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Investigating reduction of dimensionality during single-joint elbow movements: a case study on muscle synergies

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Running title: Muscle synergies for elbow movements

34 **Abstract**

35 A long standing hypothesis in the neuroscience community is that the CNS generates the
36 muscle activities to accomplish movements by combining a relatively small number of
37 stereotyped patterns of muscle activations, often referred to as “muscle synergies”. Different
38 definitions of synergies have been given in the literature. The most well-known are those of
39 synchronous, time-varying and temporal muscle synergies. Each one of them is based on a
40 different mathematical model used to factor some EMG array recordings collected during the
41 execution of variety of motor tasks into a well-determined spatial, temporal or spatio-
42 temporal organization. This plurality of definitions and their separate application to complex
43 tasks have so far complicated the comparison and interpretation of the results obtained across
44 studies, and it has always remained unclear why and when one synergistic decomposition
45 should be preferred to another one. By using well-understood motor tasks such as elbow
46 flexions and extensions, we aimed in this study to clarify better what are the motor features
47 characterized by each kind of decomposition and to assess whether, when and why one of
48 them should be preferred to the others. We found that three temporal synergies, each one of
49 them accounting for specific temporal phases of the movements could account for the
50 majority of the data variation. Similar performances could be achieved by two synchronous
51 synergies, encoding the agonist-antagonist nature of the two muscles considered, and by two
52 time-varying muscle synergies, encoding each one a task-related feature of the elbow
53 movements, specifically their direction. Our findings support the notion that each EMG
54 decomposition provides a set of well-interpretable muscle synergies, identifying reduction of
55 dimensionality in different aspects of the movements. Taken together, our findings suggest
56 that all decompositions are not equivalent and may imply different neurophysiological
57 substrates to be implemented.

58

59 **Keywords:** Muscle Synergies, Non-Negative Matrix Factorization, EMG, Elbow Rotations,
60 Dimensionality Reduction, Triphasic Pattern

61

62 Introduction

63 A large amount of studies have provided in the last two decades evidence according to
64 which the central nervous system (CNS) generates the muscle patterns necessary to achieve a
65 desired motor behaviour by combining a relatively small number of stereotyped spatial and/or
66 temporal patterns of muscle activation, often referred to as “muscle synergies” (Bizzi et al.,
67 2008). An appeal of this framework is that it suggests that the CNS may control movement
68 execution through a relatively small number of degrees of freedom.

69 Different conceptual definitions of muscle synergies have been given in the literature.
70 These in practice translate into different mathematical models used to factor
71 electromyographic (EMG) array recordings collected during the execution of variety of motor
72 tasks into different kinds of temporal, spatial or spatio-temporal organizations. Invariant
73 temporal components (or “temporal synergies”, see Ivanenko et al., 2004, 2005; Dominici et
74 al., 2011; Chiovetto et al., 2010, 2012) are defined as temporal muscle activation profiles that
75 can be simply scaled and summed together to reconstruct the actual activity of each muscle.
76 “Synchronous synergies” (Cheung et al., 2005, 2009, 2012; Ting and Macpherson, 2005;
77 Torres-Oviedo and Ting, 2007, 2010) are stereotyped co-varying groups of muscle
78 activations, with the EMG output specified by a temporal profile defining the timing of each
79 synergy during the task execution. “Time-varying synergies” (d’Avella et al., 2003, 2006,
80 2008, 2011) are genuine spatiotemporal patterns of muscle activation, with the EMG output
81 specified by the amplitude and time lag of the recruitment of each synergy.

82 Typically, previous studies about muscle synergies focused on a given decomposition
83 that was then used to investigate potential functions of muscle synergies in complex motor
84 tasks involving a large number of degrees of freedom (dof). Each of these decompositions
85 has been used successfully to identify common physiologically important factors of muscle
86 activity (Ivanenko et al., 2005; d’Avella et al., 2006; Cheung et al., 2005). The existence in
87 the literature of multiple definitions of muscle synergies and their separate application to
88 complex tasks complicates however the comparison and interpretation of the results obtained
89 across studies, and it is not always clear why and when one synergistic decomposition should
90 be preferred to another one. We propose instead here that the systematic study of the
91 application of all these decompositions to the same and simple dataset for which the
92 mechanical action of each muscle contraction is well-known would greatly help to build
93 intuition about the merit and functional interpretation of each synergistic decomposition. This
94 would moreover be beneficial to the interpretation and comparison of different studies. We
95 thus considered the extreme case of single-joint elbow movements, characterized by one
96 kinematic dof, two antagonist muscles (biceps and triceps) and four experimental tasks
97 (flexions and extensions along both the horizontal and vertical directions). We applied
98 systematically decompositions into synchronous, time varying and temporal synergies of
99 EMG data recorded during this elementary and well documented motor task (see Berardelli et
100 al., 1996 for a review), whose biomechanical and neurophysiological bases were studied
101 intensively (Gottlieb et al., 1995; Shapiro et al., 2005).

102 Our findings support the notion that each EMG decomposition provides a set of well-
103 interpretable muscle synergies, identifying reduction of dimensionality in different aspects of
104 the movements. Each temporal synergy indeed conveys information about a specific temporal
105 phase of the movement (acceleration, deceleration and stabilization). Synchronous and time-
106 varying synergies instead encode respectively the simultaneous and coordinated actions of
107 specific groups of muscles aiming to achieve a specific action goal and a task-related feature

108 of the elbow movements (specifically the direction of motion). Taken together, our findings
109 suggest that all decompositions are not equivalent and may imply different
110 neurophysiological substrates to be implemented.

111 **Material and methods**

112 **Subjects**

113 Eight healthy right-handed subjects (7 males, 1 female, ages 29 ± 4 years, mass 74 ± 9
114 kg, height 1.77 ± 0.07 m), participated voluntarily to the experiments that were all performed
115 at the Robotics, Brain and Cognitive Sciences Department at Italian Institute of Technology
116 (IIT) in Genoa (Italy). All subjects were in good health condition and had no previous
117 history of neuromuscular disease. The experiment conformed to the declaration of Helsinki
118 and informed consent was obtained from all the participants according to the protocol of the
119 ethical committee of IIT.

120 **Protocol**

121 Subjects sat on a chair with their back straight and perpendicular to the ground. They
122 were asked to perform one-shot 90 degrees elbow rotations between two reference points
123 along either a vertical and a horizontal plane (Figure 1). A total of four experimental
124 conditions were thus studied (vertical flexion, VF, vertical extension, VE, horizontal flexion,
125 HF and horizontal extension, HE). For movements along the vertical direction, the two
126 reference points were located in a vertical plane, placed laterally at approximately 10 cm
127 from the subject's movement plane. To this aim, we used a wooden hollow frame containing
128 1.5 cm-spaced thin vertical fishing wires to which fishing leads indicating the requested
129 fingertip initial position were attached. One reference point coincided with the subject's
130 fingertip position in the vertical plane when the arm was completely relaxed and extended
131 vertically with the index fingertip pointing at the ground (vertical position number 1, or VP1).
132 The second point coincided with the subject's fingertip position in the vertical plane when,
133 starting from VP1, the elbow was rotated of about 90 degrees so that at the end the forearm
134 was parallel to the ground (vertical position number 2, or VP2). The positions of the fishing
135 leads were adjusted for each subject before the initiation of the experiment, based on the
136 subject's upper arm and forearm lengths. For vertical elbow flexion subjects rotated the
137 elbow so as to move their index finger from VP1 to VP2. On the contrary, during vertical
138 elbow extension they had to move the fingertip from VP2 back to VP1. For rotation along the
139 horizontal plane subjects sat in front of a table. One reference point on the table coincided
140 with the horizontal location of the index fingertip when the upper-arm was kept horizontal
141 with respect to the ground and perpendicular to the coronal plane and the forearm flexed of
142 about 90° with respect to the upper-arm (horizontal position 1, or HP1). The second
143 reference point coincided with the fingertip location when the whole arm was completely
144 extended horizontally in front of the subjects and perpendicular to the coronal plane
145 (horizontal position 2, or HP2). After that (for each subject) HP1 and HP2 were identified,
146 their location was marked on the table by means of two small squared pieces of colored tape.
147 The table plane laid 10 cm below the plane of rotation of the arm, avoiding thus to disturb the
148 accomplishment of the movement. For horizontal elbow flexion subjects had to rotate the
149 elbow so as to move their index finger from HP1 to HP2. On the contrary, during horizontal
150 elbow extension they had to move the fingertip from HP2 back to HP1. Subjects were
151 always asked to perform fast movements (mean velocities and average peak velocities are
152 reported in Table 1 for each subject and condition). They performed 20 elbow flexion and 20

153 extensions for each plane orientation. During the experiment the wrist joint was frozen by
154 means of two light and small sticks attached to the distal part of the forearm and the proximal
155 part of the hand. At any trial repetition subjects put their index finger on the starting position.
156 The experimenter started data acquisition and gave the “go” signal. The subjects performed
157 the movement after the “go” signal and stopped on the target for about a second. Data
158 acquisition stopped automatically after two seconds. At the end of the trial the subject
159 assumed with his arm a relaxing position until the beginning of the next trial. After 20 trials
160 subjects took a pause of about 3 minutes to avoid fatigue.

161 **Apparatus**

162 During trials’ execution kinematic data were recorded by means of a Vicon (Oxford,
163 UK) motion capture system. Six passive markers were attached on subjects’ right arm (the
164 acromion process, lateral epicondyle of the humerus, the styloid process and the tip of the
165 index finger) and head (external canthus of the eye and auditory meatus). Electromyographic
166 activity of biceps brachii (Bic) and triceps longus (Tri) was monitored by means of an Aurion
167 (Milan, Italy) wireless electromyographic system. Impedance between the surface electrodes
168 was always checked not to exceed 5 K Ω : in the case of higher values, skin was rubbed by
169 means of an abrasive sponge in order to decrease it. EMG data were amplified (gain of 1000),
170 band-pass filtered (10 Hz high-pass and 1 KHz low-pass) and digitized at 1000 Hz.

