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**UNIVERSITÀ
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Doctoral Dissertation

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Doctoral Program in Bioengineering and medical and surgical sciences

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(33th Cycle)

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Muscle Synergy Assessment during Cyclic and Non-Cyclic Movements

Methodological Issues and Application-oriented Studies

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29 *Marco Ghislieri*
30
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Turin, July 12, 2021

1 Summary

2 Human locomotion and balance control are essential activities during daily life.
3 Human locomotion is one of the most complex motor tasks, due to the multiple
4 degrees of freedom of the musculoskeletal system, the several biomechanical
5 functions carried out during each gait cycle, and the high intra-cycle variability of
6 the muscle activation intervals. Balance control (e.g., single-leg stance) is also
7 important during daily living activities, as a single task as well as a component of
8 other more complex tasks, such as walking and running. It is a simple but
9 challenging condition for balance control and for this reason it is widely used for
10 training and rehabilitation programs.

11 Due to the complexity of the musculoskeletal system, the central nervous
12 system adopts different strategies to efficiently and accurately perform movements.
13 Even if there is no consensus regarding the neural strategies adopted by the central
14 nervous system to perform movements, the muscle synergy theory is the most
15 widely used in literature. According to this theory, the central nervous system
16 controls the activation level and the synchrony of small groups of muscles, rather
17 than controlling independently every single muscle involved in the movement.

18 The present doctoral thesis aims at proposing, validating, and applying novel
19 methods for extracting muscle synergies that can be applied to achieve deeper
20 insights into the neuromuscular control of human locomotion and balance control
21 tasks.

22 Considering human locomotion, two novel approaches for extracting muscle
23 synergies and selecting the optimal number of muscle synergies are proposed and
24 validated considering both physiological and pathological sample populations. The
25 novel sEMG pre-processing approach, based on the extraction of principal
26 activation intervals before muscle synergy extraction, allows a more interpretable
27 assessment of the modular organization of the central nervous system during

1 walking without any loss of information with respect to the standard pre-processing
2 approaches. Similar results are obtained considering both physiological and
3 pathological (patients affected by Parkinson’s disease) sample populations.
4 Moreover, the novel approach for selecting the optimal number of muscle
5 synergies, based on a multi-criteria decision analysis model (ELECTRE III),
6 reveals higher performance with respect to the standard threshold-dependent
7 approaches proposed in literature, considering both simulated and real sEMG data.
8 Considering balance control, instead, the modular organization during single-
9 leg stance is assessed by means of a novel approach based on the segmentation of
10 the sEMG time-instants relative to a “*well-balanced*” single-leg stance or an
11 “*unbalanced*” single-leg stance to be used as inputs of the muscle synergy
12 extraction algorithm. The consolidated approach is then applied to the analysis of
13 the modular organization of single-leg stance with and without visual feedback
14 considering a healthy sample population. The modular organization, assessed
15 through the novel approach, slightly changes when considering the eyes-closed
16 condition, revealing a reorganization of the activation levels and balance control
17 strategies.
18

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1 Chapter 1

2 Introduction

3 1.1 Human Motor Control

4 Human motor control is an outstandingly complicated system. Due to several
5 neurophysiological (e.g., motoneurons and neuromuscular junctions), anatomical
6 (e.g., muscles and joints), and kinematic (e.g., trajectories, accelerations, and
7 velocities) variables that determine the execution of the same movement [1], the
8 human motor system presents countless degrees of freedom (more than 1000).
9 Thus, several combinations of neurophysiological, anatomical, and kinematic
10 variables can be used to achieve the same movement, suggesting many possible
11 ways to perform the same motor task (e.g., walking or running).

12 In 1967, the degrees of freedom problem was assessed for the first time by
13 Nikolai Aleksandrovich Bernstein in his book “*The co-ordination and regulation*
14 *of movements*” [1]. Bernstein formally described for the first time this problem by
15 stating that:

16
17 *“The basic difficulties for co-ordination consist precisely in the extreme*
18 *abundance of degrees of freedom, with which the central nervous system [...] is*
19 *not at first in a position to deal”.*

20
21 Moreover, he gave a very modern definition of the coordination concept that
22 is still one of the supporting pillars of modern motor control theories:

23
24 *“The co-ordination of a movement is the process of mastering redundant*
25 *degrees of freedom of the moving organ, in other words its conversion to a*
26 *controllable system. More briefly, co-ordination is the organization of the control*
27 *of the motor apparatus”.*

28
29 In our body, the task of dealing with the motor system redundancy (i.e.,
30 multiple solutions for performing the same movement) is left to the Central Nervous

1 System (CNS). The CNS is composed of the brain and the spinal cord (**Figure 1.1**).
2 The latter ranges from the lower region of the brain (i.e., the medulla) to the lumbar
3 segments of the vertebral column, which encloses and protects the spinal cord. The
4 human spinal cord is composed of 31 different segments which give rise to paired
5 spinal nerves joined in the anterior and posterior roots. The posterior roots contain
6 afferent fascicles and are characterized by the presence of spinal ganglia, while
7 anterior roots contain the axons of those motor neurons (motor effector nerves) that
8 have their bodies in the ventral grey horns of the spinal cord. Finally, the
9 interneurons create reflex arcs between sensory and motor nerves, modulating the
10 information between the anterior and the posterior roots. **Figure 1.1** shows a
11 graphical representation of the central nervous system and the spinal reflex arc.

12 More specifically, 31 pairs of spinal nerves are present in the spinal cord: 8
13 pairs of cervical nerves (from C1 to C8), 12 pairs of thoracic nerves (from T1 to
14 T12), 5 pairs of lumbar nerves (from L1 to L5), 5 pairs of sacral nerves (from S1 to
15 S5), and 1 pair of coccygeal nerves. Cervical nerves are used to control muscles
16 involved in respiration and the movement of the head, neck, and arms. Thoracic

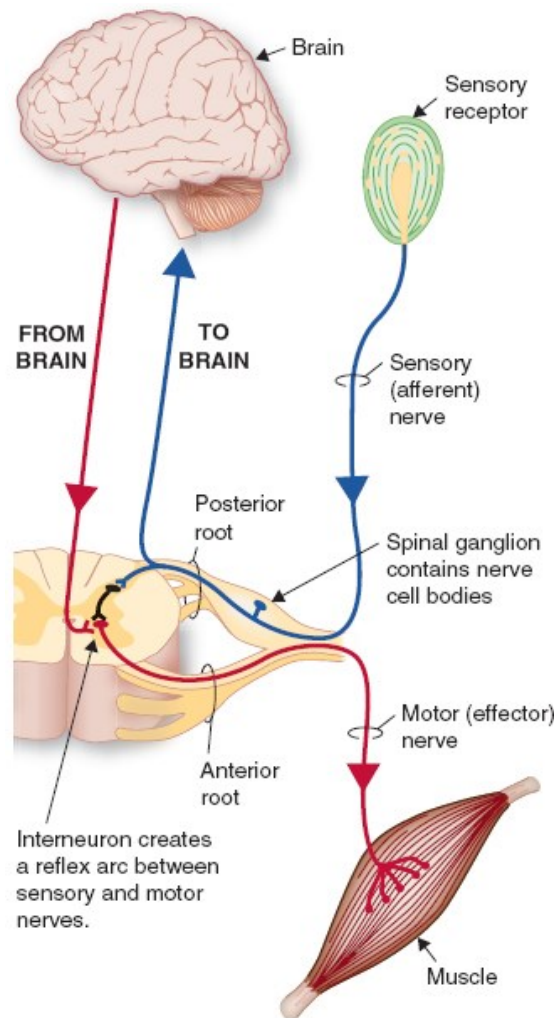


Figure 1.1 Schematic representation of the central nervous system and the spinal reflex arc. Sensory receptor generates an input signal which travels through the afferent pathway from the sensory receptor to the posterior root of the spinal cord. The motor output, instead, travels through the efferent pathway from the anterior root of the spinal cord to the muscle. The connection between the afferent (input) and efferent (output) pathways is mediated by the interneuron.

1 nerves are used to control muscles involved in finger and trunk movements. Lumbar
2 and sacral nerves are used to control locomotion, intestinal, and reproductive
3 functions.

4 Accordingly, how our CNS manages such complexity and selects only one
5 solution (i.e., a combination of neurophysiological, anatomical, and kinematic
6 variables) from all the other possibilities? How can we rapidly and accurately
7 perform movements having an outstanding amount of degrees of freedom to choose
8 from? What if our CNS, instead of choosing over multiple degrees of freedom,
9 reduces the motor control complexity by using a few combinations of low-
10 dimensional elements?

11 1.2 Muscle Synergy Theory

12 Theoretical Background

13 Despite several theoretical models (e.g., spinal force field, neuromotor
14 synergies, and unit burst generators) [2–5] have been proposed in literature over the
15 last decades, the muscle synergy theory is the one that achieved the greatest
16 consensus across researchers [6]. According to this theory, the CNS controls the
17 activation level (spatial component) and the synchrony (temporal component) of all
18 the muscles involved in a specific movement through the combination of a small
19 number of low-dimensional elements (called muscle synergies), rather than
20 controlling independently every single muscle. Moreover, some muscle synergies
21 have been demonstrated to be task-specific, while others may be shared among
22 different motor behaviors, such as walking and running [7].

23 **Figure 1.2** shows a schematic representation of the muscle synergy theory. The
24 CNS manages the multiple degrees of freedom of the motor system by controlling
25 small sets of muscles (called muscle synergies), rather than controlling every single
26 muscle involved in a specific movement, separately.

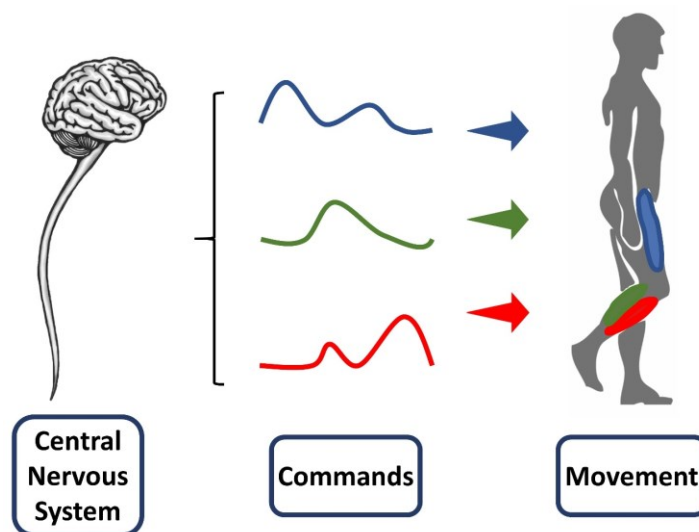


Figure 1.2 Schematic representation of the muscle synergy theory.

1 More specifically, two different definitions of the muscle synergy theory have
2 been proposed in literature: the “*time-invariant*” muscle synergies and the “*time-*
3 *varying*” muscle synergies [8]. Considering the “*time-invariant*” muscle synergies,
4 all the muscles within a specific muscle synergy are synchronously activated (i.e.,
5 there is no temporal modulation of the muscle activation levels), while, considering
6 the “*time-varying*” muscle synergies, each muscle within a specific muscle synergy
7 shows a temporal modulation over the task duration. In the present thesis, only the
8 “*time-invariant*” muscle synergy definition was considered, allowing for the
9 assessment of the temporal patterns of each muscle synergy.

10 Even if there is no consensus on the neural origin of muscle synergies [7,9,10],
11 several studies have supported the muscle synergy theory in several motor tasks,
12 both in animals and humans. Considering animal studies, several authors have
13 demonstrated a modular organization in the spinal cords and brainstems (in
14 particular in the medulla) of rats, frogs, and cats [2,6,11–14]. These studies have
15 shown that the spinal cord and brainstem express most of the muscle synergies for
16 motor behavior [3,15]. However, the job of activating and coordinating these low-
17 level motor structures (i.e., weight vectors) is left to the primary motor cortex and
18 other non-primary motor areas of the brain, such as the supplementary motor cortex
19 and premotor cortex [3]. These high-level motor structures (i.e., activation
20 coefficients) may contribute to the activation and coordination of the low-level
21 motor structures by selecting the best subset of muscle synergies for the execution
22 of a specific motor task.

23 Considering human studies, the assessment of patients affected by
24 neuromuscular or neurodegenerative diseases has contributed to the understanding
25 of muscle synergies in humans, revealing results that are in line with those found
26 considering animals. In particular, mildly-to-moderate impaired stroke survivors
27 with motor cortical lesions have shown, during voluntary arm movements, similar
28 muscle synergies between the affected and the unaffected arm, suggesting the
29 preservation of the low-level motor structures (i.e, weight vectors) [16]. However,
30 differences between the affected and the unaffected arm have been detected in the
31 temporal coordination of the muscle synergies, suggesting the presence of disrupted
32 high-level motor structures (i.e., activation coefficients) due to stroke-related
33 lesions [16].

34 The effectiveness of the muscle synergy theory in modeling the human modular
35 organization of the CNS during different motor tasks has been demonstrated in
36 several studies and research areas, such as clinics, robotics, and sport [17,18].
37 Among all the possible motor tasks, human locomotion is one of the most widely
38 studied movement to assess muscle synergies in humans. Several are the reasons
39 explaining why human locomotion is the most widely investigated motor task:

- 40 • the high number of muscles and joints involved
- 41 • the biomechanical functions carried out during each gait cycle
- 42 • is one of the most important Activities of Daily Living (ADL)
- 43 • it is easier with respect to other movements thanks to its cyclic nature
- 44 • the high cycle-by-cycle variability of the muscle activation intervals

- it can be easily performed by patients, children, and elderly that are able to independently walk without walking aids or external supports.

According to several studies, human locomotion can be modeled by a small set of muscle synergies characterized by specific biomechanical functions [19–22]. Besides inter-subject variability, the number of muscle synergies extracted and their biomechanical functions may vary depending on the type and number of muscles considered [23], the sEMG pre-processing techniques, and the factorization algorithm implemented. Nevertheless, the number of muscle synergies characterizing human locomotion is usually found equal to 4 or 5 [19,24].

Moreover, the muscle synergy theory has been recently used to gain a deeper understanding of human postural control [25–27]. The control of balance requires sensorimotor transformations that allow CNS to rapidly interpret multiple sensory input signals from all segments of the body to produce context-dependent muscle activation patterns that stabilize the body. According to several studies, the variations in muscle activation patterns evoked in response to different “controlled” perturbations (e.g., support-surface movements in the horizontal plane) could be described by a limited set of muscle synergies [25–29]. Muscle synergies that contribute to balance control have been demonstrated to be highly consistent across different balance tasks and levels of difficulty [26,30,31], suggesting that increasing the task complexity there should be only slight modifications to the basic motor control strategies involved in postural balance control. Depending on the type and the number of muscles acquired and the factorization algorithm implemented, the number of muscle synergies extracted during postural control ranges between 4 and 6.

Numerical Approaches

Muscle synergies are usually extracted from surface electromyographic (sEMG) signals acquired, during a specific motor activity, from a set of muscles that are mainly involved in the analyzed movement [6]. The factorization approach aims at identifying common sEMG patterns recorded from multiple muscles during specific motor activity.

Figure 1.3 shows the schematic representation of the muscle synergy extraction process. **Figure 1.3A** represents the sEMG signals acquired during a specific motor activity, e.g., walking, while **Figure 1.3B** shows the weight vectors and the activation coefficients extracted from the sEMG signals through factorization approaches. Finally, **Figure 1.3C** represents the activation coefficients and weight vectors that are used by the CNS to generate a specific motor activity (as described in **Figure 1.3A**).

In the last years, the modular organization of the CNS has been modeled as the linear combination of synergies as weight vectors in the space of muscle activations as defined by the following equation [32,33]:

$$M(t) = \sum_{k=1}^N C(t)_k \cdot W_k + e \quad (1.1)$$

1 where $M(t)$ is a matrix containing the original sEMG signals acquired from the
 2 observed muscles at a specific time point (t) [8], N is the number of synergies, W_k
 3 is a time-independent vector of non-negative weights (called weight vector [34]),
 4 $C(t)_k$ is a time-dependent set of non-negative muscle activation coefficients (called
 5 activation coefficients [34]), and e is the model reconstruction error [6]. According
 6 to this model, high-dimensional data, such as sEMG signals acquired from multiple
 7 muscles (**Figure 1.3A**), can be compactly represented by the linear combination of
 8 low-dimensional elements (i.e., weight vectors and activation coefficients) (**Figure**
 9 **1.3B**).

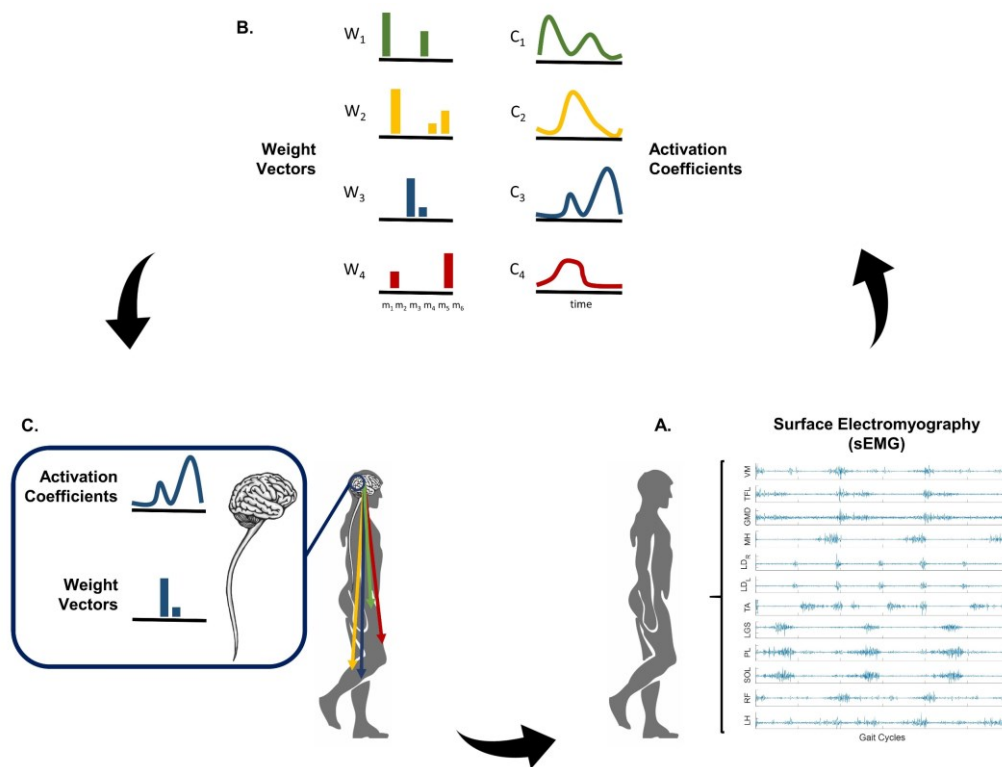


Figure 1.3 Schematic representation of the muscle synergy extraction process.

10 Several factorization methods for muscle synergy extraction have been
 11 proposed in literature. Some of the most widely used methods are the Principal
 12 Component Analysis (PCA), the Independent Component Analysis (ICA), the
 13 inverse Gaussian, and the Non-Negative Matrix Factorization (NNMF)
 14 [4,6,17,32,33,35–40]. The comparison among different factorization approaches
 15 for muscle synergy extraction reveals a high similarity in the computed muscle
 16 synergies [32,41]. However, since the nature of muscle activation is undeniably
 17 non-negative, the Non-Negative Matrix Factorization (NNMF) is one of the most
 18 widely used factorization approaches, as it does not allow for negative values.

1 According to the original definition of the NNMF approach by Daniel D. Lee
 2 and Sebastian H. Seung [41], a matrix of observations V can be decomposed in the
 3 product of two non-negative matrixes W and C as it follows (1.2):

$$V_{m \times t} \approx V_R = W_{m \times r} C_{r \times t} \quad (1.2)$$

4 where V_R represent the model reconstruction of the original observation matrix V
 5 containing the sEMG data, W is a matrix of dimension $m \times n$, and C is a matrix of
 6 dimension $n \times t$, with m being the number of observed muscles, t the number of
 7 recorded time-instants, and r the rank of factorization (i.e., the number of muscle
 8 synergies) [41].

9 **Figure 1.4** shows a schematic representation of sEMG data factorized using
 10 the NNMF approach. The matrix of observations V is graphically represented for
 11 six different muscles. Then, V can be approximated (V_R) through the linear
 12 combination of W (weight vector matrix) and C (activation coefficient matrix).
 13 Since the weight vectors are time-independent constants, W is usually represented
 14 through a bar diagram. Instead, since the activation coefficients describe the
 15 temporal modulation of the muscle synergies, C is usually represented as time-
 16 dependent curve. By multiplying and summing synergy-by-synergy the elements of
 17 W and C , it is then possible to reconstruct the matrix of observations V (representing
 18 the original sEMG data).

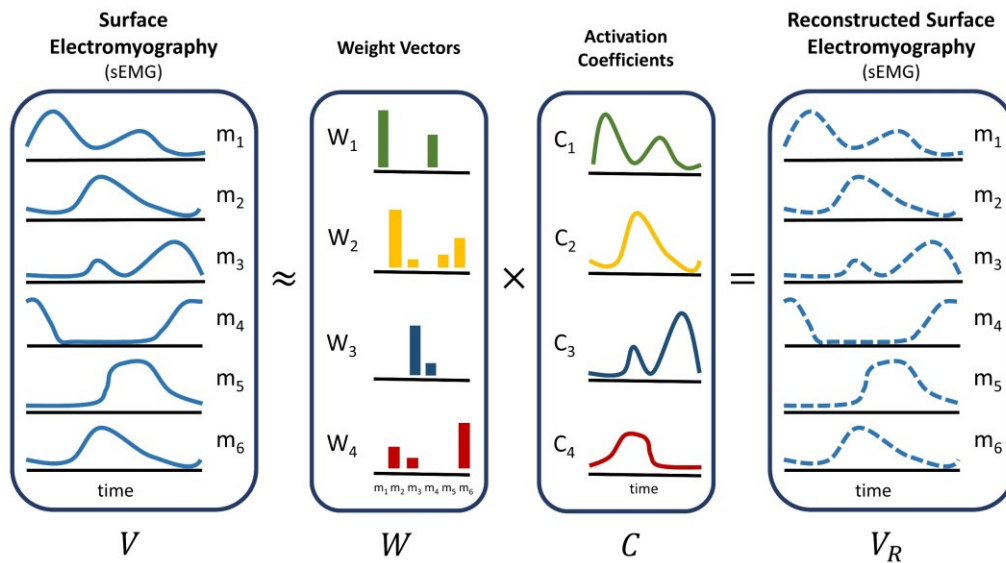


Figure 1.4 Schematic representation of sEMG data factorized through Non-Negative Matrix Factorization (NNMF) approach. Muscle activations of six different muscles (V) are compactly represented with four muscle synergies. Each weight vector (W) describes the time-independent contribution of each muscle to a specific muscle synergy. Each activation coefficient (C), instead, describes the time-dependent modulation of a specific muscle synergy. The multiplication of W and C gives an approximate reconstruction of the original sEMG data (V).

19 More specifically, the NNMF approach is an iterative process that aims at
 20 computing the W and C values that minimize the model reconstruction error (e).
 21 This optimization problem is solved by the NNMF approach by minimizing a cost
 22 function for measuring the divergence between the matrix of observations (V) and

1 the reconstructed matrix of observations (V_R) that was originally defined by Lee
 2 and Seung [42]. One of the most commonly used cost functions is the squared error
 3 (or Frobenius Norm), defined as follows:

$$F(W, C) = \|V - WC\|_F^2 \quad (1.3)$$

4 where V represents the observation matrix, W represents the weight vector matrix,
 5 and C the activation coefficient matrix.

6 An important consideration on factorization approaches is that, since none of
 7 them can automatically define the best factorization rank r (i.e., the optimal number
 8 of muscle synergies necessary to reconstruct the original sEMG data), a further step
 9 is needed to define the factorization rank that best models the original matrix of
 10 observations. To assess the reconstruction quality of the extracted muscle synergies,
 11 two metrics have been commonly used in literature: the Variance Accounted For
 12 (VAF) and the coefficient of determination R^2 . These metrics indicate to what
 13 extent the reconstructed sEMG data (obtained by multiplying W and C matrixes)
 14 are similar to the acquired sEMG signals (V). Both metrics are defined as follows:

$$VAF = \left(1 - \frac{RSS}{SST}\right) \cdot 100 = \left(1 - \frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\sum_{i=1}^n (y_i)^2}\right) \cdot 100 \quad (1.4)$$

$$R^2 = \left(1 - \frac{RSS}{SST}\right) \cdot 100 = \left(1 - \frac{\sum_{i=1}^n (y_i - \hat{y}_i)^2}{\sum_{i=1}^n (y_i - \bar{y}_i)^2}\right) \cdot 100 \quad (1.5)$$

15 where RSS represents the Residual Sum of Squares (or the sum of squared errors),
 16 SST the Sum of Squares Total (or the total sum of squares), y_i the observed data,
 17 \bar{y}_i the average of the observed data, and \hat{y}_i the estimate of the observed data through
 18 the factorization algorithm. Therefore, both metrics quantify the fraction of data
 19 variation accounted for by the muscle synergy reconstruction, but they differ in the
 20 data variation definition used for their computation. Considering R^2 metric, the data
 21 variation is computed with respect to the mean, while, for the VAF metric, the mean
 22 is not subtracted before data variation computation and, thus, the data variation is
 23 computed with respect to zero. High values of VAF or R^2 (i.e., close to 100%)
 24 indicate an accurate reconstruction of the original sEMG data, while low values
 25 (i.e., close to 0%) indicate that muscle synergies do not explain a large part of the
 26 sEMG variance. The minimum number of muscle synergies necessary to
 27 reconstruct the original sEMG data is then selected by analyzing the plot of the VAF
 28 (or R^2) versus all the tested number of muscle synergies (called VAF or R^2 curve)
 29 [16,23,32,43–47]. One of the most widely used criteria is the selection of the first
 30 point where the VAF or R^2 curve reaches a threshold that is commonly set equal to
 31 90% [24,43,44,46].

32 **Figure 1.5** shows an example of non-negative matrix factorization outcomes
 33 extracted from sEMG signals of a healthy subject during walking, where non-
 34 negative weight vectors are represented through bar diagrams and activation
 35 coefficients through curves. A large number of muscle activities (i.e., 12 muscle
 36 activities) can be modeled by a lower number of activation coefficients (C) and

1 weight vectors (W). In particular, during walking, foot plantar flexors (e.g., Lateral
 2 Gastrocnemius, Peroneus Longus, and Soleus) are used for propulsion in the second
 3 part of the stance phase, when their weight (or contribution) is highest (i.e., $W1$ in
 4 **Figure 1.5**). However, during the swing phase, their activity is not as “important”
 5 and thus their contribution is close to zero (i.e., $W2$, $W3$, $W4$, and $W5$ in **Figure**
 6 **1.5**).

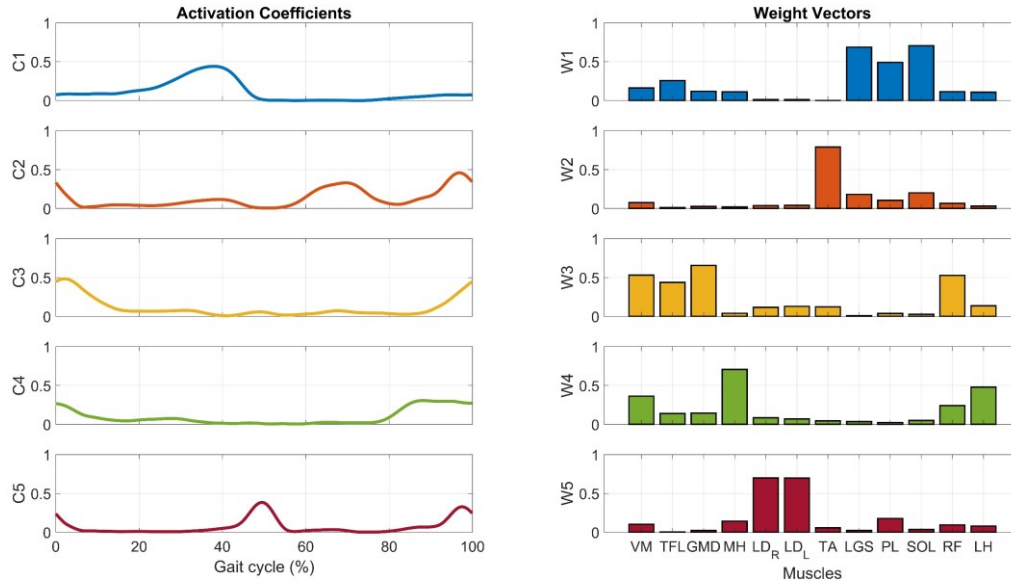


Figure 1.5 Example of muscle synergies extracted via Non-Negative Matrix Factorization (NNMF) from a set of muscles of a healthy subject during walking. Weight vectors (W) are represented through bar diagrams on an amplitude-normalized y -axis, while activation coefficients (C) are represented through curves on an amplitude-normalized y -axis. Muscle abbreviations: LD_R = right Longissimus Dorsii, LD_L = left Longissimus Dorsii, TFL = Tensor Fasciae Latae, GMD = Gluteus Medius, RF = Rectus Femoris, LH = Lateral Hamstring, MH = Medial Hamstring, VM = Vastus Medialis, LGS = Lateral Gastrocnemius, PL = Peroneus Longus, SOL = Soleus, and TA = Tibialis Anterior.

7 Another important consideration on factorization approaches is that each of
 8 them can be applied in many different ways and their outputs may vary depending
 9 on the initial data conditions. For example, several NNMF algorithms have been
 10 proposed in the last years based on different mathematical models [41,48].
 11 Additionally, the way initial random values of the NNMF approach are selected
 12 plays a role in the number and quality of the calculated muscle synergies [37].

13 The number and choice of the muscles to be acquired, the number of task
 14 repetitions, and the necessary sEMG pre-processing steps before factorization are
 15 other factors that can strongly influence the factorization output. Several studies
 16 have provided methodological recommendations to allow a more comprehensive
 17 muscle synergy analysis and guarantee comparability of the results
 18 [23,36,38,39,44,49–53]. The study by Steele *et al.* [23] demonstrated that the
 19 number and the structure of muscle synergies are dependent upon the number and
 20 choice of muscles included in the analysis. Indeed, due to constraints on time,
 21 experimental setup, and subject comfort, sEMG signals are usually measured only
 22 from a subset of muscles involved in the observed movement. In particular,
 23 researchers typically include the larger muscles thought to contribute to the
 24 observed movement and from which it is easier to acquire sEMG signals.
 25 Depending on the limb and movement observed, different numbers of muscles,

1 ranging from five to twenty-four muscles, have been measured to model motor
2 control strategies [31,54–59]. Results revealed that the average similarity of
3 synergies to the master set (e.g., muscle synergies calculated from all the acquired
4 muscles) dropped below 0.8 when fewer than eleven muscles were included in the
5 analysis. Moreover, analyses that included small subsets of muscles also
6 overestimated the variance accounted for by the synergies compared to the analysis
7 computed considering larger subsets of muscles.

8 The study by Oliveira *et al.* [38], instead, investigated the influence of data
9 structure (i.e., sEMG averaging, concatenation, and number of task repetitions) on
10 the extracted muscle synergies, providing practical guidelines on the
11 methodological aspects of muscle synergy extraction from sEMG signals during
12 locomotion. The same number of muscle synergies was extracted by NNMF from
13 all the acquired gait cycles separately, from averaging 2, 3, 5, 10, 20, and 40
14 consecutive gait cycles, and from the concatenation of the same sets of consecutive
15 gait cycles. However, the muscle synergies extracted considering the concatenation
16 of gait cycle subsets revealed a slightly reduced reconstruction accuracy compared
17 to the other two tested conditions. Nevertheless, muscle synergies extracted from
18 all the gait cycles separately or from averaging subsets of gait cycles did not account
19 for step-to-step variability.

20 Finally, in the work by Kieliba *et al.*[49], the effect of different low-pass filter
21 cut-off frequencies and amplitude-normalization techniques on muscle synergies
22 extracted from the upper limb muscles of healthy subjects during point-to-point 3D
23 reaching movements was assessed. Four different low-pass filter cut-off
24 frequencies (i.e., 0.5 Hz, 4 Hz, 10 Hz, and 20 Hz) and two different amplitude-
25 normalization techniques (i.e., maximum voluntary contraction and maximum
26 amplitude of the recorded sEMG signal) were tested. In terms of filter cut-off
27 frequencies, results showed that increasing the low-pass filter cut-off frequency had
28 the effect of decreasing the amount of variance accounted for and, perhaps, the
29 selected number of muscle synergies. Muscle synergies, instead, were not
30 significantly altered by the amplitude-normalization techniques.

31

32 **1.3 Aim of the Thesis and Organization**

33 The present doctoral thesis aims at proposing, validating, and applying novel
34 methods for extracting muscle synergies. These methods can be useful to provide
35 deeper insights into the neuromuscular control of human locomotion and single-leg
36 stance tasks. More specifically, this thesis is divided into two different sections
37 aimed at assessing the modular organization of the central nervous system during
38 cyclic (locomotion) and non-cyclic (single-leg balance) movements. In this thesis,
39 both methodological and application-oriented studies are presented and discussed.

40 **Chapter 2 - A Deep Learning Approach for Muscle Activity Detection** aims
41 at proposing a novel approach for extracting muscle activation intervals
42 (onset/offset time instants) from raw surface electromyographical (sEMG) signals
43 by using a deep learning approach based on Long Short-Term Memory (LSTM)

1 neural networks. Then, the proposed muscle activity detector will be used, in the
2 following chapter (**Chapter 3 - Motor Control Strategies during Cyclic**
3 **Movements**), as an sEMG pre-processing step before muscle synergy extraction.

4 **Chapter 3 - Motor Control Strategies during Cyclic Movements** aims at
5 assessing human motor control strategies during cyclic movements (i.e.,
6 locomotion) in both patients affected by Parkinson’s Disease and healthy subjects.
7 More specifically, two methodological studies are described. Those studies aim at
8 presenting and validating novel methods for extracting muscle synergies by
9 extracting principal activations (**Methodological Issue - Muscle Synergies**
10 **Extracted Using Principal Activations**) and for selecting the optimal number of
11 muscle synergies by applying a multi-criteria decision analysis approach
12 (**Methodological Issue – Multi-Criteria Decision Analysis for Selecting the**
13 **Optimal Number of Muscle Synergies**).

14 **Chapter 4 - Motor Control Strategies during Non-Cyclic Movements** aims
15 at assessing human motor control strategies during non-cyclic movements (i.e.,
16 single-leg stance). More specifically, one methodological and one application-
17 oriented study are described. The first one aims at presenting and validating a novel
18 method for extracting muscle synergies by distinguishing between the sEMG time-
19 instants relative to a “*well-balanced*” and an “*unbalanced*” single-leg stance
20 (**Methodological Issue – Muscle Synergy Assessment during Single-Leg**
21 **Stance**). The application-oriented study, instead, aims to assess the influence of
22 visual feedback on the modular organization of the central nervous system during
23 single-leg stance considering healthy subjects (**Application - Influence of Visual**
24 **Feedback on Muscle Synergies**).

25 **Chapter 5 - Conclusions and Perspectives** aims at summarizing the results
26 described in the previous chapters highlighting the advantages of the proposed
27 methodologies and the future perspectives.

28 Finally, in **Appendix A - Extraction of Principal and Secondary Activations**
29 **through CIMAP** chapter, a brief introduction to the Clustering for Identification
30 of Muscle Activation Patterns (CIMAP) algorithm is provided.

31 To improve the readability of the present thesis, at the beginning of each
32 chapter the muscle synergy theory, the factorization approach adopted, and the
33 sample population enrolled are briefly introduced.

34 **Figure 1.6** shows a graphical representation of the doctoral thesis’s contents.

