<th>Algorithm</th>
<th>Number of Synergies</th>
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<tbody>
<tr>
<td></td>
<td>3</td>
</tr>
</tbody>
</table>

4. CONCLUSIONS

We have presented a new NMF algorithm which incorporates C-FuzzyEn as the similarity measure in the update of the algorithm. The use of C-FuzzyEn allows us to decompose matrices whilst also preserving the complexity of our data. The preservation of this underlying dynamic structure is important when considering data for which standard measures of the variability within the data are insufficient. In particular we believe this to be the case for EMG data where small variability in the data contributes significantly to the quality and coordination of movements. We have highlighted the performance of the algorithm in comparison to a standard NMF algorithm for a representative EMG data set. The results illustrate that even small improvements in performance can significantly change the representation of the data. While we have shown the applicability of this algorithm for the factorization of EMG data sets, making it ideally suited to the extraction of muscle synergies, we also believe the algorithm could find application in other areas where the data under consideration contains small but significant fluctuations.

5. REFERENCES