171

172

173 **Data pre-processing**

174 Data were analysed off-line using customized software written in Matlab (Mathworks,
175 Natick, MA). Kinematic data were low-pass filtered (Butterworth filter, cut-off frequency of
176 20 Hz). The angular displacement of the elbow was computed starting from the markers’
177 spatial positions. Elbow angular velocity of rotation was obtained by numerical
178 differentiation of the angular position. Mean and peak angular velocities were computed for
179 each trial. The mean velocity was computed as the mean value of the angular velocity over
180 the movement duration. The time instants of movement initiation (t_0) and end (t_f) were
181 defined respectively as the instants at which the bell-shaped angular velocity profile of the
182 elbow exceeded and dropped below 5% of its peak value. For the EMG analysis, muscle
183 signals were full-wave rectified, normalized in amplitude with respect to their maximum
184 value recorded across all trials and conditions and low-pass filtered once more with a zero-lag
185 Butterworth filter (cut-off frequency 5 Hz). The filtered EMG signals relative to each trial
186 and comprised between 100 ms before t_0 and t_f were normalized to a standard time window of
187 200 samples. By considering 100 ms before movement initiation we wanted to include in the
188 analysis any kind of anticipatory activity associated with the movement. To identify specific
189 invariant patterns characterizing the EMG activities of the different subjects, two versions of
190 non-negative matrix factorization were applied to the low-pass filtered EMGs. The standard
191 NMF algorithm (Lee and Seung, 1999) was used to identify both temporal components and
192 synchronous synergies.

193 *Temporal synergies (or temporal components)*

194 NMF was applied to the matrix \mathbf{M} of the EMG signals (size m by T , where m is the number
 195 of muscles signals and T the number of time samples), providing two matrices \mathbf{U} and \mathbf{C} (of
 196 dimension respectively m by N_c and N_c by T , where N_c is the number of temporal
 197 components) such that, at the time instant t , it results

$$198 \quad \mathbf{M}(t) = \sum_{i=1}^{N_c} \mathbf{U}_i \mathbf{C}_i(t) + \text{residuals} \quad (1)$$

199
200
201

202 where \mathbf{U}_i indicates the i -th column of the matrix \mathbf{U} and $\mathbf{C}_i(t)$ the i -th element of the column
 203 vector $\mathbf{C}(t)$. Note the number of muscles m indicates the number of muscles recorded
 204 during one single experimental trial. When considering multiple trials the matrix \mathbf{M} was
 205 obtained by concatenating vertically the matrices of the single trials.

206 *Synchronous synergies.*

207 NMF was applied to the transpose matrix \mathbf{M}' of \mathbf{M} , providing thus two matrices \mathbf{V} and \mathbf{W}
 208 (this time of dimension respectively T by N_s and N_s by m , where N_s is the number of
 209 synchronous synergies) such that

$$210 \quad \mathbf{M}'(t) = \sum_{i=1}^{N_s} \mathbf{V}_i(t) \mathbf{W}_i + \text{residuals} \quad (2)$$

211
212
213

214 where $\mathbf{V}_i(t)$ indicates the i -th element of the row vector $\mathbf{V}(t)$ and \mathbf{W}_i the i -th row of the
 215 matrix \mathbf{W} . Note that in (1) the j -th row of the matrix \mathbf{M} results from the linear combination of
 216 the rows of the matrix \mathbf{C} scaled by the scalar coefficients of the j -th row of the matrix \mathbf{U} .
 217 Each row of \mathbf{C} therefore contains one temporal component. In (2), conversely, the j -th row of
 218 \mathbf{M}' is obtained by combining linearly the rows of \mathbf{W} scaled by the coefficients of the j -th row
 219 of \mathbf{V} . Each row of \mathbf{W} therefore, of dimension 1 by N_s , represents a vector of muscle
 220 activations, i.e. a synchronous synergy. Note also that, because of the constraints imposed by
 221 NMF on parameters, all the entries of the matrices \mathbf{U} , \mathbf{V} , \mathbf{C} , and \mathbf{W} are non-negative. Even in
 222 this case, when considering multiple trials before applying NMF the transposed of the
 223 matrixes of the single trials were concatenated vertically.

224 *Time-varying synergies.*

225 We applied a customized version of standard NMF and that was developed by d'Avella and
 226 colleagues (2003, 2005). Similarly to standard NMF all the identified parameters are non-
 227 negative, but temporal shifts of the synergies are also allowed so that each column vector of
 228 \mathbf{M} at the instant t is the following relationship is such that

$$229 \quad \mathbf{M}(t) = \sum_{i=0}^{N_t} c_i \mathbf{w}_i(t - \tau_i) + \text{residuals} \quad (3)$$

230
231
232

233 where N_t is the number of time-varying synergies and the c_i and τ_i are respectively the scaling
 234 coefficient and the time delay associated the synergy w_i . The algorithm by d'Avella and
 235 colleagues requires specifying the temporal duration of each time-varying synergy. In this
 236 study the time duration of each synergy was set, for each subject, as long as the time duration
 237 of the whole trial after time standardization (200 samples). Note that the residuals in (1), (2)
 238 and (3) decrease as the number of synergies increase. In case of multiple trials, the matrixes
 239 of the single trials were concatenated horizontally.

240 *Selection of the number of synergies to be included in the EMG decomposition and their*
 241 *significance.*

242 In (1), (2) and (3) the numbers of muscle synergies (N_c , N_s and N_t) are free parameters of the
 243 analysis that can be set arbitrarily by the experimenter. Here, it was decided to set in all the
 244 three cases the number of synergies according to a criterion based on the computation of the
 245 variance accounted for (VAF) as a function of N_c , N_s and N_t . The VAF was defined as it
 246 follows

$$247 \quad \text{VAF} = 100 \cdot (1 - (\|\mathbf{M} - \mathbf{D}\|^2 / \|\mathbf{M} - \text{mean}(\mathbf{M})\|^2)) \quad (4)$$

248

249 where \mathbf{D} is the matrix of the reconstructed EMG obtained by using a certain number of
 250 synergies and $\text{mean}()$ is an operator that compute a matrix of the same size of the matrix \mathbf{M}
 251 and whose rows are equal point by point to the mean values of the corresponding rows of \mathbf{M} .
 252 The number of synergies was determined as the number of components at which the graph of
 253 the cumulative VAF presented a considerable change of slope (an “elbow”) and after which
 254 the slope of the graph became constant (Ferré, 1995). The exact point of change was
 255 quantitatively determined by using a linear regression procedure already used in literature
 256 (Cheung et al., 2005, 2009; d'Avella et al., 2006; Chiovetto et al., 2010, 2012). We computed
 257 a series of linear regressions, starting from a regression on the entire cumulative VAF curve
 258 and progressively removing the smallest value of number of component from the regression
 259 interval. We then compute the mean square residual error of the different regressions and
 260 selected the number of optimal synergies the first number whose corresponding error was
 261 smaller than 10^{-3} . To minimize the probability to find local minima, we always ran NMF
 262 25 different times on the same data set and consider as valid solution that provided the lowest
 263 reconstruction error between original and reconstructed error. To test the robustness and
 264 generality of the synergies extracted from each data set we exploited the two following cross-
 265 correlation procedures. We divided each data set in 5 parts of the same size. Since every data
 266 set consisted of the EMG activities of the Bic and Tri muscles collected during 20 repetitions
 267 of the same movement accomplished by one subject, each part consisted of the EMG
 268 activities of four trials. We then chose randomly 4 parts to use as training data set and one
 269 part as test data set. We extracted the synergies from the training data set and used them to
 270 reconstruct the activations of the test data set. We used the original and reconstructed test
 271 data sets to compute the VAF to draw the graph of the cumulative VAF. We also used the
 272 synergies extracted from each subject to reconstruct the EMG data sets of all the other
 273 subjects and assessed the level of reconstruction goodness by computing the VAF. For all
 274 cases, we verified that the extracted synergies did not result from a bias associated with the
 275 extraction methods by running a simulation. For each subject and decomposition, we
 276 compared the VAF values for the reconstruction of the experimental data obtained by
 277 combining the identified synergies with the VAF values of the reconstruction of random,

278 structureless data reconstructed by combination of the synergies identified from those
 279 artificial data. Such data sets were generated by reshuffling the samples of each muscle
 280 independently in each trials of each subject. Reshuffled data were then low-pass filtered (5
 281 Hz cutoff). For each one of the actual data set we simulated 50 artificial data sets and
 282 extracted the synergies by using the same procedure used for the observed data. We estimated
 283 the significance by computing the 95th percentile of the VAF distribution for simulated data.

284 *Similarity of synergies across subjects.*

285 The similarity between synergies of different subjects was quantified by computing their
 286 scalar products. For synchronous synergies and temporal components we proceeded as
 287 follows. For all possible pairs of normalized synergies of two different subjects the
 288 corresponding scalar products were computed. Note that, by definition, such a product can
 289 only adopt values ranging between 0 and 1. The pair with highest similarity was selected and
 290 the corresponding synergies were removed from the two groups of synergies. The similarities
 291 between the remaining synergies were then computed, and the best matching pair of
 292 synergies was selected and removed from the original and reconstructed model. This
 293 procedure was iterated until all synergies were matched. To compute the similarity between
 294 time-varying synergies the procedure was very similar to the one just described above with
 295 the only difference that in the last case, before computing the scalar product, the matrices of
 296 the synergies were first rearranged by disposing the entries of the matrices in form of vectors.
 297 The similarity between synergies was then quantified by computing the maximum of the
 298 scalar products over all possible time delays of the second synergy with respect to the first.
 299 To assess however the significance of the values of similarity provided by the scalar products
 300 we defined a similarity index (S) between two synergies. This index, ranging from 0
 301 (similarity at chance level) and 1 (perfect matching of the synergies) was defined as follows

$$302 \quad S = (S_{data} - S_{chance}) / (1 - S_{chance}) \quad (5)$$

303 Where s_{data} is the scalar product between two synergies extracted from the actual data and
 304 s_{chance} is the mean scalar product between 200 pairs of random synergies. We generated the
 305 artificial synergies by resampling randomly from the distribution of the activation amplitude
 306 of each muscle in the data set from which the synergies were extracted and constructed
 307 sequences of random data with the same length of the extracted synergies. Artificial data
 308 were then low-pass filtered to match the smoothness of the actual data.