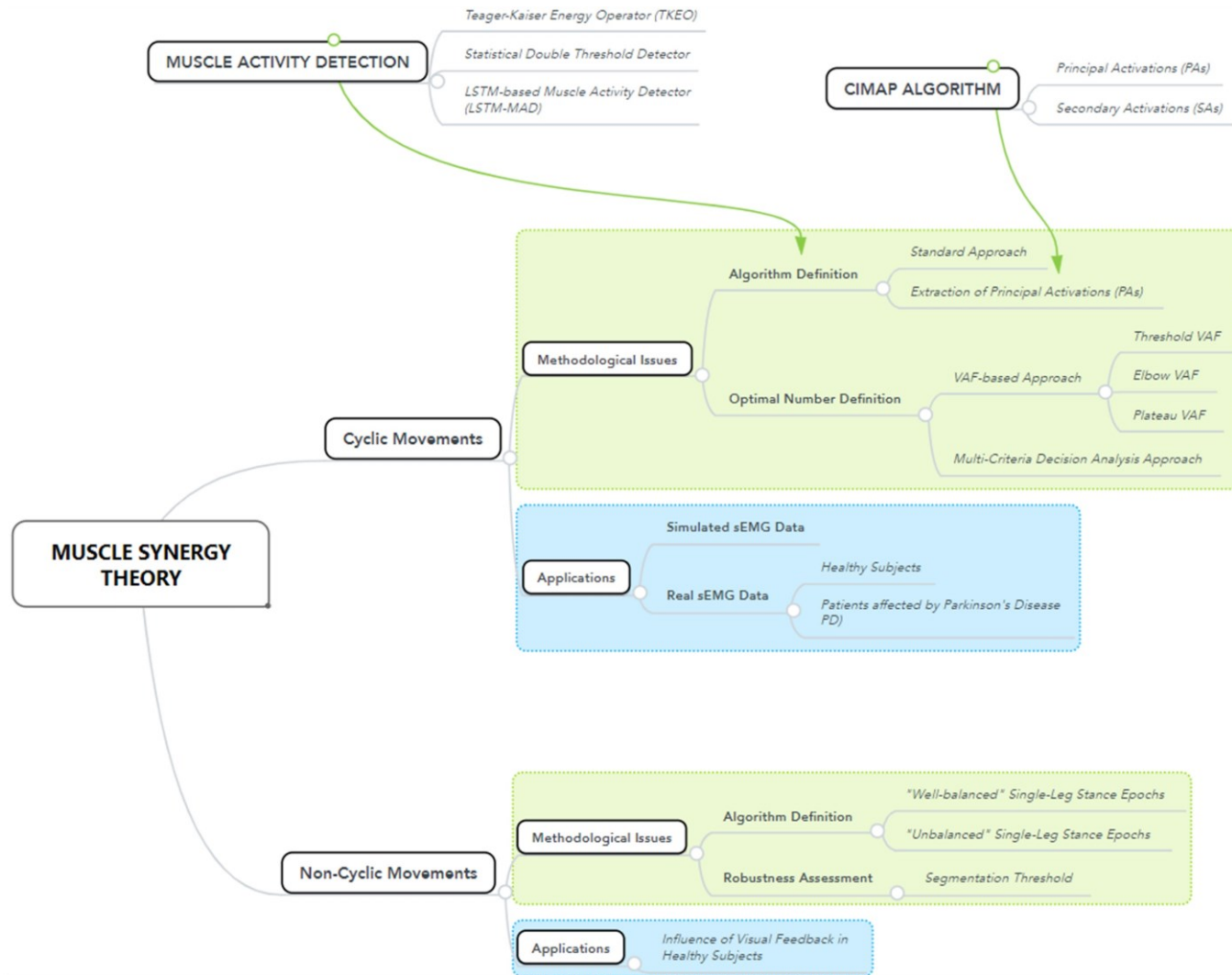


Figure 1.6 Graphical representation of the doctoral thesis's contents.

1 Chapter 2

2 A Deep Learning Approach for 3 Muscle Activity Detection

4 Determining muscle activation intervals (onset/offset time-instants) during
5 human movements is of great interest in different research fields, such as gait
6 analysis, myoelectric control of prostheses, and pre-processing of muscle synergy
7 extraction.

8 In the following chapter, the methodological issues related to muscle activation
9 interval detection are presented and discussed. More specifically, a novel approach
10 for muscle activity detection, based on LSTM neural networks, is proposed and its
11 performance is compared against two of the most widely used approaches: the
12 Teager-Kaiser Energy Operator (TKEO) detector and the statistical double
13 threshold detector.

14 2.1 Introduction

15 Dynamic muscle activity can be non-invasively investigated by means of
16 surface electromyography (sEMG). Determining the start (onset) and end (offset)
17 instants of muscle activations during human movements is of great interest in
18 different research fields including gait analysis [60], motor rehabilitation and sport
19 science [61], myoelectric control of prostheses [62], human-machine interaction
20 [63], design of biofeedback systems [64], and pre-processing of muscle synergy
21 extraction [46,65,66]. In particular, the accurate temporal analysis of muscle
22 activation in terms of burst onset, duration of the activation interval, and burst
23 offset, can be useful in the assessment of the altered locomotion patterns of
24 orthopedic and neurological patients [67].

25 A classical way to detect the timing of muscle activations from sEMG signals
26 is using a double threshold detector, such as the statistical detector by Bonato *et al.*
27 [68], specifically developed for gait analysis. However, this detector requires, as a
28 necessary input parameter, to set the first (amplitude) threshold that depends on the

1 background noise level. Furthermore, to fine-tune the second threshold, it is
2 important to estimate the signal-to-noise ratio (*SNR*), e.g., through the algorithm
3 described in Ref. [69].

4 Alternative methods, such as deep learning approaches, are being explored to
5 perform sEMG-based pattern recognition [70–79]. Muscle activity detection is
6 somewhat easier with respect to a pattern-recognition problem. Indeed, the focus is
7 not on classifying different movements, but simply detecting the presence or
8 absence of muscle activation. Exploiting artificial intelligence, such as a Recurrent
9 Neural Network (RNN), has been a winning strategy in a wide variety of different
10 applications and might be explored also for our problem. RNN is a powerful
11 learning algorithm inspired by the biological neural networks that constitute the
12 human brain, and it is trained to present to the network a large number of labeled
13 “examples” [80]. Long Short-Term Memory (LSTM) neural networks are a widely
14 used type of RNNs specifically designed to recognize patterns and time-
15 dependencies in sequential data, such as numerical time series, texts, and audio
16 tracks [81]. These neural networks were first introduced by Hochreiter and
17 Schmidhuber in 1997 [82] and represent an extension of the Recurrent Neural
18 Networks (RNNs), allowing for a better assessment of the time-dependencies in
19 long sequential data. Actually, LSTM neural networks represent the state of art in
20 natural language processing and speech recognition problems [83].

21 The aim of this chapter is to assess the applicability of a novel approach for
22 muscle activity detection, based on LSTM neural networks, specifically developed
23 to overcome the limitations of the standard approaches. The performance of the
24 LSTM-based Muscle Activity Detector (LSTM-MAD) are evaluated and compared
25 against a standard approach (Teager-Kaiser Energy Operator, TKEO) [84,85], and
26 a statistical approach (Statistical Double Threshold Detector) [68] in terms of
27 Precision, Recall, F1-score, and Jaccard similarity index, both on simulated and real
28 sEMG signals.

2.2 Materials and Methods

First, a dataset of simulated sEMG signals was built to assess the applicability of the LSTM-based approach to muscle activity detection and to compare its performance with respect to a standard and a statistical approach. For the standard approach, the Teager-Kaiser Energy Operator (TKEO) followed by the linear envelope extraction was implemented [85], while for the statistical approach it was implemented the double threshold statistical detector proposed in Ref. [68]. **Figure 2.1** represents the block diagram of the procedure followed in this chapter.

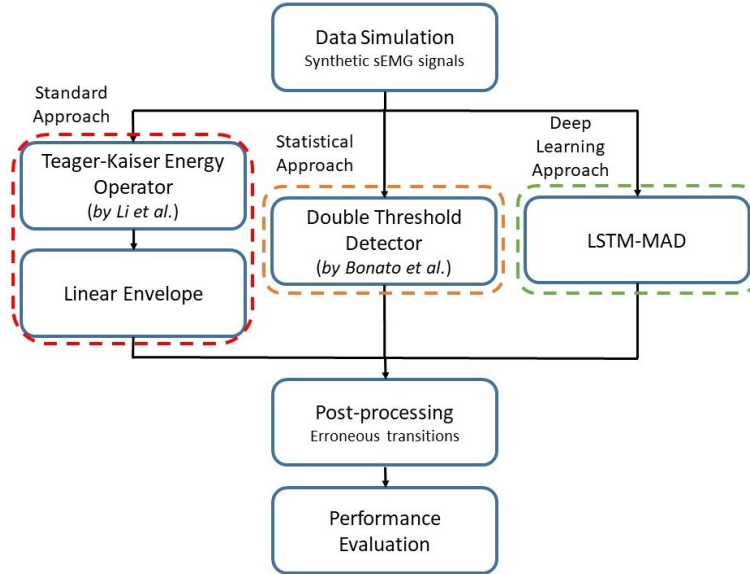


Figure 2.1 Schematic representation of the procedure followed to assess the performance of the new LSTM-MAD (LSTM-based Muscle Activity Detector) with respect to standard and statistical approaches for muscle activity detection.

Second, the LSTM-based approach was optimized on real sEMG signals, acquired in previous studies from lower limb and trunk muscles during gait, to emphasize its advantages in gait analysis.

1. Simulated Data

The sEMG signals acquired during cyclic movements, such as gait, can be modeled by the superimposition of two different contributions: (i) the electrical activity ($s(t)$) generated by each muscle during the contraction and (ii) the background noise ($n(t)$) mainly generated by the neighboring muscles and, to a lesser extent, by the acquisition system electronics. Under the hypothesis of cyclic contractions, the sEMG signal can be defined as a cyclostationary process [86] and, therefore, described through the superimposition of two different stationary processes [69]:

- i. The muscle activity ($s(t)$) modeled as a Gaussian process with zero-mean and variance σ_s^2 , as described in (2.1):

$$s(t) \in N(0, \sigma_s^2) \quad (2.1)$$

1 where σ_s was set equal to $10^{(SNR/20)} \cdot 1\mu V$.

2 *ii.* The background noise ($n(t)$) modeled as a zero-mean Gaussian process with
3 variance σ_n^2 , as described in (2.2):

$$n(t) \in N(0, \sigma_n^2) \quad (2.2)$$

4 where σ_n was set equal to $1\mu V$.

5 Each simulated sEMG signal was finally modeled as the superimposition of
6 two stationary processes: the background noise process only in correspondence of
7 the time-instants in which the muscle is non-active (OFF state) and a second
8 stationary process $x(t)$ in correspondence of the time-instants in which the muscle
9 is active (ON state) given by the superimposition of the muscle activity (2.1) and
10 the background noise (2.2) Gaussian processes as follows (2.3).

$$x(t) = n(t) + s(t) \quad (2.3)$$

11
12 Each realization of the muscle activity process $s(t)$ was simulated assuming a
13 time period of 1s (gait cycle) and a sampling frequency of 1 kHz. Physiological
14 muscle activity was modeled by time-windowing the Gaussian process $s(t)$ through
15 a single truncated Gaussian function centered at 50% of the gait cycle. Different
16 standard deviations (σ), and time supports ($2\alpha\sigma$) of the truncated Gaussian function
17 have been considered to simulate sEMG signals similar to those observed in leg,
18 thigh, and trunk muscles during gait. More specifically, three different values of the
19 standard deviation ($\sigma = 50, 100, 150 \text{ ms}$) and four different values of the time
20 support $2\alpha\sigma$ (with $\alpha = 1, 1.5, 2, 2.4$) have been tested [68]. Each muscle activity
21 was then band-pass filtered through a 4th order Butterworth digital filter with a
22 lower cut-off frequency of 10 Hz and a higher cut-off frequency of 450 Hz to
23 simulate the frequency spectrum of real sEMG signals [87]. Then, the background
24 noise process ($n(t)$) was superimposed. Nine different values of Signal-to-Noise
25 Ratio (SNR) were simulated ($SNR = 3, 6, 10, 13, 16, 20, 23, 26, 30 \text{ dB}$) [68]. For
26 each triplet of σ , α , and SNR values, 100 different realizations (N) have been
27 simulated and, therefore, a dataset composed by 10800 different realizations (3
28 standard deviations \times 4 time supports \times 9 $SNRs$ \times 100 realizations) was built.

29 **Figure 2.2** represents an example of a simulated sEMG signal with the
30 superimposition of the truncated Gaussian function used to model a physiological
31 muscle activity.

32 The time-instants relative to each simulated muscle activity ($s(t)$) were defined
33 by a binary mask (BM_{sim}) that was set equal to 1 in correspondence of the time-

1 instants in which the truncated Gaussian assumed values higher than 0, and it was
2 set equal to 0 otherwise.

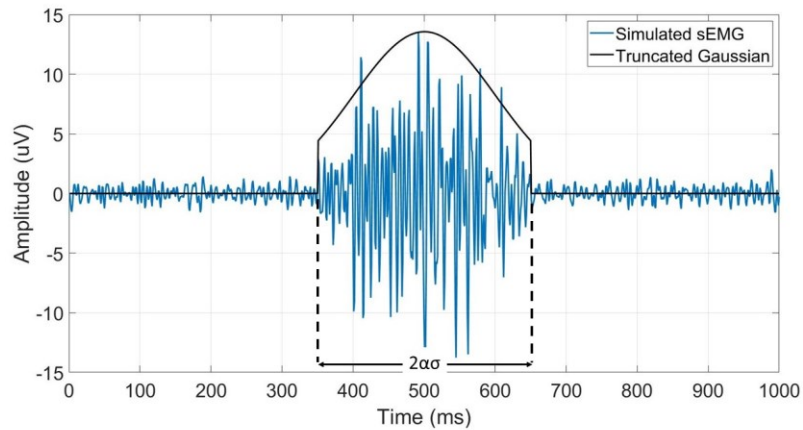


Figure 2.2 Example of a simulated sEMG signal (blue line) with the indication of the truncated Gaussian function (black line) used for the simulation of the muscle activity. The *SNR* is set equal to 20 dB, the standard deviation of the truncated Gaussian (σ) is equal to 100 ms, and the time support (2σ) is obtained for $\alpha=1.5$.

3 2. Real Data

4 Gait data were acquired from 20 subjects at PoliToBIOMed Lab (Politecnico
5 di Torino, Turin, Italy) to test the performance of the three different approaches
6 when applied to real sEMG signals. Subjects were randomly selected from our
7 database to include both healthy individuals, and patients affected by neurological
8 or orthopedic pathologies. This non-homogeneous group of subjects was
9 specifically chosen to verify that the algorithm works under different conditions. In
10 particular, 8 out of 20 subjects were healthy adults (age: 38.0 ± 13.1 years, height:
11 164.9 ± 5.4 cm, weight: 65.4 ± 21.2 kg) [46], 6 were patients with unilateral total
12 hip prosthesis (age: 73.8 ± 8.4 years, height: 175.5 ± 7.6 cm, weight: 86.8 ± 16.3
13 kg) [88], and the other 6 were patients affected by idiopathic normal pressure
14 hydrocephalus (age: 75.7 ± 6.3 years, height: 170.5 ± 6.3 cm, weight: 72.5 ± 10.4
15 kg) [89].

16 Signals were recorded through a multichannel acquisition system (STEP32,
17 Medical Technology, Italy), specifically developed for statistical gait analysis
18 [46,90,91]. Surface electromyographic (sEMG) signals were acquired through
19 active probes (configuration: single differential, size: $19 \text{ mm} \times 17 \text{ mm} \times 7 \text{ mm}$, Ag-
20 disks diameter: 4 mm, interelectrode distance: 12 mm, gain: variable in the range
21 from 60 dB to 86 dB) placed over the following 4 muscles of the lower limb: Rectus
22 Femoris (RF), Lateral Hamstring (LH), Lateral Gastrocnemius (LGS), and Tibialis
23 Anterior (TA). The dominant lower limb was analyzed for healthy subjects, while
24 the most affected limb was selected for pathological subjects.

25 For each subject, 5 gait cycles were randomly selected from the whole walking
26 task to build the real dataset. Therefore, a dataset composed of 400 different sEMG
27 signals (20 subjects \times 5 gait cycles \times 4 muscles) was obtained. The time-instants
28 relative to each real-muscle activity were manually segmented by three expert
29 operators. More specifically, a binary mask (BM_{real}) was set equal to 1 in

1 correspondence of the time-instants in which the majority of the expert operators
2 (at least two out of three) detected a muscle activity and to 0 otherwise.

3 The real dataset described and used in this chapter to train and test the LSTM-
4 MAD can be freely found online at Zenodo (doi: 10.5281/zenodo.4391062) in MAT
5 and CSV format.

6 3. *Standard Approach: TKEO and Linear Envelope*

7 One of the most commonly applied standard approaches for muscle activity
8 detection is the Teager-Kaiser Energy Operator (TKEO) followed by the linear
9 envelope extraction. This approach has been demonstrated to increase the accuracy
10 of the simple linear envelope approach [84,85].

11 More specifically, a single threshold was applied to the filtered sEMG signals
12 after the computation of the TKEO (ψ), defined as in (2.4):

$$\psi_{x(n)} = x(n)^2 - x(n+1)x(n-1) \quad (2.4)$$

13 where x represents the sEMG time-series and n the sample number. The single
14 threshold (Thr) was defined as described in (2.5):

$$Thr = \mu_n \pm j \cdot \sigma_n \quad (2.5)$$

15 where μ_n , σ_n and j represent the mean of the background noise, the standard
16 deviation of the background noise, and a multiplicative constant, respectively. In
17 this chapter, the constant j was set equal to 7 as suggested in Ref. [84],[92]. Since
18 the average (μ_n) and the standard deviation (σ_n) of the background noise are
19 required as inputs of this approach, time-instants corresponding to the background
20 noise were automatically selected considering a 100-ms window before the onset
21 of the simulated and segmented muscle activities, for the simulated and real sEMG
22 signals, respectively.

23 The output of this detector was finally defined as a binary mask ($Output_{TKEO}$)
24 as it follows:

- 25 • $Output_{TKEO} = 1$, if $\psi_{x(n)} \geq Thr$
- 26 • $Output_{TKEO} = 0$, if $\psi_{x(n)} < Thr$.

27

28 4. *Statistical Approach: Double Threshold Statistical Detector*

29 The statistical approach used in this chapter is the double threshold statistical
30 detector proposed in Ref. [68]. The algorithm, specifically developed for statistical
31 gait analysis, operates on raw sEMG signals and, hence, it does not require any pre-
32 processing step (e.g., envelope or TKEO computation). The computation steps used
33 for the double threshold detector are the following:

1 *i.* An auxiliary sequence z_i is computed from the raw sEMG signals as the sum
2 of the squared values of two successive samples (2.6):

$$z_i = x_i^2 + x_{i+1}^2 \quad (2.6)$$

3 where x_i and x_{i+1} represent two consecutive samples of the sEMG time series

4
5 *ii.* A first threshold ζ is applied on a sliding detection window defined by m
6 consecutive samples of the auxiliary sequence z_i

7
8 *iii.* A muscle activation is detected if at least r_0 (second threshold) out of m
9 consecutive samples of the detection window are above the first threshold ζ .

10 Considering the computation steps above mentioned, the double threshold
11 statistical detector has a more elaborate structure compared to single threshold
12 approaches. In particular, muscle activations are detected when at least r_0 (second
13 threshold) out of m consecutive values of the auxiliary sequence (z_i) cross the first
14 threshold (ζ). Accordingly, this detector is defined by three different input
15 parameters: (*i*) the first (or amplitude) threshold (ζ), (*ii*) the length of the
16 observation window (m), and (*iii*) the second (or temporal) threshold. As suggested
17 in Ref. [68], the length of the observation window (m) and the second threshold r_0
18 were set equal to 5 and 1, respectively.

19 The output of this detector was finally defined as a binary mask ($\text{Output}_{\text{Stat}}$) as
20 it follows:

- 21 • $\text{Output}_{\text{Stat}} = 1$, if $z_i \geq \zeta$ for at least r_0 out of m samples
- 22 • $\text{Output}_{\text{Stat}} = 0$, otherwise.

23

24 5. Deep Learning Approach: LSTM-MAD

25 An LSTM model is generally composed by the following architecture:

- 26 *i.* An *input sequence layer*
- 27 *ii.* One or more *LSTM layers* used to learn the time-dependencies within the
28 sequential data
- 29 *iii.* A *fully connected layer* used to convert the output size of the previous
30 layers into the number of classes to be recognized
- 31 *iv.* A *softmax layer* used to compute the belonging probability to each class
- 32 *v.* A *classification output layer* used to compute the cost function.

33 In this chapter, several LSTM neural network models were tested to assess the
34 applicability of this approach for muscle activity detection during gait, considering
35 sEMG data without any pre-processing step. To define the best LSTM model for

1 muscle activity detection, the entire dataset of simulated (and real) sEMG signals
2 was divided into 3 different sets: training set (70%), validation set (15%), and test
3 set (15%), respectively. The training set was used to train the LSTM model, while
4 the validation set (or “development” set) was used to evaluate the network
5 performances and to avoid the overfitting of the training data. More specifically,
6 the validation set was used to stop training automatically when the validation
7 accuracy stopped increasing to avoid overfitting [80]. Finally, the test set was used
8 for the final validation and the comparison of LSTM-MAD with the other two
9 detectors.

10 Using the Deep Learning Toolbox of MATLAB[®] release R2020b (The
11 MathWorks Inc., Natick, MA, USA), 720 different LSTM neural networks were
12 tested. All the LSTM models had a sequence *input layer* consisting of 1 unit
13 (dimension of a single simulated sEMG signal), and a *fully connected output layer*
14 consisting of 2 units (number of classes to be recognized). Different numbers of
15 LSTM layers (n), numbers of hidden units for each layer (n_{units}), values of learning
16 rate (α), and values of drop period (δ) were tested to achieve the LSTM architecture
17 with the highest performance. More specifically, two different numbers of LSTM
18 layers ($n = 1, 2$), nine different numbers of hidden units in each LSTM layer
19 ($n_{units} = 100, 125, 150, 175, 200, 225, 250, 275, \text{ and } 300$), five different learning
20 rates ($\alpha = 0.01, 0.015, 0.02, 0.025, \text{ and } 0.03$), and eight different drop rate values
21 ($\delta = 10, 15, 20, 25, 30, 35, 40, \text{ and } 45$) were tested [80]. The adaptive moment
22 (ADAM) optimization algorithm was adopted in this work to train all the tested
23 LSTM models [93]. The performance of each LSTM model was assessed
24 considering the simulated (or real) test set by computing the overall classification
25 accuracy, defined as the number of correctly classified sEMG samples normalized
26 to the total number of sEMG samples within the test set.

27 The training process was performed on a workstation with a 3.2 GHz six-core
28 CPU, 32 GB of RAM memory, and a 64-bit Windows operating system.

29 The BM_{sim} , extracted from the truncated Gaussian functions, and the BM_{real} ,
30 manually defined by the expert operators, were used to compute the target (ground
31 truth) of the LSTM model for the simulated and real datasets, respectively.

32 The output of the LSTM approach was computed as a binary mask ($\text{Output}_{\text{LSTM-}}$
33 MAD) that was defined as it follows:

- 34 • $\text{Output}_{\text{LSTM-MAD}} = 1$, if the sEMG time-instant was classified as muscle
35 activity (class 1)
- 36 • $\text{Output}_{\text{LSTM-MAD}} = 0$, if the sEMG time-instant was classified as
37 background noise (class 0).

38 39 6. Post-processing

1 A post-processing step was applied to the output of each detector (standard,
2 statistical, and deep learning approach) to reject the erroneous transitions due to the
3 stochastic nature of the sEMG signal. Since it is generally accepted that a muscle
4 activation shorter than 30 ms does not affect the kinetics and the kinematics of gait
5 [94], all the muscle activations lasting less than 30 ms were discarded [68]. This
6 concept is illustrated in **Figure 2.3**. In particular, **Figure 2.3A** shows a sample
7 realization of a simulated sEMG signal modulated by a truncated Gaussian function
8 ($SNR = 16$ dB, $\sigma = 100$ ms, and $\alpha = 1.5$). **Figure 2.3B** represents the output of the
9 standard approach ($Output_{TKEO}$) without any post-processing step, while **Figure**
10 **2.3C** shows the effect of the post-processing on the detector's output.

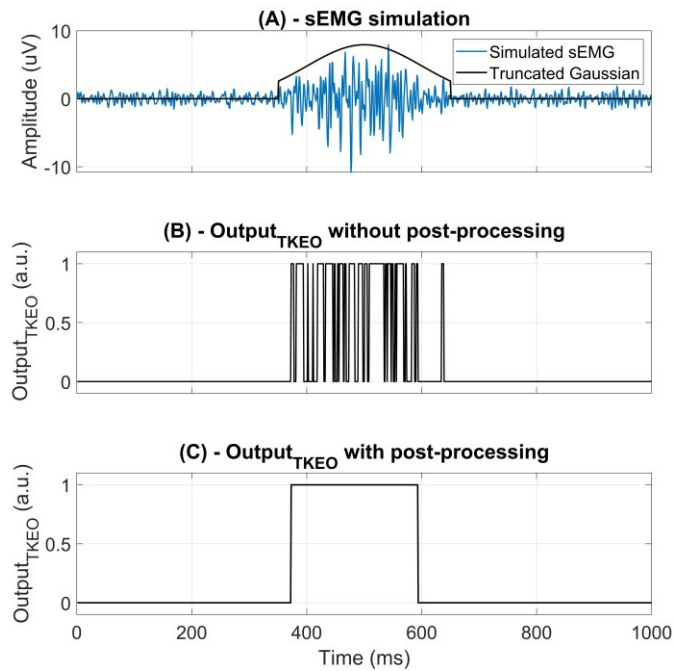


Figure 2.3 Example of post-processing applied to the standard approach output ($Output_{TKEO}$). (A) Sample realization of a simulated sEMG signal (blue line) with the superimposition of the truncated Gaussian function used to modulate the muscle activity (black line). The SNR is set equal to 20 dB, standard deviation of the truncated Gaussian (σ) is 100 ms and the multiplicative constant (α) of the time support is 1.5. (B) Output of the standard approach ($Output_{TKEO}$) without any post-processing. (C) Output of the standard approach after rejecting all the activations shorter than 30 ms (post-processing step).

11 7. Performance Evaluation

12 The muscle activations detected by the three different approaches ($Output_{TKEO}$,
13 $Output_{Stat}$, and $Output_{LSTM-MAD}$) were quantitatively compared in terms of (i)
14 Precision, (ii) Recall, (iii) F1-score, and (iv) Jaccard similarity index. More
15 specifically, the indexes were defined as it follows:

$$Precision = \frac{TP}{TP + FP} \quad (2.7)$$

$$Recall = \frac{TP}{TP + FN} \quad (2.8)$$

$$F1 \text{ score} = \frac{2 \times (Recall \times Precision)}{(Recall + Precision)} \quad (2.9)$$

$$Jaccard = \frac{|Output_i \cap BM|}{|Output_i \cup BM|} \quad (2.10)$$

1 where TP represents the True Positive (i.e., number of sEMG time-instants
 2 correctly classified by the detectors as muscle activity), FN describes the False
 3 Negative (i.e., number of sEMG time-instants incorrectly classified by the detectors
 4 as background noise), and FP represents the False Positive (i.e., number of sEMG
 5 time-instants incorrectly classified by the detectors as muscle activity). Finally,
 6 $Output_i$ represents the binary mask computed by the i -th detector ($i = 1$: Standard
 7 approach with TKEO, $i = 2$: Statistical approach with the double-threshold detector,
 8 $i = 3$: New approach with LSTM-MAD), and BM represents the ground truth of the
 9 simulated (or real) data.

10 8. *Effect of SNR on Muscle Activity Detection*

11 To assess the effect of the SNR on the performance of the three muscle activity
 12 detectors, the performance parameters described above were computed on the
 13 simulated test set, separately for each of the nine SNR values ($SNR =$
 14 $3, 6, 10, 13, 16, 20, 23, 26, 30$ dB).

15 9. *Statistical Analysis*

16 A one-way repeated-measures analysis of variance (ANOVA) followed by
 17 *post-hoc* analysis with Bonferroni adjustment for multiple comparisons was
 18 performed to assess significant differences in the performance of the three
 19 approaches, setting the significance level (α) equals to 0.05.

20 **2.3 Results**

21 First, the results supporting the applicability of the LSTM approach for muscle
 22 activity detection (LSTM-MAD) are presented, considering simulated sEMG
 23 signals. Second, the performance of the three detectors (standard TKEO, double-
 24 threshold statistical detector, and the newly introduced LSTM-MAD) on simulated
 25 sEMG signals are compared, highlighting the effect of the SNR . Finally, LSTM-
 26 MAD is validated on real data.

27 1. *Simulated Data - LSTM Model Definition*

28 The best LSTM model was selected among all the tested networks as the one
 29 with the highest overall classification accuracy on the test set, discarding those
 30 networks with a difference between the training and validation accuracy higher than
 31 4% (to avoid overfitting of the training data).

1 **TABLE 2.1** shows the properties of the LSTM model that achieved the highest
 2 overall classification accuracy ($96.8\% \pm 4.3\%$) on the test set.

TABLE 2.1
 PROPERTIES OF THE BEST LSTM MODEL

<i>LSTM Layers</i>	<i>PROPERTIES</i>
Sequence input layer	1 input feature
LSTM layer #1	275 hidden units, Bi-directional sequence-to-sequence architecture
LSTM layer #2	138 hidden units, Bi-directional sequence-to-sequence architecture
Fully connected layer	2 units
Softmax layer	Softmax activation function (threshold = 0.5)
Classification output layer	2 classes (1 = muscle activity, 0 = background noise)

3 2. *Simulated Data – Performance Evaluation*

4 The performances of the three different muscle activity detectors were assessed
 5 and compared in terms of (i) Precision, (ii) Recall, (iii) F1-score, and (iv) Jaccard
 6 similarity index:

- 7 *i. Precision:* An average precision of 0.92 ± 0.10 , 0.98 ± 0.09 , and 0.95 ± 0.08
 8 was found on the simulated test set, considering the standard, the statistical,
 9 and the deep learning approach, respectively. One-way ANOVA followed by
 10 *post-hoc* analysis revealed significant differences between each pair of
 11 detectors ($p < 0.0001$)
- 12 *ii. Recall:* On average, a recall of 0.86 ± 0.20 , 0.53 ± 0.39 , and 0.96 ± 0.08 was
 13 found on the simulated test set, considering the standard, the statistical, and
 14 the deep learning approach, respectively. One-way ANOVA followed by
 15 *post-hoc* analysis revealed significant differences between each pair of
 16 detectors ($p < 0.0001$)
- 17 *iii. F1-score:* On average, an F1-score of 0.87 ± 0.14 , 0.76 ± 0.25 , and $0.95 \pm$
 18 0.06 was found on the simulated test set, considering the standard, the
 19 statistical, and the deep learning approach, respectively. One-way ANOVA
 20 followed by *post-hoc* analysis revealed significant differences between each
 21 pair of detectors ($p < 0.0001$)
- 22 *iv. Jaccard similarity index:* An average Jaccard index of 0.80 ± 0.19 , $0.52 \pm$
 23 0.38 , and 0.91 ± 0.10 was found on the simulated test set, considering the
 24 standard, the statistical, and the deep learning approach, respectively. One-
 25 way ANOVA followed by *post-hoc* analysis revealed significant differences
 26 between each pair of detectors ($p < 0.0001$).

1 **Figure 2.4A** compares the performance of the three detectors tested (standard
 2 TKEO [84], statistical double-threshold detector [68], and LSTM-MAD), based on
 3 the four parameters introduced above. The average values and standard errors of
 4 the parameters were estimated on the simulated dataset. Asterisks highlight
 5 statistical differences between each pair of detectors.

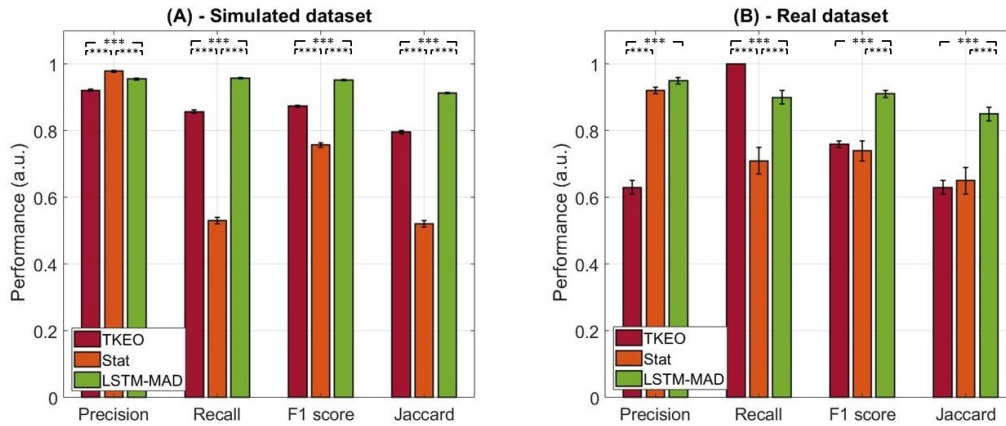


Figure 2.4 Comparison of the performance of the three muscle activity detectors (standard TKEO by Li *et al.* [24], statistical double-threshold detector by Bonato *et al.* [10], and LSTM-MAD) estimated considering (A) the simulated and (B) the real dataset. Average values and standard errors are represented. Statistically significant differences are indicated by asterisks (***) $p < 0.0001$.

6 3. Simulated Data – Effect of SNR on Muscle Activity Detection

7 The effect of the *SNR* on the detectors' performance was assessed by
 8 computing the performance parameters (Precision, Recall, F1-score, and Jaccard
 9 similarity index), separately for each simulated *SNR*-value.

10 **Figure 2.5** represents, for each muscle activity detector, the average values
 11 (and standard errors) of Precision (**Figure 2.5A**), Recall (**Figure 2.5B**), F1-score
 12 (**Figure 2.5C**), and Jaccard similarity index (**Figure 2.5D**), for each simulated
 13 *SNR*-value.

14 For all the parameters, LSTM-MAD revealed a higher performance consistency
 15 across the different *SNR* values, suggesting a lower effect of *SNR* on muscle
 16 activity detection with respect to the other two approaches. The approach more
 17 affected by the *SNR* was the statistical double threshold detector, with an evident
 18 decrease of the performances for simulated sEMG signals with *SNR*-values lower
 19 than 20 dB.

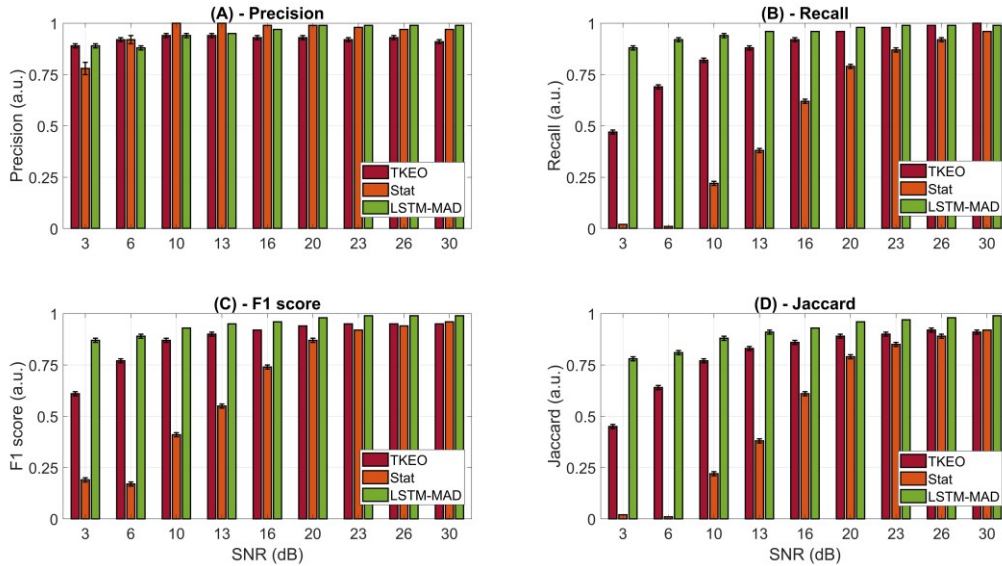


Figure 2.5 Values of (A) Precision, (B) Recall, (C) F1 score, and (D) Jaccard similarity index, averaged on the simulated test set, for each value of **SNR** and for each muscle activity detector. Each colored bar represents the average performance of a specific detector (TKEO by Li *et al.* [24] in red, double-threshold statistical detector by Bonato *et al.* [10] in orange, and LSTM-MAD in green). Error bars represent the standard errors.

1 4. Real Data

2 The same procedure described for the simulated dataset was followed to
 3 determine the best LSTM model considering real sEMG data. In the following, the
 4 results obtained from the real dataset for the three different approaches are detailed.

5 5. Real Data – LSTM Model Definition

6 The same 720 different LSTM neural networks considered for the simulated
 7 dataset were tested. The best LSTM model that achieved the highest overall
 8 classification accuracy ($90.1\% \pm 14.28\%$) on the test set had the same architecture
 9 and properties as the one selected considering the simulated data (see **TABLE 2.1**).
 10 **Figure 2.6** shows an example of a real sEMG signal acquired from the TA muscle
 11 of a healthy subject with the superimposition of the ground truth (BM_{real}) and the
 12 outputs of the three detectors ($Output_{TKEO}$, $Output_{Stat}$, and $Output_{LSTM-MAD}$).