309

310 **Results**

311 To compare systematically the results provided by different synergistic
 312 decompositions when characterizing the same EMG data set, we recorded EMGs during a
 313 series of elbow rotations and then we extracted and compared synchronous, time-varying and
 314 temporal muscle synergies.

315 To illustrate the data, we begin by showing in Figure 2A the EMGs recorded during a
 316 typical trial accomplished by one subject and relative to an elbow flexion in the horizontal
 317 plane. Consistent with previous literature (Berardelli et al., 1996), such a movement is
 318 characterized by a sequence of three EMG bursts: an initial burst of the agonist muscle

319 having the goal of providing the propulsive force to accelerate the movement, followed by a
320 second burst of the antagonist to decelerate the movement and a third burst of the agonist to
321 dampen the oscillation that other appears at the end of the movement. The latter final
322 corrective action is also reflected in the final overshoot of the finger velocity profile. This
323 sequence of bursts of activity was found also for elbow extension in the horizontal plane and
324 flexion and extension in the vertical one (Figure 2B).

325 We then considered the extraction of synergies from these data. The first interesting
326 question is how many synergies of each type are needed to describe the data. The number of
327 synergies to consider was determined, for each subject and type of decomposition, from the
328 dependence of the percentage of VAF (see Methods) upon the number of synergies. The
329 latter curves are plotted in Figure 2 for each type of synergy factorization and for each
330 subject. The VAF curves in each decomposition were very similar across subjects. While for
331 the temporal and time-varying decomposition we could extract up to 6 synergies (Figure 2A
332 and 2C) we found that, when referring to a synchronous synergistic decomposition, two
333 synergies were enough to account for 100% of the variance associated with the original data.
334 We thus did not extract a number of synergies higher than two. In Figure 3B, however, we
335 reported an amount of variance equal to 100% even for $N = 3,4,5$ and 6, to make Figure 3B
336 graphically coherent with the other two panels, i.e. Figure 3A and 3B.

337 Figure 3A reports the VAF dependence upon the number of extracted temporal
338 synergies. For all subject, the VAF reached a high value when including 3 synergies, and the
339 linear interpolation algorithm that we used (see Methods) indicated that in all subjects 3
340 temporal synergies were sufficient to explain the vast majority of the variance (with
341 additional temporal synergies generated by the NMF algorithm adding only a very small
342 fraction of the total variance). The VAF curves for synchronous (Figure 3B) and time-varying
343 (Figure 3C) synergies show that, for each individual subject, only two synergies were instead
344 required to account for the variance of the EMG data.

345 After having individuated their number, we next considered the shapes of the
346 synergies extracted by each decomposition. Figure 4A reports the shapes of the three
347 temporal synergies extracted from the EMGs of a typical subject (LA). The three temporal
348 components clearly remind of the triphasic organization presented in Figure 2. Each temporal
349 component is characterized by one major bump. The first temporal synergy can be interpreted
350 as the component contributing the most to the modulation of the first burst of the agonist
351 muscle during movement accomplishment: the second as the first burst of the antagonist; and
352 the third as the second burst of the agonist. Note that the third temporal synergy shows an
353 initial deactivation before the occurrence of the main peak. This initial part of the synergy can
354 be associated to the antagonist deactivation, prior to movement initiation, of the anti-
355 gravitational muscles during rotation along the vertical plane. The combination coefficients in
356 Figure 4B (averaged across the repetitions of each kind of movement) show the contribution
357 of each component to the activity of each muscle. Consistently with a triphasic pattern, it is
358 evident that the first component is contributing more to the activity of the biceps during VF
359 and HF; conversely, it contributes more to the activation of the triceps in VE and HE.
360 Similarly the second temporal synergy is more active for the muscles opposing the actions
361 exerted by the muscles activated by the first components. Thus for HF and VF movements
362 the coefficients of the triceps are higher than those of the biceps. Whereas for VE the
363 coefficient of the biceps is higher than that of the biceps, for HE movements however the
364 level of the coefficients of the two antagonist muscles is approximately the same. The
365 coefficients show then that, in all movements, the third component is contributing to the

366 activations of both muscles in approximately equivalent proportion, in order to compensate
367 for overshoots or to increase the joint stiffness by co-activating opposing muscles.

368 There are two points that need to be remarked. First of all in the pre-processing step
369 all the EMG signals of each muscle were normalized with respect to the maximum value that
370 was recorded for that muscle across all trials. Such a procedure may consequently lead to a
371 partial loss of information about the relationship among the EMG amplitudes of different
372 muscles monitored within the same trial. Moreover, trials were normalized in duration, which
373 may introduce some supplementary temporal variability when merging all trials together to
374 extract synergies. These can explain why the average coefficients of biceps and triceps
375 relative to temporal synergy 2 in Figure 4B had approximately the same value for condition
376 HE, differently from the expectation according to which the coefficient of the biceps should
377 have appeared much larger than that of the triceps. According to the triphasic strategy,
378 indeed, it should have been expected the second component to contribute mainly to the
379 activation of biceps muscle which, in HE, is devoted to exert the antagonist role.

380 In addition, it is important to note that the number of identified temporal synergies is
381 three, which is higher than the number of degrees-of-freedom to control (one joint angle, two
382 muscles). This may look at first as a useless increase of complexity. However, the strength of
383 a triphasic strategy in a single-joint motor task lies likely in its flexibility and power of
384 generalization. Indeed, similar triphasic muscle organizations were found characterizing also
385 arm raising (Friedli et al., 1984), rapid voluntary body sway (Hayashi 1998) and whole-body
386 reaching (Chiovetto et al., 2010, 2012) motor tasks. In accordance with this premise, one can
387 note that the four tasks were all executed through a triphasic motor pattern. While previous
388 studies mainly demonstrated the powerfulness of the synergy idea to reduce the
389 dimensionality of motor control and execution, our results show in addition that temporal
390 synergies present marked functional features.

391 Figure 5A depicts the two synchronous synergies extracted from the EMGs of a
392 typical subject (LA). Each synergy is characterized by the activation of one single muscle.
393 Due to their antagonist nature, biceps and triceps therefore were found to share no common
394 level of activation. Note that, although such a result may seem trivial in a two dimensional
395 space, we might have obtained a pair of linearly independent vectors characterized by
396 noticeable activity of both muscles. In Figure 5B the temporal evolution of the scaling
397 coefficients averaged across movement repetition are shown for each muscle and each
398 movement. Note how, within each movement condition, the activities of the agonist and
399 antagonist muscles are always characterized by one main burst in agreement with a classic
400 triphasic pattern. Only for the first coefficient relative to HF movements the second burst is
401 not clearly visible, this being very likely due to the averaging procedure.

402 Finally, the two time-varying synergies are shown in Figure 6A. They were characterized by
403 one single burst for each muscle, one for the biceps and one of the triceps. The two synergies
404 differed however for the temporal order in which the two burst occurred: whereas the burst of
405 the biceps anticipated the burst of the triceps in the first time-varying synergy, the order of
406 the peaks was reversed in the second one. The average scaling coefficients and temporal
407 delays corresponding to each synergy are shown in Figure 6B-C. Note that also in this case,
408 the contribution of each synergy to the EMG activity of each movement is consistent with the
409 biomechanical feature of the movement itself. Thus time-varying synergy 1, in which the
410 biceps is activated first, contributes more to HF and VF movements, while time-varying
411 synergy 2, in which the triceps is activated first, contributes more to HE and VE movements.

412 In sum, we found that each kind of muscle decomposition provided a set of interpretable
 413 synergies. Each temporal component described a temporal phase of the movement. Each
 414 synchronous synergy described the simultaneous and coordinated action of a group of
 415 muscles (only one in our case) aiming to achieve a specific action goal. Each time-varying
 416 synergy related instead to a specific task-related variable (specifically a direction of motion).

417 We used the synergies extracted from each subject to reconstruct the EMG data of
 418 each one of the others and assessed the percentage of VAF. The results are reported in forms
 419 of confusion matrices (Figure 7). The average percentage of VAF computed across subjects
 420 was $90 \pm 7 \%$ when temporal synergies were extracted and used for reconstruction, and $87 \pm$
 421 4 for the data sets reconstructed by using the time varying synergies. These values were
 422 found to be significant and did not result from a bias built in the extraction methods. The
 423 average 95th percentile of the distribution of VAF values obtained from the reconstructions
 424 of the simulated data were indeed much lower of the ones obtained from the reconstruction of
 425 the actual data, respectively 17.6% and 39.3% when data were decomposed according to
 426 the temporal and time-varying synergistic decompositions. The synchronous case was not
 427 considered given the features of the extracted sources and the fact that with such synergies a
 428 perfect match of the actual data could always be achieved.

429 We quantified how much the synergies illustrated in Figure 4, 5 and 6 for one single
 430 subject were representative also of the synergies extracted from the EMG activity of the other
 431 subjects. To this purpose we computed the average scalar products and similarity indices
 432 between groups of synergies belonging to different participants. For the temporal
 433 components, the average scalar product was $s = 0.93 \pm 0.01$, $s = 1 \pm 0$ for the synchronous
 434 synergies and $s = 0.91 \pm 0.05$ for the time-varying ones. The scalar products across subjects
 435 of synchronous synergies were always equal to 1 because for all the subjects the same set of
 436 synchronous synergies was always identified, in which only one single muscle was recruited
 437 at a time. Note that in this case also the similarity index S is always automatically equal to 1.
 438 The mean S values computed between groups of synergies extracted from different subjects
 439 are plotted in Figure 8. On average $S = 0.86 \pm 0.06$ for the groups of temporal synergies and
 440 $S = 0.85 \pm 0.11$ for the time-varying synergies. Note that in both cases the average similarity
 441 index was much higher than 0 (chance level). In sum, all synergies decompositions show a
 442 very high degree of robustness across subjects.

443 Discussion

444 We used NMF-based methods to extract three different kinds of muscle synergies
 445 from the EMG activity of two antagonist muscles during the accomplishment of single-joint
 446 elbow rotations along both the horizontal and vertical planes. By using a well-understood
 447 motor task, we aimed to clarify better what are the motor features characterized by each kind
 448 of decomposition and to assess whether, when and why one of them should be preferred to
 449 another. We found well-defined interpretable results for each of the EMG signals
 450 decomposition considered. This allow us to discuss more in detail about what motor features
 451 each kind of muscle synergy decomposition encodes and, consequently, to explain why
 452 sometimes the extraction of one type of synergy may be more meaningful than another one.