13 6. Real Data – Performance Evaluation

14 The performance of the three different muscle activity detectors on real sEMG
 15 signals were assessed considering the same four parameters described for the
 16 simulated dataset:

- 17 *i. Precision:* An average precision of 0.63 ± 0.15 , 0.92 ± 0.11 , and 0.95 ± 0.11
 18 was found on the real test set, considering the standard, the statistical, and the

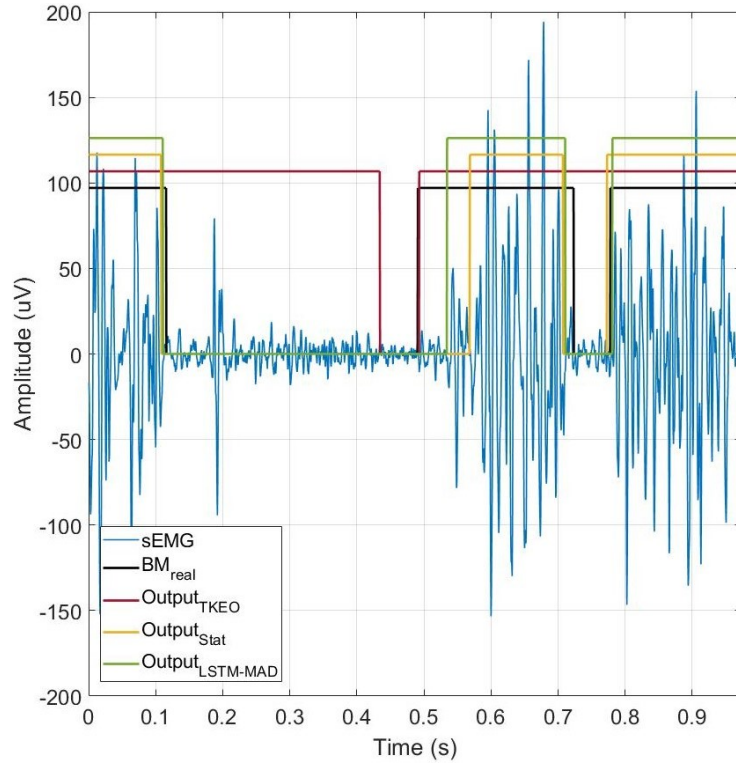


Figure 2.6 Example of a real sEMG signal (blue line) acquired from the Tibialis Anterior (TA) muscle of a healthy subject of the sample population. The output of the standard (red line), statistical (orange line), and LSTM-MAD approach (green line) are represented along with the indication of the ground truth (black line) manually segmented by expert operators. All the muscle activity shorter than 30 ms are rejected by means of the post-processing step for all the approaches.

- 1 deep learning approach, respectively. One-way ANOVA followed by *post-*
2 *hoc* analysis revealed significant differences between the standard and the
3 statistical approach ($p < 0.0001$), and between the standard and the deep
4 learning approach ($p < 0.0001$), while no difference was found between the
5 statistical and the deep learning approach
- 6 *ii. Recall:* On average, a recall of 1.00 ± 0.00 , 0.71 ± 0.31 , and 0.90 ± 0.15 was
7 found on the real test set, considering the standard, the statistical, and the deep
8 learning approach, respectively. One-way ANOVA followed by *post-hoc*
9 analysis revealed significant differences between each pair of detectors ($p <$
10 0.0001)
- 11 *iii. F1-score:* On average, an F1-score of 0.76 ± 0.11 , 0.74 ± 0.26 , and $0.91 \pm$
12 0.11 was found on the real test set, considering the standard, the statistical,
13 and the deep learning approach, respectively. One-way ANOVA followed by
14 *post-hoc* analysis revealed significant differences between the standard and
15 the deep learning approach ($p < 0.0001$), and between the statistical and the
16 deep learning approach ($p < 0.0001$), while no difference was found between
17 the standard and the statistical approach
- 18 *iv. Jaccard similarity index:* An average Jaccard index of 0.63 ± 0.15 , $0.65 \pm$
19 0.27 , and 0.85 ± 0.16 was computed from the real test set, considering the
20 standard, the statistical, and the deep learning approach, respectively. One-
21 way ANOVA followed by *post-hoc* analysis revealed significant differences

1 between the standard and the deep learning approach ($p < 0.0001$), and
2 between the statistical and the deep learning approach ($p < 0.0001$), while no
3 difference was found between the standard and the statistical approach.

4 Considering the real test set, **Figure 2.4B** compares the performance of the
5 three detectors, based on the four parameters introduced above. The average values
6 and standard errors of the parameters are reported, as well as asterisks to highlight
7 statistical differences between each pair of detectors.

8 **2.4 Discussion and Conclusions**

9 This chapter demonstrates that the detection of muscle activity during gait can
10 be successfully performed by means of a novel approach, based on a Long Short-
11 Term Memory (LSTM) Recurrent Neural Network (RNN). The newly introduced
12 muscle activity detector (LSTM-MAD) is able to effectively separate muscle
13 activation intervals from background noise, with an overall classification accuracy
14 of 97% (considering simulated data).

15 Results showed that the proposed detector outperforms current methods used
16 to discern muscle activity. In particular, LSTM-MAD clearly exhibits better
17 performance than both the alternative approaches tested (standard approach using
18 the Teager-Keiser Energy Operator (TKEO) [84], and statistical double-threshold
19 detector [68]). Indeed, improved performance was demonstrated both on simulated
20 and real sEMG signals.

21 Considering simulated signals, all the four “performance” parameters
22 introduced (Precision, Recall, F1-score, and Jaccard similarity index) showed
23 remarkably higher values for LSTM-MAD when compared to the statistical
24 detector. Furthermore, three out of four parameters (Recall, F1-score, and Jaccard
25 similarity index) displayed greater values for LSTM-MAD when compared to the
26 TKEO detector, and only a slightly worst precision was found. However, it should
27 be noticed that while LSTM-MAD shows an excellent balance between Precision
28 and Recall, the same cannot be said for the TKEO detector, which displays a very
29 high Precision (i.e., low number of false positive), but a very low Recall (i.e., a high
30 number of false-negative). In other words, the TKEO detector demonstrates a
31 reduced probability of detection and an increased number of false-negative
32 compared to the LSTM-MAD. By the way, the “optimal balance” between
33 Precision and Recall is incorporated in the definition of the F1-score, which is
34 broadly used in literature specifically to take into account the balance between
35 Precision and Recall by considering both the number of false-positive and the
36 number of false-negative.

37 Considering real signals, all the four “performance” parameters showed
38 remarkably higher values for LSTM-MAD with respect to the TKEO detector.
39 Furthermore, three out of four parameters (Precision, F1-score, and Jaccard
40 similarity index) displayed greater values for LSTM-MAD when compared to the
41 TKEO detector. Only the Recall was higher in the TKEO detector with respect to
42 our approach. Again, it should be noted that LSTM-MAD is characterized by an

1 excellent balance between Precision and Recall, while this is not true for the TKEO
2 detector, which is characterized by a higher probability of detection and, hence, by
3 a higher number of false-positive (i.e., lower precision) compared to the LSTM-
4 MAD. Indeed, the TKEO detector shows a very good Recall to the detriment of a
5 very poor Precision. The statistical detector, instead, shows different behavior,
6 revealing a very high Precision (similar to the Precision obtained considering the
7 LSTM-MAD approach) to the detriment of a very poor Recall. In other words, the
8 statistical detector demonstrates a reduced probability of detection and an increased
9 number of false-negative compared to the LSTM-MAD.

10 Overall, LSTM-MAD revealed a smaller variability in the detector's
11 performance, especially compared to the TKEO approach. Although a thorough
12 analysis of this aspect is beyond the scope of this work, this reduced variability can
13 be qualitatively appreciated in **Figure 2.4**, when comparing the small error bars
14 obtained for LSTM-MAD compared to those obtained for TKEO.

15 Furthermore, the novel approach introduced in this chapter revealed increased
16 robustness of the detector's performance with respect to the effect of the signal-to-
17 noise ratio (*SNR*), suggesting the applicability of the LSTM-MAD to a wider range
18 of noise conditions compared to the other two tested approaches. Indeed, while it is
19 evident that decreasing *SNR* inevitably diminishes the detection performance of
20 each approach, LSTM-MAD is the least affected one (see **Figure 2.5**). In particular,
21 focusing on the parameters Recall, F1-score, and Jaccard similarity index, we found
22 a conspicuous worsening of the performance of the statistical double-threshold
23 detector with decreasing *SNR*. The situation is even more dramatic considering the
24 TKEO detector. On the contrary, our detector LSTM-MAD shows a limited
25 worsening of the performance with decreasing *SNR*. Indeed, even at very low *SNR*
26 values (e.g., 3 or 6 dB), the performance of LSTM-MAD never degrades too much
27 (Recall is always greater than 0.88, F1-score is always greater than 0.87, and
28 Jaccard similarity index is always greater than 0.78). For what concerns the
29 parameter Precision, none of the three detectors showed a drastic decrease of the
30 performance with decreasing *SNR* ratio. The analysis about how a poor *SNR* can
31 eventually degrade detectors' performance was carried out on simulated signals
32 only. This was chosen to study the above-described phenomenon in a more
33 controlled condition, i.e., to have a precise knowledge (*a priori*) about the *SNR*
34 itself (since sEMG signals were simulated at each specific *SNR* level). Indeed,
35 considering real signals one would have needed to apply some additional algorithm
36 to estimate the *SNR*.

37 A further valuable attribute and distinctive quality of the LSTM-MAD
38 approach is that it does not require any additional input parameter. The proposed
39 algorithm directly works on "raw" sEMG signals, the only pre-processing step
40 being the usual passband filtering (e.g., between 10 Hz and 450 Hz), applied to all
41 the three approaches, in the same manner. On the contrary, the statistical double-
42 threshold detector requires, as a necessary input parameter, the knowledge of the
43 background-noise power. Furthermore, the estimation of the *SNR* is also usually
44 required to properly fine-tune the algorithm parameters. The estimation of

1 background-noise power and *SNR*-ratio is usually obtained by analyzing 30 s-
2 windows of sEMG signal. However, since LSTM-MAD does not require any
3 additional input parameter (e.g., background-noise power or *SNR* estimation), it is
4 intrinsically more adaptable to eventual *SNR* variations arising during signal
5 acquisition. Furthermore, differently from the TKEO approach, it does not require
6 the computing of a linear envelope.

7 In conclusion, a Long-Short-Term-Memory approach for muscle activity
8 detection was proposed and validated (both on simulated and real sEMG signals).
9 The presented approach clearly outperforms previous detectors, and it is robust
10 even when applied to signals with low to medium signal-to-noise ratio. Therefore,
11 it may be considered a valuable tool, in all the applications requiring an accurate
12 and effective recognition/distinction of muscle activity from background noise.

1 Chapter 3

2 Motor Control Strategies during 3 Cyclic Movements

4 In this chapter, two fundamental methodological issues in the assessment of
5 motor control strategies through the muscle synergy theory are discussed: the
6 sEMG pre-processing technique used to extract muscle synergies and the criterion
7 used to select the optimal number of muscle synergies. First, an innovative sEMG
8 pre-processing approach for muscle synergy extraction during gait is presented and
9 compared with respect to a “standard” pre-processing approach (**Methodological**
10 **Issue - Muscle Synergies Extracted Using Principal Activations**). Then, to
11 overcome the limitations of the *VAF*- or R^2 -based methods for the selection of the
12 optimal number of muscle synergies, a novel approach, based on Multi-Criteria
13 Decision Aiding (MCDA), is assessed and validated considering both simulated
14 and real data (**Methodological Issue – Multi-Criteria Decision Analysis for**
15 **Selecting the Optimal Number of Muscle Synergies**).

16 **Methodological Issue - Muscle Synergies Extracted Using** 17 **Principal Activations**

18 Some of the results presented in this paragraph are reproduced and modified
19 from an article published in *IEEE Transactions on Neural Systems and*
20 *Rehabilitation Engineering* (Volume: 28, Issue: 2, Feb. 2020) by Ghislieri *et al.*
21 The final authenticated version of the manuscript is available online at
22 <http://dx.doi.org/10.1109/TNSRE.2020.2965179>.

23 **3.1 Introduction**

24 According to the muscle synergy theory, the Central Nervous System (CNS)
25 controls small groups of muscles (called *muscle synergies*) rather than coordinating
26 and activating every single muscle involved in a specific motor task. The

1 effectiveness of the muscle synergy theory in modeling the modular organization
2 of the CNS during different motor tasks has been demonstrated in several studies
3 and research areas, such as clinics, robotics, and sport [17,18,27,47,95,96].

4 More specifically, human locomotion is one of the most widely studied and
5 complex motor tasks, due to the multiple degrees of freedom of the skeletal muscle
6 system, the several biomechanical functions carried out during each gait cycle [97],
7 and the high intra-cycle variability of the muscle activation intervals [98]. Previous
8 studies demonstrated that human locomotion can be modeled by a small set of
9 muscle synergies characterized by specific biomechanical functions [19–22]. The
10 number of muscle synergies extracted and their biomechanical functions may vary
11 depending on the type and the number of muscles acquired [23], the sEMG pre-
12 processing techniques, and the factorization algorithm implemented. On average,
13 five muscle synergies are needed to properly describe human locomotion [19,24].

14 Muscle synergies during human locomotion are often extracted from surface
15 electromyography (sEMG) signals by applying the Non-Negative Matrix
16 Factorization (NNMF) algorithm [28,41]. Before computing the factorization, the
17 acquired sEMG signals are usually pre-processed to extract their envelopes. In the
18 last years, several pre-processing approaches were proposed in literature for muscle
19 synergy extraction during cyclic movements [19,23,27,28,41,55,99]. For example,
20 in the study by Clark *et al.* [55] the acquired sEMG signals were previously high-
21 pass filtered at 40 Hz through a zero-lag 4th-order digital Butterworth filter,
22 demeaned, full-cycle rectified, and smoothed by means of a zero-lag 4th-order
23 Butterworth digital filter with a cut-off frequency of 4 Hz. Finally, to facilitate the
24 comparison among different muscles and different motor conditions, the filtered
25 sEMG signals were amplitude-normalized with respect to the global maximum of
26 the signal acquired from each recorded muscle. In the work by Steele *et al.* [23]
27 sEMG data were band-pass filtered with a lower cut-off frequency of 20 Hz and a
28 higher cut-off frequency of 400 Hz, rectified and low-pass filtered at 10 Hz.

29 The application of these *standard* pre-processing techniques may not be
30 sufficient to fully understand the motor control mechanism due to the high intra-
31 cycle variability of the sEMG activation intervals [100]. Statistical Gait Analysis
32 (SGA) [60] was introduced to deal with this issue by selecting only the
33 representative gait patterns. It was successfully applied to the study of the
34 frequency-of-occurrence of muscle activation modalities [100], muscle activation
35 timing [101], and co-contractions [102]. Moreover, through the application of the
36 Clustering for Identification of Muscle Activation Patterns (CIMAP) algorithm
37 [103], [104], it is possible to define the principal and secondary muscle activations.
38 Principal activations (PAs) are defined as those activations that are strictly
39 necessary to accomplish the motor task, while secondary activations (SAs) may
40 have an auxiliary function, such as providing corrections to motion and body
41 segment posture. The CIMAP algorithm was successfully applied to the study of
42 gait asymmetry of healthy, orthopedic, and neurological patients [67,105,106].

43 The aim of this work is to assess how the combined application of the SGA and
44 CIMAP (i.e., extraction of PAs) techniques can be used to overcome the limitations
45 of the *standard* pre-processing algorithms in terms of intra-subject consistency,

1 robustness, and interpretability of muscle synergies during gait in both
2 physiological and pathological populations.

3 **3.2 Materials and Methods**

4 *1. Sample Population and Experimental Protocol*

5 Gait data from twenty-two patients affected by Parkinson's disease (7 females
6 and 15 males, age: 58.9 ± 8.1 years, height: 174.4 ± 9.4 cm, weight: 76.5 ± 13.1 kg)
7 and twenty-two healthy subjects (18 females and 4 males, age: 39.2 ± 17.0 years,
8 height: 165.2 ± 8.2 cm, weight: 60.9 ± 17.5 kg) were acquired and analyzed at
9 PoliToBIOMed Lab (Politecnico di Torino, Turin, Italy). Patients affected by
10 Parkinson's Disease (PD) were enrolled from those eligible for Deep Brain
11 Stimulation (DBS) at the Stereotactic and Functional Neurosurgery Unit of the
12 University of Turin (Italy). The inclusion criteria were:

- 13 *i.* Diagnosis of PD, according to the UK Brain Bank guidelines
- 14 *ii.* Good response to Levodopa
- 15 *iii.* Age at surgery below 70 years
- 16 *iv.* Absence of freezing of gait and postural instability unresponsive to
17 pharmacological therapy
- 18 *v.* Ability to walk independently for a few minutes without walking aids or
19 external supports, at least within the best-on pharmacological time-
20 window.

21 PD patients affected by other co-morbidities that could potentially affect gait
22 performance were excluded from the study.

23 Gait and balance performance were rated using the Unified Parkinson's Disease
24 Rating Motor Subscale (UPDRS-III) [107]. For the enrolled PD population, the
25 average UPDRS-III score was equal to 17.1 ± 0.2 and 44.6 ± 12.3 considering *ON*
26 and *OFF* condition, respectively.

27 Considering the healthy subjects, none of them reported lower limb injuries or
28 had neurological or musculoskeletal disorders that could compromise their gait
29 performance.

30 The experimental protocol consisted of a 5-minute walk at self-selected speed,
31 back and forth on a 9-m straight walkway. PD patients were acquired during their
32 best-on conditions. **Figure 3.1** shows a schematic representation of the walking
33 path. The protocol conformed to the Helsinki declaration on medical research

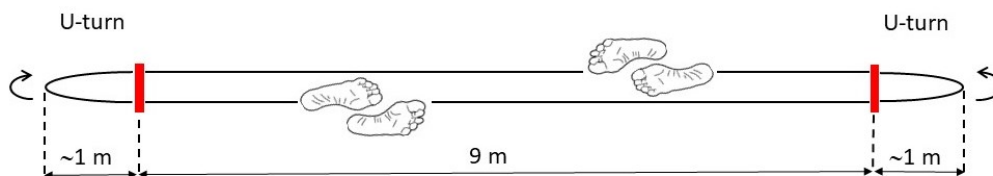


Figure 3.1 Schematic representation of the walking path. Subjects walked back and forth, without interruptions, along a straight path of 9 m, for approximately 5 minutes. U-turns were excluded by the analysis.

1 involving human subjects and was approved by the Ethics Committee of the A.O.U.
 2 Città della Salute e della Scienza di Torino - A.O. Ordine Mauriziano - A.S.L. “Città
 3 di Torino” (No. 0092029).

4 **TABLE 3.1** represents the anthropometric characteristics of the PD and healthy
 5 populations analyzed in this chapter.

TABLE 3.1
 ANTHROPOMETRIC CHARACTERISTICS OF PD AND HEALTHY POPULATIONS.

	Age (years)	Weight (kg)	Height (cm)	UPDRS-III (ON condition)	UPDRS-III (OFF condition)	Disease duration (years)
PD	58.9 ± 8.1	76.5 ± 13.1	174.4 ± 9.4	17.1 ± 0.2	44.6 ± 12.3	12.8 ± 3.7
Healthy	39.2 ± 17.0	60.9 ± 17.5	165.2 ± 8.2	N/A	N/A	N/A

Values of parameters are reported as mean ± standard deviation over the population.

UPDRS-III: Unified Parkinson’s Disease Rating Motor Subscale

N/A: Not Applicable

6 2. Data Acquisitions

7 A multichannel system specifically developed for statistical gait analysis
 8 (STEP32, Medical Technology, Italy) was used to acquire gait data. The following
 9 signals were simultaneously recorded:

- 10 *i.* Surface electromyographic (sEMG) signals, by means of active probes
 11 (configuration: single differential, size: 19 mm × 17 mm × 7 mm, Ag-
 12 disks diameter: 4 mm, interelectrode distance: 12 mm, gain: variable in
 13 the range from 60 dB to 86 dB)
- 14 *ii.* Foot-switch signals (size: 10 mm × 10 mm × 0.5 mm, activation force: 3
 15 N) to detect gait phases
- 16 *iii.* Knee joint kinematics signals in the sagittal plane by means of electro-
 17 goniometers (accuracy: 0.5°).

18 The sEMG active probes were placed over the following 12 muscles of the
 19 dominant lower limb and trunk (bilaterally): right and left Longissimus Dorsii
 20 (LD_R, LD_L), Tensor Fasciae Latae (TFL), Gluteus Medius (GMD), Rectus Femoris
 21 (RF), Lateral Hamstring (LH), Medial Hamstring (MH), Vastus Medialis (VM),
 22 Lateral Gastrocnemius (LGS), Peroneus Longus (PL), Soleus (SOL) and Tibialis
 23 Anterior (TA). Foot-switches were positioned beneath the heel, the first, and fifth
 24 metatarsal heads, bilaterally. Electro-goniometers were positioned on the lateral
 25 aspect of the knee joint, bilaterally.

26 An example of sensor placement is shown, for a representative healthy subject
 27 of the sample population, in **Figure 3.2**.

28 Signals were acquired with a sampling frequency of 2000 Hz, converted by a
 29 12-bit analog to digital converter, and sent to a PC for real-time representation. The
 30 acquired signals were then imported into the MATLAB[®] release 2020b (The
 31 MathWorks Inc., Natick, MA, USA) to be processed by means of custom routines.

32 3. Data Processing

33 Before muscle synergy extraction, the acquired sEMG signals were pre-
 34 processed considering two different approaches:

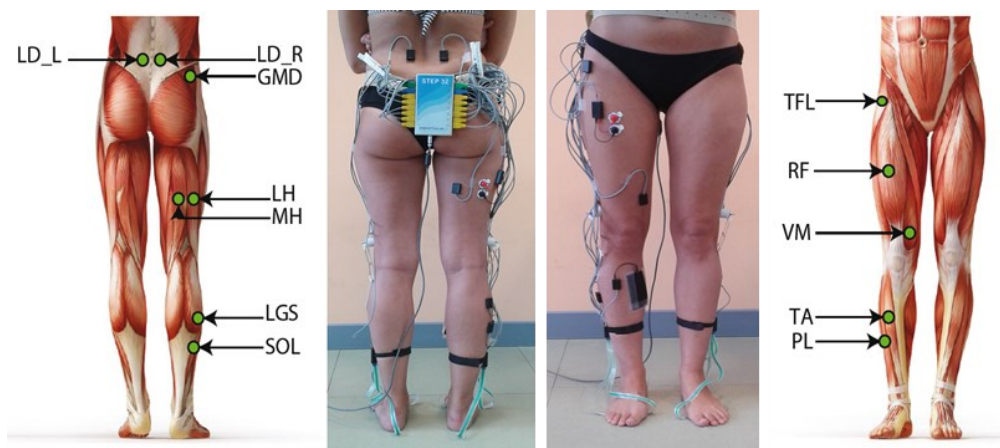


Figure 3.2 Sensor placement for a healthy subject of the sample population. sEMG active probes are placed over the principal muscle of the dominant lower limb and the trunk (bilaterally). Foot-switches are placed beneath the heel, the first, and the fifth metatarsal heads to detect gait phases and time-segment gait cycles. Electro-goniometers are positioned on the lateral aspect of the knee joint.

- 1 *i.* the *standard* approach, in which all sEMG time-instants were considered
- 2 for muscle synergy extraction
- 3 *ii.* the *novel* approach in which only the principal activation intervals of the
- 4 sEMG signal were considered.

5 **Figure 3.3** shows the workflow of the two approaches implemented for the

6 extraction of the muscle synergies.

7 *Gait Cycle Segmentation and Normalization*

8 In the last years, several approaches have been proposed to time-segment

9 sEMG signal [108–110]. In this study, the foot-switch signals were used to time-

10 segment and classify the gait cycles according to the foot-floor contact sequence.

11 Only the HFPS gait cycles, defined as the physiological sequence of Heel Contact

12 (H), Flat Foot Contact (F), Push Off (P), and Swing (S), were considered, discarding

13 those characterized by atypical gait cycles [108]. Only the gait cycles belonging to

14 the rectilinear path (see **Figure 3.1**) were analyzed, removing those corresponding

15 to direction changes at the beginning and at the end of the walkway (including

16 deceleration before and acceleration after the U-turn) [11]. Finally, each segmented

17 gait cycle was normalized in amplitude with respect to its global maximum, time-

18 normalized to 1000 samples [55], and all gait cycles selected were concatenated in

19 a single vector [111].

20 *Extraction of Principal Activations (PAs) through CIMAP algorithm*

21 The muscle activation onset/offset timing was computed from the sEMG

22 signals by means of a muscle activity detector based on LSTM neural networks

23 (LSTM-MAD). See **A Deep Learning Approach for Muscle Activity Detection**

24 chapter for further details on the LSTM-MAD approach. The optimized version of

25 the CIMAP (Clustering for Identification of Muscle Activation Patterns) [104]

26 algorithm was then applied to the muscle activation intervals to select PAs.

27 Briefly, PAs are defined as those muscle activations that are strictly necessary

28 to accomplish a specific biomechanical task and describe the fundamental

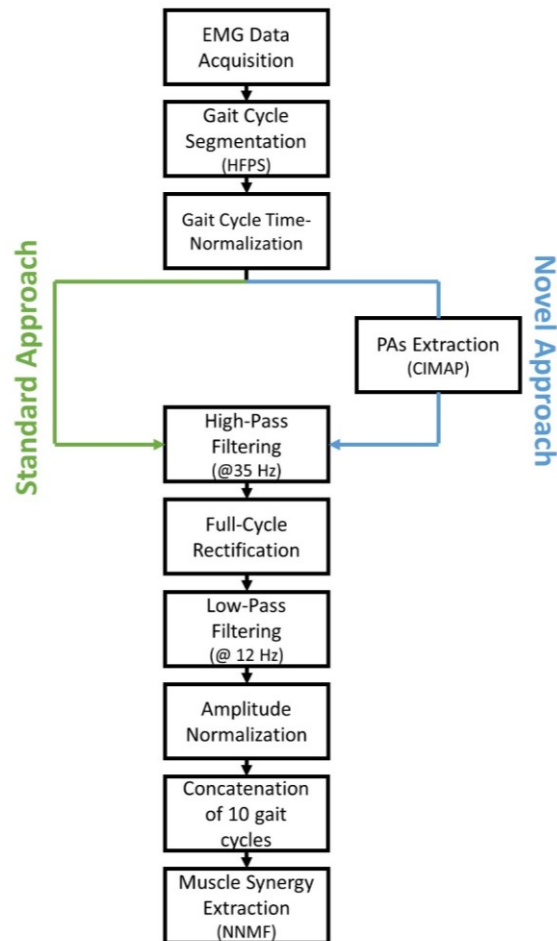


Figure 3.3 Workflow of the two approaches implemented for the muscle synergy extraction. Reprinted from “*Muscle Synergies Extracted Using Principal Activations: Improvement of Robustness and Interpretability*” by Ghislieri *et al.* (2020).

1 activation intervals of a specific muscle. The CIMAP algorithm, based on
 2 hierarchical clustering, groups together the gait cycles sharing similar sEMG onset-
 3 offset patterns. For each cluster, the cluster prototype is defined as the median
 4 timing pattern. Considering only the most representative clusters (i.e., those
 5 containing a significant number of elements compared to the total number of gait
 6 cycles acquired during the task), PAs are computed as the intersection of the
 7 representative clusters’ prototypes [104]. Then, for each observed muscle, a single
 8 PA was obtained. PA was defined as a 1000-samples binary mask that was set to 1
 9 in correspondence of the time-instants in which a principal activation was detected
 10 and to 0 otherwise.

11 **Figure 3.4** represents an example of application of the CIMAP algorithm to
 12 sEMG gait data acquired during the walking task of a healthy subject of the sample
 13 population from the PL and GMD muscles. **Figure 3.4A** represents normalized
 14 activation intervals for the various gait cycles, grouped in clusters sharing similar
 15 activation timings. Each orange interval represents the prototype of a representative
 16 cluster. **Figure 3.4B** depicts how PAs are obtained from the intersection of the
 17 cluster prototypes.

18 The extraction of the PA intervals from the original sEMG signals of a specific

1 muscle was performed by windowing all the time-normalized gait cycles through
2 the correspondent binary mask.

3 See **Extraction of Principal and Secondary Activations through CIMAP**
4 chapter for additional details on the optimized version of the CIMAP algorithm.

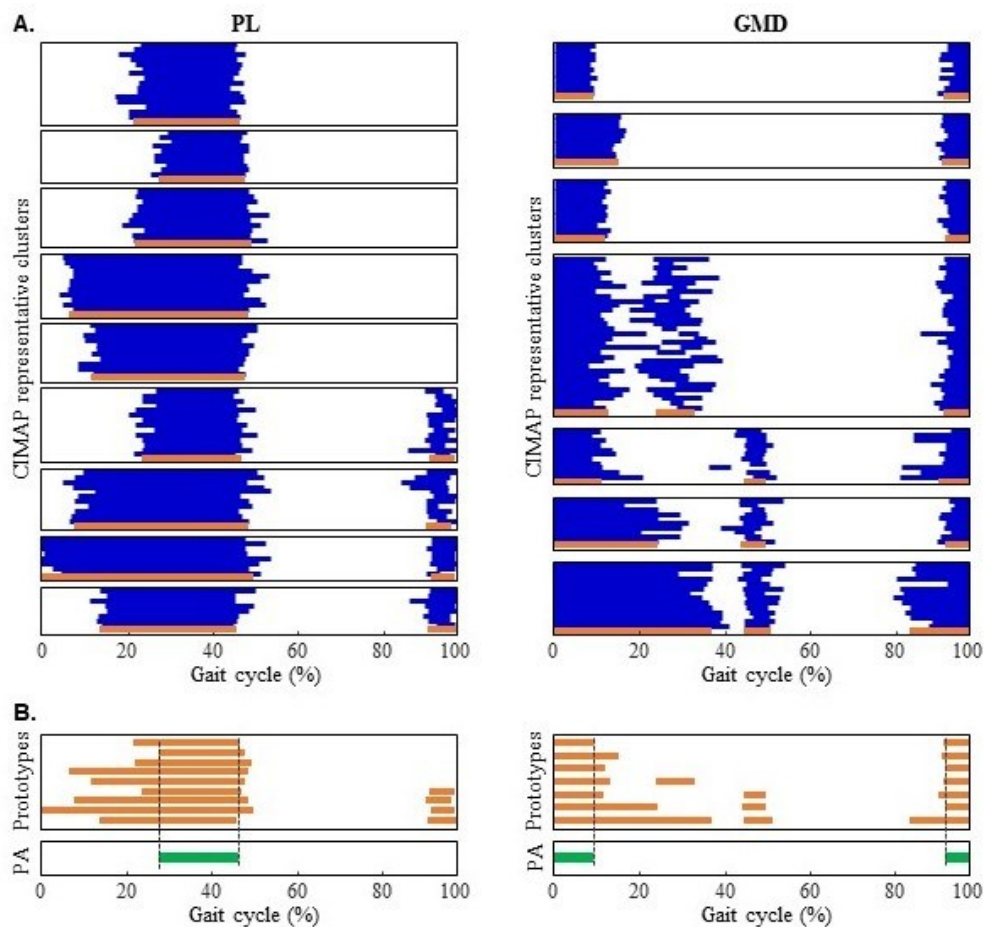


Figure 3.4 Application of the optimized version of the CIMAP algorithm to sEMG signals acquired from the PL (left) and GMD (right) muscles of a healthy subject of the sample population during a walking task. (A) Blue lines represent the cluster elements (muscle activation intervals computed through LSTM-MAD algorithm) normalized into 1000 time points with respect to the gait cycle duration, while orange intervals represent the prototypes of each cluster (computed as median of the elements that belong to the same cluster). (B) Principal activation (PA) is represented in green and is defined as the intersection of all the cluster's prototypes. Reprinted from “*Muscle Synergies Extracted Using Principal Activations: Improvement of Robustness and Interpretability*” by Ghislieri *et al.* (2020).

5 *Muscle Synergies Extraction and Sorting*

6 The sEMG signals were high-pass filtered at 35 Hz by means of an 8th-order
7 Butterworth digital filter to remove motion artifact, demeaned, and full-cycle
8 rectified to obtain a non-negative signal. Then, the rectified signals were low-pass
9 filtered by means of a 5th order digital Butterworth filter with a cut-off frequency
10 of 12 Hz to obtain the sEMG envelope. The signals were then normalized in
11 amplitude with respect to the global maximum of each muscle to ensure an equally
12 weighted contribution of each acquired muscle in the muscle synergy extraction
13 process.

14 The normalized envelopes were divided into groups of 10 concatenated gait

1 cycles (called subgroups) allowing for muscle synergy assessment over the entire
 2 walk duration [112]. For each subgroup, muscle synergies were extracted from the
 3 filtered sEMG signals by means of the Non-Negative Matrix Factorization (NNMF)
 4 algorithm. This algorithm models the original sEMG signals (M) as the linear
 5 combination of the time-independent muscle synergy weights (W) and the time-
 6 dependent activation coefficients (C) [34] as described in (3.2):

$$M(t) = \sum_{k=1}^N C(t)_k \cdot W_k + e \quad (3.2)$$

7 where N represents the optimal number of muscle synergies needed to properly
 8 model the original sEMG data. The weight vector W_k describes the contribution of
 9 each observed muscle to the k -synergy, the activation coefficient vector
 10 $C(t)_k$ represents the time-dependent modulation of the muscles recruited in the k -
 11 synergy and e represents the prediction error of the factorization algorithm. The
 12 MATLAB[®] function “*nnmf*” was used to apply the NNMF algorithm, setting the
 13 routine’s input parameters as follows: multiplicative update as factorization
 14 algorithm, $1e^{-6}$ as function and search tolerance, 50 as the number of factorization
 15 replicates, and 1000 as the maximum number of factorization iterations.

16 The first algorithm initialization was performed differently for the weight
 17 vector W_k and the activation coefficient vector $C(t)_k$. The C matrix was initialized
 18 with values randomly selected from a uniform distribution in the range $[0, 1]$. To
 19 improve the performance of the factorization algorithm and the accuracy in the
 20 reconstruction of the original sEMG data, a sparseness constraint was imposed in
 21 the initialization of the W matrix [37]. In particular, W matrix was first initialized
 22 with values randomly chosen from a uniform distribution in the range $[0, 0.05]$,
 23 then one random element of each W_k vector was set to a value selected from a
 24 uniform distribution in the range $[0.7, 0.8]$. Therefore, only one muscle for each k -
 25 synergy has a significant contribution, obtaining an extremely sparse NNMF
 26 initialization [37].

27 To explore different solutions of the NNMF algorithm, the factorization
 28 process was run many times on the same gait data, changing the number of muscle
 29 synergies (N) in the range $[1, 8]$. For each value of N , the reconstruction quality
 30 was assessed by means of the total Variance Accounted For ($tVAF$), defined as the
 31 uncentered Pearson’s correlation coefficient expressed in percentage (3.3):

$$tVAF = \left(1 - \frac{\sum_{k=1}^m (M_k - M_k^R)^2}{\sum_{k=1}^m M_k^2} \right) \cdot 100 \quad (3.3)$$

32 where m represents the number of muscles observed, $\sum_{k=1}^m (M_k - M_k^R)^2$ describes
 33 the sum of the squared errors between the original (M_k) and reconstructed (M_k^R)
 34 sEMG signals of the k -synergy, and $\sum_{k=1}^m M_k^2$ represents the total sum of squared
 35 M_k values with respect to zero. The optimal number of muscle synergies needed to
 36 properly model the sEMG signals of the i -th subgroup ($N_{90,i}$) was selected by
 37 choosing the smallest number of synergies (N) granting a $tVAF$ value equal to or

greater than 90% [55]. Since each subgroup could be described by a different number of muscle synergies, the final number of synergies (N_{90}) was then selected, for each subject, as the mode of the numbers of muscle synergies computed on each 10-gait-cycle subgroup ($N_{90,i}$).

To represent the synergy output in the range [0, 1], the weight vectors W_k were normalized in amplitude with respect to their global maximum. Then, the activation coefficient vectors $C(t)_k$ were multiplied by the correspondent normalized values.

To sort the muscle synergies in the same order for each subgroup and for each subject, the k -means algorithm was applied to the W matrix [22]. Clustering was performed by means of the MATLAB[®] routine “*kmeans*” using the following input parameters: N_{90} as number of k -means clusters, 1000 as maximum number of iterations, 15 as number of replicates, and cosine as distance metric. The activation coefficients matrix C was ordered accordingly.

14 *Muscle Synergy Analysis*

The number of muscle synergies, the intra-subject consistency, the robustness, and the interpretability were used in the following to quantitatively compare the muscle synergies extracted from the PD and healthy populations considering the *standard* and the novel approach.

19 *i. Number of Muscle Synergies*

As stated above, the optimal number of muscle synergies needed to properly reconstruct the sEMG signals of the i -th subgroup ($N_{90,i}$) was selected by choosing the smallest number of synergies (N) granting a $tVAF \geq 90\%$. The final number of synergies (N_{90}) was then selected, for each subject, as the mode of the number of muscle synergies computed on each 10-gait-cycle subgroup.

26 *ii. Intra-Subject Consistency*

The intra-subject consistency of the muscle synergies among the 10-gait-cycle subgroups was evaluated by computing the Cosine Similarity (CS) [30] of the previously sorted weight vectors W_k and activation coefficient vectors $C(t)_k$, separately. The CS between the previously sorted weight vectors belonging to the i - and j -th subgroup of the k -synergy was defined as the normalized scalar product between the vectors expressed in percentage, as described in (3.4) and (3.5):

$$CS_{W,k}^{i,j} = \left(\frac{W_k^i \cdot W_k^j}{\|W_k^i\| \|W_k^j\|} \right) \cdot 100 \quad (3.4)$$

$$CS_{C,k}^{i,j} = \left(\frac{C_k^i \cdot C_k^j}{\|C_k^i\| \|C_k^j\|} \right) \cdot 100 \quad (3.5)$$

where $CS_{W,k}^{i,j}$ and $CS_{C,k}^{i,j}$ represent the cosine similarity computed between the weight vectors W_k and the activation coefficients $C(t)_k$ of the i - and j -th subgroup, respectively. The CS values range between 0 (no similarity) and 1 (complete similarity).