453 In some previous studies (Tresch et al., 2006; Ivanenko et al., 2005) different
 454 unsupervised learning algorithms were applied to the same data set to verify the
 455 independence of the synergies from the particular technique used for their identification, or to
 456 test the superiority of an algorithm with respect to another one. In such studies however, all

457 the algorithms used always relied on the same generative model, i.e. on the same definition of
458 synergy. To our knowledge, this is the first study comparing synchronous, time-varying and
459 temporal muscle synergies extracted from the same data set. Hence it offers the possibility to
460 gain novel insights into the benefits provided by the different modular decompositions. Our
461 choice of an elementary motor task for which most of the neuromuscular functions are well-
462 understood, made the interpretations of various synergies as transparent as possible.

463 The results that we presented revealed that in all the cases NMF led to the
464 identification of interpretable muscle synergies. The extraction of synchronous synergies
465 yielded two primitives, each one characterized by the activation of only one of the two
466 muscles, indicating that biceps and triceps (respectively flexor and extensor of the elbow
467 joint) assumed independent levels of activation; in other words their activation waveforms
468 did not, in general, co-vary in time. This might look like a trivial result given the small
469 number of muscles considered and in view of antagonist nature of the two muscles during
470 elbow rotations. However, following the generic definition of a muscle synergy as a group of
471 muscles working together to achieve a common goal, it may appear surprising to find that the
472 two main muscles controlling the task performance are not synergistic. However, the
473 definition of synergies can be restated as groups of muscles acting at one or multiple joints to
474 achieve a specific motor function (in our case the motor function could be simply flexing or
475 extending the arm; in other terms, accelerate or decelerate the arm). From this point of view,
476 our interpretation is in agreement with other previous studies considering more complex
477 movements and a larger number of muscles. Similarly to us, for instance, the synergies
478 extracted by Cheung and colleagues (2009) from the EMG activations of sixteen elbow and
479 shoulder muscles of subjects performing a set of arm movements in space can be easily split
480 in two groups: one encompassing synergies in which the most active muscles are flexor and
481 another one in which extensor muscles are instead dominating (see Cheung et al. 2009, their
482 Figure 3A). Also in this case, therefore, the goal associated with each synergy was to flex or
483 extend the arm. By extension, this may suggest that muscles belonging to the same
484 synchronous synergy share similarities with respect to their biomechanical function for the
485 movement to be performed. Synchronous synergies were shown however encoding also other
486 kinds of functional goals, or “strategies”. Torres-Oviedo and Ting (2007) extracted
487 synchronous synergies from a set of leg and trunk muscles during a postural task and found
488 synergies characterized mainly by activation of either ankle or knee muscle. These synergies
489 resulted therefore in producing muscle activation patterns associated with two well-known
490 postural strategies, usually referred to as “hip” and “ankle” strategies, which were previously
491 deeply described in human postural control (Horak and Macpherson, 1996).

492 When extracting temporal muscle components the application of NMF provided a
493 decomposition based on three temporal synergies. Each one of them was found playing a
494 well-determined functional role during movement accomplishment, in agreement with the
495 three movement phases present in the classical triphasic pattern (see Berardelli et al., 1996,
496 for a review relative to elbow and wrist movements). The three phases can be resumed as
497 follows: a first phase (coinciding with the first agonist EMG burst) to provide the impulsive
498 force to initiate the movement, a second phase (antagonist burst) dedicated to halt the
499 movement at the desired end-point and a third phase (coinciding with the second agonist
500 burst) to dampen out the oscillations which might occur at the end of the movement.
501 Although in a single-joint motor task such a triphasic strategy may look like a useless
502 increase of complexity due to the fact that the number of synergies is higher than the
503 number of muscles to control, its strength lies likely in its flexibility and power of
504 generalization. Indeed, similar muscle organizations were found characterizing also arm

505 raising (Friedli et al., 1984), rapid voluntary body sway (Hayashi 1998) and whole-body
506 reaching (Chiovetto et al., 2010, 2012) motor tasks. Along with the need of reducing
507 movement complexity by reducing the number of degrees of freedom (number of muscles),
508 the decomposition of EMG activations based on the definition of temporal synergies
509 showed that at some extent even the temporal dimension of the movement is a source of
510 complexity that could be controlled and simplified by the CNS. These findings also pose the
511 question of the neural implementation of this kind of temporal synergies. For single-joint
512 rotations, Irlbacher et al. (2006) showed that the bursts composing the triphasic pattern were
513 triggered in cascade with the possibility for the second burst to depend partly on what
514 occurred during the first burst and not as a complete undividable sequence. This is
515 compatible with the extraction of three temporal synergies to account for the control of
516 elbow rotations across several conditions. However, this asks the question whether there are
517 indeed three 'spinal' temporal patterns recruited by different premotor drives or if the same
518 temporal pattern is recruited by a delayed sequence of premotor drives. Interestingly, this
519 idea of time shifts is present in the time-varying model of muscle synergies, which might
520 have solved this issue.

521 We found that two time-varying muscle synergies could account quite well for the
522 EMG activity associated with elbow movements. Each synergy was characterized by two
523 main bursts of activation for both the biceps and triceps, whereas the time of occurrence of
524 their peaks was inverted in the two synergies. While the burst of the biceps in the first
525 synergy of Figure 6A occurs for first and may be thought to contribute therefore to start
526 elbow flexion and the burst of the triceps to brake it, in the second synergies to role of the
527 two muscles is inverted and the synergy is consistent with the pattern associated with an
528 elbow extension. The two synergies seem therefore to intrinsically encode the direction of
529 motion, or in other words, the motor task, and therefore may allow a hierarchical control of
530 movements, in which task goals are only needed to be specified to generate complete muscle
531 patterns. This finding is coherent with the results presented in previous investigations
532 regarding arm movements (d'Avella et al. 2006, 2008, 2011) in which, even when a larger
533 number of muscles was taken into account in the analysis, time-varying synergies were
534 found to be directionally tuned, so that they resulted active only when the movements
535 occurred in well-determined directions. We also stress the subtle difference between the
536 interpretation of time-varying synergies and synchronous synergies: with the first time-
537 varying synergy only flexions can be performed (maybe varying its speed or amplitude
538 depending the way it is recruited). In contrast, the first synchronous synergy can be used for
539 both flexion (to accelerate) and extensions (to decelerate), showing that both representations
540 encode divergent aspects of the movements data set.

541 The use of very simple motor tasks characterized by well-known triphasic pattern allows us
542 to evaluate some pros and cons of each of the decompositions used in this study. Previous
543 works demonstrated that, in a triphasic pattern, the time of activation of the antagonist muscle
544 is controlled independently by the cerebellum (Manto et al., 1995). Other studies (Cheron and
545 Godaum, 1986) also reported that the timing of the antagonist burst onset increases with the
546 movement amplitude, whereas the one of the agonist does not. Our results showed that
547 neither the temporal synergistic decomposition nor the time-varying one can capture such
548 timing features. In the first case, indeed, each one of the three bumps of Figure 4 is invariant
549 in time and cannot be shifted temporally. This makes impossible to model the inter-trial
550 variability of the onset of the antagonist muscle. Rather, each bump represents the average
551 temporal evolution of the corresponding bursts across all trials. In the second case,
552 differently, in each of the time-varying synergies that we identified from the experimental

553 data set, the time lag between the activation of the two antagonist muscles is constant. This
554 prevents the possibility, when reconstructing the data, to vary from trial to trial the time
555 interval between the activations of the agonist and antagonist muscles, as observed in human
556 subjects. Different considerations can instead be made for the results associated with the
557 synchronous decomposition. As each synergy that was identified from the data is responsible
558 for the recruitment of one single muscle indeed, the activation profile of each muscle can be
559 set arbitrarily and independently for each trial. This allows therefore not only to model
560 independently the times of activation of each burst in each trial, but also their amplitudes, in
561 agreement with other experimental observations. Hannaford and colleagues (1985)
562 demonstrated indeed that the first agonist burst is not modified by the vibration of the agonist
563 muscle. In contrast the amplitude of the second agonist burst is increased and the vibration of
564 the antagonist muscle increases the amplitude of the antagonist burst. Similarly to the
565 synchronous one, even the temporal decomposition is suitable to capture such features of the
566 amplitudes in the reconstructed data, as it allows the separate scaling of each one of the three
567 identified bumps. The time-varying decomposition, on the contrary, introduces instead by
568 construction a correlation between the amplitudes of the different muscles.

569 It was demonstrated that discrete movements regulated by a triphasic pattern may present an
570 oscillatory component in the neural command (see for instance Cheron & Godaux, 1986).
571 Very recently, it was also shown by the analysis of the dynamical structure of reaching
572 movement that non-periodic movement such as the one presented here contains a strong
573 rhythmic structure (Churchland et al, 2012). In this study the authors proved that, although
574 EMG responses do not themselves exhibit state-space rotations, EMG can however be
575 constructed from underlying rhythmic components. It makes thus sense to wonder which one
576 of the decomposition methods that we investigated can be more useful or complementary for
577 the understanding of the oscillatory nature of the control of movement. Each model might
578 indeed provide a set of synergies revealing specific oscillatory features underlying the EMGs.
579 In this framework, synchronous components cannot be of help, as they carry spatial and not
580 temporal information. Interesting results might instead be provided by drawing the phase
581 plots associated with each temporal component or with the activity of each muscle trace in a
582 time-varying synergy. In case the plots presented evident rotations indeed, the hypothesis put
583 forward by Cheron and Godaux and later by Churchland and colleagues would be
584 strengthened. In the contrary case, however, the results obtained by these authors would not
585 be discredited, as the absence of rhythmic features in the components might instead be due to
586 the incapability of the synergy models to account for such features correctly.