1 *iii. Robustness*

2 The robustness of the muscle synergies among different subgroups of 10 gait
3 cycles was assessed through the Cross-Variance Accounted For [111] defined as in
4 (3.6):

$$CrossVAF^{i,j} = \left(1 - \frac{\sum_{k=1}^m (M_k^i - M_k^{R,j})^2}{\sum_{k=1}^m (M_k^i)^2} \right) \cdot 100 \quad (3.6)$$

5 where M_k^i and $M_k^{R,j}$ represent the original and the reconstructed sEMG signals of
6 the k -muscle for the i - and j -th subgroup, respectively.

7 This parameter assesses how well the muscle synergies extracted for the i -th
8 subgroup reconstruct the sEMG signals that belong to the j -th subgroup. For each
9 subject, the average *CrossVAF* value was computed over all the possible couples
10 of 10-gait-cycle subgroups without considering the main diagonal of the *CrossVAF*
11 matrix. The average *CrossVAF* can assume values ranging from 0% to 100%,
12 corresponding to a low or high correlation between the reconstructed and the
13 original sEMG signals from different subgroups, respectively.

14

15 *iv. Interpretability*

16 Previous studies demonstrated that five muscle synergies are usually needed to
17 properly describe human locomotion [19,24]. Each of these muscle synergies can
18 be characterized by a specific biomechanical function. These biomechanical
19 functions were generally assigned to each muscle synergy by applying an amplitude
20 threshold to the weight vectors ($W_k > 0.5$) in order to define the prevailing muscles
21 contributions [24]. Moreover, the profile of the activation coefficients $C(t)_k$ was
22 usually observed to define the gait cycle phase in which the muscle synergy is
23 mainly activated, according to the functional gait cycle phases proposed by Perry
24 *et al.* [97]. In the study by Rimini *et al.* [24] 5 biomechanical functions common to
25 all the analyzed subjects during the walking task were discovered. TABLE 3.2
26 reports the description of each biomechanical function and the muscles mainly
27 involved in each function ($W_k > 0.5$).

TABLE 3.2
BIOMECHANICAL FUNCTIONS OF THE MUSCLE SYNERGIES DURING GAIT.

Function	Involved muscles	Biomechanical function
F1	TFL, GMD	Hip joint stabilization during heel strike and load acceptance phase.
F2	LGS, PL, SOL	Propulsion at the mid and terminal stance.
F3	TA	Forefoot clearance control during the swing phase and foot control during the first rocker.
F4	MH, LH	Leg deceleration at the end of the swing phase.
F5	LD _R , LD _L	Control of the trunk position in the frontal plane at heel strike.

Reprinted from "Muscle Synergies Extracted Using Principal Activations. Improvement of Robustness and Interpretability" by Ghislieri *et al.*, 2020

28 For example, the biomechanical function F2 is used to generate the propulsion
29 and requires, among the observed muscles, the involvement of the LGS, PL, and
30 SOL muscles (the other acquired muscles are not directly involved during
31 propulsion).

1 The muscle synergy Interpretability (I) was evaluated considering the average
 2 of the muscle synergy weights that are not directly involved in the k -synergy
 3 biomechanical function as described in (3.7):

$$I = \frac{1}{N} \sum_{k=1}^N ((1 - \overline{W}_k') \cdot 100) \quad (3.7)$$

4 where N represents the final number of muscle synergies needed to describe the
 5 motor task and \overline{W}_k' is the average contribution of the muscles not directly involved
 6 in the biomechanical task described by the k -synergy. A muscle synergy can be
 7 considered more easily interpretable when the averages of the muscle synergy
 8 weights not directly involved in the specific biomechanical function are close to
 9 zero ($\overline{W}_k' \cong 0$), while it can be considered less interpretable when they have values
 10 comparable with the weights of the muscles directly involved in the biomechanical
 11 function ($\overline{W}_k' \cong 1$). The I values are expressed in percentage, and range between
 12 0% and 100%, suggesting low and high interpretability, respectively.

13 *Statistical Analysis*

14 To determine if there is a statistically significant difference, in the walking
 15 speed, between healthy and PD subjects, the Wilcoxon signed-rank test was used,
 16 setting the significance level (α) at 0.05. Two-way repeated measures analysis of
 17 variance (ANOVA) followed by *post-hoc* analysis with Tukey's adjustment for
 18 multiple comparisons was used to test differences in muscle synergy results
 19 between Group (Healthy and PD) and Approach (Standard and PAs). The effect
 20 size of the statistically significant differences was calculated by means of the
 21 Hedges' g including the correction for small sample sizes [113]. The significance
 22 level (α) was set equal to 0.05. The statistical analysis was carried out using the
 23 Statistical and Machine Learning Toolbox of MATLAB®.

24 **3.3 Results**

25 During the experimental sessions, an average speed of 1.2 ± 0.1 m/s and $1.0 \pm$
 26 0.2 m/s and a dataset of 156 ± 25 and 129 ± 37 *typical* gait cycles (i.e., gait cycles
 27 characterized by the HFPS foot-floor contact sequence) were assessed for the
 28 healthy and PD population, respectively. On average, the Wilcoxon test revealed a
 29 slightly significant decrease ($p = 0.04$) in the walking speed of PD patients with
 30 respect to the healthy subjects.

31 In the following, the results obtained using the *standard* and the novel approach
 32 (entailing the extraction of PAs) were compared in terms of (i) number of muscle
 33 synergies, (ii) intra-subject consistency, (iii) robustness, and (iv) interpretability.

34 *i. Number of Muscle Synergies*

35 On average, an optimal number of muscle synergies of 4.8 ± 0.1 and 5.0 ± 0.1
 36 was found on the healthy population, considering the *standard* and the novel

1 approach, respectively. Considering the PD population, an average number of 4.7
 2 ± 0.1 and 5.1 ± 0.1 muscle synergies was computed considering the *standard* and
 3 the novel approach, respectively. Two-way repeated measures ANOVA revealed a
 4 statistically significant increase ($p = 0.002$, $g = 0.69$) in the optimal number of
 5 muscle synergies computed considering the novel approach with respect to the
 6 *standard* one, while no statistically significant differences ($p = 0.83$) were assessed
 7 between healthy and PD populations. No significant interaction effects were
 8 detected between Group and Approach.

9 The average number of muscle synergies values are reported in **TABLE 3.3**,
 10 with the indication of the statistically significant differences ($p < 0.05$).

TABLE 3.3
 MUSCLE SYNERGY RESULTS ON HEALTHY SUBJECT AND PARKINSON DISEASE (PD) PATIENTS

	HEALTHY SUBJECTS		PD PATIENTS		2-WAY ANOVA (P-VALUE)	
	Standard	PAs	Standard	PAs	Group	Approach
Number of muscle synergies	4.8 \pm 0.1	5.0 \pm 0.1	4.7 \pm 0.1	5.1 \pm 0.1	0.83	0.002
Intra-subject consistency						
• Weight vectors	0.99 \pm 0.01	0.99 \pm 0.01	0.98 \pm 0.02	0.98 \pm 0.02	0.003	0.13
• Activation coefficients	0.99 \pm 0.01	0.99 \pm 0.01	0.98 \pm 0.01	0.98 \pm 0.02	0.005	0.24
Robustness (%)	80.7 \pm 0.7	80.6 \pm 1.2	79.1 \pm 0.7	79.6 \pm 0.7	0.13	0.82
Interpretability (%)	90.6 \pm 4.4	93.6 \pm 3.9	90.1 \pm 2.6	93.7 \pm 1.7	0.79	< 0.0001

Values of parameters are reported as mean \pm standard deviation over the population.

Statistically significant differences ($p < 0.05$) between Group or Approach are represented in bold.

11 **Figure 3.5** and Error! Reference source not found. report the weight vectors
 12 and the activation coefficients extracted from a representative healthy subject and
 13 PD patient of the sample population, respectively, using the two pre-processing
 14 techniques: (

15 **Figure 3.5A** and Error! Reference source not found.A) *standard* approach and
 16 (

17 **Figure 3.5B** and Error! Reference source not found.B) the novel approach
 18 (extraction of PAs). No significant differences both in the final number of muscle
 19 synergies and in their composition were revealed.

20 *ii. Intra-Subject Consistency*

21 Considering the activation coefficients $C(t)_k$, two-way repeated-measures
 22 ANOVA revealed a statistically significant decrease ($p = 0.005$, $g = 0.83$) in the
 23 intra-subject consistency of the PD population with respect to the healthy
 24 population. No statistically significant differences ($p = 0.24$) between Approach
 25 and no significant interaction effects between Group and Approach were detected.

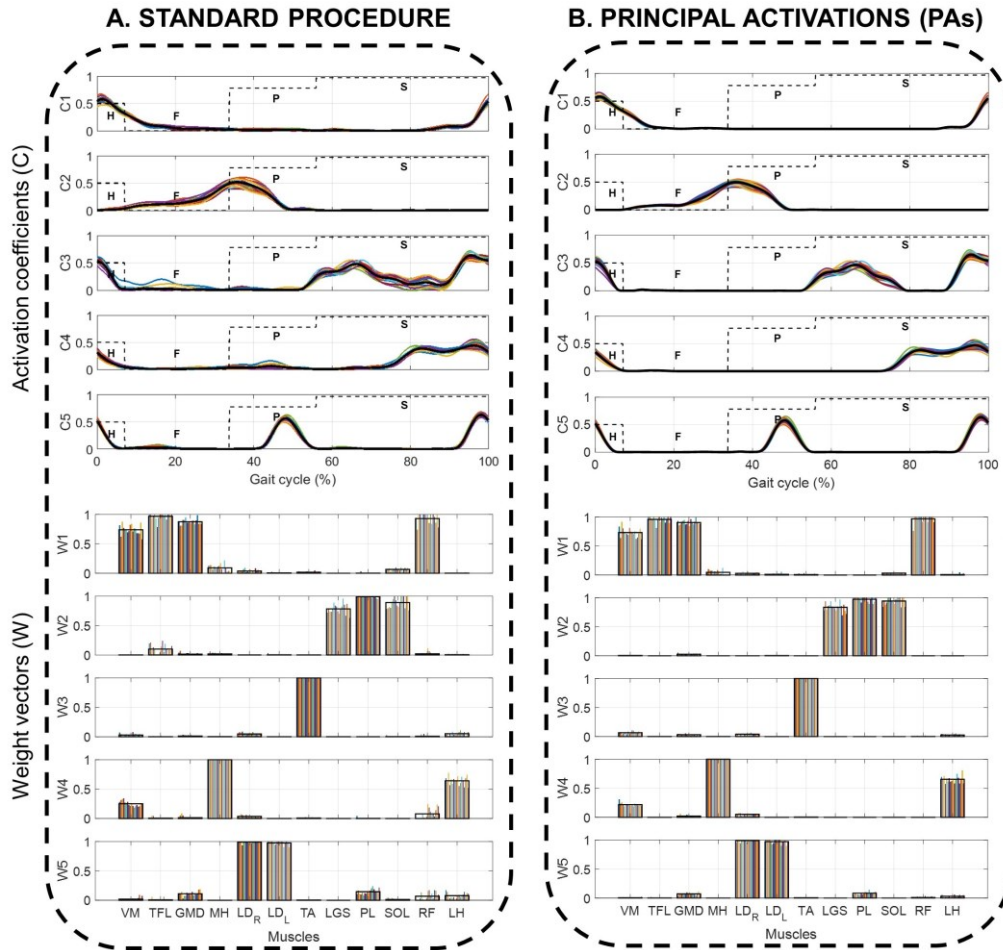


Figure 3.5 Example of activation coefficients C_k and weight vectors W_k extracted from a representative healthy subject of the sample population using the two different processing techniques: (A) standard approach, and (B) novel approach (extraction of PAs). Each colored line (or colored vertical bar) represents C_k (or W_k) extracted from a single 10-gait-cycle subgroup, while black lines (or top of black rectangles) represent the average C_k (or W_k) computed over subgroups. The dotted lines, in the C_k -plots, represent the mean foot-switch signal with the indication of the 4 gait phases: Heel contact (H), Flat foot contact (F), Push off (P) and Swing (S).

1 Considering the weight vectors W_k , two-way ANOVA revealed a statistically
 2 significant decrease ($p = 0.003$, $g = 0.66$) in the intra-subject consistency of the PD
 3 population with respect to the healthy population. No statistically significant
 4 differences ($p = 0.13$) between Approach and no significant interaction effects
 5 between Group and Approach were detected.

6 The average intra-subject consistency values are reported in **TABLE 3.3**, with
 7 the indication of the statistically significant differences ($p < 0.05$).

8 **Figure 3.7A** shows the boxplots of the intra-subject consistency computed on
 9 the healthy and PD populations using the two approaches.

10 *iii. Robustness*

11 On average, a robustness value of $80.7\% \pm 0.7\%$ and $80.6\% \pm 1.2\%$ was found
 12 on the healthy population, considering the *standard* and the novel approach,
 13 respectively. Considering the PD population, instead, an average robustness value
 14 of $79.1\% \pm 0.7\%$ and $79.6\% \pm 0.7\%$ was computed considering the *standard* and

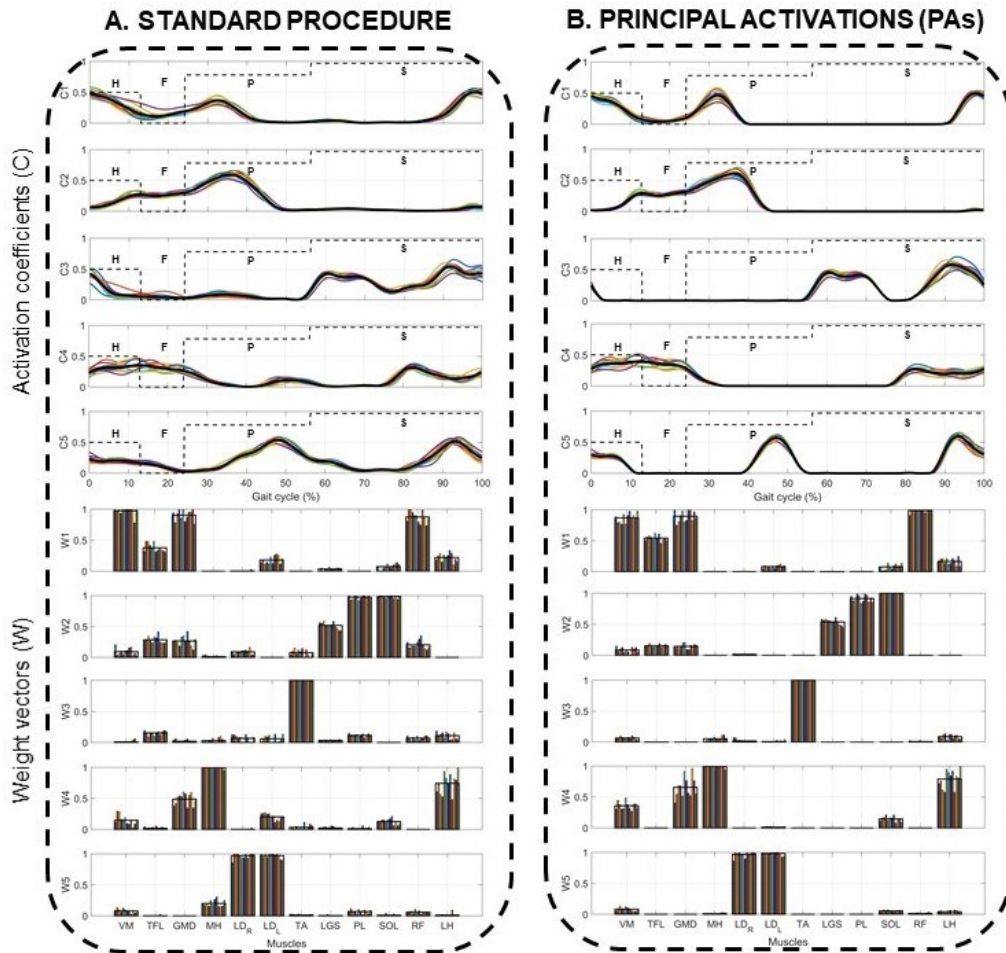


Figure 3.6 Example of activation coefficients C_k and weight vectors W_k extracted from a representative PD patient of the sample population using the two different processing techniques: (A) standard approach, and (B) novel approach (extraction of PAs). Each colored line (or colored vertical bar) represents C_k (or W_k) extracted from a single 10-gait-cycle subgroup, while black lines (or top of black rectangles) represent the average C_k (or W_k) computed over subgroups. The dotted lines, in the C_k -plots, represent the mean foot-switch signal with the indication of the 4 gait phases: Heel contact (H), Flat foot contact (F), Push off (P) and Swing (S).

1 the novel approach, respectively. Two-way repeated-measures ANOVA revealed
 2 no statistically significant differences in the muscle synergy robustness between
 3 Group ($p = 0.13$) and between Approach ($p = 0.82$). No significant interaction
 4 effects were detected between Group and Approach.

5 The average robustness values are reported in TABLE 3.3, with the indication
 6 of the statistically significant differences ($p < 0.05$).

7 **Figure 3.7B** represents the boxplots of the $CrossVAF$ computed on the healthy
 8 and PD populations considering the two processing approaches.

9 *iv. Interpretability*

10 On average, a muscle synergy interpretability value of $90.6\% \pm 4.4\%$ and
 11 $93.6\% \pm 3.9\%$ was found on the healthy population, considering the *standard* and
 12 the novel approach, respectively. Considering the PD population, instead, an
 13 average interpretability value of $90.1\% \pm 2.6\%$ and $93.7\% \pm 1.6\%$ was computed
 14 considering the *standard* and the novel approach, respectively. Two-way repeated-

1 measures ANOVA reveals a statistically significant increase ($p < 0.0001$, $g = 1.01$)
 2 in the interpretability of the muscle synergies extracted considering the novel
 3 approach with respect to those extracted considering the *standard* one. No
 4 statistically significant differences ($p = 0.79$) were assessed between Group
 5 (healthy and PD) and no significant interaction effects were detected between
 6 Group and Approach.

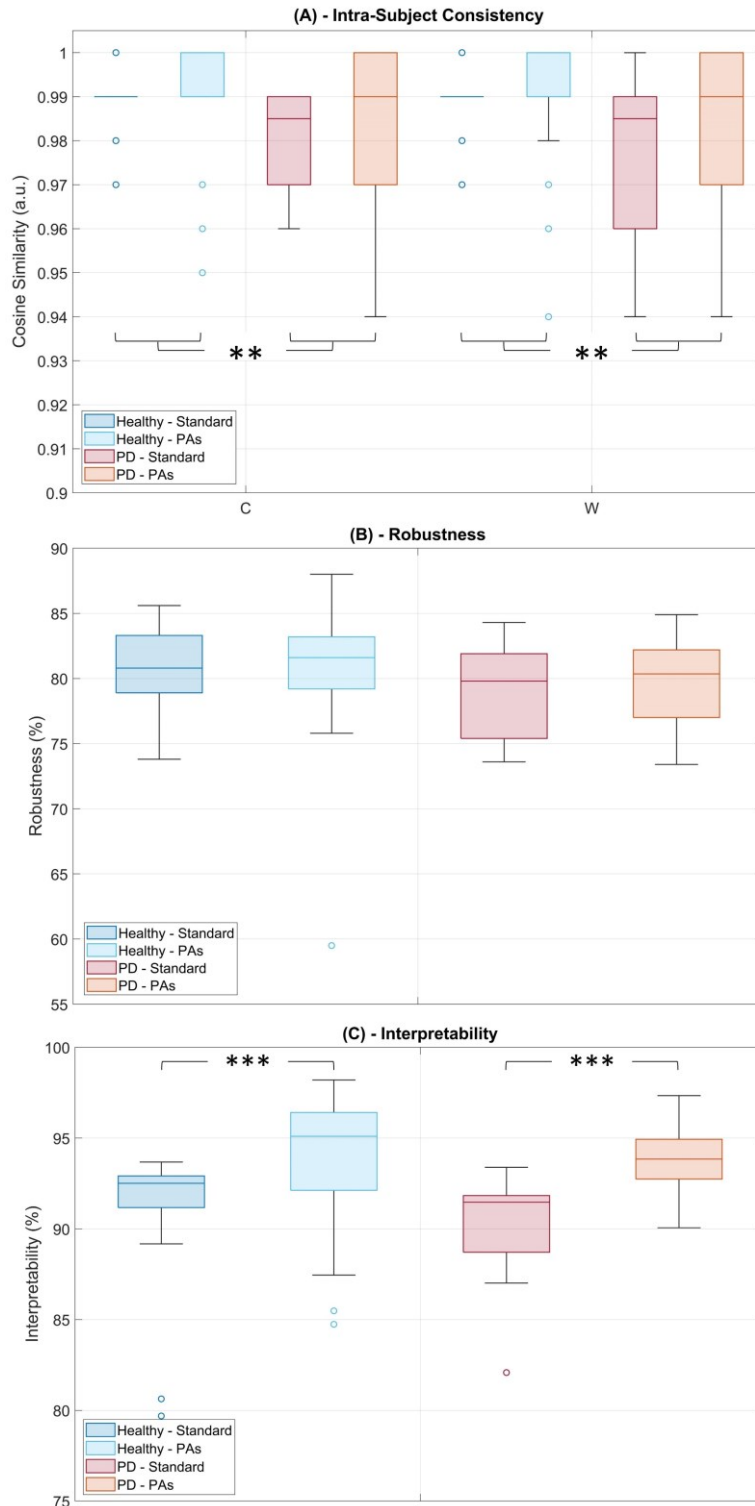


Figure 3.7 Boxplots of (A) intra-subject consistency, (B) robustness, and (C) interpretability used to compare the standard and the novel approach for muscle synergy extraction. Outliers are indicated by circles. Single and double asterisks represent p -values lower than 0.05 and 0.01, respectively.

1 The average interpretability values are reported in **TABLE 3.3**, with the
2 indication of the statistically significant differences ($p < 0.05$).

3 **Figure 3.7C** shows the boxplots of muscle synergy interpretability computed on
4 the healthy and PD populations considering both the pre-processing techniques.

5 **3.4 Discussion and Conclusions**

6 Although the effectiveness of the muscle synergy theory in modeling the
7 modular organization of the CNS during motor tasks is well known in literature
8 [7,24,27,30,47,55], there are no shared standards for sEMG pre-processing before
9 muscle synergy extraction. The pre-processing techniques previously used in
10 literature (“*standard*” approaches) for muscle synergy extraction generally
11 considered the whole sEMG time-instants as input of the factorization algorithm
12 [19,23,28,55]. However, these standard approaches may be influenced by the high
13 cycle-by-cycle variability of the sEMG activation patterns [100]. Therefore, these
14 approaches may not be sufficient to fully understand the motor control strategies
15 during human locomotion.

16 The combination of SGA and CIMAP (with the extraction of the principal and
17 secondary muscle activation intervals) algorithms as a pre-processing step before
18 muscle synergy extraction may provide a valuable tool to understand how a set of
19 muscle synergies are recruited and modulated by the central nervous system under
20 different conditions. In particular, the ability to distinguish between the principal
21 and the secondary muscle activation intervals allows for separately assessing
22 human motor control strategies adopted during gait cycle phases characterized by
23 highly consistent and repeatable activation intervals (PAs) and during gait cycle
24 phases characterized by extemporary and lowly repeatable activation intervals
25 (SAs), respectively. Moreover, the selection of the PAs from the whole sEMG
26 signal may help the interpretation of muscle activation patterns by reducing the
27 intra-subject variability of the activation intervals. In this chapter, only the
28 applicability of the selection of PAs as a pre-processing step before muscle synergy
29 extraction was tested, leaving the study of motor control strategies during secondary
30 muscle activations to future studies.

31 As stated before, the extraction of the PAs allows evaluating only the
32 “*necessary*” muscle activations, discarding those with auxiliary function [105],
33 such as those providing corrections to cyclic motion and body segment posture
34 (whose muscle synergies can be investigated through the selection of the SAs from
35 the whole sEMG signals). According to this approach, only the sEMG time-instants
36 in correspondence of PAs are considered as inputs of the factorization algorithm
37 [24,104], while the remaining sEMG time-instants are set to zero.

38 To assess the differences in muscle synergies between the newly proposed
39 approach and the standard one, the muscle synergies extracted from 22 healthy
40 subjects and 22 patients affected by Parkinson’s disease during a walking task have
41 been compared in terms of number of muscle synergies, intra-subject consistency,
42 robustness, and muscle synergy interpretability.

43 In terms of the number of muscle synergies, both approaches accurately

1 reconstruct the original sEMG signals acquired from both the sample populations
2 through five muscle synergies. These muscle synergies are very similar both in their
3 composition (temporal modulation of the activation coefficients and weighted
4 contribution of the observed muscles) and in the biomechanical functions described
5 by each of them. In particular, the same five biomechanical functions described in
6 Ref. [24] can be associated with the muscle synergies extracted considering both
7 the standard and the novel approach. Even if a statistically significant difference in
8 age between PD patients and control subjects was detected, a few studies, such as
9 those published by Monaco *et al.*[20,114], demonstrated that neither temporal
10 patterning of activation coefficients nor muscles' contribution to each muscle
11 synergy were significantly affected by aging. Moreover, the purpose of this chapter
12 was to assess the applicability and the performance of the novel approach
13 considering both physiological and pathological conditions rather than assess the
14 difference in motor control strategies between the two sample populations.

15 In terms of intra-subject consistency and robustness, the standard and the novel
16 approach reveal similar average values of intra-subject consistency and robustness
17 of muscle synergies considering both healthy and pathological subjects, suggesting
18 high repeatability of the motor control strategy among the 10-gait-cycle subgroups
19 and high robustness of muscle synergy results, independently from the pre-
20 processing technique used. However, statistically significant ($p < 0.05$) differences
21 in weight vector and activation coefficient intra-subject consistencies between the
22 PD patients and healthy controls have been detected considering both the standard
23 and the novel approach. This finding is consistent with a previous study by Allen
24 *et al.* [115] focused on assessing the improvement of gait and balance function in
25 PD patients after short-term rehabilitation through the muscle synergy theory.
26 Results revealed that muscle synergies were more consistently recruited
27 immediately after rehabilitation, suggesting more reliable and repeatable motor
28 output. Furthermore, intra-subject consistency has been demonstrated to be a more
29 sensitive marker of improved motor performance compared to the number of
30 muscle synergies. Indeed, only a modest reduction in muscle synergy number has
31 been detected in the PD population compared to the control subjects.

32 In terms of interpretability of the muscle synergies, the novel approach
33 outperforms the standard one. Indeed, results suggest that the extraction of the PAs
34 allows for obtaining higher interpretability of the muscle synergies considering both
35 healthy ($90.6 \% \pm 4.4 \%$ vs. $93.6 \% \pm 3.9 \%$) and pathological patients ($90.1\% \pm$
36 2.6% vs. $93.7 \% \pm 1.7 \%$) compared to the standard approach, thus providing a more
37 clear assessment of the modular organization of the CNS during gait. Moreover, the
38 possibility to extract more distinctly organized (or interpretable) muscle synergies
39 may be useful to make easier the assignment of a specific biomechanical function
40 to each muscle synergy.

41 The analysis of the final number and the composition of the obtained muscle
42 synergies revealed no loss of information due to the extraction of the PAs,
43 compared to the standard approach. Moreover, the higher performance in terms of
44 muscle synergy interpretability obtained considering the novel approach
45 demonstrates that the extraction of the PAs may successfully improve the muscle

1 synergy analysis during gait in both healthy and pathological subjects.

2 The number and composition of muscle synergies can also be strongly
3 influenced by other steps of the muscle synergy extraction process
4 [23,36,38,39,44,49–53], such as the sEMG pre-processing (e.g., low-pass filtering
5 techniques) [38,49–51,53] and the number and choice of muscles acquired [23].
6 However, the focus of this chapter was on developing and validating an approach
7 for muscle synergy extraction based on the combined application of SGA and
8 CIMAP (with the extraction of PAs) algorithms, and not on evaluating the effect of
9 different filtering techniques or subsets of acquired muscles on the extraction of the
10 muscle synergies. Indeed, the same number of acquired muscles (i.e., 12 lower limb
11 and trunk muscles) and filtering steps (i.e., high-pass filter at 35 Hz, full-cycle
12 rectification, and low-pass filter at 12 Hz) were considered for both the tested
13 approaches (standard and novel approach), following the common
14 recommendations and guidelines for muscle synergy extraction.

15 In this chapter, an innovative pre-processing technique for muscle synergy
16 extraction was proposed combining Statistical Gait Analysis (SGA) and a clustering
17 algorithm (CIMAP) that allows for obtaining the principal activations (PAs). To
18 properly compute PAs, sEMG signals must be recorded for at least 3 minutes during
19 gait. More specifically, sEMG signals used in this chapter for muscle synergy
20 extraction were acquired during 5-minute walking tasks. Signal recording during a
21 “long” physiological walk is needed to be able to collect at least 100 - 200 typical
22 gait cycles required for PAs computation. This requirement does not limit the
23 feasibility and applicability of the proposed methodology to pathological
24 populations. Indeed, gait analysis is commonly used to quantitatively assess
25 patients’ locomotion performance only in those patients able to independently walk,
26 for some minutes, without walking aids or external supports. In the past, several
27 studies demonstrated the feasibility of gait data acquisition, during recording
28 sessions lasting 3 minutes, in patients suffering from different neurological
29 conditions, e.g., Normal Pressure Hydrocephalus [89], mild ataxia [116], and
30 cerebral palsy [117].

31 In this chapter, the CIMAP algorithm was used to select PAs and discard
32 secondary activations, suggesting that muscle synergies are better understood when
33 considering only PAs. This is in line with a previous work in which the selection of
34 PAs was used to define a robust asymmetry index, based on sEMG activity during
35 locomotion [67]. In particular, the study by Castagneri *et al.* [67] provided a
36 validation both on healthy and pathological populations.

37 The approach proposed in this chapter was validated considering both healthy
38 and pathological populations, for whom the assessment of motor control through
39 muscle synergies may be of the uttermost importance [45,115]. However, when
40 considering pathological populations, also secondary activations (SAs) might be
41 fundamental in the interpretation of the results. Indeed, discarding the effect of
42 auxiliary functions included in the secondary activations might bias the
43 interpretation of pathological types of behavior, instead of improving it.
44 Nevertheless, it should be noticed that the opportunity to separate principal from
45 secondary activations, provided by the CIMAP algorithm, does not preclude

1 studying also secondary (auxiliary) activations. Therefore, future developments of
2 the proposed approach will include the analysis of muscle synergies extracted from
3 a pathological sample population considering the principal and the secondary
4 activations, separately.

5 In conclusion, the results presented in this chapter demonstrate that the
6 extraction of the principal activations can be successfully used as a pre-processing
7 step before muscle synergy extraction, allowing a more interpretable assessment of
8 the modular organization of the CNS during a walking task without any loss of
9 information.

1 Methodological Issue – Multi-Criteria Decision Analysis 2 for Selecting the Optimal Number of Muscle Synergies

3

4 3.5 Introduction

5 One of the main challenges for researchers and clinicians interested in human
6 motion control is to understand how the CNS controls a complex system, such as
7 the musculoskeletal system, during different movement tasks. The complexity of
8 the musculoskeletal system lies in its redundancy, due to the countless number of
9 muscles and joints involved and the many different ways that muscles can be
10 enrolled to execute the same motor task.

11 One of the most widely accepted theory for the assessment of human motor
12 control is the muscle synergy theory. Muscle synergies are a valuable tool to
13 quantitatively and non-invasively assess how the CNS can manage the high
14 complexity and redundancy of the motor control system. Applications range from
15 the medical field (e.g., monitoring of patients suffering from neurological,
16 neurodegenerative diseases [18,43], or joint disorders [118]), to the rehabilitation
17 field (e.g., pre-/post-treatment comparisons [119]), to the robotic field (e.g., control
18 of robotic devices or exoskeletons [17,120]). According to this theory, muscles
19 involved in a specific movement are activated in small groups, commonly referred
20 to as muscle synergies [121]. Therefore, the CNS can reduce the complexity of the
21 motor control system thanks to the activation of multiple muscles with a single
22 control signal. Previous studies have demonstrated that muscle activity during a
23 variety of movement tasks, such as walking, running, and cycling, can be
24 successfully described through a small set of muscle synergies responsible for
25 specific biomechanical functions [19]. In these studies, muscle synergies are
26 extracted from surface electromyography (sEMG) signals acquired from sets of
27 muscles during a variety of motor tasks, using different matrix factorization
28 algorithms, such as “standard” Non-Negative Matrix Factorization (NNMF)
29 [41,122], Gaussian NNMF (GNNMF), inverse Gaussian NNMF (IGNNMF)
30 [36,123], and principal or independent component analysis. Among all the
31 factorization algorithms, NNMF is the most widely used algorithm for muscle
32 synergy extraction, even if it was demonstrated that similar results can be obtained
33 by applying other approaches [32,40]. The NNMF algorithm models the original
34 sEMG data as a linear combination of muscle synergy weight vectors and activation
35 coefficients, whose dimensions depend on the number of muscle synergies needed
36 to properly reconstruct the motor task.

37 In the last years, the optimal number of muscle synergies (N_{opt}) has been
38 proposed as a meaningful feature for the analysis of motor control strategies in
39 pathological populations. In the study by Allen *et al.* [124], a reduced number of
40 muscle synergies has been demonstrated in patients affected by brain injury with
41 respect to a healthy population while executing walking tasks. Similar results have
42 been obtained in the work of Rodriguez *et al.* [45], in which a decreased

1 neuromuscular complexity during gait has been assessed in patients affected by
2 Parkinson's disease. These studies suggest that the number of muscle synergies and
3 their composition could be correlated with the motor control capacity and its
4 reduction in pathological conditions [33,55].

5 Despite the growing interest in the assessment of muscle synergy number, there
6 is a lack of standardized criteria or shared guidelines for computing the optimal
7 number of muscle synergies (N_{opt}) from sEMG recordings, revealing a low
8 reproducibility and comparability of muscle synergy results obtained from different
9 studies. One of the most widely used methods is the application of a cut-off
10 threshold based on the Variance Accounted For (VAF) [23,24,43–
11 46,112,118,120,125,126], defined as the uncentered Pearson's correlation
12 coefficient between the original and the reconstructed sEMG signals. More
13 specifically, according to this method, N_{opt} is defined as the smallest number of
14 synergies that ensures a VAF value above a cut-off threshold (threshold criterion).
15 The VAF threshold is commonly set equal to 90%
16 [23,24,43,44,46,112,118,120,125,126], but several other studies considered
17 threshold values in the range from 80% to 95% [45]. Discordance over the VAF
18 threshold values significantly intensifies the confusion over the selection of the
19 optimal number of muscle synergies and reduces the comparability with other
20 studies. To a lesser extent, N_{opt} was selected by means of the coefficient of
21 determination R^2 [16,31,34,47,121,122,127,128], setting a cut-off threshold for the
22 mean squared error between the R^2 curve and the linear fit. In the studies by Sawers
23 *et al.* [129] and by Cheung *et al.* [31], bootstrapping techniques have been used to
24 resample multiple times the sEMG data and compute VAF and R^2 curves,
25 respectively. The optimal number of muscle synergies was then selected as the
26 value at which the 95% confidence interval exceeds 90% VAF (or R^2).

27 To the best of the author's knowledge, only a few works have already explored
28 alternatives to VAF or R^2 curves, using the variability of muscle synergies among
29 task cycles [130], a task decoding-based metric [131], likelihood ratio tests, Bartlett
30 Akaike, Bayesian, and Laplacian information criteria [32,132–135].

31 In the last years, Multi-Criteria Decision Analysis (MCDA) or Multi-Criteria
32 Decision-Making (MCDM) approaches have been proposed in the field of
33 operational research, in addition to Cost-Benefit Analysis (CBA), to support
34 rigorous decision-making by explicitly taking into account multiple criteria [136].
35 Applications range from the medical field [137,138] to the transport field [139,140],
36 to the environment and water management [141,142]. The selection of the optimal
37 number of muscle synergies can be considered itself a multi-criteria approach,
38 where several criteria (e.g., reconstruction accuracy and muscle synergy
39 consistency) should be addressed to obtain the best representation of the motor
40 system.

41 This chapter aims at assessing the applicability of a novel approach for the
42 selection of the optimal number of muscle synergies (N_{opt}), based on an MCDA
43 approach, specifically developed to overcome the limitations of the standard VAF -
44 based approaches. The performance of the MCDA-based approach is evaluated and

1 compared against standard approaches (Threshold *VAF*, Elbow *VAF*, and Plateau
2 *VAF*) in terms of estimation bias and mean absolute error (*MAE*), both on simulated
3 and real sEMG signals.