587 We have in this discussion tried to provide evidence that the simple results that we
588 found for the simple movement and system considered in this study might very likely hold
589 also for more complex behaviours involving the action of large number of muscles. We think
590 therefore that, in general, each kind of muscle synergy may encode a different motor feature.
591 Specifically, temporal components encode different temporal phases of the movement, each
592 one playing a specific functional role. Synchronous synergies encode the simultaneous and
593 coordinated actions of specific groups of muscles aiming to achieve a specific motor function
594 (e.g. accelerate the body toward the target). Finally, time-varying synergies encode high-level
595 task-related functions (in this case the direction of motion). This suggests that the type of
596 factorization to be chosen in each condition depends on which of these aspects the study
597 intends to reveal. Note however that each type of synergies may not always characterize
598 uniquely only one single motor feature, mainly because two or more variables may be
599 correlated. Thus, for instance, the direction of motion can be inferred also from the amplitude
600 of the scaling coefficients relative to temporal components (Figure 4B) once the action

601 exerted by the muscles in known, or the triphasic temporal organization can be also reflected
602 in the temporal evolution of the scaling coefficients in Figure 5B.

603 We conclude by stressing that a unifying synergy extraction method capturing all
604 those aspects at once could simplify the interpretation of future works. If all these
605 representations of synergies are simultaneously valid, then a more general model on the top
606 of them should exist. Used systematically, such a model could allow better comparisons and
607 interpretations of muscle synergy studies in more complex motor tasks.

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691 Captions

692 **Figure 1: Sketch of the experimental paradigm.** Subjects sat on a chair and had to
693 accomplish flexions or extensions of the elbow along both the vertical (V) and horizontal (H)
694 plane.

695 **Figure 2: Typical EMG and kinematic associated with the experimental paradigm.** A.
696 EMG traces of individual rapid flexion and extension movements of the elbow in a normal
697 subject. In all conditions the triphasic pattern results clearly present. B. From the top, angular
698 elbow displacement and velocity associated with one typical elbow flexion in the horizontal
699 plane are respectively depicted, along with the EMG activities of Bic and Tric muscles. In the
700 two panels at the bottom, the smoothest lines represent the envelopes of the rectified EMGs
701 and were obtained by low-pass filtering the rectified EMG at 5 Hz, the spikiest ones at 20 Hz.
702 Clearly, different filtering frequencies do not modify the main temporal features of the
703 signals.

704 **Figure 3: Levels of approximation as a function of the number of synergies.** A.
705 Percentage of VAF as a function of the number of temporal synergies. B. Percentage of VAF
706 as a function of the number of synchronous synergies. C. Percentage of VAF as a function of
707 the number of time-varying synergies. Each coloured line is associated to a specific subject
708 (see most right panel), which in the figure is identified by the initials of his first and last

709 name. In all the three panels the vertical arrows indicate the number of primitives at which
710 the curves satisfy the linear regression criterion to choose the number of primitives (see
711 Methods). These points are invariant across subjects and coincide, in most of the cases, with
712 the points at which the curves present an “elbow” and start becoming straight.

713 **Figure 4: Identified temporal synergies.** A. Temporal components extracted from one
714 typical subject (LA), ordered according to the time of the occurrence of their main peaks. B.
715 Corresponding scaling coefficients.

716 **Figure 5: Identified synchronous synergies.** A. Synchronous synergies identified from one
717 typical subject (LA). B. Temporal evolution of the corresponding scaling coefficients

718 **Figure 6: Identified time-varying synergies.** A. Time-varying synergies extracted from the
719 EMG activity of one typical subject (LA). B. Corresponding scaling coefficients. C.
720 Corresponding temporal delays.

721 **Figure 7: Cross-validation results.** A. Percentage of VAF for the reconstruction of the
722 actual EMG data set of one subject by using the temporal synergies identified from the data
723 sets of the other subjects. VAF values along each row are associated with the reconstruction
724 of the data of one single subject. B. Percentage of VAF for the reconstruction of the actual
725 EMG data set of one subject by using the time-varying synergies identified from the data sets
726 of the other subjects.

727 **Figure 8: Average level of similarity between groups of synergies identified from the**
728 **EMG data of the 8 subjects that participated to the experiment.** A. Similarity between
729 groups of temporal synergies. B. Similarity between groups of time-varying synergies. The
730 average level of similarity between synchronous synergies is not shown as the identified set
731 of synchronous synergies was the same across all subjects.

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742 **Table 1: Average mean and peak angular velocities.** For each movement and subject the
 743 average velocities (\pm standard deviation) are reported. Averages and standard deviations were
 744 computed over all trials repetitions.

		HF	HE	VF	VE
Peak vel. (rad/s)	AL	8.86 \pm 0.91	-8.95 \pm 0.77	10.81 \pm 1.12	-11.65 \pm 2.37
	AR	11.78 \pm 0.97	-9.84 \pm 1.29	12.51 \pm 0.89	-12.55 \pm 1.11
	CA	9.69 \pm 1.35	-8.26 \pm 0.92	9.63 \pm 1.66	-9.08 \pm 2.59
	MA	7.09 \pm 0.99	-7.45 \pm 1.83	7.18 \pm 0.83	-7.92 \pm 1.42
	FR	4.85 \pm 0.41	-7.37 \pm 0.77	5.84 \pm 1.07	-5.29 \pm 0.57
	FA	11.32 \pm 1.01	-10.35 \pm 1.03	9.72 \pm 1.24	-12.04 \pm 1.30
	GI	7.96 \pm 0.93	-8.17 \pm 0.89	9.99 \pm 0.69	-9.28 \pm 1.48
	LA	8.56 \pm 2.10	-9.70 \pm 0.68	10.19 \pm 0.69	-12.63 \pm 0.93
Mean vel. (rad/s)	AL	3.26 \pm 1.00	-3.47 \pm 0.74	4.31 \pm 0.68	-4.06 \pm 0.98
	AR	3.55 \pm 0.34	-2.92 \pm 0.83	3.95 \pm 0.56	-3.69 \pm 0.76
	CA	3.86 \pm 0.81	-3.65 \pm 0.42	3.53 \pm 0.60	-3.06 \pm 0.64
	MA	3.01 \pm 0.37	-2.68 \pm 0.49	3.00 \pm 0.43	-2.81 \pm 0.47
	FR	2.04 \pm 0.31	-1.61 \pm 0.78	2.56 \pm 0.35	-2.18 \pm 0.47
	FA	4.10 \pm 0.92	-2.15 \pm 0.54	3.68 \pm 0.64	-3.54 \pm 0.71
	GI	3.41 \pm 0.37	-3.65 \pm 0.29	3.82 \pm 0.63	-3.58 \pm 0.54
	LA	3.24 \pm 1.89	-3.57 \pm 0.60	3.66 \pm 0.70	-3.47 \pm 0.54

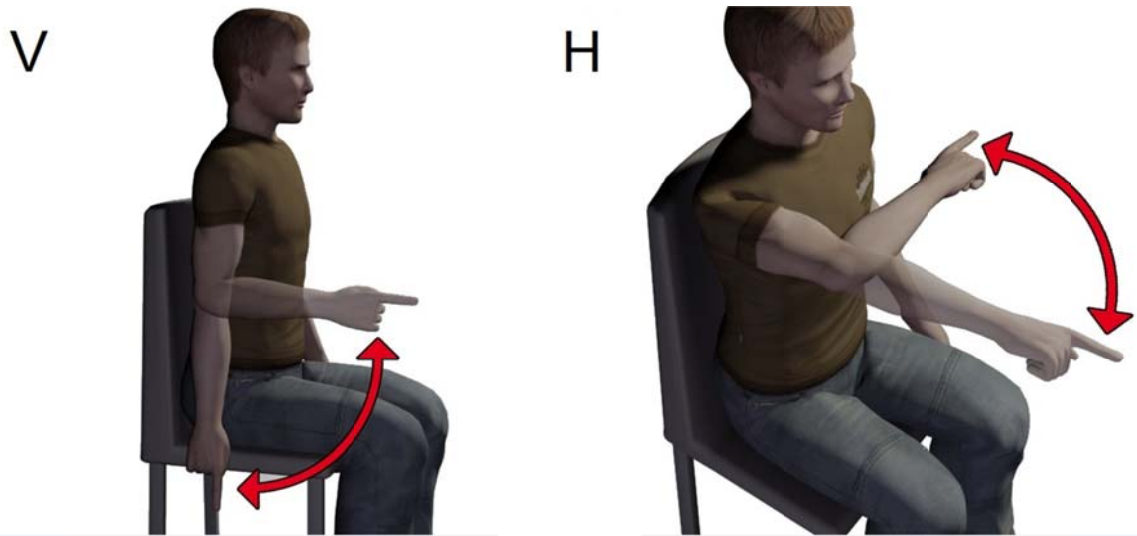
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Figure 1

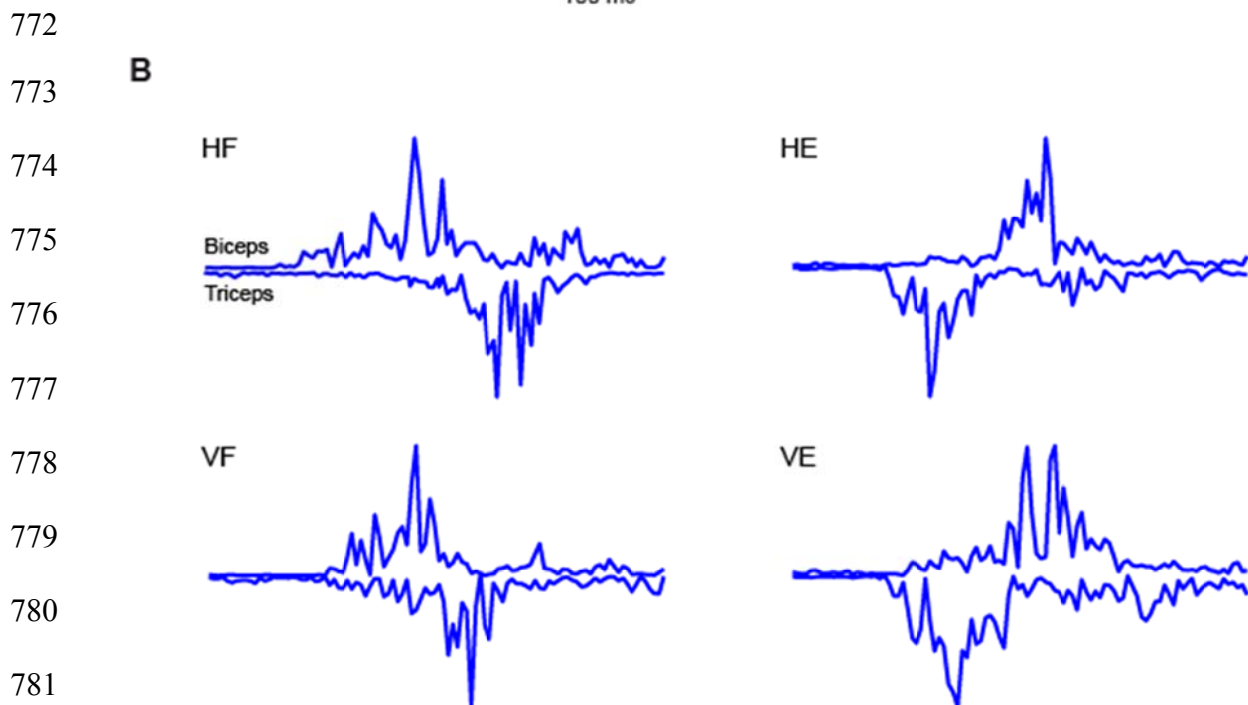
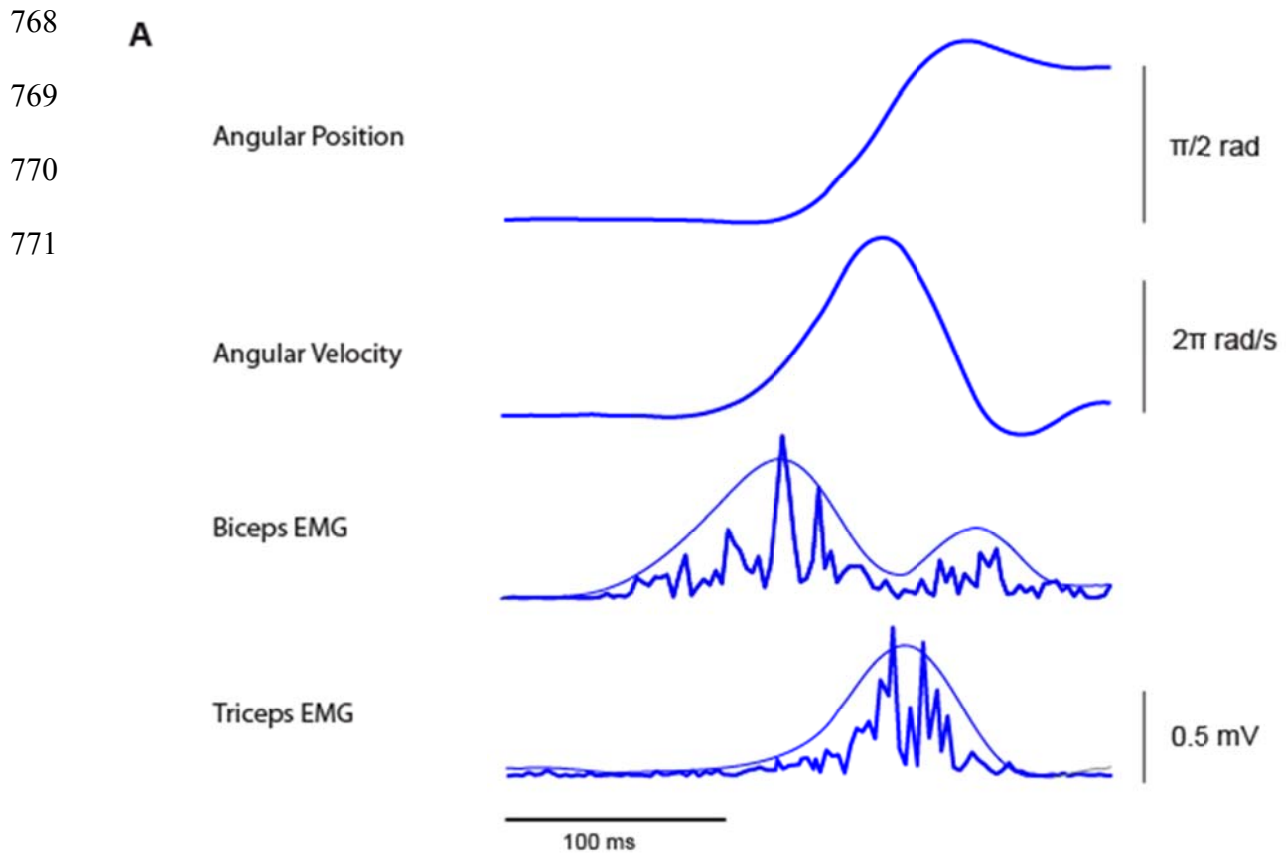


Figure 2

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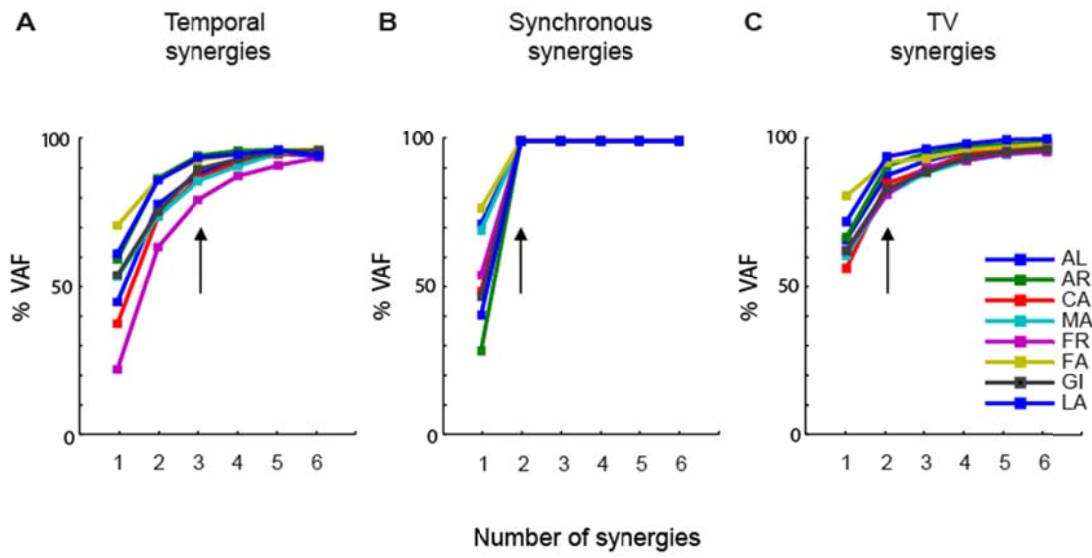
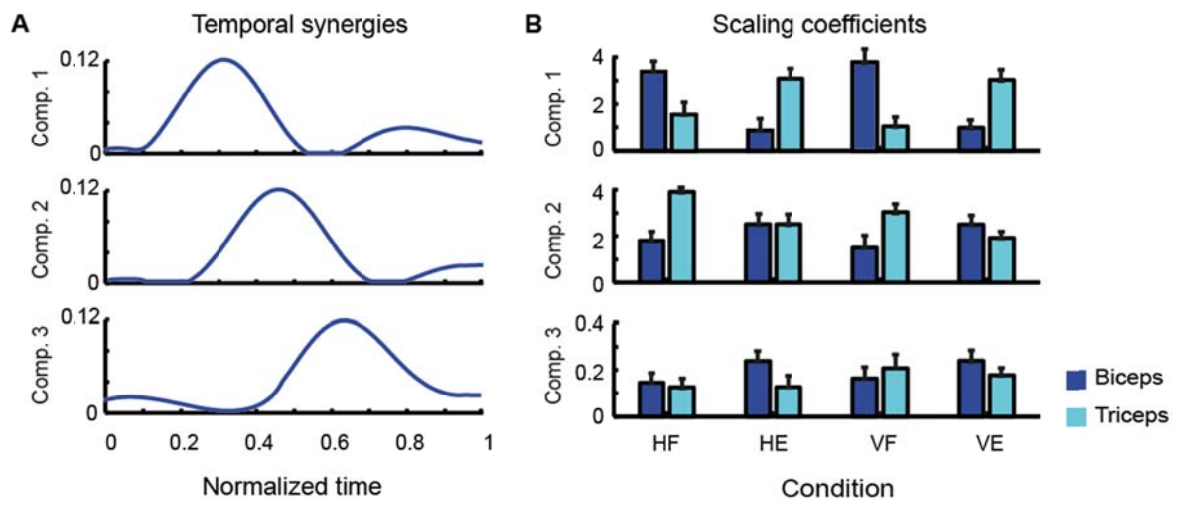


Figure 3



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Figure 4

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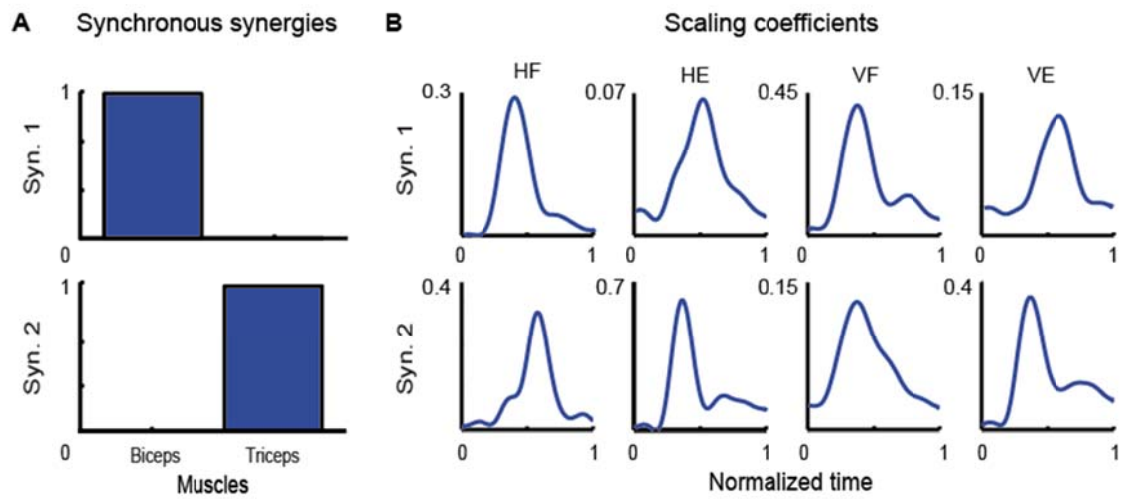
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Figure 5