4 **3.6 Materials and Methods**

5 A dataset of simulated and real sEMG signals was used to assess the
6 applicability of the MCDA-based approach for selecting the optimal number of
7 muscle synergies (N_{opt}) and to compare its performance to the principal *VAF*-based
8 approaches [23,24,43–46,112,118,120,125,126].
9

10 *1. Real Data*

11 Gait data acquired from 12 subjects were retrospectively analyzed to test the
12 performance of the four different approaches (Threshold *VAF*, Elbow *VAF*, Plateau
13 *VAF*, and MCDA-based approach) when applied to real sEMG signals. Subjects
14 were randomly selected from the PoliToBIOMed Lab (Politecnico di Torino, Turin,
15 Italy) database to include both healthy individuals and patients affected by
16 Parkinson’s Disease (PD). This non-homogeneous group of subjects was
17 specifically chosen to verify that the algorithm could work under different
18 conditions.

19 In particular, 6 out of 12 subjects were patients affected by PD (age: 55.5 ± 5.7
20 years, height: 171.1 ± 9.1 cm, weight: 76.7 ± 13.2 kg, UPDRS-III [96] during *OFF*
21 condition: 45.8 ± 13.1 , UPDRS-III during *ON* condition: 16.2 ± 6.5), while the
22 remaining 6 subjects were healthy controls (age: 52.3 ± 10.0 years, height: $171.3 \pm$
23 7.2 cm, weight: 73.0 ± 8.7 kg). PD patients were enrolled from those eligible for
24 Deep Brain Stimulation (DBS) at the Stereotactic and Functional Neurosurgery
25 Unit of the University of Turin (Italy). Moreover, none of the healthy subjects
26 reported lower limb injuries or had neurological or musculoskeletal disorders that
27 could compromise their gait performance.

28 Gait data were recorded through the STEP32 system (Medical Technology,
29 Italy)[46,90,91]. Surface electromyographic (sEMG) signals were acquired through
30 active probes (configuration: single differential, size: $19 \text{ mm} \times 17 \text{ mm} \times 7 \text{ mm}$, Ag-
31 disks diameter: 4 mm, interelectrode distance: 12 mm, gain: variable in the range
32 from 60 dB to 86 dB) placed over the following 12 muscles of the lower limb: right
33 and left Longissimus Dorsii (LD_R , LD_L), Tensor Fasciae Latae (TFL), Gluteus
34 Medius (GMD), Rectus Femoris (RF), Lateral Hamstring (LH), Medial Hamstring
35 (MH), Vastus Medialis (VM), Lateral Gastrocnemius (LGS), Peroneus Longus
36 (PL), Soleus (SOL) and Tibialis Anterior (TA). The dominant lower limb was
37 analyzed for healthy subjects, while the most affected limb was selected for
38 pathological subjects. **Figure 3.2** shows an example of sensor placement for a
39 representative subject of the healthy population.

40 Subjects were asked to perform a 5-minute walk at a self-selected speed, back
41 and forth on a 9-m straight walkway (**Figure 3.1**). The protocol conformed to the
42 Helsinki declaration on medical research involving human subjects and all the

1 volunteers signed a written informed consent to participate in the study before each
2 experimental session.

3 The reconstructed sEMG envelopes were computed, for each subject and each
4 number of muscle synergies in the range [1, 8], as the linear combination of the
5 weight vectors (W) and the activation coefficients (C) extracted from the real data
6 through the NNMF algorithm (**Equation 1.2**).

8 2. Simulated Data

9 The sEMG signal during cyclic contractions, such as those observed during
10 walking, can be modeled as the superimposition of two different contributions: (i)
11 the muscle activity and (ii) the background noise generated by the activity of the
12 neighboring muscles and electronics. Thus, the sEMG signal can be defined as a
13 cyclostationary process [86] and modeled as the superimposition of two different
14 stationary processes [69]:

- 15 i. Muscle activity (s) defined as a zero-mean Gaussian process with variance
16 σ_s^2 :

$$s(t) \in N(0, \sigma_s^2) \quad (3.8)$$

17 where $\sigma_s = 10^{\left(\frac{SNR}{20}\right)} \cdot 1 \mu V$.

- 18 ii. Background noise (n) defined as a Gaussian process with zero-mean and
19 variance σ_n^2 :

$$n(t) \in N(0, \sigma_n^2) \quad (3.9)$$

20 where $\sigma_n = 1 \mu V$.

21 Similarly to what was performed in previous studies[37,125,143], simulated
22 sEMG signals were generated starting from real data, defining *a priori* the optimal
23 number of muscle synergies (N_{opt}). More specifically, the muscle activity of twelve
24 muscles was simulated by time-windowing twelve Gaussian processes (s) by means
25 of different time supports. Each time support was defined as the reconstructed
26 sEMG envelope calculated from the muscle synergies extracted from the real
27 data[143], selecting *a priori* the number of muscle synergies (N_{opt}) in the range
28 between 4 and 6 synergies (i.e., numbers of muscle synergies usually selected
29 during walking in healthy subjects). Then, twelve background noise processes (n)
30 were superimposed to simulate the crosstalk of the neighboring muscles. Eight
31 different values of SNR were simulated ($SNR = 6, 10, 13, 16, 20, 23, 26, 30$ dB) to
32 assess the impact of SNR on the approaches' performance.

33 The simulated sEMG signals were then band-pass filtered through a 4th order
34 Butterworth digital filter with a lower cut-off frequency of 10 Hz and a higher cut-
35 off frequency of 450 Hz. The steps for the computation of the simulated sEMG
36 signals are represented in **Figure 3.8** for the Tibialis Anterior (TA) muscle. First,
37 muscle synergies (W and C) were extracted from the real data of a representative
38 PD subject of the sample population (**Figure 3.8a**). Second, the reconstruction of
39 the TA envelope was calculated as $W_{TA} \cdot C$ (**Figure 3.8b**). Third, a simulated sEMG
40 signal without additive noise is generated by time-windowing a Gaussian process
41 through the TA envelope (**Figure 3.8c**). Finally, noise is superimposed to the

1 previous signal to simulate the crosstalk from the neighboring muscles (**Figure**
 2 **3.8d**).

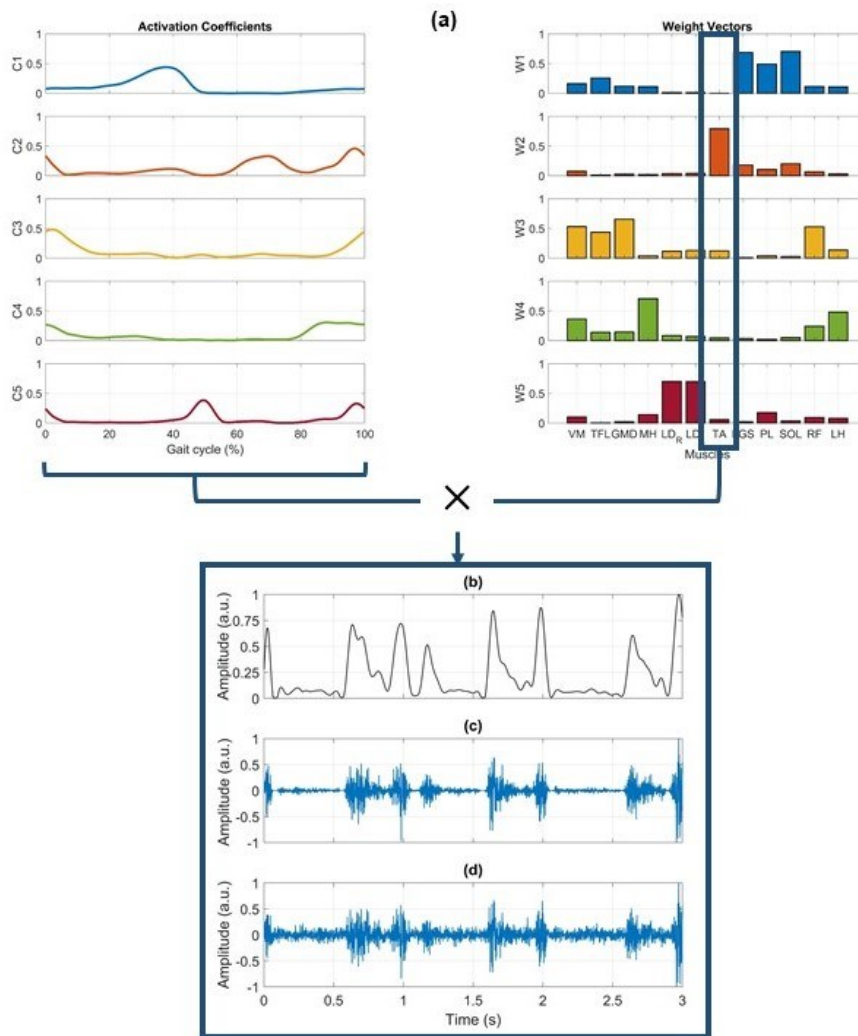


Figure 3.8 Example of the generation of simulated sEMG signal for the Tibialis Anterior (TA) muscle: the first step is (a) the extraction of muscle synergies (W and C) from the real data of a representative subject, the second is (b) the reconstruction of the TA envelope (obtained as $W_{TA} * C$). Then, (c) a simulated sEMG signal without additive noise is generated. Finally, noise is added to the previous signals. An example of a simulated sEMG signal with $SNR = 16$ dB is shown in (d).

3 Therefore, for each of the 12 muscles mentioned above, a dataset composed of
 4 288 simulated sets ($12 \text{ subjects} \times 8 \text{ SNRs} \times 3 N_{opt}$) was built.

5

6 3. Muscle Synergy Extraction

7 Before muscle synergy extraction, both the simulated and real sEMG signals
 8 were pre-processed to obtain the sEMG envelopes.

9 Considering the real data, sEMG signals were first time-segmented into gait
 10 cycles, according to the foot-floor contact sequence. Second, only the HFPS gait
 11 cycles, defined as the physiological sequence of Heel Contact (H), Flat Foot
 12 Contact (F), Push Off (P), and Swing (S), were considered, discarding those
 13 characterized by atypical gait cycles (i.e., non-HFPS gait cycles) [108]. Third only
 14 the gait cycles belonging to the rectilinear path (see **Figure 3.1**) were analyzed,

1 removing those corresponding to direction changes at the beginning and at the end
2 of the walkway (including deceleration before and acceleration after the U-turn)
3 [24]. Finally, each segmented gait cycle was time-normalized to 1000 samples [55],
4 and all gait cycles selected were concatenated in a single vector [24].

5 Both simulated and real sEMG signals were then high-pass filtered through an
6 8th order Butterworth digital filter with a cut-off frequency of 35 Hz, demeaned,
7 full-cycle rectified, and low-pass filtered at 12 Hz through a 5th order Butterworth
8 digital filter to obtain the sEMG envelopes. Each sEMG envelope was then
9 normalized in amplitude with respect to its global maximum to ensure an equally
10 weighted contribution of each acquired muscle in the muscle synergy extraction
11 process. The amplitude-normalized envelopes were divided into groups of 10
12 concatenated gait cycles (called subgroups) allowing for muscle synergy
13 assessment over the entire walk duration [24,112].

14 Muscle synergies were then extracted for each subgroup of 10 gait cycles
15 [24,46,112] by means of the NMF algorithm [32,41]. The NMF algorithm
16 describes the original sEMG signals as a linear combination of muscle synergy
17 weight vectors (W_k) and activation coefficients ($C(t)_k$), as detailed in (3.2). More
18 specifically, the weight vector W_k describes the time-independent contribution of
19 each observed muscle to the k -synergy, while the activation coefficient vector
20 $C(t)_k$ represents the time-dependent modulation of the muscles recruited in the k -
21 synergy. The MATLAB[®] function “*nmmf*” was used to apply the NMF algorithm,
22 setting the routine’s input parameters as follows: multiplicative update as
23 factorization algorithm, $1e^{-6}$ as function and search tolerance, 50 as the number of
24 factorization replicates, and 1000 as the maximum number of factorization
25 iterations.

26 The first algorithm initialization was performed differently for the weight
27 vector W_k and the activation coefficient vector $C(t)_k$. The C matrix was initialized
28 with values randomly selected from a uniform distribution in the range [0, 1]. To
29 improve the performance of the factorization algorithm and the accuracy in the
30 reconstruction of the original sEMG data, a sparseness constraint was imposed in
31 the initialization of the W matrix [37]. In particular, W matrix was first initialized
32 with values randomly chosen from a uniform distribution in the range [0, 0.05],
33 then one random element of each W_k vector was set to a value selected from a
34 uniform distribution in the range [0.7, 0.8]. Therefore, only one muscle for each k -
35 synergy has a significant contribution, obtaining an extremely sparse NMF
36 initialization [37].

37 To sort the muscle synergies in the same order for each subgroup, the k -means
38 algorithm was applied to the W matrix [22]. Clustering was performed by means of
39 the MATLAB[®] routine “*kmeans*” using the following input parameters: N as
40 number of k -means clusters, 1000 as maximum number of iterations, 15 as number
41 of replicates, and cosine as distance metric. The activation coefficients matrix C
42 was ordered accordingly.

To explore different solutions of the NMF algorithm, the factorization process was run many times on the same gait data, changing the number of muscle synergies (N) in the range [1, 8].

4. Optimal Number Selection: VAF-base Methods

Variance Accounted For (VAF) is widely used in literature to compute the reconstruction accuracy of a factorization algorithm [23,24,43–46,112,118,120,125,126]. VAF is defined as the uncentered Pearson's correlation coefficient expressed in percentage, as it follows:

$$VAF = \left(1 - \frac{\sum_{k=1}^m (M_k - M_k^R)^2}{\sum_{k=1}^m M_k^2} \right) \cdot 100 \quad (3.10)$$

where m represents the number of muscles observed, $\sum_{k=1}^m (M_k - M_k^R)^2$ describes the sum of the squared errors between the original (M_k) and reconstructed (M_k^R) sEMG signals of the k -synergy, and $\sum_{k=1}^m M_k^2$ represents the total sum of squared M_k values with respect to zero. The optimal number of muscle synergies (N_{opt}) is often selected by applying different approaches on the plot of the VAF versus the number of muscle synergies (called VAF curve).

Figure 3.8 shows an example of VAF curves extracted from (**Figure 3.8A**) real and (**Figure 3.8B**) simulated sEMG signals.

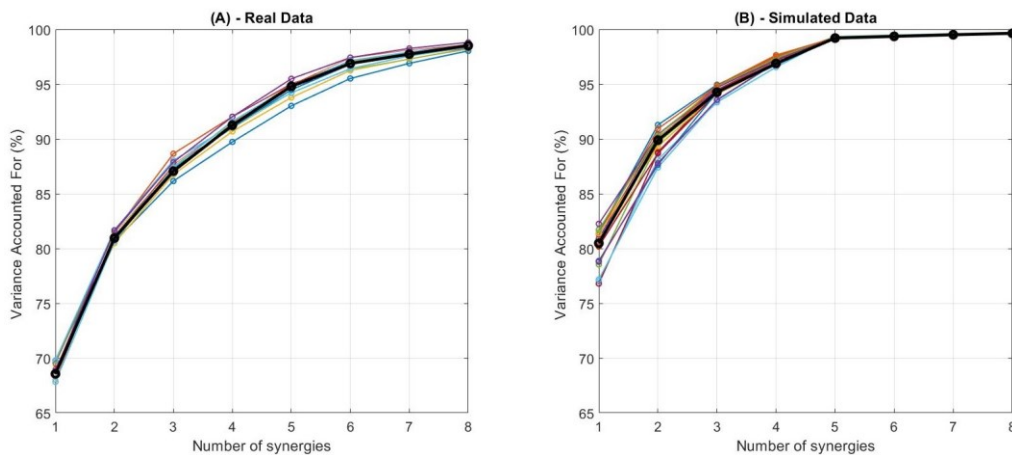


Figure 3.8 Example of VAF curves extracted from (A) real and (B) simulated sEMG signals. For each tested number of synergies, the Variance Accounted For (VAF) measures the correlation between the original and the reconstructed sEMG data. Each colored line represents a VAF curve extracted from a single 10-gait-cycle subgroup, while black lines represent the average VAF curve computed over subgroups.

In this chapter, the performance of the MCDA-based approach is quantitatively compared with respect to three standard VAF -based approaches: (i) Threshold VAF (T_{VAF}), (ii) Elbow VAF (E_{VAF}), and (iii) Plateau VAF (P_{VAF}) methods.

- i. *Threshold VAF* (T_{VAF}): this approach defines N_{opt} as the first number of synergies granting a VAF value above a heuristically defined cut-off threshold. This threshold is commonly set at 90% VAF [23,24,43,44,46,112,118,120,125,126].

- 1
- 2 *ii. Elbow VAF (E_{VAF}):* this approach defines N_{opt} as the number of synergies
- 3 at which the *VAF* curve achieves the highest curvature (“elbow”) [32].
- 4 More specifically, slope changes are detected by computing the curvature
- 5 for every three consecutive *VAF* curve points (i.e., the first curvature is
- 6 computed considering $N = 1, 2,$ and 3 ; the second curvature considering
- 7 $N = 2, 3, 4,$ etc.). Finally, the optimal number of muscle synergies is
- 8 defined as the number of synergies in correspondence of the highest
- 9 curvature among the computed ones.
- 10
- 11 *iii. Plateau VAF (P_{VAF}):* this approach defines N_{opt} as the number of
- 12 synergies beyond which the *VAF* curve reaches a plateau. More
- 13 specifically, the *VAF* curve is iteratively fitted by means of the least-
- 14 squares linear regression, first considering all the *VAF* curve points, and
- 15 then moving the fit window toward the right side of the *VAF* curve by
- 16 excluding the smallest number of muscle synergies. The first fit iteration
- 17 considers all the *VAF* curve points (e.g., N ranges from 1 to 8) for the
- 18 linear regression, the second fit iteration considers only the last seven
- 19 *VAF* curve points (N ranges from 2 to 8), and so on until only the last two
- 20 *VAF* curve points are considered. For each fit, the Mean Squared Error
- 21 (MSE) was computed and compared with respect to a heuristically
- 22 defined threshold, that in this study was set equal to 10^{-5} [16,47]. Finally,
- 23 the optimal number of muscle synergies is defined as the first number of
- 24 synergies at which the linear fit produces a $MSE \leq 10^{-5}$.
- 25

26 5. *Optimal Number Selection: MCDA-based Method*

27 Multi-Criteria Decision Analysis (MCDA) was developed in the field of

28 operational research and consists of a family of methods that aim at selecting the

29 optimal solution (among all the possible alternatives) by explicitly taking into

30 account multiple criteria. To find a reliable and systematic solution to the muscle

31 synergy number problem, moving from single criterion approaches (e.g., *VAF*- or

32 R^2 -based approaches) to a multicriteria approach, an MCDA model is herein

33 proposed to obtain a systematic and reliable approach to criteria modeling and

34 alternatives evaluation.

35 To define the best MCDA model, the following four main steps were

36 considered:

- 37 *i.* The definition of the alternatives set
- 38 *ii.* The definition of the evaluation criteria
- 39 *iii.* The choice and definition of the multicriteria aggregation procedure

40

41 *Definition of the alternatives set*

42 The alternatives considered in the MCDA model were defined as the different

43 ranks of the factorization process (r), i.e., all the possible solutions explored by the

44 NNMF algorithm. Since the factorization is usually repeated several times on the

1 same data, changing the factorization rank (i.e., the number of muscle synergies N)
 2 between 1 and 8, eight different alternatives were considered in the decision-
 3 making process. Thus, each alternative corresponds to a different number of muscle
 4 synergies.

5

6 *Definition of the evaluation criteria*

7 The main objective of the MCDA approach was to find a robust and repeatable
 8 approach for the selection for the optimal number of muscle synergies, moving from
 9 single criterion approaches, such as VAF - or R^2 -base approaches, to a multicriteria
 10 approach. To define the evaluation criteria, this main objective was divided into
 11 four sub-objectives:

- 12 *i.* Maximize the reconstruction accuracy of the factorization algorithm
- 13 *ii.* Maximize the muscle synergy consistency
- 14 *iii.* Minimize the muscle synergy redundancy
- 15 *iv.* Maximize the factorization model robustness.

16 As shown in **Table 3.4**, these four sub-objectives were translated into six
 17 evaluation criteria to measure the degree to which an alternative attains each sub-
 18 objective.

TABLE 3.4
 OBJECTIVE, SUB-OBJECTIVES, AND EVALUATION CRITERIA

<i>SUB-OBJECTIVES</i>	<i>EVALUATION CRITERIA (CRITERION WEIGHT)</i>
<i>Maximize the reconstruction accuracy of the factorization algorithm</i>	Evaluation of Variance Accounted For (10%)
<i>Maximize the muscle synergy consistency</i>	Evaluation of Weight Vector Intra-Cluster Consistency (18%) Evaluation of Activation Coefficient Intra-Cluster Consistency (18%)
<i>Minimize the muscle synergy redundancy</i>	Evaluation of Weight Vector Inter-Cluster Similarity (17%) Evaluation of Activation Coefficient Inter-Cluster Similarity (17%)
<i>Maximize the factorization model robustness</i>	Evaluation of Cross-Variance Accounted For (20%)

19 In the following, the six evaluation criteria are detailed:

- 20 *i.* *Evaluation of Variance Accounted For (VAF)*: this criterion quantifies the
 21 reconstruction accuracy of the factorization results and it is defined as the
 22 uncentered Pearson’s correlation coefficient, expressed in percentage, as
 23 defined in **Equation (3.10)**. A high value of VAF (high accuracy) is
 24 desired to obtain an accurate reconstruction accuracy.
- 25
- 26 *ii.* *Evaluation of the Weight Vector and Activation Coefficient Intra-Cluster*
 27 *Consistency (ICC)*: these criteria quantify the consistency of weight

vectors (ICC_W) and activation coefficients (ICC_C) over time (i.e., over the different 10-gait-cycle subgroups) by computing the cosine similarity (CS) [30] of the previously sorted weight vectors W_k (3.11) and activation coefficient vectors $C(t)_k$ (3.12), separately. More specifically, CS was computed between the weight vector (or activation coefficient) belonging to the i -th subgroup and the cluster centroid, defined as the average over all the 10-gait-cycle subgroups, of the k -synergy as described in (3.4) and (3.12):

$$CS_{W,k}^i = \left(\frac{W_k^i \cdot \bar{W}_k}{\|W_k^i\| \|\bar{W}_k\|} \right) \cdot 100 \quad (3.11)$$

$$CS_{C,k}^i = \left(\frac{C_k^i \cdot \bar{C}_k}{\|C_k^i\| \|\bar{C}_k\|} \right) \cdot 100 \quad (3.12)$$

where $CS_{W,k}^i$ and $CS_{C,k}^i$ represent the CS computed between the weight vectors W_k of the i -th subgroup and the cluster centroid of the k -synergy and between the activation coefficients $C(t)_k$ of the i -th subgroup and the cluster centroid of the k -synergy, respectively. The CS values range between 0 (meaning no similarity) and 100 (meaning complete similarity). Then, ICC was defined, separately for each k -synergy, as the average of the $CS_{W,k}$ and $CS_{C,k}$ values for the weight vectors and the activation coefficients, separately. Notice that the average operator is always applied across 10-gait-cycle subgroups. Finally, the “ min ” function is used to select the most variable muscle synergy (“worst” condition), obtaining a single ICC value for each number of muscle synergies N . The ICC value ranges from 0 (i.e., completely different muscle synergy across subgroups) to 100 (i.e., perfectly repeatable muscle synergy among the different subgroups). A high value of ICC (high consistency) is desired to be able to assess repeatable muscle synergies.

iii. *Evaluation of the Weight Vector and Activation Coefficient Inter-Cluster Similarity (ICS)*: these criteria quantify the level of similarity of weight vectors (ICC_W) and activation coefficients (ICC_C) belonging to different muscle synergies by computing the cosine similarity (CS) [30]. For each number of synergies ($N > 1$), and for each synergy i (with $i = 1, \dots, N$), CS was computed between the cluster centroids (defined as the average over all the 10-gait-cycle subgroups) of different muscle synergies for the weight vectors and the activation coefficients, separately. Finally, the “ max ” function is used to select the most similar muscle synergy (“worst” condition), obtaining a single ICS -value for each number of muscle synergies N as detailed in (3.13) and (3.14):

$$ICS_W = \max \left(\frac{\bar{W}^i \cdot \bar{W}^k}{\|\bar{W}^i\| \|\bar{W}^k\|} \right) \quad (3.13)$$

$$ICS_C = \max \left(\frac{\bar{C}^i \cdot \bar{C}^k}{\|\bar{C}^i\| \|\bar{C}^k\|} \right) \quad (3.14)$$

where \bar{W}^i and \bar{W}^k represent the average weight vectors (cluster centroid) computed across subgroups for the i - and k -synergy, respectively. Instead, \bar{C}^i and \bar{C}^k represent the average activation coefficient vectors (cluster centroid) computed across subgroups for the i - and k -synergy, respectively. ICS values can range from 0 to 1: $ICS = 0$ indicates a complete dissimilarity, while $ICS = 1$ indicates a complete similarity between \bar{W}^i and \bar{W}^k or between \bar{C}^i and \bar{C}^k . A small value of ICS (low similarity) is desired to avoid redundant information.

iv. *Evaluation of the Muscle Synergy Robustness*: this criterion quantifies the robustness of the factorization model over different 10-gait-cycle subgroup through the Cross-Variance Accounted For (*CrossVAF*) [111], defined as it follows:

$$CrossVAF^{i,j} = \left(1 - \frac{\sum_{k=1}^m (M_k^i - M_k^{R,j})^2}{\sum_{k=1}^m (M_k^i)^2} \right) \cdot 100 \quad (3.15)$$

where M_k^i and $M_k^{R,j}$ represent the original and the reconstructed sEMG signals of the k -muscle for the i - and j -th 10-gait-cycle subgroup, respectively. This parameter is used to assess how well the muscle synergies extracted for the i -th subgroup reconstruct the sEMG signals that belong to the j -th subgroup. The average *CrossVAF* value was then computed over all the possible couples of 10-gait-cycle subgroups. The average *CrossVAF* can assume values ranging from 0% (i.e., low correlation between the original and the reconstructed sEMG signals) to 100% (i.e., a high correlation between the original and the reconstructed sEMG signals). A high value of *CrossVAF* (high robustness) is desired to assess a robust factorization model.

Selection and Definition of the Multicriteria Aggregation Procedure

Although the problem addressed in the present chapter can be described as a selection problem, herein it was addressed as a ranking problem, in which the alternatives are ranked from the best to the worst with the possibility of *ex-aequo* and/or incomparability between alternatives.

The multicriteria aggregation procedure ELECTRE III (ELimination Et Choix Traduisant la REalité or ELimination and Choice Expressing the REality) [144–147] was used to rank the alternatives from the best to the worst based on their performance on the six evaluation criteria previously defined (**Table 3.4**). More specifically, ELECTRE-III [144–147] is a partially compensatory ordinal method specifically developed for ranking problems. Considering the decision-making process addressed in this chapter, it is desired to minimize the compensation effect

1 and penalize unbalanced alternatives, i.e., alternatives with a mix of very good and
 2 very bad evaluations. ELECTRE III is based on 4 basic outranking relations
 3 between alternatives [147]:

- 4 • *Indifference*: it represents a condition where there are clear and positive
 5 reasons that justify the equivalence between the two alternatives A and B
- 6 • *Strict preference*: it represents a condition where there are clear and positive
 7 reasons in favor of one of the two alternatives
- 8 • *Weak preference*: it represents a condition where there are clear and positive
 9 reasons in favor of one of the two alternatives, but they are insufficient to
 10 justify a *strict preference* relation
- 11 • *Incomparability*: it represents a condition where there are no clear and
 12 positive reasons that would justify any of the above-mentioned relations.

13 The outranking of an alternative A by an alternative B (defined by assigning to
 14 the pair of alternatives A and B one and only one of the four basic outranking
 15 relations above defined) is determined by concordance and discordance tests. In
 16 particular, for each pair of alternatives, the concordance and discordance indexes
 17 are computed on the basis of the evaluation matrix, i.e., the performance of each
 18 alternative on each criterion. **Table 3.5** represents an example of an evaluation
 19 matrix, used as input of the ELECTRE III model, computed from real sEMG signals
 20 of a PD subject of the sample population during a walking task.

TABLE 3.5
 EVALUATION MATRIX OF A PD PATIENT OF THE SAMPLE POPULATION.

Alternatives	Evaluation Criteria					
	<i>VAF</i> (%)	<i>ICC_w</i> (a.u.)	<i>ICC_c</i> (a.u.)	<i>ICS_w</i> (a.u.)	<i>ICS_c</i> (a.u.)	<i>CrossVAF</i> (%)
1 Synergy	76.37	1.00	1.00	NaN ¹	NaN ¹	72.14
2 Synergies	85.42	1.00	0.99	0.42	0.61	78.12
3 Synergies	89.95	0.99	0.99	0.38	0.69	79.16
4 Synergies	93.55	0.99	0.99	0.35	0.70	80.61
5 Synergies	95.62	0.98	0.99	0.38	0.78	80.83
6 Synergies	97.08	0.99	0.99	0.34	0.93	80.17
7 Synergies	98.00	0.75	0.89	0.36	0.92	80.02
8 Synergies	98.68	0.94	0.98	0.53	0.94	79.73

¹ Inter-cluster similarity cannot be computed for a number of synergy $N = 1$.

21 The concordance index of a pair of alternatives A and B is used to validate the
 22 assertion that “ A outranks B ” and to measure the credibility of this assertion. From
 23 a mathematical point of view, it is defined as the weighted sum of the criteria on
 24 which A is evaluated at least as good as B . In particular, the importance weight for
 25 each criterion were defined for each of the six criteria using the revised Simon’s
 26 procedure [148] as shown in brackets in **Table 3.4**.

27 The discordance index for the alternatives A and B on criterion j , instead, is
 28 used to assess the degree to which criterion j goes against the assertion that “ A
 29 outranks B ”. Finally, a ranking of the alternatives from the best one to the worst

1 one was obtained from the strict and weak outranking relations. **Figure 3.9** shows
2 an example of outranking graph obtained considering real sEMG signals of a PD
3 subject of the sample population during walking.

4 In the outranking graph, when two alternatives are represented at the same rank,
5 it means that they are incomparable (e.g., “6 Synergies” and “8 Synergies”
6 alternatives in **Figure 3.9**) or equivalent.

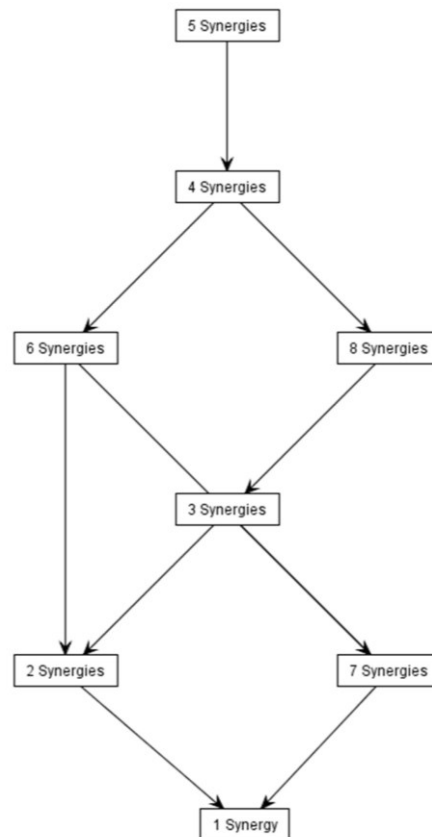


Figure 3.9 Outranking graph extracted through the ELECTRE III methods considering a PD subject of the sample population. Alternatives are ranked from the best one (top) to the worst (bottom) one. The “5 Synergies” alternative was selected as the best alternative. Therefore, the optimal number of muscle synergies was defined equal to 5.

7 ELECTRE-III model was implemented and validated using MATLAB[®] release
8 2020b (The MathWorks Inc., Natick, MA, USA), modifying the MCDA-ULaval
9 open-source software developed by professor Irène Abi-Zeid of Laval University
10 of Québec, Canada. The original MCDA-ULaval open-source software, developed
11 in Java, can be freely found online at the following address:
12 <https://cersvr1.fsa.ulaval.ca/mcda-ulaval/?q=en>.

13

14 6. Performance Evaluation

15 The optimal numbers of muscle synergies computed through the four different
16 approaches (T_{VAF} , E_{VAF} , P_{VAF} , and ELECTRE III) were quantitatively compared in
17 terms of (i) estimation bias and (ii) Mean Absolute Error (MAE), both for the

1 simulated and real datasets. The *MAE* was used to quantify how far a given
 2 approach deviates from the ground truth (i.e., the correct number of muscle
 3 synergies), while the estimation bias was used to know whether an approach goes
 4 wrong by defect or excess. More specifically, the indexes were defined as it follows:

$$bias = \frac{1}{n} \sum_{i=1}^n (\hat{N}_i - N_{opt,i}) \quad (3.16)$$

$$MAE = \frac{1}{n} \sum_{i=1}^n (|\hat{N}_i - N_{opt,i}|) \quad (3.17)$$

5 where $N_{opt,i}$ is the optimal number of muscle synergies for the i -th subject (ground
 6 truth) and \hat{N}_i is the estimated optimal number of muscle synergies identified by one
 7 of the tested approaches.

8 To be able to know which number of muscle synergies should be considered
 9 optimal (N_{opt}), ground truth was defined for both the simulated and real signals.
 10 Considering the simulated data, the ground truth was known *a-priori* based on the
 11 sEMG signal simulation process. Considering the real data, instead, the judgment
 12 of two expert operators (Mr. Marco Ghislieri and Prof. Valentina Agostini) was
 13 used to define the ground truth. Their judgment was performed blind to the details
 14 of the ELECTRE III algorithm as well as to the results of the various methods
 15 tested. For each real signal, they analyzed the muscle synergy plots considering
 16 different numbers of muscle synergies N (in the range between 1 and 8 muscle
 17 synergies) and they chose - separately - the number they considered as correct
 18 (N_{opt}), based on their knowledge of motor control strategies, muscle synergy
 19 analysis, and gait biomechanics. More specifically, the operators used basically the
 20 same criteria to identify the correct number of muscle synergies. They considered
 21 the consistency in time of the motor control strategies (muscle synergies are
 22 expected not to change too much over task duration) and the biomechanical task
 23 associated to each muscle synergy by assessing which number of synergies better
 24 represent a set of well-distinguished functions and avoiding the presence of
 25 redundant information (i.e., avoiding having two identical or “too similar” muscle
 26 synergies). However, expert judgment is subjective, at least to some extent.
 27 Cohen’s kappa statistic [149] was used to compute the degree of agreement between
 28 raters. In case of disagreement, the two expert operators discussed the discordant
 29 cases to achieve a common ground truth.

30 7 *Effect of SNR on Muscle Synergy Number Selection*

31 To assess the effect of the *SNR* on the performance of the four approaches, the
 32 performance parameters described above were computed on the simulated dataset,
 33 separately for each of the eight *SNR* values ($SNR = 6, 10, 13, 16, 20, 23, 26, 30$
 34 dB).

35 8 *Statistical Analysis*

1 One-way repeated-measures analysis of variance (*ANOVA*) followed by *post-*
2 *hoc* analysis with Tukey's adjustment for multiple comparisons was performed to
3 assess statistically significant differences in the performance of the four tested
4 approaches for selecting the optimal number of muscle synergies, considering both
5 simulated and real data. The effect size of the statistically significant differences
6 was calculated through the Hedges' *g* including the correction for small sample sizes
7 [113]. The significance level (α) was set equal to 0.05.

8 **3.7 Results**

9 First, the results supporting the applicability of the ELECTRE III approach for
10 selecting the optimal number of muscle synergies are presented, considering
11 simulated sEMG signals. Second the performance of the four approaches (T_{VAF} ,
12 E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III) on simulated sEMG signals
13 are compared, highlighting the effect of the *SNR*. Finally, ELECTRE III is
14 validated on real data.

15 *Simulated Data*

16 The performance of the four different approaches were assessed and compared
17 in terms of (i) estimation bias and (ii) Mean Absolute Error (*MAE*):

- 19 i. *Estimation bias*: On average, an estimation bias of -2.0 ± 1.6 muscle
20 synergies, -0.4 ± 1.1 muscle synergies, 1.7 ± 0.7 muscle synergies, and -0.6
21 ± 0.9 muscle synergies was found on the simulated dataset, considering the
22 T_{VAF} , the E_{VAF} , the P_{VAF} , and the MCDA approach, respectively. One-way
23 ANOVA followed by *post-hoc* analysis revealed significant differences
24 between each pair of approaches ($p < 0.001$)
- 25 ii. *Mean Absolute Error*: On average, a *MAE* of 2.0 ± 1.6 muscle synergies,
26 0.4 ± 1.1 muscle synergies, 1.7 ± 0.7 muscle synergies, and 0.8 ± 0.8 muscle
27 synergies were found on the simulated dataset, considering the T_{VAF} , E_{VAF} ,
28 P_{VAF} , and the ELECTRE III approach, respectively. One-way ANOVA
29 followed by *post-hoc* analysis revealed significant differences between each
30 pair of approaches ($p < 0.001$)

1 **Figure 3.10** compares the performance of the four approaches tested (T_{VAF} ,
 2 E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III), based on the two parameters
 3 introduced above. The average values and standard errors of the parameters were
 4 estimated on the simulated dataset.

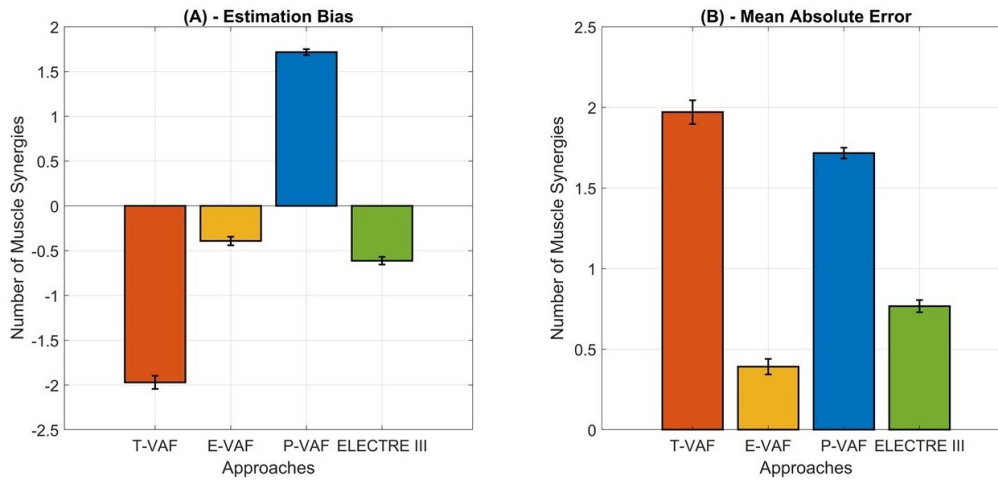


Figure 3.10 Comparison of the performance of the four tested approaches (T_{VAF} , E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III approach) estimated considering (A) the estimation bias and (B) the Mean Average Error (*MAE*). Average values are represented with the indication of the standard errors. Considering both parameters, statistically significant differences were detected between each pair of approaches.

5

6 *Effect of SNR on Muscle Synergy Number Selection*

7 The effect of the *SNR* on the approaches' performance was assessed by
 8 computing the "performance" parameters (estimation bias and mean absolute
 9 error), separately for each simulated *SNR*-value.

10 **Figure 3.11** represents, for each tested approach, the average values (and
 11 standard errors) of estimation bias (**Figure 3.11A**) and *MAE* (**Figure 3.11B**), for
 12 each simulated *SNR*-value.

13 For both parameters, threshold-independent approaches (E_{VAF} and ELECTRE
 14 III) revealed a higher performance across the different *SNR* values, suggesting a
 15 lower effect of *SNR* on the selection of the optimal number of muscle synergies
 16 with respect to the other two threshold-dependent approaches (T_{VAF} and P_{VAF}).
 17 Even if the E_{VAF} approach revealed the highest performance in terms of estimation
 18 bias and *MAE*, the approach less affected by the *SNR* was the ELECTRE III,
 19 revealing similar performance across the different values of *SNR*. The approach
 20 more affected by the *SNR*, instead, was the T_{VAF} , with an evident decrease in the
 21 performance for simulated sEMG signals with *SNR*-values lower than 16 dB. On
 22 average, a significant decrease in the performance of all the tested approaches was
 23 detected for simulated sEMG signals with *SNR* lower than 16 dB.

24 *Real Data*

25 The performance of the four different approaches on real sEMG signals were
 26 assessed with respect to the ground truth by considering the same two parameters
 27 described for the simulated dataset. The inter-rater agreement, computed by means
 28 of Cohen's kappa, was equal to 0.5, suggesting a moderate agreement between the

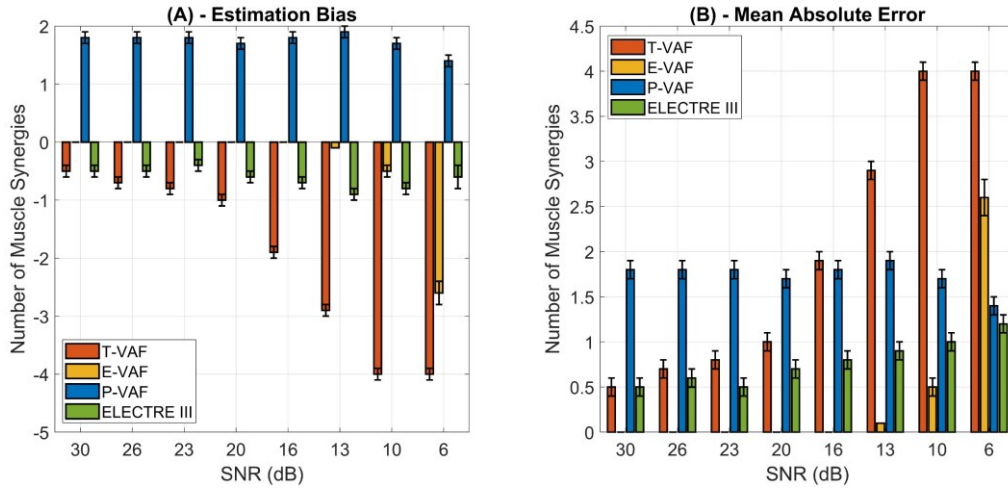


Figure 3.11 Comparison of the performance of the four tested approaches (T_{VAF} , E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III approach) estimated considering (A) the estimation bias and (B) the Mean Average Error (MAE). Average values are represented with the indication of the standard errors.

1 two expert operators. In the following, the results obtained considering the two
 2 above-mentioned parameters are detailed:

- 3 *i. Estimation bias:* On average, an estimation bias of 0.9 ± 0.3 muscle
 4 synergies, -0.8 ± 0.2 muscle synergies, -0.8 ± 0.2 muscle synergies, and
 5 0.0 ± 0.3 muscle synergies was found on the real dataset, considering the
 6 T_{VAF} , the E_{VAF} , the P_{VAF} , and the ELECTRE III approach, respectively.
 7 One-way ANOVA followed by *post-hoc* analysis revealed statistically
 8 significant differences between the T_{VAF} and E_{VAF} approaches ($p = 0.003$,
 9 $g = 5.88$) and between the T_{VAF} and P_{VAF} approaches ($p = 0.001$, $g = 6.25$),
 10 while no differences were found between the ELECTRE III approach and
 11 the VAF-based approaches.
- 12 *ii. Mean Absolute Error:* An average MEA of 1.1 ± 0.3 muscle
 13 synergies, 0.8 ± 0.2 muscle synergies, 0.8 ± 0.2 muscle synergies, and 0.5 ± 0.3
 14 muscle synergies was computed from the real dataset, considering the
 15 T_{VAF} , the E_{VAF} , the P_{VAF} , and ELECTRE III approach, respectively. One-
 16 way ANOVA followed by *post-hoc* analysis revealed no statistically
 17 significant differences between the four tested approaches ($p > 0.05$).

18 Considering the real sEMG data, **Figure 3.12** compares the performance of the
 19 four approaches (T_{VAF} , E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III
 20 approach), considering the estimation bias (**Figure 3.12A**) and MAE (**Figure**
 21 **3.12B**) parameters previously defined. The average values and standard errors of
 22 the computed parameters are reported, as well as asterisks to highlight statistical
 23 differences ($p < 0.05$) between each pair of approaches.

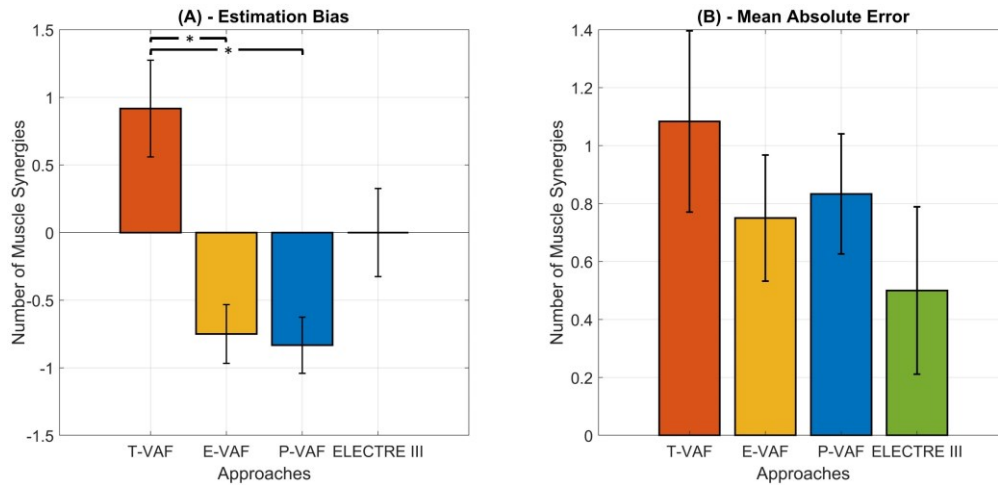


Figure 3.12 Comparison of the performance of the four tested approaches (T_{VAF} , E_{VAF} , P_{VAF} , and the newly introduced ELECTRE III approach) estimated considering (A) the estimation bias and (B) the Mean Average Error (MAE). Average values are represented with the indication of the standard errors. Statistically significant differences ($p < 0.05$) are highlighted by means of asterisks (*).

1 3.8 Discussion and Conclusions

2 In literature, the muscle synergy theory has been proposed to assess the modular
3 organization of the CNS during different movement tasks, such as walking and
4 running, in both physiological and pathological populations. In the last years,
5 several studies demonstrated that the optimal number of muscle synergies can be a
6 meaningful feature for the assessment of the motor control complexity in patients
7 affected by neuromuscular or neurodegenerative diseases, such as brain injury
8 [124] and Parkinson’s disease [45]. According to these studies, the number of
9 muscle synergies and their composition could be correlated with the motor control
10 capacity and its reduction in pathological conditions [33]. Despite the growing
11 interest in the assessment of the optimal number of muscle synergies, there is a lack
12 of standardized methods for the precise identification of the number of synergies,
13 making comparisons between studies and cohorts difficult. The number of muscle
14 synergies characterizing a given movement, indeed, significantly varies within and
15 across studies, even for healthy individuals. Hence, as muscle synergy analysis is
16 growing in popularity in motor control neuroscience and rehabilitation engineering,
17 how to improve and, hopefully, to standardize the selection of the optimal number
18 of muscle synergies are timely and important questions.

19 The approach currently accepted and most widely used by researchers for the
20 identification of the optimal number of muscle synergies is based on the application
21 of an arbitrary cut-off threshold based on the Variance Accounted For (VAF)
22 [23,24,43–46,112,118,120,125,126], defined as the uncentered Pearson’s
23 correlation between the original and the reconstructed sEMG signals. More
24 specifically, the optimal number of muscle synergies is defined as the first number
25 of synergies that produces a VAF value equal to or higher than an arbitrary cut-off
26 threshold. This cut-off threshold is usually set at 90%
27 [23,24,43,44,46,112,118,120,125,126], but several other studies considered

1 threshold values that range from 80% to 95% [45]. Several are the disadvantages
2 of this *VAF*-based approach. First, the discordance over the *VAF* threshold values
3 significantly intensifies the confusion over the selection of the optimal number of
4 muscle synergies and reduces results comparability between studies. Second, even
5 small variations in the cut-off threshold value could significantly change the final
6 results in terms of number of muscle synergies. Third, there is no evidence why the
7 same cut-off threshold value should be applied for all individuals and cohorts.
8 Finally, since the *VAF* is defined as the uncentered Pearson's correlation, different
9 values of mean activity computed from different muscles may strongly affect the
10 selection of the optimal number of muscle synergies.

11 To the best of the author's knowledge, few other studies have proposed
12 alternatives to the *VAF* threshold approach. In the study by Kim *et al.*[130], a novel
13 approach was proposed to enhance precision and reliability in synergy number
14 identification based on *k*-means clustering and intra-cluster correlation analysis.
15 However, this approach was tested considering sEMG data acquired only from
16 eight leg muscles of a small sample population (9 healthy subjects) during treadmill
17 walking. Delis *et al.*[131], instead, developed a more "physiological" approach,
18 introducing a task decoding-based metric during an arm pointing task.

19 In this chapter, a novel approach for choosing the optimal number of muscle
20 synergies is proposed and validated on both simulated and real sEMG data. The
21 novel ELECTRE III approach has potential advantages with respect to the
22 commonly used *VAF*-based threshold methods, such as the use of minimization and
23 maximization approaches rather than an arbitrary threshold and the applicability to
24 a variety of different motor tasks. It relies on six parameters directly computed from
25 muscle synergy results: the reconstruction accuracy, weight vector and activation
26 coefficient intra-cluster consistency, weight vector and activation coefficient inter-
27 cluster similarity, and muscle synergy robustness. The applicability of the proposed
28 approach was first checked considering simulated sEMG data. Then, its
29 performance was quantitatively compared against three widely used *VAF*-based
30 approaches (T_{VAF} , E_{VAF} , and P_{VAF}), in terms of estimation bias (*Bias*) and Mean
31 Absolute Error (*MAE*). Finally, ELECTRE III is validated on real data.

32 Considering the simulated sEMG signals, the two threshold-independent
33 approaches (E_{VAF} and ELECTRE III) outperform the other two threshold-
34 dependent methods (T_{VAF} and P_{VAF}) in terms of both estimation bias and mean
35 absolute error. In particular, E_{VAF} outperform all the other approaches, revealing
36 high performance considering both the estimation bias and the mean absolute error.
37 These results confirm the hypothesis that the threshold-dependent approaches could
38 be more affected by even small changes in the sEMG characteristics that could
39 modify the *VAF* curve, such as the SNR, compared to the threshold-independent
40 approaches. Indeed, considering the effect of the SNR on the approaches'
41 performance, the most affected ones were, as expected, the threshold-dependent
42 approaches (i.e., T_{VAF} and P_{VAF}), revealing a significant decrease in the
43 performance for simulated sEMG signals with SNR-values lower than 16 dB.
44 ELECTRE III approach, instead, revealed similar performance across the tested

1 *VAF* values, suggesting to be less affected by the *SNR* with respect to the other
2 three approaches.

3 Considering real sEMG signals, results similar to those of the other three
4 methods were achieved, applying the ELECTRE III approach, in terms of
5 estimation bias and mean absolute error. However, it could be noticed that there are
6 some clear differences in the performance of the threshold-dependent approaches
7 between the simulated and the real sEMG signals. In particular, when considering
8 the simulated data, T_{VAF} and P_{VAF} approaches underestimate and overestimate the
9 number of muscle synergies, respectively, while a complementary behavior is
10 assessed considering the real data. This is probably due to the nature of the
11 simulated dataset: the simulated signals may be less complex to factorize and, then,
12 the *VAF* assumes higher values already at small numbers of synergies. Indeed, the
13 estimation bias values show that the T_{VAF} always goes wrong by defect.

14 Therefore, the results presented in this chapter demonstrated the applicability
15 of the proposed *MCDA*-based approach for the selection of the optimal number of
16 muscle synergies, showing equal or higher performance with respect to the other
17 three single-criterion methods. However, even if the ELECTRE III revealed lower
18 performance with respect to the E_{VAF} approach, the former can represent a valid
19 alternative for the selection of the muscle synergy number, moving from single
20 criterion approaches, such as *VAF*- or R^2 -base approaches, to a multicriteria
21 approach.

22 The *MCDA*-based approach has several advantages with respect to the
23 commonly used *VAF*-based methods, such as the use of maximization and
24 minimization approaches rather than arbitrary thresholds and its applicability to
25 sEMG signals acquired from different sets of muscles and during different motor
26 tasks (contrary to the study by Delis *et al.* [131]). Instead, one of the limitations of
27 the proposed approach is that it requires several 10-gait-cycle subgroups to extract
28 the evaluation matrix. This requirement does not limit the feasibility and
29 applicability of the proposed methodology to pathological populations. Indeed, gait
30 analysis is commonly used to quantitatively assess patients' locomotion
31 performance only in those patients able to independently walk, for some minutes,
32 without walking aids or external supports. In the past, several studies demonstrated
33 the feasibility of gait data acquisition, during recording sessions lasting 3 minutes,
34 in patients suffering from different neurological conditions, e.g., normal pressure
35 hydrocephalus[89], mild ataxia[116], and cerebral palsy[117].

36 The author is aware that the number of muscle synergies can be also strongly
37 influenced by other steps in the muscle synergy extraction process
38 [23,36,38,39,44,49–53], such as the sEMG collection and pre-processing (e.g., low-
39 pass filtering techniques) [49–51,53], the number and choice of muscles acquired
40 [23], and sEMG data preparation before factorization (e.g., averaging or
41 concatenating sEMG epochs) [38]. However, the focus of this chapter is on
42 developing an approach that can be applied subsequent to a factorization algorithm
43 to select the optimal number of muscle synergies and not on evaluating the effect
44 of different pre-processing techniques on the identification of the synergy number.

1 Future developments of the proposed approach will include the assessment of
2 the MCDA-based approach on larger and different cohorts (for age or pathological
3 condition) and during different motor tasks, the optimization of the approach by
4 tuning the importance weights of each evaluation criterion, the evaluation of
5 different sets of evaluation criteria, and the evaluation of the effect of the number
6 of 10-gait-cycle subgroup per subject on the ELECTRE III performance.

7 In conclusion, a multicriteria approach for the identification of the optimal
8 number of muscle synergies was proposed and validated on simulated and real
9 sEMG signals. The proposed approach outperforms the other threshold-dependent
10 *VAF*-based approaches in terms of estimation bias and mean absolute error and
11 overcomes their limitations, being independent of arbitrary cut-off thresholds.
12 Moreover, it may help the standardization of motor control studies based on muscle
13 synergies among different research laboratories.

1 Chapter 4

2 Motor Control Strategies during 3 Non-Cyclic Movements

4 In the following chapter, the methodological issues in the assessment of motor
5 control strategies during single-leg stance tasks are discussed. First, a novel pre-
6 processing approach for the selection of sEMG epochs corresponding to “*well-*
7 *balanced*” and “*unbalanced*” single-leg stance is assessed and validated
8 (**Methodological Issue – Muscle Synergy Assessment during Single-Leg**
9 **Stance**). Second, differences in terms of balance control strategies between “*well-*
10 *balanced*” and “*unbalanced*” single-leg stance epochs are analyzed by using the
11 muscle synergy theory (**Methodological Issue – Muscle Synergy Assessment**
12 **during Single-Leg Stance**). Then, to investigate muscle synergies in lower limb
13 and back muscles during a single-leg stance task with and without visual
14 information, the proposed approach is assessed considering a sample population of
15 healthy young individuals (**Application - Influence of Visual Feedback on**
16 **Muscle Synergies**).

17 **Methodological Issue – Muscle Synergy Assessment during** 18 **Single-Leg Stance**

19 Some of the results presented in this paragraph are reproduced and modified
20 from an article published in *IEEE Transactions on Neural Systems and*
21 *Rehabilitation Engineering* (Volume: 28, Issue: 12, Dec. 2020) by Ghislieri *et al.*
22 The final authenticated version of the manuscript is available online at
23 <http://dx.doi.org/10.1109/TNSRE.2020.3030847>.

24 **4.1 Introduction**

25 The upright stance is a useful test condition to assess motor skills and evaluate
26 proprioception and coordination with applications both in clinics and sport. Several

1 balance conditions can be studied to assess postural sway, such as Double-Leg
2 Stance (DLS), tandem, semi-tandem, and Single-Leg Stance (SLS) [150]. Each of
3 these balance tasks is characterized by different motor control strategies and levels
4 of difficulty in carrying out the task. In particular, SLS or unipedal stance requires
5 the maintenance of the upright stance using a single limb, and it may be challenging
6 in patients affected by Chronic Ankle Instability (CAI) [151–153]. Moreover,
7 considering a specific balance task, different conditions of visual and
8 somatosensory integrations can be tested [154]. Typically, along with the eyes open
9 (EO) condition, in which the subject exploits the visual feedback to maintain
10 balance, a condition with eyes closed (EC) is also studied to evaluate the effect of
11 visual deprivation on postural balance control [155–157].

12 The study of muscle synergies is facing new challenges in the field of postural
13 balance analysis. To the best of the author’s knowledge, the analysis of muscle
14 synergies during balance tasks is mainly focused on the evaluation of balance
15 recovery after a “controlled” perturbation [26,158–161]. More specifically, a few
16 studies have demonstrated that muscle synergies are highly consistent across
17 different balance tasks and levels of difficulty [26,28,162]. This suggests that,
18 increasing the task complexity, there should be only slight modifications to the
19 basic motor control strategies involved in postural balance control. However, no
20 studies have assessed the motor control strategies adopted to maintain balance
21 during SLS. One possible reason may be the difficulty to select epochs of sEMG
22 signals when the subject firmly maintains unipedal stance. Indeed, it is important
23 to separate sEMG epochs in which balance is properly maintained from those in
24 which a slight disequilibrium occurs. Therefore, this chapter aims at defining a
25 robust approach for the segmentation of the sEMG time-instants relative to a “*well-*
26 *balanced*” (WB) SLS or an “*unbalanced*” (UB) SLS to be used as inputs of the
27 muscle synergy extraction algorithm. The proposed approach might help the
28 interpretation of muscle synergies in the SLS task.

29 **4.2 Materials and Methods**

30 *1. Sample Population*

31 A sample population of twenty-two healthy subjects (11 females and 11 males;
32 age: 24 ± 3 years; height: 175.7 ± 9.6 cm; weight: 65.9 ± 12.2 kg) was enrolled in
33 the study. None of the volunteers reported lower limb injuries or had
34 neurological/musculoskeletal disorders that could compromise the execution of the
35 balance task. All the subjects were right-limb dominant, according to the preferred
36 lower limb to start walking. This study was reviewed and approved by the Ethics
37 Committee of the Area Vasta Emilia Centro della Regione Emilia Romagna (CE
38 AVEC 193/2019/Sper/IOR approved on October 4, 2019). All participants signed
39 written informed consent for the protocol before each experimental session, and all
40 the acquisitions were performed in accordance with the Declaration of Helsinki.

41 *2. Experimental Protocol*

1 The subjects were asked to perform a SLS task twice in two different
2 conditions: eyes open (EO) and eyes closed (EC). The order condition was
3 randomized for each subject by tossing a coin. In particular, for each condition,
4 subjects were asked to perform a transition from double-leg stance (DLS) to single-
5 leg stance (SLS), to maintain SLS for at least 30 seconds, and then to return back
6 to double-leg stance. If the subject failed to maintain the SLS balance for at least
7 30 s and required to land on both feet during the task, the test was stopped and
8 repeated until the minimum time duration was reached. The test was performed on
9 a firm surface (force plate), with the subject keeping the arms straight at the sides.
10 **Figure 4.1** shows a schematic representation of the experimental protocol.

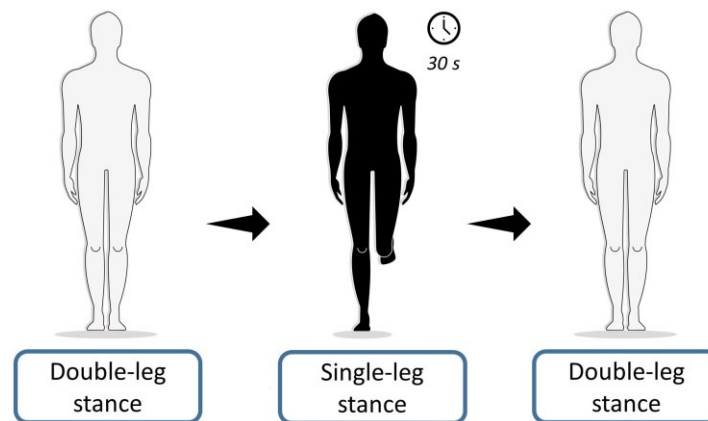


Figure 4.1 Schematic representation of the experimental protocol. Subjects were asked to perform a transition from double-leg stance (DLS) to single-leg stance (SLS), maintaining SLS for at least 30 seconds, and the returning back to DLS to end the task. Reprinted from “*Muscle Synergy Assessment During Single-Leg Stance*” by Ghislieri *et al.* (2020).

11 3. Data Acquisitions

12 During each experimental session, sEMG, foot-switch, and ground reaction
13 force signals were simultaneously acquired. sEMG signals were recorded through
14 active probes (FREEEMG 1000, BTS Bioengineering, Milan, Italy) placed over the
15 main muscles of the lower limb and the trunk. Foot-switch signals (FREEEMG
16 1000 – Footswitch Kit, BTS Bioengineering, Milan, Italy) were acquired to detect
17 the onset/offset timing of the SLS task, while ground reaction forces were detected
18 through a force plate (Dynamic Walkway P6000, BTS Bioengineering, Milan,
19 Italy) to assess body sway.

20 More specifically, sEMG active probes were placed over the following 13
21 muscles of the dominant lower limb (the one sustaining the SLS) and the trunk
22 bilaterally: right Longissimus Dorsii (LD_R), left Longissimus Dorsii (LD_L), Gluteus
23 Medius (GMD), Rectus Femoris (RF), Lateral Hamstring (LH), Medial Hamstring
24 (MH), Vastus Medialis (VM), Vastus Lateralis (VL), Lateral Gastrocnemius
25 (LGS), Peroneus Longus (PL), Peroneus Brevis (PB), Soleus (SOL), and Tibialis
26 Anterior (TA). These signals were acquired at a sampling frequency of 1 kHz.

1 The foot-switch sensor was placed under the first metatarsal head of the non-
2 dominant foot (corresponding to the left foot, for each subject of the sample
3 population).

4 All the acquired signals were then imported into MATLAB[®] release R2019b
5 (The MathWorks Inc., Natick, MA, USA) to be offline processed through custom
6 routines.

7 **Figure 4.2** shows a schematic representation of the acquisition system
8 composed of the sEMG active probes placed over the main muscles of the dominant
9 lower limb and the trunk, the foot-switch sensor mounted on the contralateral side
10 to detect the SLS timing, and the force plate used to assess body sway.

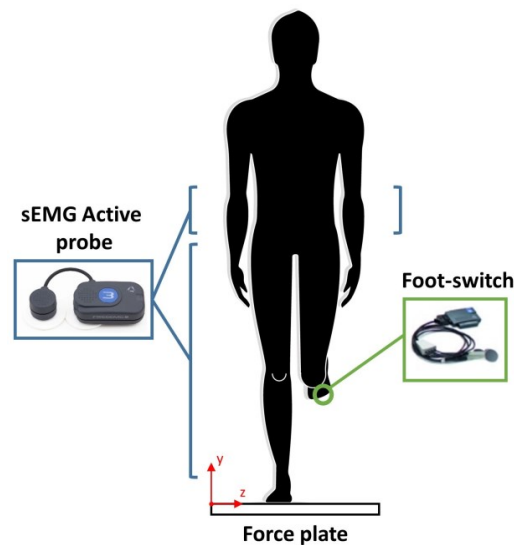


Figure 4.2 Schematic representation of the acquisition system. sEMG active probes were placed over the main muscles of the dominant lower limb (sustaining the SLS) and the trunk. A foot-switch was placed beneath the first metatarsal head of the contralateral foot (left side for all the subjects of the sample population) to detect the SLS timing. A force plate was used to assess body sway during SLS. Reprinted from “*Muscle Synergy Assessment During Single-Leg Stance*” by Ghislieri *et al.* (2020).

11 4. Segmentation of WB and UB SLS

12 The signals acquired from the foot-switch sensor placed beneath the
13 contralateral foot (left) and the ground reaction force acquired from the force plate
14 were used to segment the sEMG time-instants relative to WB or UB SLS.

15 In particular, the foot-switch signal was used to detect the time-instants when
16 the subjects performed the transition from DLS to SLS and vice versa. First, the
17 foot-switch signal was normalized in amplitude between 0 (open foot-switch, foot
18 raised from the floor) and 1 (closed foot-switch, foot on the floor). Second, the onset
19 of the SLS task was established 5 seconds after the 1-to-0 transition, while the offset
20 was established 5 seconds before the 0-to-1 transition, discarding from the analysis
21 DLS-to-SLS and SLS-to-DLS transitions.

22 To distinguish between WB and UB sEMG epochs during SLS, the ground
23 reaction force acquired through the force plate was used. The x -axis of the ground
24 reaction force is aligned to the antero-posterior (AP) direction, the y -axis is aligned
25 to the down-top vertical direction, and the z -axis is aligned to the medio-lateral
26 (ML) direction. Due to the high correlation between the planar components (AP

1 and ML) and the vertical component of the ground reaction force, only AP and ML
 2 were considered to segment WB and UB sEMG epochs, discarding the vertical
 3 component of the force [150].

4 Both the ground reaction force components (AP and ML) were first low-pass
 5 filtered through a 5th order Butterworth digital filter with a cut-off frequency of 10
 6 Hz [163,164]. Then, the resultant force (F_{res}) was computed as described in (4.1):

$$F_{res} = \sqrt{F_{AP}^2 + F_{ML}^2} \quad (4.1)$$

7 where F_{AP} and F_{ML} represent the AP and ML components of the low-pass filtered
 8 ground-reaction force, respectively.

9 The Root-Mean-Square (RMS) of the resultant reaction force ($F_{res_{RMS}}$) was
 10 computed by windowing the signal into 1s-epochs without overlap to ensure a
 11 sufficient number of samples to be used as input of the muscle synergy extraction
 12 algorithm. Finally, an adaptive threshold (Th_c) was applied to the $F_{res_{RMS}}$ signal
 13 to detect the sEMG time-instants relative to a WB or UB SLS, defined as in (4.2):

$$Th_c = mean(F_{res_{RMS}}) + c \cdot std(F_{res_{RMS}}) \quad (4.2)$$

14 where c is a multiplicative constant.

15 Due to the high inter-subject variability of the $F_{res_{RMS}}$ signals and strong
 16 correlation between the ground reaction forces and several anthropometric
 17 parameters (e.g., body weight, body height, and leg length), an adaptive threshold
 18 was preferable to a fixed threshold to properly detect well-balanced and unbalanced
 19 SLS epochs according to the distribution of the ground reaction forces of each
 20 subject.

21 **Figure 4.3** represents the segmentation masks used to select the “*well-*
 22 *balanced*” and “*unbalanced*” SLS epochs from a healthy subject of the sample
 23 population, considering three different values of the multiplicative constant c ($c =$
 24 0.5 , $c = 1.0$, $c = 1.5$). These c -values have been chosen to achieve a sufficient
 25 number of sEMG time-instants for muscle synergy extraction (for both WB and UB
 26 epochs).

27 More specifically, each Segmentation Mask (SM) was a binary mask defined
 28 as it follows:

- 29 • SM = 1, if $F_{res_{RMS}} \leq Th_c$ (WB epochs)
- 30 • SM = 0, if $F_{res_{RMS}} > Th_c$ (UB epochs).

31

32 where Th_c represents the adaptive threshold value given the value of the
 33 multiplicative constant c .

34 The sEMG epochs relative to a “*well-balanced*” and “*unbalanced*” SLS were
 35 then segmented using the above defined SM.

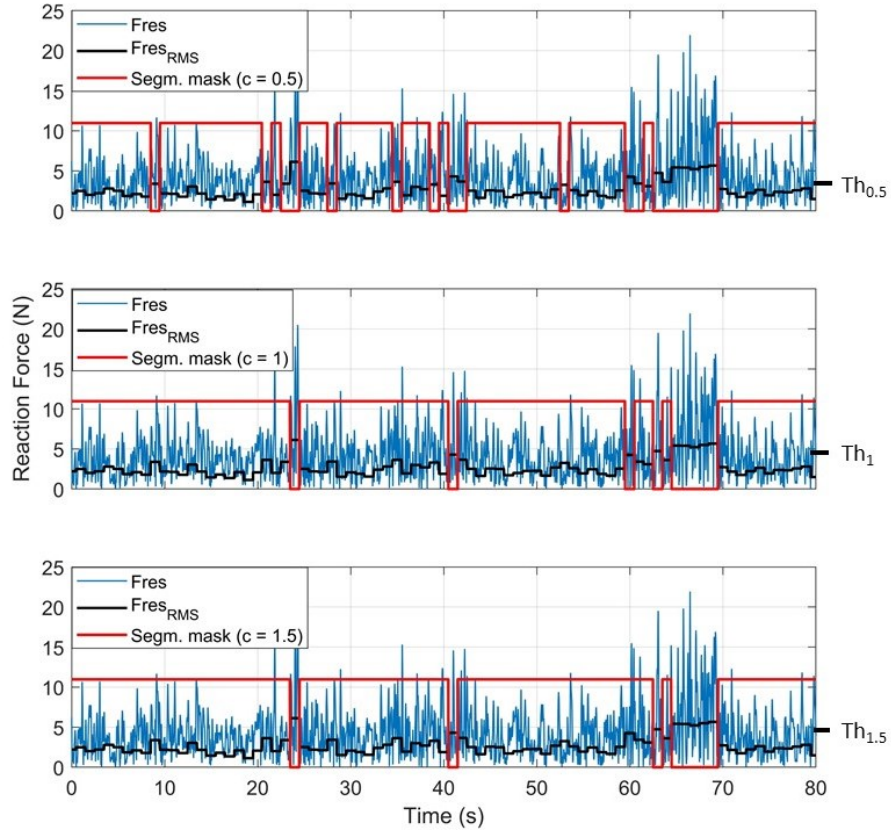


Figure 4.3 Segmentation masks used to select the “well-balanced” and “unbalanced” SLS epochs for a representative subject of the sample population, considering three different values of the multiplicative constant c . In blue it is represented the resultant reaction force (F_{res}) during SLS task, in black the RMS of the resultant reaction force ($F_{res_{RMS}}$), and in red the segmentation mask computed considering each of the three different values of c . Each segmentation mask assumes values equal to 1 in correspondence of “well-balanced” SLS epochs and to 0 otherwise (“unbalanced” SLS epochs). Reprinted from “Muscle Synergy Assessment During Single-Leg Stance” by Ghislieri *et al.* (2020).

1 5. Muscle Synergy Extraction and Number Selection

2 After sEMG segmentation, muscle synergies were extracted and properly
 3 sorted for the WB and UB SLS epochs, separately. First, the segmented sEMG
 4 signals were high-pass filtered at 35 Hz through an 8th order Butterworth digital
 5 filter to attenuate motion artifacts. Second, signals were demeaned and full-wave
 6 rectified to obtain non-negative signals. Third, rectified sEMG signals were low-
 7 pass filtered at 12 Hz through a 5th order low-pass Butterworth digital filter [165].
 8 Fourth, each envelope was normalized in amplitude with respect to its global
 9 maximum to ensure equally weighted contributions of all the acquired muscles in
 10 the muscle synergy extraction process [165].

11 Afterward, muscle synergies were extracted from the amplitude-normalized
 12 sEMG envelopes by means of the NNMF algorithm [28,41]. The NNMF
 13 decomposes the original sEMG envelope matrix ($M(t)$) as the linear combination
 14 of two different components: the time-dependent activation coefficients ($C(t)$) and
 15 the time-independent weight vectors (W) [34] as described in (3.2)

16 The activation coefficient vector $C(t)_k$ describes the modulation over time of
 17 the muscles enrolled in the k -synergy (the temporal component of the motor

1 control), while the weight vector W_k represents the contribution of each muscle to
 2 the k -synergy (the spatial component of the motor control).

3 The MATLAB[®] function “*nnmf*” was used to factorize the original sEMG
 4 envelope matrix, setting the routine’s input parameters as detailed in TABLE 4.1.
 5 The input parameters used in this study were optimized in previous works focused
 6 on muscle synergy extraction during gait [46,66]. To explore different solutions of
 7 the NNMF algorithm, the “*nnmf*” function was run several times on the same sEMG
 8 data, changing the number of muscle synergies (N) from 1 to 8.

TABLE 4.1
 INPUT PARAMETERS OF THE MATLAB[®] ROUTINE “*NNMF*” USED FOR MUSCLE SYNERGY EXTRACTION

Parameters	Values
Algorithm	multiplicative update
Function tolerance	$1e^{-6}$
Number of replicates	50
Number of iterations	1000

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9 The reconstruction accuracy of the original sEMG envelope matrix ($M(t)$) was
 10 assessed for each number of muscle synergies (N) by computing the total Variance
 11 Accounted For ($tVAF$), defined as the uncentered Pearson’s correlation coefficient
 12 expressed in percentage (3.3).

13 The optimal number of muscle synergies (N_{opt}) was first selected by choosing
 14 the least number of muscle synergies ensuring $tVAF \geq 90\%$ (global criterion) [55].
 15 Second, considering the number of muscle synergies selected according to the
 16 global criterion, the Variance Accounted For (VAF) was also computed for each of
 17 the acquired muscles. If $VAF \geq 75\%$ for each of the 13 muscles (local criterion), it
 18 was concluded that no additional muscle synergies were needed to reconstruct the
 19 original sEMG envelopes. Otherwise ($VAF < 75\%$), the number of muscle
 20 synergies (N) was incremented until all the muscles achieved a VAF value equal
 21 to or greater than 75% [28,166].

22 To graphically represent the muscle synergies, the weight vectors (W) were
 23 normalized in amplitude in the range $[0, 1]$ with respect to their global maximum.
 24 Then, the activation coefficient vectors ($C(t)$) were multiplied by the correspondent
 25 normalized values.

26 Since the factorization algorithm may return W and $C(t)$ in a different order
 27 for each subject or test condition, proper sorting was required. For this purpose, a
 28 k -means clustering algorithm was applied to the weight vectors (W) [22]. The
 29 clustering algorithm was set considering N_{opt} as number of k -means clusters, 1000
 30 as maximum number of iterations, 15 as number of replicates, and cosine similarity
 31 as distance metric. Activation coefficients were then sorted accordingly.