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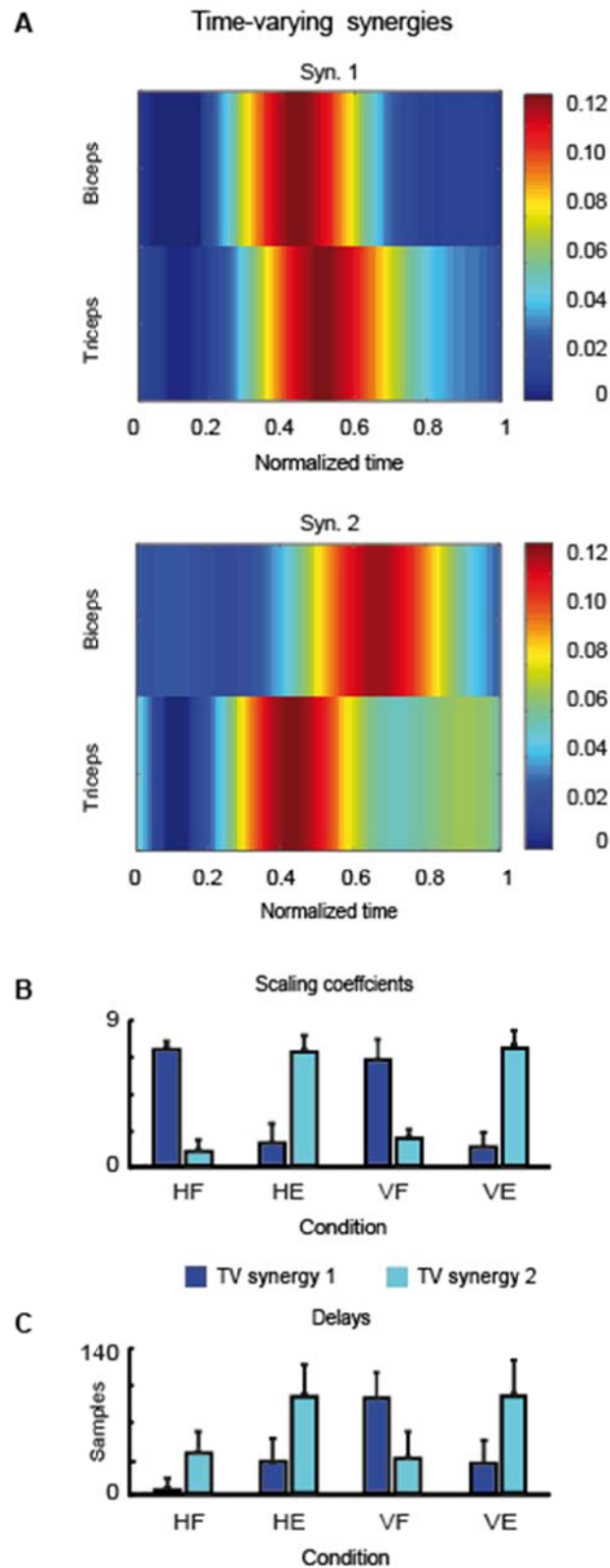


Figure 6

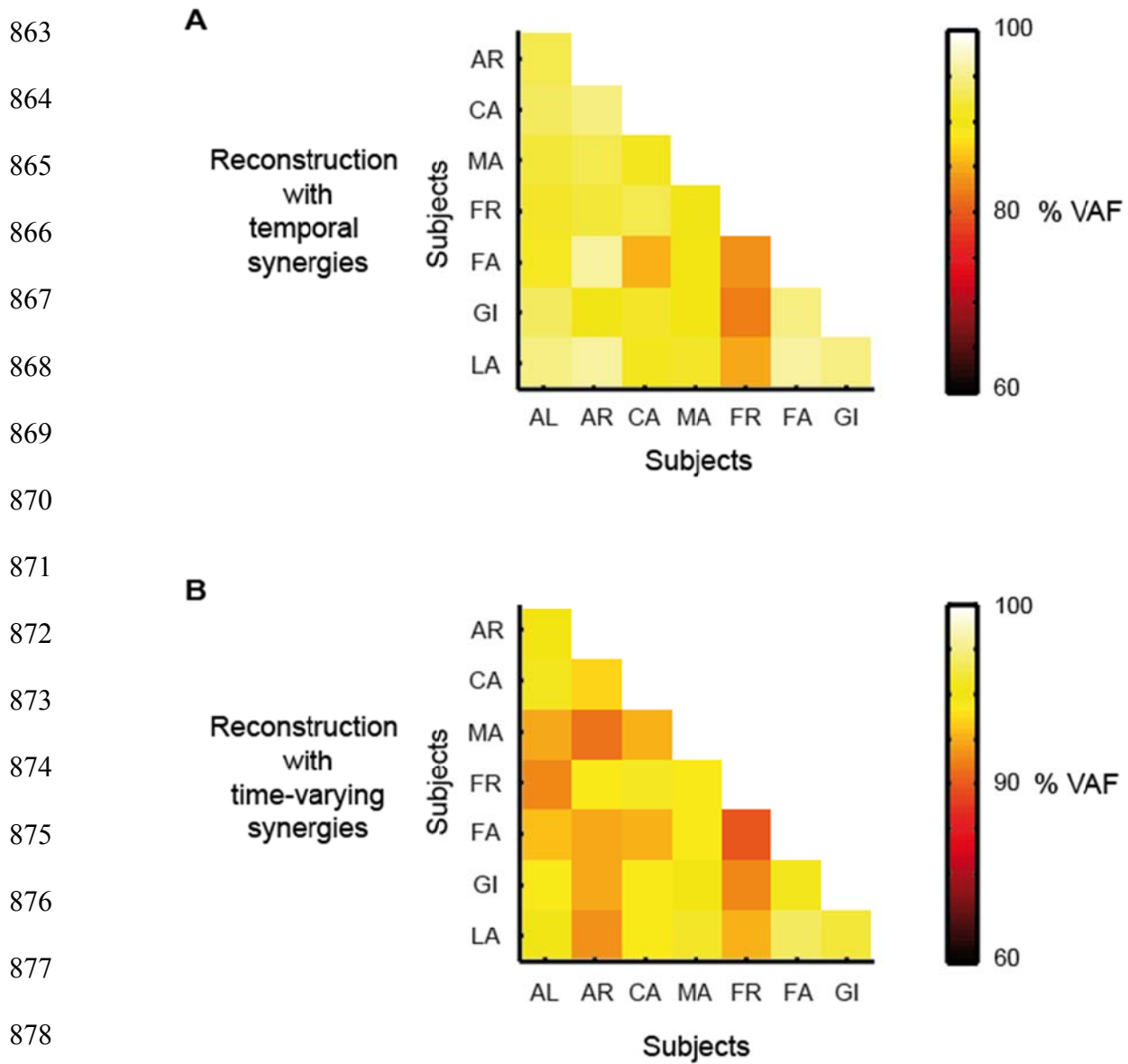
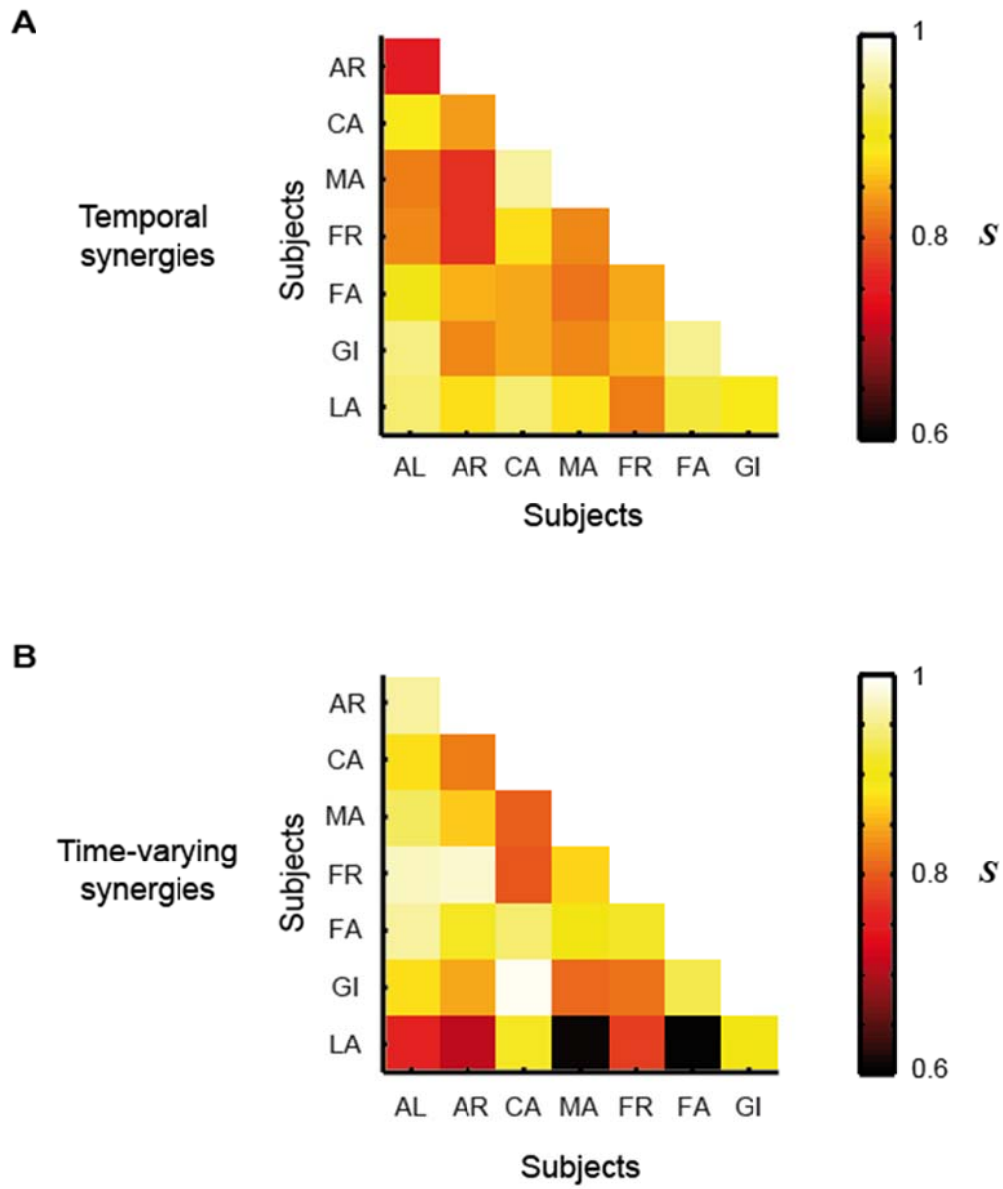