32 6. Robustness Assessment

33 To assess the robustness of the proposed approach with respect to the
 34 segmentation threshold Th_c , muscle synergies were extracted from sEMG
 35 envelopes relative to WB or UB SLS epochs, considering 3 different values of the
 36 multiplicative constant c : $c = 0.5$, $c = 1.0$, and $c = 1.5$.

1 Afterward, the muscle synergies extracted considering the three different
 2 thresholds $Th_{0.5}$, $Th_{1.0}$, and $Th_{1.5}$ were quantitatively compared in terms of (i) the
 3 consistency of the optimal number of muscle synergies (N_{opt}), and (ii) the
 4 similarity of weight vectors estimated through Pearson's correlation coefficient (R).

5 7. Muscle Synergy Analysis

6 To prove the necessity to segment WB or UB SLS epochs, the correspondent
 7 muscle synergies were extracted (setting $c = 1.0$) and compared one to each other
 8 in terms of (i) the optimal number of muscle synergies (N_{opt}), (ii) the average
 9 recruitment level of the activation coefficient vectors ($Recr$), and (iii) the balance
 10 control strategies (S).

11 i. Optimal Number of Muscle Synergies (N_{opt})

12 The optimal number of muscle synergies (N_{opt}) necessary to accurately model
 13 the original sEMG signals was selected by choosing the smallest number of
 14 synergies which guarantees $tVAF \geq 90\%$ (global criterion) and $VAF \geq 75\%$
 15 (local criterion) for each of the acquired muscles [28,55,166].

16 ii. Average Recruitment Level ($Recr$)

17 Since no typical cyclostationary processes can be assessed during the SLS task,
 18 any direct or morphological interpretation of the activation coefficient vectors
 19 $C(t)_k$ is difficult. Hence, the average recruitment level ($Recr_k$) was considered to
 20 quantitatively compare muscle synergy activation coefficient vectors ($C(t)$) [28].
 21 The $Recr_k$ was computed as the average (over time) of the activation coefficient
 22 vector $C(t)_k$ of the k -synergy.

23 iii. Balance Control Strategy (S)

24 Starting from the balance task performed and the recorded muscles, three
 25 different balance control strategies can be identified: (a) ankle control, (b) knee
 26 control, and (c) hip/trunk control [26]:

- 27 a. The ankle control strategy (S_{ankle}) is mainly characterized by the activation
 28 of PL, PB, TA, LGS, and SOL muscles.
- 29 b. The knee control strategy (S_{knee}) is mainly characterized by the activation
 30 of VM, VL, and RF muscles.
- 31 c. The hip/trunk control strategy (S_{hip}) is mainly characterized by the
 32 activation of LH, MH, GMD, LD_R, and LD_L muscles.

33 The average weight vector (W_k) across those muscles belonging to the same
 34 balance control strategy was computed to quantitatively assess each balance control
 35 strategy, as detailed in (4.3):

$$S_{j,k} = \frac{\sum_{i=1}^m W_{k,i}}{m} \quad (4.3)$$

1 where $S_{j,k}$ represents the j -th balance control strategy ($j=1$: ankle control; 2: knee
 2 control; 3: hip/trunk control) for the k -synergy, and m represents the number of
 3 muscles enrolled in the j -th balance control strategy.

4 Finally, only a single balance control strategy (S_k) was associated with each
 5 muscle synergy by calculating the highest $S_{j,k}$ among those computed for the same
 6 k -synergy, as detailed in (4.4):

$$S_k = \max (S_{1k}, S_{2k}, S_{3k}) \quad (4.4)$$

7 where k can assume value from 1 to 3 ($S_1 = S_{\text{ankle}}$, $S_2 = S_{\text{knee}}$, $S_3 = S_{\text{hip}}$).

8 8. Statistical Analysis

9 To assess the robustness of the proposed approach with respect to the
 10 segmentation threshold, a two-way analysis of variance (ANOVA) followed by
 11 *post-hoc* analysis with Bonferroni adjustment for multiple comparisons was
 12 performed.

13 First, the Lilliefors test was used to test the hypothesis of normality of the
 14 distribution, setting the significance level (α) equals to 0.05. Second, according to
 15 the results of the Lilliefors test, the Wilcoxon signed-rank test ($\alpha = 0.05$) or the two-
 16 tailed paired Student's t -test ($\alpha = 0.05$) was performed to assess significant changes
 17 in the optimal number of muscle synergies, in the average recruitment levels, and
 18 in the balance control strategies considering different sEMG epochs (WB and UB).

19 The effect size of the statistically significant differences was calculated by
 20 means of the Hedges' g including the correction for small sample sizes [113].

21 4.3 Results

22 Results revealed, on average, a similar number of WB and UB SLS epochs (8
 23 ± 3 WB/UB epochs), but different epoch durations were measured. Considering the
 24 EO condition, the average WB and UB epoch durations were equal to 70.5 ± 9.8 s
 25 and 10.3 ± 2.9 s, respectively, while considering the EC condition, they were equal
 26 to 42.8 ± 27.3 s and 5.8 ± 3.8 s.

27 First, the results related to the robustness assessment of the segmentation
 28 threshold were presented. Second, the muscle synergies obtained considering the
 29 WB and UB SLS epochs are quantitatively compared to justify the separation into
 30 WB and UB epochs of SLS.

31 1. Robustness of the Segmentation Threshold

32 All the tested segmentation thresholds ($Th_{0.5}$, $Th_{1.0}$, and $Th_{1.5}$) required the same
 33 number of muscle synergies (N_{opt}) to accurately model the original sEMG signals
 34 with a $tVAF \geq 90\%$ and a $VAF \geq 75\%$ for each of the acquired muscles. In
 35 particular, considering WB SLS epochs, 4 muscle synergies were needed to
 36 reconstruct the sEMG signals for all the tested thresholds, both in EO and EC
 37 conditions. The same results were obtained considering UB epochs.

Moreover, results revealed high values of the Pearson’s correlation coefficients between each pair of thresholds. Considering the WB SLS epochs, no statistically significant differences in terms of weight-vector correlation among the 3 thresholds ($p = 0.17$) and the 2 tested conditions ($p = 0.87$) were detected. Considering the UB SLS epochs, no statistically significant differences among thresholds ($p = 0.28$) were detected, while a significant decrease ($p = 0.01$, $g = 0.47$) in the weight correlation was detected in the EO condition with respect to the EC condition.

TABLE 4.2 represents the average Pearson’s correlation coefficients (R) between each pair of thresholds, separately for WB and UB SLS epochs. Results reveal a very high similarity among the muscle synergy weight vectors extracted considering the 3 thresholds, both in WB and UB epochs.

Considering the high similarity of the weight vectors and the high consistency of the optimal number of muscle synergies obtained from the 3 thresholds, the multiplicative constant c of the segmentation threshold was set equal to 1.0.

2. “Well-balanced” and “Unbalanced” Single-Leg Stance

No statistically significant differences were found in terms of the number of muscle synergies between WB and UB SLS epochs. Considering the EO condition, 4 muscle synergies were necessary to reconstruct the original sEMG signals with an average $tVAF$ value of $93.0\% \pm 1.2\%$ for WB, and $93.1\% \pm 1.3\%$ for UB, respectively. For the EC condition, similar results were obtained. More specifically, 4 muscle synergies were extracted with an average $tVAF$ value of $92.6\% \pm 1.5\%$ for WB, and $92.8\% \pm 1.3\%$, for UB, respectively.

TABLE 4.2
WEIGHT VECTOR CORRELATION (R) AVERAGED ON THE SAMPLE POPULATION

SLS epochs		Pearson’s Correlation Coefficient (R) (mean \pm standard deviation)		
		$Th_{0.5}$ vs Th_1	$Th_{0.5}$ vs $Th_{1.5}$	Th_1 vs $Th_{1.5}$
<i>Well-balanced</i>	EO	0.91 ± 0.16	0.92 ± 0.14	0.95 ± 0.11
	EC	0.91 ± 0.15	0.89 ± 0.17	0.97 ± 0.08
<i>Unbalanced</i>	EO	0.84 ± 0.19	0.75 ± 0.20	0.80 ± 0.20
	EC	0.88 ± 0.17	0.86 ± 0.17	0.90 ± 0.18

$Th_{0.5}$: first threshold with $c=0.5$; $Th_{1.0}$: second threshold with $c=1.0$; $Th_{1.5}$: third threshold with $c=1.5$. EO: Eyes Open; EC: Eyes Closed.

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Figure 4.4 shows muscle synergy results, averaged over the sample population, extracted considering the EO condition (**Figure 4.4A**) and the EC condition (**Figure 4.4B**), respectively. Moreover, for each balance condition, the muscle synergies extracted considering WB and UB SLS epochs are compared. More specifically, **Figure 4.4** represents, for each k -synergy, both the average recruitment level ($Recr_k$) and the weight vector (W_k).

Wilcoxon signed-rank test revealed a statistically significant increase in the average recruitment levels of three out of four muscle synergies extracted considering UB epochs with respect to those extracted considering WB epochs, during both EO and EC conditions. **TABLE 4.4** details the average recruitment levels

1 of each muscle synergy with the indication of the statistically significant differences
 2 between WB and UB SLS epochs (indicated by asterisks).

3 In **Figure 4.4A** and **Figure 4.4B**, statistically significant changes ($p < 0.05$) of
 4 the average recruitment levels and the muscles' contribution to each muscle synergy
 5 are indicated by asterisks during the EO and EC conditions, respectively.

TABLE 4.4
 RECRUITMENT LEVELS (*Recr*) AVERAGED
 ON THE SAMPLE POPULATION

Recruitment Levels		Average Recruitment Levels (<i>Recr</i>) (mean \pm standard deviation)	
		<i>Well-balanced (WB)</i>	<i>Unbalanced (UB)</i>
EO	<i>Recr</i> ₁	0.20 \pm 0.06*	0.24 \pm 0.05*
	<i>Recr</i> ₂	0.19 \pm 0.05**	0.24 \pm 0.05**
	<i>Recr</i> ₃	0.22 \pm 0.06	0.24 \pm 0.06
	<i>Recr</i> ₄	0.15 \pm 0.04*†	0.17 \pm 0.05*‡
EC	<i>Recr</i> ₁	0.17 \pm 0.05***	0.25 \pm 0.05***
	<i>Recr</i> ₂	0.19 \pm 0.07	0.22 \pm 0.06
	<i>Recr</i> ₃	0.19 \pm 0.06*	0.22 \pm 0.07*
	<i>Recr</i> ₄	0.19 \pm 0.05****	0.23 \pm 0.05****

*Recr*_k represents the average recruitment level of the *k*-synergy ($k = 1,2,3,4$). The asterisk (*) indicates a statistically significant difference ($p < 0.05$) between well-balanced (WB) and unbalanced (UB) epochs of single-leg stance (SLS), while the dagger (†) between eyes open (EO) and eyes closed (EC) conditions. Single, double, and triple asterisks (or daggers) represent *p*-values lower than 0.05, 0.01, and 0.001, respectively. Reprinted from “*Muscle Synergy Assessment During Single-Leg Stance*” by Ghislieri *et al.* (2020).

TABLE 4.3
 BALANCE CONTROL STRATEGIES (*S*) AVERAGED
 ON THE SAMPLE POPULATION

Balance Control Strategies		Average Balance Control Strategies (<i>S</i>) (mean \pm standard deviation)	
		<i>Well-balanced (WB)</i>	<i>Unbalanced (UB)</i>
EO	Ankle	0.37 \pm 0.11*†	0.43 \pm 0.11*
	Knee	0.63 \pm 0.24†	0.73 \pm 0.22
	Hip/Trunk	0.51 \pm 0.14	0.56 \pm 0.16
EC	Ankle	0.43 \pm 0.12†	0.40 \pm 0.12
	Knee	0.76 \pm 0.20†	0.79 \pm 0.24
	Hip/Trunk	0.50 \pm 0.20	0.49 \pm 0.08

Average balance controls of each of the three identified strategies. The asterisk (*) indicates a statistically significant difference ($p < 0.05$) between well-balanced (WB) and unbalanced (UB) epochs of single-leg stance (SLS), while the dagger (†) between eyes open (EO) and eyes closed (EC) conditions. Single, double, and triple asterisks (or daggers) represent *p*-values lower than 0.05, 0.01, and 0.001, respectively. Reprinted from “*Muscle Synergy Assessment During Single-Leg Stance*” by Ghislieri *et al.* (2020).

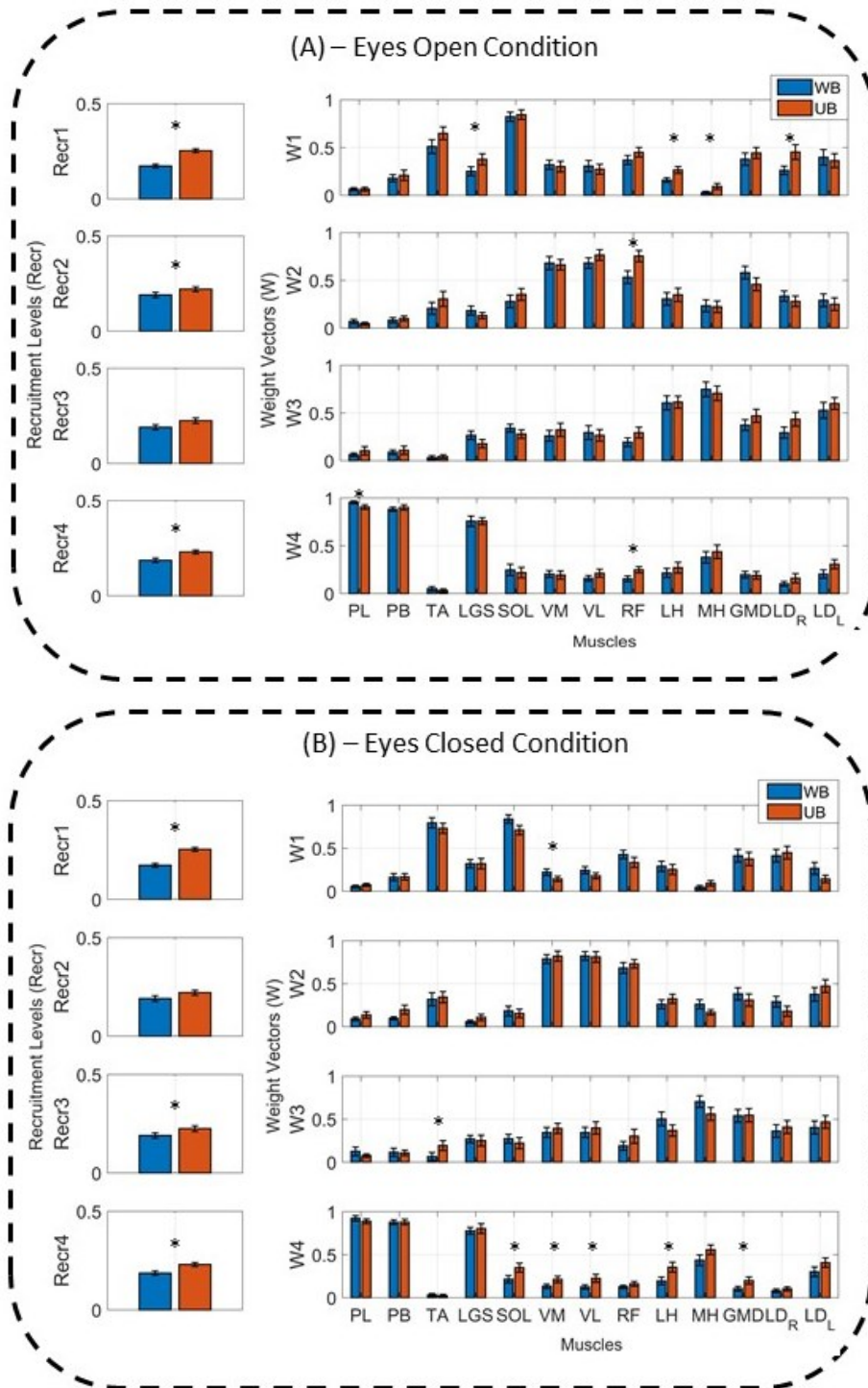


Figure 4.4 Comparison of the muscle synergies extracted during “well-balanced” (WB) and “unbalanced” (UB) epochs of single-leg stance (SLS), for both the eyes open (EO) and eyes closed (EC) conditions. In both panel (A) and (B), the colored vertical bars represent the average recruitment levels $Recr_k$ (on the left) and weight vectors W_k (on the right) of the k -synergy, over the sample population, with the superimposition of the standard error (black lines). The asterisk (*) indicates a statistically significant difference ($p < 0.05$). Reprinted from “Muscle Synergy Assessment During Single-Leg Stance” by Ghislieri *et al.* (2020).

1 According to equation (4.4), the ankle control strategy (S_{ankle}) can be mainly
 2 associated to the first and the fourth muscle synergies, the knee control strategy
 3 (S_{knee}) to the second muscle synergy, and the hip/trunk control strategy (S_{hip}) to
 4 the third muscle synergy. Considering the EO condition, results revealed a
 5 statistically significant increase ($p = 0.05$, $g = 0.13$) of the ankle control strategy

1 recruitment during the UB SLS epochs with respect to the WB ones. No statistically
2 significant differences between WB and UB SLS epochs were found considering
3 the knee and hip/trunk balance control strategies, mainly associated with the third
4 and fourth muscle synergies. No statistically significant differences in terms of
5 balance control strategies were detected between WB and UB SLS epochs. **TABLE**
6 **4.3** shows the values of the balance control strategies, averaged over the sample
7 population, with the indication of the statistically significant changes between WB
8 and UB epochs of SLS (indicated by asterisks).

9 Results presented in this chapter reveal differences both in terms of average
10 recruitment levels and balance control strategies between muscle synergies
11 extracted from WB and UB SLS epochs, justifying the proposed segmentation
12 approach as a necessary pre-processing procedure to properly assess the motor
13 control strategies and to help the interpretation of the muscle synergies during SLS.

14 **4.4 Discussion and Conclusions**

15 The muscle synergy theory was widely used in literature to assess human motor
16 control strategies during different motor tasks, such as gait and balance exercise. In
17 the last years, muscle synergy assessment during different balance conditions was
18 mainly focused on the evaluation of balance recovery after a “controlled”
19 perturbation [26,158–161], rather than on the maintenance of the balance condition
20 itself. The segmentation approach proposed in this chapter can be a useful tool for
21 the assessment of the motor control strategies adopted to maintain single-leg stance,
22 distinguishing between “*well-balanced*” (WB) and “*unbalanced*” (UB) SLS
23 epochs.

24 The robustness of the segmentation approach with respect to the threshold
25 selection was verified obtaining high consistency of the optimal number of muscle
26 synergies and high similarity of the weight vectors (Pearson’s correlation
27 coefficient R ranges from 0.75 to 0.97) across the three different values of the
28 adaptive segmentation threshold. Muscle synergies extracted from WB and UB SLS
29 epochs are consistent with previous studies [26,28] both in terms of activation
30 coefficients and weight vectors. However, due to the reduced complexity of the
31 balance exercise analyzed in this chapter, a fewer number of muscle synergies was
32 needed to properly reconstruct the original sEMG signals with respect to the
33 previous studies, in which SLS was assessed after the application of
34 multidirectional perturbations.

35 The number and the composition of the muscle synergies revealed no
36 statistically significant differences between the WB and UB SLS epochs. However,
37 differences in terms of average recruitment levels and balance control strategies
38 were detected, suggesting a slightly different motor control strategy between WB
39 and UB SLS epochs. More specifically, UB SLS epochs require on average higher
40 recruitment levels with respect to the WB SLS epoch, considering both the EO and
41 EC conditions, and an increased ankle control strategy, in the EO condition. These
42 differences justify the necessity to distinguish between WB and UB single-leg
43 stance epochs before muscle synergy extraction. Results presented in this chapter

1 are in line with those of previous studies in which the effect of the task complexity,
2 postural configuration, and loading conditions on motor control strategies were
3 assessed [26,30,31]. According to these studies, our central nervous system recruits
4 similar muscle synergies rather than generating a completely new set of muscle
5 synergies for each balance condition. Indeed, only slight modifications to the basic
6 motor control strategies (number and composition of the muscle synergies)
7 involved in SLS were detected, while differences in terms of average recruitment
8 levels and balance control strategies were assessed between EO and EC conditions.
9 Considering the WB and UB SLS epochs, the EC condition requires a higher
10 average recruitment level of the fourth muscle synergy (mainly associated with the
11 ankle control strategy), and an increased level of ankle and knee control strategies,
12 suggesting higher recruitment of the muscle synergies controlling the distal
13 muscles.

14 This study was conducted on a sample population of young healthy subjects
15 that are able to maintain the upright stance on one foot, even with their eyes closed,
16 for at least 30 s. This may represent a first limitation of the study, since it may be
17 difficult to apply the same experimental protocol also to elderly or pathological
18 populations affected by severe balance impairments. Moreover, muscle synergy
19 results obtained (similarity between motor control strategies adopted during WB
20 and UB SLS epochs), can be biased by the specific population considered. Different
21 results might be obtained in populations with diminished equilibrium skills or
22 severe balance impairments. Another limitation of this study is that it focused only
23 on balance strategies during SLS maintenance, without analyzing transition tasks.
24 The analyzed signals started 5 seconds after the first (DLS-to-SLS) transition and
25 stopped 5 seconds before the second (SLS-to-DLS) transition. Therefore, our
26 findings on motor control strategies adopted during SLS (excluding the transitions)
27 cannot be extended to task transitions.

28 In conclusion, results demonstrate that the proposed segmentation approach is
29 robust with respect to the selection of the segmentation threshold and can be
30 successfully used as a pre-processing step before muscle synergy extraction,
31 allowing for a better assessment of the modular organization of the central nervous
32 system during the maintenance of the single-leg stance. Further studies will focus
33 on the application of this approach to sEMG signals acquired from subjects affected
34 by chronic ankle instability (CAI) during SLS, to assess its applicability in
35 pathological conditions.

36

1 Application - Influence of Visual Feedback on Muscle 2 Synergies

3

4 4.5 Introduction

5 The ability to maintain single-leg stance is essential during daily living
6 activities, as a single task as well as a component of other more complex tasks, such
7 as walking and running. It is a simple but challenging condition for balance control
8 and for this reason it is widely used for training and rehabilitation [167,168]. In
9 research and clinical practice, it is widely used as a testing task as it allows
10 quantifying balance alterations and deficits of the single lower limb otherwise
11 concealed during the performance of double leg tasks [169–174].

12 From a physiological point of view, the single-leg stance can be considered as
13 a high demanding postural task for neuromuscular and central nervous systems
14 requiring an efficient integration of somatosensory, visual, and vestibular
15 information with the aim to orchestrate a continuous and effective motor response
16 to manage a reduced base of support [175]. The effectiveness of postural control
17 has been usually expressed by means of mechanical parameters such as the center
18 of pressure (COP), joints, or body segment displacement [176–178]. Previous
19 literature studies have reported the essential role of the ankle joint for postural
20 stabilization in particular when tasks show an increase in instability, as in the
21 transition from double- to single-limb stance [179] or from stable to unstable
22 surfaces [170]. When the ankle movements are not sufficient to guarantee balance,
23 the involvement of more proximal joints and body segments has been reported
24 [170,180]. Further, an increase in the instability during stance tasks has been also
25 reported in case of a number of pathological conditions [172,181,182] and in case
26 of abnormal sensitive information [172,183,184]. Above all, it has been shown that
27 vision has a key role in posture control and that the lack of visual feedback or
28 abnormal visual feedback leads to peculiar adaptations in mechanical parameters
29 featuring postural tasks [155,185].

30 Even if mechanical parameters, such as COP or joint displacement, are useful
31 to quantify instability during postural tasks, they do not give adequate information
32 on motor control. Essential information for motor control assessment comes from
33 the analysis of muscle activations, which mediates CNS control and mechanical
34 expression of movement.

35 While a wide number of studies investigated multi-muscles activations during
36 double limb stance, in the transition from double to single stance, or during various
37 stance tasks in response to sudden perturbations [26,158,159,186,187], less is
38 known about quiet single-limb stance. Few studies focused on ankle/foot muscle
39 activations, given their important role as previously described [188,189].

40 However, the investigation of muscle activations around a single joint is
41 reductive, since it is well known that CNS organizes motor response to a given task
42 in terms of muscle synergies [28,190]. This means that CNS coordinates the

1 activation of a set of muscles that are synergistic for a given task, or a number of
2 similar tasks [26].

3 To the best of the author's knowledge, the muscle synergies used for balance
4 control during the maintenance of single-leg stance (SLS) conditions are not
5 known. Since this task is largely used, discovering the muscle synergies adopted by
6 healthy individuals is essential to address future research, such as training,
7 rehabilitation, and functional assessment both in healthy and pathological
8 individuals. The aim of this chapter is to investigate muscle synergies in lower limb
9 and back muscles during a single-leg stance task, with and without visual
10 information, in healthy young individuals.

11 **4.6 Materials and Methods**

12 *1 Participants*

13 Eleven male participants (age: 23.9 years \pm 2.2 years; height: 182 cm \pm 8.4 cm;
14 body mass: 74.5 kg \pm 10.8 kg) and eleven female participants (age: 24.5 years \pm 2.9
15 years; height: 169 cm \pm 5.8 cm; body mass: 57.2 kg \pm 6.5 kg) were recruited to
16 participate in the study. Inclusion criteria were *a*) age between 20 and 35 years, *b*)
17 physical activity level of 2 and 3 according to the Saltin and Grimby scale [191],
18 thus excluding sedentary individuals and competitive athletes, and *c*) absence of
19 known neurological diseases. Exclusion criteria were *a*) previous injuries or
20 surgery, and *b*) abnormalities in lower limb and foot joints.

21 Each participant signed informed consent before participating in the study. The
22 study was conducted in accordance with the Declaration of Helsinki and received
23 ethical approval from the Ethics Committee of the Area Vasta Emilia Centro della
24 Regione Emilia Romagna (CE AVEC 193/2019/Sper/IOE approved on October 4,
25 2019).

26 *2 Experimental Protocol and Data Analysis*

27 Participants were asked to stand barefoot on a force platform (Dynamic
28 Walkway P6000, BTS Bioengineering, Milan, Italy) with the dominant limb and to
29 maintain the contralateral knee joint flexed at approximately 90°. They were asked
30 to look forward, to maintain upper limbs aligned to the trunk, and to remain as still
31 as possible for at least 30 seconds (see **Figure 4.5**). Minimal arms movements were
32 allowed, however, participants were asked to minimize them as much as possible.
33 They performed the task in both eyes open (EO) and eyes closed (EC) conditions.
34 These two conditions were performed in a random order for each subject (by tossing
35 a coin) and with two minutes of rest between trials. Muscle activations were
36 recorded from 13 muscles of the lower limb and trunk by means of
37 electromyography wireless probes (BTS FreeEMG 1000, BTS Bioengineering,
38 Milan, Italy) fixed on sEMG electrodes (Ag/AgCl) applied over Tibialis Anterior
39 (TA), Peroneus Longus (PL), Peroneus Brevis (PB), Soleus (SOL), Lateral
40 Gastrocnemius (LGS), Vastus Medialis (VM), Vastus Lateralis (VL), Rectus
41 Femoris (RF), Lateral Hamstring (LH), Medial Hamstring (MH), Gluteus Medius

1 (GMD), right Longissimus Dorsii (LD_R) and left Longissimus Dorsii (LD_L) in
 2 accordance with SENIAM recommendations [192]. To reduce the skin impedance,
 3 before electrode application, the skin area was shaved and cleaned with ethyl
 4 alcohol. A footswitch (FSW) was placed under the first metatarsal head of the non-
 5 dominant foot. Force platform, sEMG, and FSW signals were part of the same
 6 integrated system and were recorded with a 1000 Hz sampling rate.

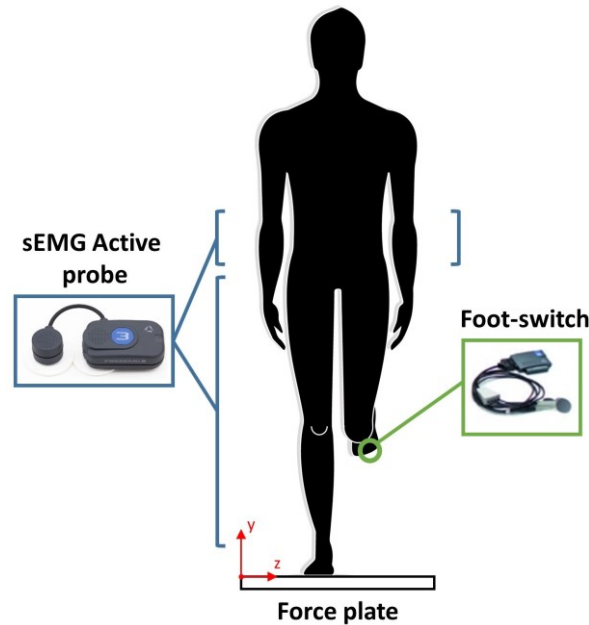


Figure 4.5 Schematic representation of the acquisition system. sEMG probes are positioned over the main muscles of the dominant lower limb (sustaining the single-leg stance) and the trunk. A foot-switch is placed under the first metatarsal head of the contralateral foot to detect the onset/offset timing of SLS. A force plate is used to assess body sway during SLS. Reprinted from “Muscle Synergy Assessment During Single-Leg Stance” by Ghislieri *et al.* (2020).

7 3 Muscle Synergy Analysis

8 Muscle synergies extracted from the segmented sEMG signals during the two
 9 different task conditions (EO and EC) were quantitatively compared in terms of (i)
 10 the optimal number of muscle synergies (N_{opt}), (ii) the average recruitment levels
 11 ($Recr$), and (iii) balance control strategies (S).

- 12 *i. Optimal number of muscle synergies (N_{opt}):* the optimal number of muscle
 13 synergies (N_{opt}) was selected for each subject and task condition by choosing
 14 the smallest number of muscle synergies (N) which guarantees $tVAF \geq 90\%$
 15 (global criterion) and $VAF \geq 75\%$ (local criterion) for each of the acquired
 16 muscles.
- 17 *ii. Average recruitment levels ($Recr$):* since no typical cyclostationary processes
 18 can be assessed during single-leg stance, it is difficult to compute average
 19 activation coefficients over the sample population. Thus, the activation
 20 coefficients ($C(t)$) were compared in terms of average recruitment level
 21 ($Recr_k$), defined as the average (over time) of each activation coefficient vector
 22 $C(t)_k$ [27,28]. The average recruitment level values range between 0 (no

1 recruitment) and 1 (maximum recruitment) and quantify how much a specific
2 muscle synergy is activated in the execution of the task.
3 *iii. Balance control strategies (S):* considering the task performed and the acquired
4 muscles, three different balance control strategies were identified: (a) ankle
5 control, (b) knee control, and (c) hip/trunk control. The ankle control strategy
6 (S_{ankle}) was mainly related to the activation of 5 muscles of the leg (PL, PB,
7 LGS, TA, and SOL), the knee control strategy (S_{knee}) to the activation of 3
8 muscles of the shank (VM, VL, and RF), and the hip/trunk control strategy (S_{hip})
9 to the activation of 5 muscles of the proximal lower limb and the trunk (LH,
10 MH, GMD, LD_L, and LD_R). Balance control strategies were quantified for each
11 k -synergy by computing the average weight vector (W_k) over the muscles
12 belonging to the same balance control strategy, obtaining three balance control
13 strategy values for each muscle synergy. A single balance control strategy was
14 finally associated with each muscle synergy by selecting the highest balance
15 control strategy among the three computed.

16

17 4 Statistical Analysis

18 To assess statistically significant changes in the optimal number of muscle
19 synergies considering the two different task conditions (EO and EC), the hypothesis
20 of normality of the distribution was firstly tested through the Lilliefors test. If the
21 normality hypothesis was rejected, the Wilcoxon signed-rank test was performed,
22 otherwise, a two-tailed paired Student's t-test was performed. Two-way ANOVA
23 for repeated measures followed by post-hoc analysis with Tukey adjustment for
24 multiple comparisons was performed to evaluate the differences between conditions
25 (OE and CE) and muscle synergies (factors: condition and synergies), for both the
26 average recruitment levels ($Recr$) and balance control strategies (S). For the weight
27 vectors (W), an analogous two-way ANOVA was applied to evaluate the
28 differences between conditions and muscles. The effect size of the statistically
29 significant differences was calculated by means of the Hedges' g including the
30 correction for small sample sizes [113].

31 All the levels of significance (α) were set equal to 0.05. The statistical analysis
32 was carried out using the Statistical and Machine Learning Toolbox of MATLAB®.

33 4.7 Results

34 In the following, the muscle synergy results are computed considering the two
35 different single-leg stance conditions (EO and EC). More specifically, muscle
36 synergies were compared in terms of (i) the optimal number of muscle synergies,
37 (ii) average recruitment levels, and (iii) balance control strategies.

38 *i. Optimal number of muscle synergies (N_{opt})*

39 The application of the Wilcoxon signed-rank test revealed no statistically
40 significant differences ($p = 0.52$) in the optimal number of muscle synergies
41 (N_{opt}) between the EO and EC conditions. In particular, 4 muscle synergies

1 were needed to accurately model the motor control strategies during both the
 2 EO and EC conditions.

3 **Figure 4.6** shows the muscle synergies, averaged over the sample
 4 population, extracted from the two different task conditions: EO represented in
 5 blue and EC in red. More specifically, for each muscle synergy, the recruitment
 6 levels $Recr_k$ (on the left) and the weight vectors W_k (on the right) are
 7 represented. The asterisk (*) indicates statistically significant differences
 8 between conditions (repeated measures ANOVA, $p < 0.05$), both for the
 9 average recruitment levels and weight vectors.

10 ii. *Average recruitment levels (Recr)*

11 A statistically significant decrease ($p = 0.02$, $g = 3.35$) in the average
 12 recruitment level of the third muscle synergy extracted during the EC condition
 13 (0.17 ± 0.01) with respect to the EO condition (0.21 ± 0.01) was assessed. No
 14 statistically significant differences were detected considering the remaining
 15 three muscle synergies between EO and EC conditions, suggesting no changes
 16 in the recruitment levels of those synergies due to the loss of visual feedback.

17 **Figure 4.6** shows the average recruitment levels (on the left), over the
 18 sample population, extracted during EO and EC single-leg stance conditions.

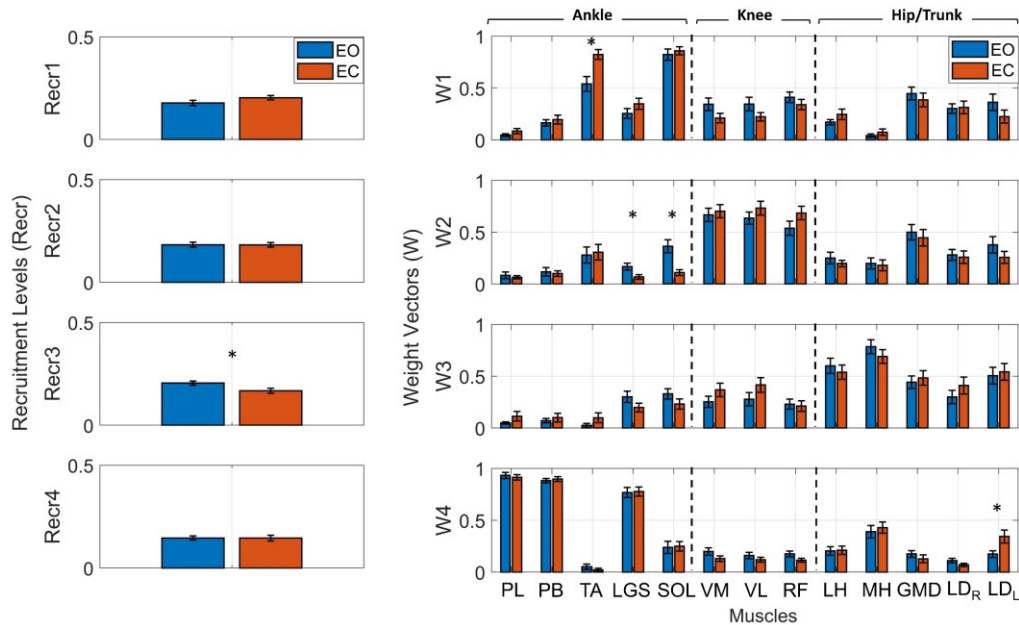


Figure 4.6 Comparison of muscle synergies extracted during eyes open (EO) and eyes closed (EC) single-leg stance conditions. Color vertical bars represent average recruitment levels $Recr_k$ (on the left) and weight vectors W_k (on the right) of the k -synergy, over the sample population, with the superimposition of the standard error (black lines). The asterisk (*) represents a statistically significant difference between conditions, in the weight vectors and average recruitment levels.

19 iii. *Balance control strategies (S)*

20 As shown in **Figure 4.6**, the first and the fourth muscle synergies can be
 21 mainly associated with an ankle control strategy, since the muscles mainly
 22 enrolled are those belonging to the leg (PL, PB, LGS, and SOL), the second
 23 muscle synergy to a knee control strategy and the third muscle synergy to a

1 hip/trunk control strategy. ANOVA for repeated measures revealed a
 2 statistically significant increase ($p = 0.03$, $g = 0.78$) of the ankle control
 3 strategies (S_{ankle}) during the EC condition (0.52 ± 0.06) with respect to the EO
 4 condition (0.47 ± 0.06). No additional statistically significant differences were
 5 detected considering the remaining two balance control strategies (S_{knee} and
 6 S_{hip}) between conditions.

7 **TABLE 4.5** represents the values of the balance control strategies, averaged
 8 over the sample population, with the indication of the statistically significant
 9 changes between EO and EC conditions.

TABLE 4.5
 BALANCE CONTROL STRATEGIES (S) AVERAGED ON THE SAMPLE POPULATION

BALANCE CONTROL STRATEGIES	AVERAGE BALANCE CONTROL STRATEGIES (S) (MEAN \pm STANDARD DEVIATION)		
	<i>EO</i>	<i>EC</i>	<i>ANOVA (P-VALUE)</i>
S_{ankle}	0.47 ± 0.06	0.52 ± 0.06	0.03
S_{knee}	0.61 ± 0.24	0.71 ± 0.25	0.22
S_{hip}	0.53 ± 0.15	0.53 ± 0.17	0.89

S_{ankle} : ankle control strategy, S_{knee} : knee control strategy, and S_{hip} : hip/trunk control strategy.

10 4.8 Discussion and Conclusions

11 Since the work by Horak and Nashner (1986), it is widely recognized the
 12 essential role of the ankle joint in the control of upright stance and for the
 13 maintenance of posture when balance is challenged by perturbations of the
 14 supporting surface. In these circumstances, muscles around the ankle joint provide
 15 the first activation strategy for balance maintenance [180]. In this chapter, no
 16 perturbations were applied to the supporting surface, and participants were required
 17 to maintain a quiet single-leg stance (SLS). The key role of the ankle joint for
 18 posture control in a quiet stance is confirmed by the observation of two ankle-
 19 dominant muscle synergies adopted by the participants in this study. The first ankle-
 20 dominant synergy (W_1) is mainly featured by the Tibialis Anterior (TA) and the
 21 Soleus (SOL) muscle activation. The second ankle dominant muscle synergy (W_4)
 22 is mainly featured by Peroneus Longus (PL), Peroneus Brevis (PB), and
 23 Gastrocnemius Lateralis (LGS) muscle activations. These two synergies may
 24 reflect the activations related to antero-posterior sway and medio-lateral sway,
 25 respectively, which may occur during a SLS task. In particular the co-activation of
 26 antagonist muscles (TA and SOL) might represent a strategy to cope with reduced
 27 base of support, with the aim to reduce movement variability and maintaining
 28 stability. Previous studies found an increase in TA and SOL activations in particular
 29 in older adults to compensate for reduced vision [193] or decreased tendon stiffness
 30 [194], and both in children and elderly which showed a diminished postural
 31 steadiness when compared with young adults [195].

32 Literature reports that as more difficult the task becomes as higher is the
 33 involvement of more proximal joints for the maintenance of balance, in particular

1 the hip [170,180]. In experimental settings, the difficulty of the task is usually
2 increased by increasing the magnitude of a perturbation, by decreasing the
3 supporting surface, or by changing the features of the supporting surface
4 [28,159,170,187]. For example, it has been reported that by moving from a stable
5 to an unstable surface, the angular displacement of the ankle was stable across all
6 the testing conditions, with the knee and hip displacement arising when the
7 difficulty of the task was higher [170,196].

8 In this chapter, the difficulty of the task was not modified throughout the
9 experiment and the support surface was not unstable. However, standing on a single
10 limb might be considered a *per se* difficult task because of the reduced base of
11 support in comparison to the common double-leg stance (DLS) condition. Usually,
12 when the support base is reduced, a precaution strategy consisting of moving
13 forward the center of mass is adopted to avoid falling backward. This condition
14 may justify the existence of the hip/trunk balance control strategy (characterized by
15 the activation of the LH, MH, GMD, LDL, and LDR muscles) adopted by all the
16 volunteers during the SLS condition. It is reasonable to think that the participants
17 of the present study used the hip/trunk control strategy (W_3) to compensate for the
18 ankle dorsiflexion used to move forward the center of mass to manage the reduced
19 base of support. It should be also mentioned that in a condition of quiet stance the
20 co-existence of the hip control strategy (W_2) and the ankle control strategy (W_1 and
21 W_4) has been reported [196], highlighting that the two strategies are not different
22 entities, but one predominates depending upon the task and conditions of the
23 environment.

24 Previous literature has reported an increase in the knee joint displacement
25 during SLS conditions when the difficulty of the task increased, i.e., the surface
26 became less stable [170]. Thus, in this study, the knee synergy was probably used
27 when the ankle synergy was not effective for the maintenance of balance, but the
28 condition did not require yet the involvement of the hip or the back synergies. These
29 results highlight the fine coordination between ankle muscles and quadriceps
30 muscle. This was especially observed in the closed-eyes condition when the lack of
31 visual information led to an increase in the difficulty of the task. In fact, it was
32 observed a significantly lower activation of the SOL and LGS muscles when the
33 knee-dominant synergy was used.

34 The results presented in this chapter show that 4 muscle synergies are needed
35 during single-leg stance during eyes open (EO) as well as eyes closed (EC)
36 conditions. In addition, there is no difference in the average recruitment levels
37 between the EO and EC conditions, except for the hip/trunk muscle synergy, which
38 showed a decrease in activation during the EC condition compared to the EO
39 condition. At the same time, an increase in the ankle balance control strategy was
40 found in the EC condition compared to the EO condition. However, despite some
41 differences in the EC condition compared to the EO condition, the number of
42 synergies used is the same between the conditions, as well as the average
43 recruitment levels. This is in accordance with previous literature reporting the
44 stability of muscle synergies adopted between tasks with different visual feedback
45 conditions [187,197]. It has been shown that the lack of visual feedback does not

1 affect muscle synergies, since standing postural control mostly relies on
2 proprioceptive feedback [187,197]. Furthermore, the results of previous
3 investigations show that proprioceptive disturbance, but not visual disturbances,
4 affected muscle synergies [187] and increases body sway [197]. Regarding the
5 results of the present chapter, it can be thus speculated that the lack of visual
6 information affects the modulation of muscle activations without altering the
7 number and composition of the muscle synergies adopted.

8 Moreover, the results of this chapter suggest that muscle synergies could be not
9 exclusively managed throughout a feedforward control, but can be modulated with
10 a feedback control based on the signals arising from sensory receptors, with the aim
11 to correct movement errors which may occur in some circumstances. It is likely to
12 think that the maintenance of the single-limb stance in this study was controlled
13 with pre-programmed muscles synergies. However, the difficulty of the task
14 leading to continuous losses and recovery of balance, probably need a continuous
15 movement correction based on a feedback control relying on information arising
16 from sensory receptors. Animal studies have reported organized patterns of muscles
17 activations in response to focal stimulation of the spinal cord [11], thus suggesting
18 that a feedback control may be launched at spinal level in response to specific
19 sensory stimuli to modulate the centrally organized synergy recruitment. It is likely
20 to think that similar patterns may regulate muscle synergies also in humans.

21 To the best of the author's knowledge, this is the first study investigating muscle
22 synergies deputed to the maintenance of posture during a single-leg stance task, in
23 an EO and EC condition. Due to the large use of this kind of task in clinical practice,
24 both for rehabilitation and functional assessment, as well as in sport practice for
25 training and testing, the results of the present study give important information on
26 motor control of this kind of task in healthy individuals. Future studies should
27 investigate muscle synergies also in other populations to investigate the effects of
28 orthopedic and neurologic pathologies on muscle synergies, as well as the effect of
29 rehabilitation and training.

30 The main limitation of this study is that only healthy young individuals were
31 recruited, and thus the results cannot be generalized to all healthy individuals.
32 Future studies should identify muscle synergies used for single-leg stance also in
33 other age groups. A second limitation of the study is that muscle synergies for the
34 transition between double- and single-limb stance (and vice versa) were not
35 analyzed, thus the results of the present study have to be considered exclusively for
36 steady single-limb stance tasks.

37 In conclusion, the single-leg stance is featured by four major muscle synergies,
38 two ankle-dominant, one knee-dominant, and one hip/trunk-dominant. The lack of
39 visual feedback did not affect the number of synergies used. In general, an increase
40 of activation of the ankle muscles and a decrease in the recruitment of the hip/trunk
41 synergy was observed in the absence of visual information in comparison to the
42 normal vision condition. Future studies should investigate muscle synergies during
43 single-leg stance also in other age groups, and it seems of high clinical relevance to
44 investigate synergies on orthopedic and neurologic patients to address clinical
45 practice and rehabilitation interventions.

1 Chapter 5

2 Conclusions and Perspectives

3 With this doctoral thesis, the author endeavored to provide a deeper insight into
4 the neuromuscular control of human movements, focusing on two essential daily
5 life activities: human locomotion and balance control.

6 In the first part of the thesis, it was assessed the modular organization of the
7 CNS during cyclical movements (i.e., human locomotion), using the widely used
8 muscle synergy theory. The first step was to develop a reliable and accurate method
9 for extracting muscle activation intervals during cyclical movements to be used as
10 a pre-processing step before muscle synergy extraction (**A Deep Learning**
11 **Approach for Muscle Activity Detection**). First, the applicability of the proposed
12 approach was assessed considering simulated sEMG data and, then, its performance
13 was compared against other standard muscle activity detectors using real sEMG
14 signals both in physiological and pathological conditions. The second step was to
15 methodologically assess two of the main weaknesses of the muscle synergy theory:
16 the sEMG pre-processing steps before factorization and the selection of the optimal
17 number of muscle synergies (i.e., the factorization rank). Considering the sEMG
18 pre-processing weakness, a novel pre-processing approach based on the extraction
19 of the principal activations before muscle synergy extraction was proposed and
20 validated on real sEMG data, allowing a more interpretable assessment of the
21 modular organization of the CNS during a walking task without any loss of
22 information (**Methodological Issue - Muscle Synergies Extracted Using**
23 **Principal Activations**). Considering the synergy number selection weakness, a
24 novel method, based on multi-criteria decision analysis, for the selection of the
25 optimal number of muscle synergies was proposed to overcome the limitations of
26 the standard *VAF*-based approaches. Following an approach similar to the one
27 proposed for the muscle activity detector, the applicability of the approach was first
28 assessed considering simulated sEMG data and, then, its performance was
29 compared against other standard *VAF*-based approaches using real sEMG signals
30 both in physiological and pathological conditions (**Methodological Issue – Multi-**

1 **Criteria Decision Analysis for Selecting the Optimal Number of Muscle** 2 **Synergies).**

3 In the second part of the thesis, instead, it was assessed the modular
4 organization of the CNS during non-cyclical movements (i.e., single-leg stance)
5 through the muscle synergy theory. As in the first part of the thesis, the first step
6 was to develop a reliable and accurate method for the segmentation of the sEMG
7 time-instants relative to a “*well-balanced*” single-leg stance or an “*unbalanced*”
8 single-leg stance to be used as inputs of the muscle synergy extraction algorithm
9 (**Methodological Issue – Muscle Synergy Assessment during Single-Leg**
10 **Stance**). Subsequently, the consolidated method was applied to the analysis of the
11 modular organization of single-leg stance with and without visual feedback
12 considering a healthy sample population (**Application - Influence of Visual**
13 **Feedback on Muscle Synergies**).

14 In the following paragraphs, the practical implications of the studies presented
15 in this doctoral thesis will be discussed, also addressing their limitations and the
16 scientific perspectives that emerged from this work.

17 **Muscle Activity Detection**

18 Determining the onset and offset time-instants of muscle activations during
19 human movements is of great interest in different research fields including gait
20 analysis, motor rehabilitation, sport science, myoelectric control of prostheses,
21 human-machine interaction, design of biofeedback systems, and pre-processing of
22 muscle synergy extraction. More specifically, the accurate temporal analysis of
23 muscle activation in terms of burst onset, duration of the activation interval, and
24 burst offset, can be useful in the assessment of the altered locomotion patterns of
25 orthopedic and neurological patients.

26 The LSTM-based muscle activity detector presented in the first study of this
27 thesis can be considered a valuable tool in all the applications requiring an accurate
28 and effective recognition/distinction of muscle activity from background noise. In
29 particular, it significantly outperforms previous detectors (such as the statistical
30 double threshold detector and Teager-Kaiser Energy Operator detector) in terms of
31 detection accuracy, it directly works on “raw” sEMG signals (does not require any
32 additional input parameters), and it is robust even when applied to signals with low
33 to medium signal-to-noise ratio.

34 Future works will be focused on testing the proposed approach also on sEMG
35 signals acquired from subjects with musculoskeletal or neurodegenerative disorders
36 other than Parkinson’s disease and on developing a novel formulation of this
37 approach for real-time muscle activity detection.

38 This novel muscle activity detector is currently being applied in a number of
39 projects investigating muscle activity during walking. First, thanks to the
40 collaboration with the Stereotactic and Functional Neurosurgery Unit of the
41 University of Turin (Italy), it is being applied to assess the effect of Deep Brain
42 Stimulation (DBS) neurosurgery on the muscle activity of patients affected by
43 Parkinson’s disease during walking by analyzing the onset, the offset, and the

1 duration of each muscle activation within a gait cycle. Second, the detector is
2 currently being used as a pre-processing step before the application of the CIMAP
3 (Clustering for Identification of Muscle Activation Patterns) algorithm, allowing
4 the extraction of principal and secondary muscle activation intervals from sEMG
5 signals during cyclic movements. Finally, thanks to the collaboration with the
6 Department of Control and Computer Engineering (DAUIN) of Politecnico di
7 Torino, the proposed muscle activity detector is being used to control a haptic
8 exoskeleton for postural rehabilitation of patients with locomotor impairments.
9 More specifically, the muscle activation intervals extracted through the LSTM-
10 MAD from sEMG signals are currently being used to predict the angular positions
11 of leg joints and control the exoskeleton.

12 **Muscle Synergies during Walking**

13 Human locomotion is one of the most widely studied and complex motor tasks,
14 due to the multiple degrees of freedom of the skeletal muscle system, the several
15 biomechanical functions carried out during each gait cycle, and the high cycle-by-
16 cycle variability of muscle activation intervals. Several studies already
17 demonstrated that human locomotion can be modeled by a small set of muscle
18 synergies characterized by specific biomechanical functions. Despite the growing
19 interest in the assessment of the modular organization of the CNS during walking
20 through the muscle synergy theory, there is a lack of standards and common
21 procedures for the sEMG pre-processing steps and the extraction of muscle
22 synergies (e.g., selection of the optimal number of muscle synergies).

23 In the present doctoral thesis, these two weaknesses of the muscle synergy
24 theory were assessed by presenting two novel methodological approaches that
25 could provide a deeper insight into human motor control.

26 Considering the former, an innovative pre-processing technique for muscle
27 synergy extraction was proposed combining Statistical Gait Analysis (SGA) and
28 CIMAP algorithm, allowing for obtaining the principal and secondary activations.
29 Results presented in this thesis demonstrated that the extraction of the principal
30 activations only can be successfully used as a pre-processing step before muscle
31 synergy extraction, allowing a more interpretable assessment of the modular
32 organization of the CNS during a walking task in both physiological and
33 pathological populations without any loss of information. Even if long-lasting
34 signal recordings (at least 2-3 minutes are required to obtain a sufficient number of
35 typical gait cycles) are needed to be able to properly compute the principal
36 activations, this requirement does not limit the feasibility and applicability of the
37 proposed approach to pathological populations. Indeed, gait analysis is commonly
38 used to quantitatively assess patients' locomotion performance only in those
39 patients able to independently walk, for some minutes, without walking aids or
40 external supports. One possible limitation of the present approach is that only
41 principal activation intervals were used to extract muscle synergies, since the
42 assessment of the variability expressed by the muscle synergy analysis could be
43 particularly significant when analyzing sEMG data acquired from subjects affected

1 by movement disorders. To overcome this limitation, future studies will focus on
2 muscle synergies extracted from the secondary activations (SAs), i.e., those muscle
3 activations that occur only in some gait cycles and could have an auxiliary function
4 in motor control, such as to provide corrections to motion and body segment
5 posture.

6 Considering the latter weakness of the muscle synergy theory, a novel
7 ELECTRE III approach for choosing the optimal number of muscle synergies was
8 proposed and validated on both simulated and real sEMG data. Even if this
9 approach does not outperform the other tested threshold-independent method (i.e.,
10 elbow VAF method), several are the potential advantages of the ELECTRE III
11 approach compared to the commonly used *VAF*-based threshold methods, such as
12 the use of minimization and maximization approaches rather than an arbitrary
13 threshold, the use of a multi-criteria decision-making approach based on different
14 parameters computed from the muscle synergy results (not only *VAF* values), and
15 the applicability to a variety of different motor tasks. Future works will be focused
16 on optimizing the proposed approach by considering additional evaluation criteria
17 or changing their importance weights and on applying the ELECTRE III approach
18 to the studies described in Chapter 3 to assess the robustness of the proposed results
19 with respect to the approach for selecting the optimal number of muscle synergies.

20 Both these approaches are currently being applied in two different projects
21 investigating motor control strategies of pathological populations during walking.
22 First, thanks to the collaboration with the Stereotactic and Functional Neurosurgery
23 Unit of the University of Turin (Italy), they are being applied to assess the changes
24 in motor control strategies of patients affected by Parkinson's Disease (PD) due to
25 Deep Brain Stimulation (DBS) neurosurgery. More specifically, this study aims at
26 assessing differences in the number and composition of the muscle synergies
27 extracted from 30 PD patients during a 5-minutes lasting walking task at a self-
28 selected speed before and after the DBS neurosurgery. Second, the effect of
29 cognitive dual-task (i.e., verbal fluency and music-based dual tasks) while walking
30 on PD motor control strategies is currently being studied through the muscle
31 synergy theory considering the same sample population above-mentioned.

32 **Muscle Synergies during Single-Leg Stance**

33 The ability to maintain the single-leg stance is essential during daily living
34 activities, as a single task as well as a component of other more complex tasks, such
35 as walking and running. It is a simple but challenging condition for balance control
36 and for this reason it is widely used for training and rehabilitation. In research and
37 clinical practice, it is widely used as a testing task as it allows to quantify balance
38 alterations and deficits of the single lower limb otherwise concealed during the
39 performance of double leg tasks. In the last years, muscle synergy assessment
40 during different balance conditions (e.g., single-leg stance) was mainly focused on
41 the evaluation of balance recovery after a "controlled" perturbation, rather than on
42 the maintenance of the balance condition itself. Thus, the segmentation approach
43 proposed in this thesis aims at providing a useful tool for the assessment of the

1 motor control strategies adopted to maintain the single-leg stance, distinguishing
2 between “*well-balanced*” and “*unbalanced*” single-leg stance epochs.

3 Results demonstrated that the proposed segmentation approach is robust
4 concerning the selection of the segmentation threshold and can be successfully used
5 as a pre-processing step before muscle synergy extraction, allowing a better
6 assessment of the modular organization of the central nervous system during the
7 maintenance of the single-leg stance. Moreover, it was applied for the assessment
8 of the effect of visual feedback on muscle synergies extracted from healthy subjects
9 during unipedal stance, revealing a high consistency of motor controls strategies
10 adopted by healthy subjects with and without visual feedback.

11 One of the advantages of the proposed approach is the possibility to assess
12 motor control strategies used by subjects to maintain the balance condition, instead
13 of evaluating balance recovery after a “controlled” perturbation. Instead, the fact
14 that the proposed method was validated on a sample population of young healthy
15 subjects that can maintain the single-leg stance for at least 30 s may represent a first
16 limitation of the study, since it may be difficult to obtain similar task durations
17 considering elderly or pathological populations affected by severe balance
18 impairments. Moreover, different muscle synergy results might be obtained in
19 populations with diminished equilibrium skills or severe balance impairments.

20 Currently, the proposed muscle synergy extraction approach is being applied in
21 a project that aims at assessing the modular organization of CNS during single-leg
22 stance in pathological conditions. More specifically, thanks to the collaboration
23 with the Rizzoli Orthopedic Institute of Bologna (Italy), it is being applied to assess
24 muscle synergies extracted during single-leg stance tasks from patients affected by
25 Chronic Ankle Instability (CAI).

1 **Appendix A**

2 **Extraction of Principal and** 3 **Secondary Activations through** 4 **CIMAP**

5 **A.1 Introduction**

6 Instrumented gait analysis is a widely used approach to quantitatively and non-
7 invasively assessed the physiological and pathological motor functions during
8 human locomotion [198]. In particular, the analysis of muscle activation patterns is
9 of great interest, in research as well as in clinical practice, to study the altered
10 locomotion patterns of orthopaedic and neurological patients or to assess the
11 efficacy of rehabilitation programs. Muscle activation patterns are computed from
12 muscles' electrical activity acquired by means of surface electromyography
13 (sEMG) during cyclic movements. Then, for each movement cycle (e.g., each gait
14 cycle), muscle activation patterns are computed determining the timing of muscle
15 activations (burst onset, duration, and end). However, the high cycle-to-cycle
16 variability of sEMG activations intervals makes it difficult to obtain a synthetic and
17 clear representation of the patient's motor functions [98,101]. **Figure A.1**
18 represents an example of sEMG signals acquired from an 11-years-old child during
19 three different gait cycles from the Lateral Hamstring (LH) muscle.

20 To overcome this limitation, Statistical Gait Analysis (SGA) [60] and
21 Clustering for Identification of Muscle Activation Patterns (CIMAP) [103,104,199]
22 algorithms were recently introduced. Statistical Gait Analysis allows an easier
23 interpretation and comparison of muscle activation intervals through the acquisition
24 and processing of a large number of gait cycles. CIMAP algorithm, indeed, enables
25 the grouping into clusters of gait cycles characterized by similar muscle activation
26 patterns. By combining SGA and CIMAP algorithms, principal activations (PAs)
27 and secondary activations (SAs) can be computed from the sEMG signals
28 [24,65,106]. Principal activations are defined as those muscle activations that are

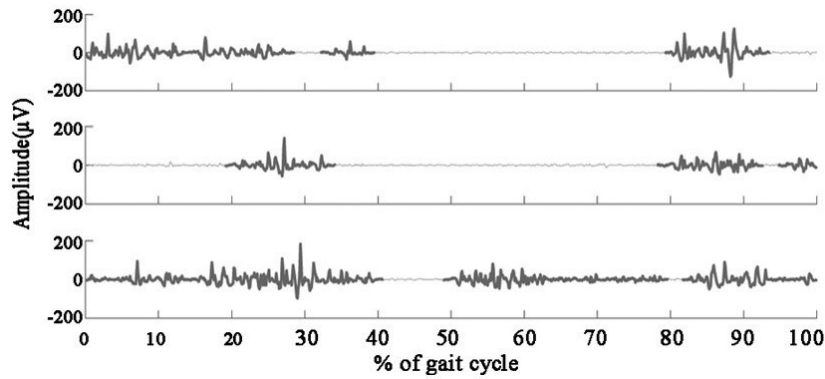


Figure A.1 Example of sEMG signals acquired from an 11-years-old healthy child during gait from Lateral Hamstring (LH) muscle.

1 necessary for the execution of a specific motor task and they describe the essential
 2 contribution of a specific muscle to the movement. The concept of PAs is
 3 complementary to the one of secondary activation, which are muscle activations
 4 that occur only in some gait cycles and could have an auxiliary function in motor
 5 control (e.g., to provide corrections to motion and body segment posture).

6 The extraction of principal and secondary muscle activations as a pre-
 7 processing step may significantly improve the analysis of sEMG signals, reducing
 8 the cycle-to-cycle variability and simplifying the understanding of muscle
 9 activation patterns both in physiological and pathological conditions.

10 This chapter aims at briefly describing the processing steps needed to compute
 11 the PAs and SAs from sEMG signals.

12 **A.2 Extraction of Principal and Secondary Activations**

13 Before computing principal and secondary activations, sEMG signals were pre-
 14 processed as it follows:

- 15 i. Muscle activation intervals were detected through the LSTM-MAD
 16 algorithm described in 0. The detector's output is computed as a binary
 17 mask that was set equal to 1 in correspondence of the sEMG time-
 18 instants classified as muscle activity and to 0 otherwise (background
 19 noise)
- 20 ii. Time-segmentation of muscle activation intervals into gait cycles using
 21 foot-switch signal [108]
- 22 iii. Time-normalization of each gait cycle into 1000 time points.

23 Principal and secondary activations were computed from the muscle activation
 24 intervals, separately for each muscle, applying the optimized version of the CIMAP
 25 algorithm [104]. **Figure A.2** shows an example of the application of the CIMAP
 26 algorithm to sEMG signals acquired from a healthy subject during gait from
 27 Peroneus Longus (PL) and Gluteus Medius (GMD) muscles.

28 The CIMAP algorithm, based on agglomerative hierarchical clustering, groups
 29 together the gait cycles sharing similar muscle activation intervals. Each significant
 30 cluster is characterized by an element (called prototype) that is representative of all

1 the elements (gait cycles) belonging to the same cluster. Prototypes are computed
 2 as the median value of the elements belonging to the same cluster. **Figure A.2A**
 3 represents in blue the time-normalized activation intervals for each gait cycle,
 4 grouped in significant clusters sharing similar activation intervals. Each orange
 5 interval represents the prototype of a significant cluster.

6 Starting from a number of clusters equals to the number of gait cycles analyzed
 7 (each cluster contains a single element), the CIMAP algorithm iteratively merges
 8 the two “closest” clusters until a single cluster, containing all the gait cycles, is
 9 obtained. The complete linkage method is used to select the two “closest” clusters
 10 to be merged. The implemented linkage method uses the farthest distance between
 11 every pair of elements in the two considered clusters as merging criterion. During
 12 the linkage process, the distance between each couple of elements is assessed using
 13 the Manhattan distance. Hence, the dendrogram is constructed.

14 The final number of clusters (cutoff rule) was selected to achieve:

- 15 i. Clusters characterized by a comparable number of elements (C) within
 16 a significant cluster
- 17 ii. Small intra-cluster variability (ICV), computed as the Euclidian
 18 distance between each element of the cluster and the corresponding
 19 cluster’s prototype.

20 More specifically, the optimized version of CIMAP algorithm defines a cutoff
 21 rule based on three different criteria. First, the differences of inter-cluster Euclidian
 22 distances ($Diff_k$) between two consecutive dendrogram iterations (k) are
 23 computed. Second, three cutoff points are computed based on the following three
 24 criteria:

- 25 i. First dendrogram iteration k in which $Diff_k \geq mean(Diff_k)$, where
 26 $mean(Diff_k)$ is the average of $Diff_k$ computed over all the iterations
- 27 ii. First dendrogram iteration k in which $Diff_k \geq mean(Diff_k) + 1 \cdot$
 28 $std(Diff_k)$, where $mean(Diff_k)$ and $std(Diff_k)$ are the average and
 29 the standard deviation of $Diff_k$ computed over all the iterations,
 30 respectively
- 31 iii. Starting from the last dendrogram iteration and stepping backwards, the
 32 first iteration in which $Diff_k$ series stop decreasing monotonically
 33 (after a moving average filtering step).

34 Finally, since the three cutoff criteria may result in different numbers of final
 35 clusters, the best cutoff was automatically identified using an index that takes into
 36 account both the intra-cluster variability (ICV) and the number of elements within
 37 each significant cluster (C_i), as described in (A.1):

$$Cutoff = \frac{\sum_{i=1}^n ICV_i / n}{\sum_{i=1}^n |C_i|} \quad (\text{A.1})$$

38 where n represents the number of significant clusters, $|C_i|$ is the number of cycles
 39 belonging to the i -th cluster, and ICV_i represents the intra-cluster variability of the

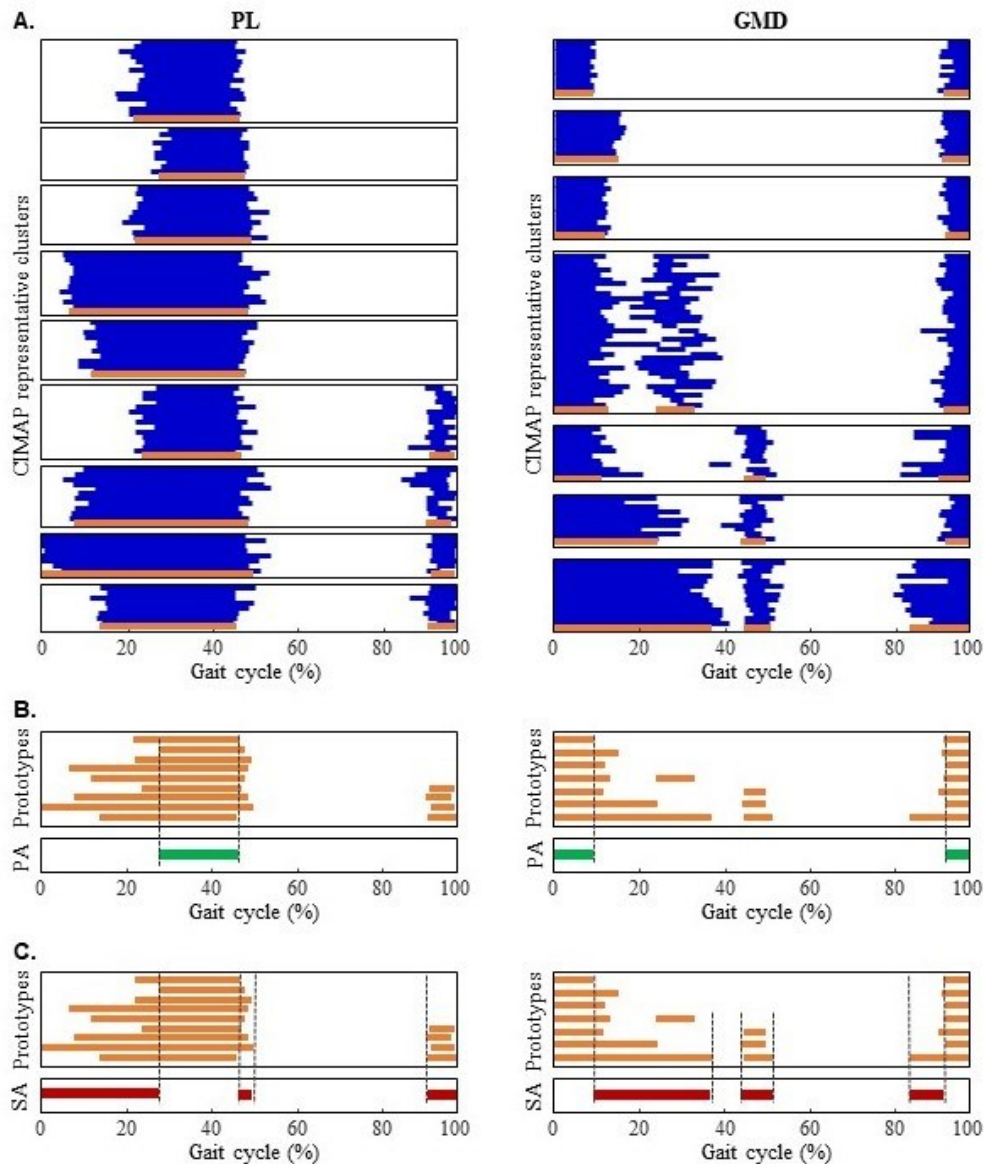


Figure A.2 Example of application of the CIMAP algorithm to sEMG signals acquired from a healthy subject during gait from PL (left) and GMD (right) muscles. (A) Blue intervals represent the cluster elements (muscle activation intervals computed through LSTM-MAD algorithm) normalized into 1000 time points with respect to the gait cycle duration, while orange intervals represent the prototypes of each cluster (computed as median of the elements that belong to the same cluster). (B) Principal activation (PA) is represented in green and is defined as the intersection of all the cluster's prototypes. (C) Secondary activation (SA) is represented in red and is defined as all the time-instants that are classified as muscle activity and are not PAs. Reprinted from “How to Improve Robustness in Muscle Synergy Extraction” by Ghislieri *et al.*

1 i -th cluster. Low values of the *Cutoff* index correspond to clusters with a small
 2 intra-cluster variability and a high number of elements belonging to each significant
 3 cluster.

4 After the definition of the final number of clusters, the principal and secondary
 5 activations are computed, separately for each muscle, from the prototypes of the
 6 significant clusters. More specifically, the PAs are defined as the intersection of all
 7 the significant clusters' prototypes, while SAs are defined as all the time-instants
 8 classified as muscle activations, but excluded from the PAs, since they occur only

1 in some gait cycles. For each observed muscle, a single principal (or secondary)
2 activation interval was obtained and defined as a 1000-samples binary mask.

3 **Figure A.2B** and **Figure A.2C** depict how the PAs and the SAs are obtained
4 from the significant clusters' prototypes, respectively.

5 For each gait cycle, the extraction of the PAs (or SAs) from the original sEMG
6 signals of a specific muscle was performed by time-windowing all the time-
7 normalized gait cycles through the corresponding binary mask that was set equals
8 to 1 in correspondence of the time-instants defined as PAs (or SAs) and to 0
9 otherwise (no PAs or SAs computed).

10 **Figure A.3** shows an example of principal and secondary activation extraction
11 performed on PL and GMD muscles acquired from a healthy subject during a single
12 gait cycle.

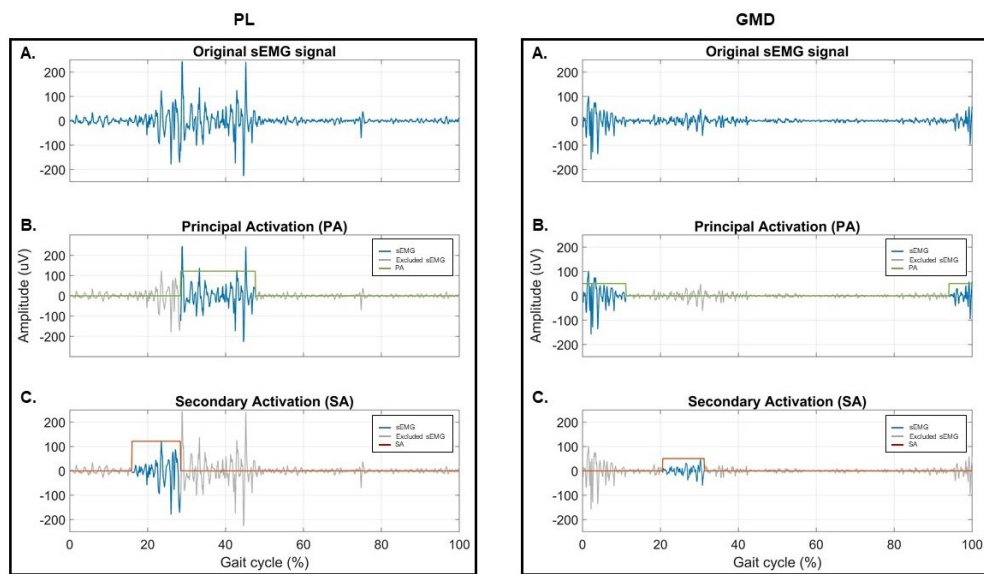


Figure A.3 Extraction of principal and secondary activations from PL (left) and GMD (right) muscles of a healthy subject during a walking task. (A) Original sEMG signals time-normalized with respect to the duration of the gait cycle. (B) Extraction of the principal activation (in green) from the original sEMG signal with the indication of the excluded sEMG time-instants in gray. (C) Extraction of the secondary activation (in red) from the original sEMG signals with the indication of the excluded sEMG time-instants in gray.

13

1 **Appendix B**

2 **List of Abbreviations**

3 The following abbreviations are used in the manuscript:

4		
5	ANOVA	Analysis of Variance
6	CAI	Chronic Ankle Instability
7	CBA	Cost-Benefit Analysis
8	CIMAP	Clustering for Identification of Muscle Activation Pattern
9	CNS	Central Nervous System
10	DBS	Deep Brain Stimulation
11	ELECTRE	ELimination Et Choix Traduisant la REalité
12	GMD	Gluteus Medius
13	ICV	Intra-Cluster Variability
14	LD _L	Left Longissimus Dorsii
15	LD _R	Right Longissimus Dorsii
16	LGS	Lateral Gastrocnemius
17	LH	Lateral Hamstring
18	LSTM	Long-Short Term Memory neural network
19	LSTM-MAD	Long-Short Term Memory for Muscle Activity Detection
20	MAE	Mean Absolute Error
21	MCDA	Multi Criteria Decision Analysis
22	MCDM	Multi Criteria Decision-Making
23	MH	Medial Hamstring
24	PAs	Principal Activations
25	PB	Peroneus Brevis
26	PL	Peroneus Longus
27	RF	Rectus Femoris
28	RMS	Root Mean Squared
29	RNN	Recurrent Neural Network
30	SAs	Secondary Activations
31	sEMG	Surface Electromyography signals
32	SGA	Statistical Gait Analysis
33	SNR	Signal-to-Noise Ratio

1	SOL	Soleus
2	TA	Tibialis Anterior
3	TKEO	Teager-Kaiser Energy Operator
4	UPDRS	Unified Parkinson's Disease Rating Score
5	VAF	Variance Accounted For
6	VL	Vastus Lateralis
7	VM	Vastus Medialis

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