Figure 7



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Figure 8

Figure 1.JPEG

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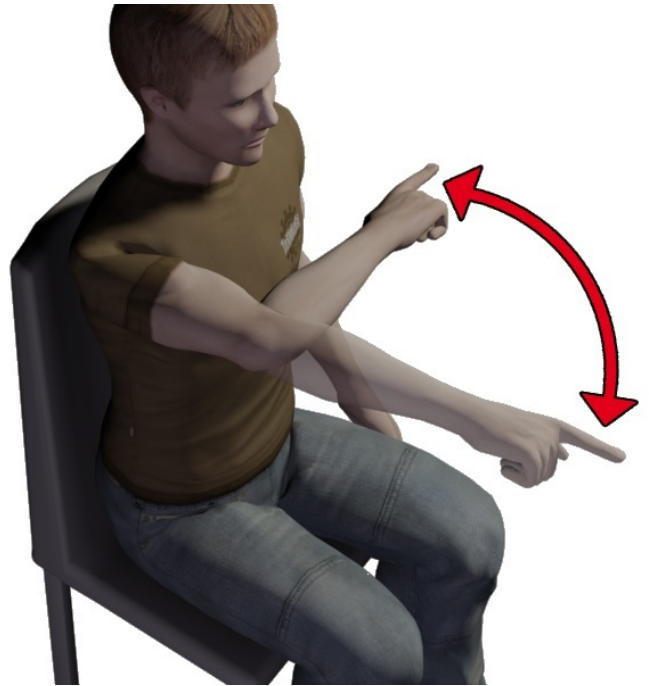
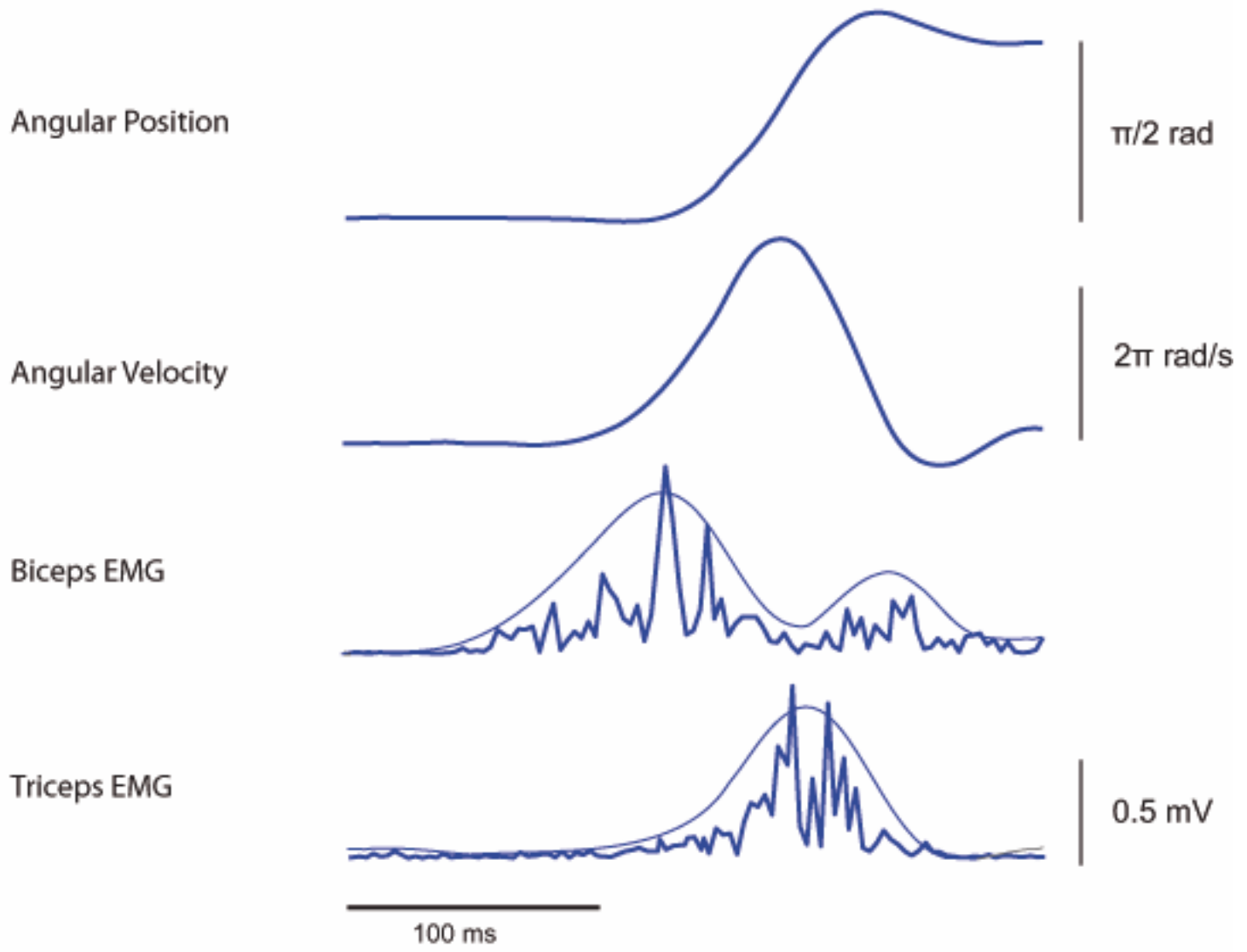


Figure 2.TIF

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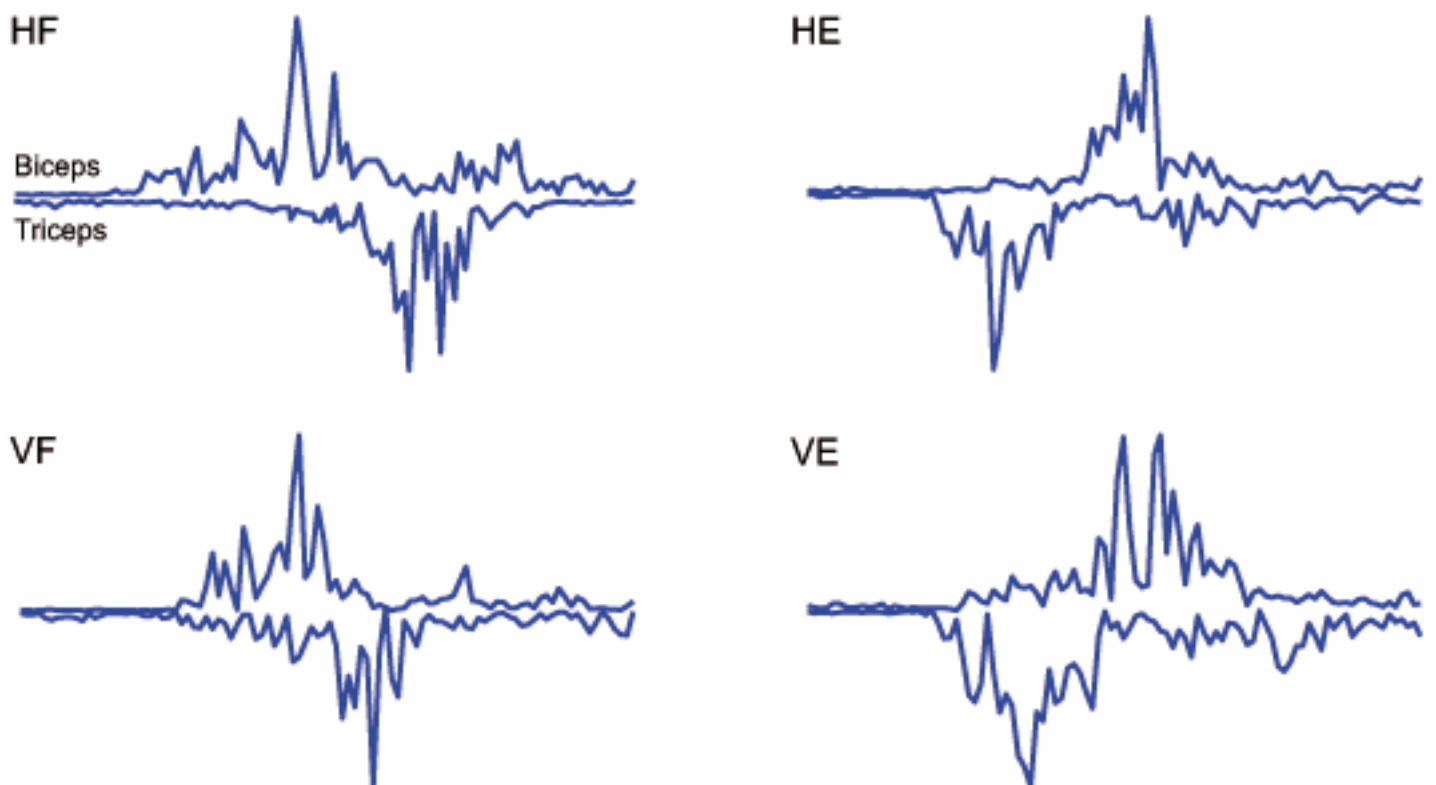


Figure 4.TIF

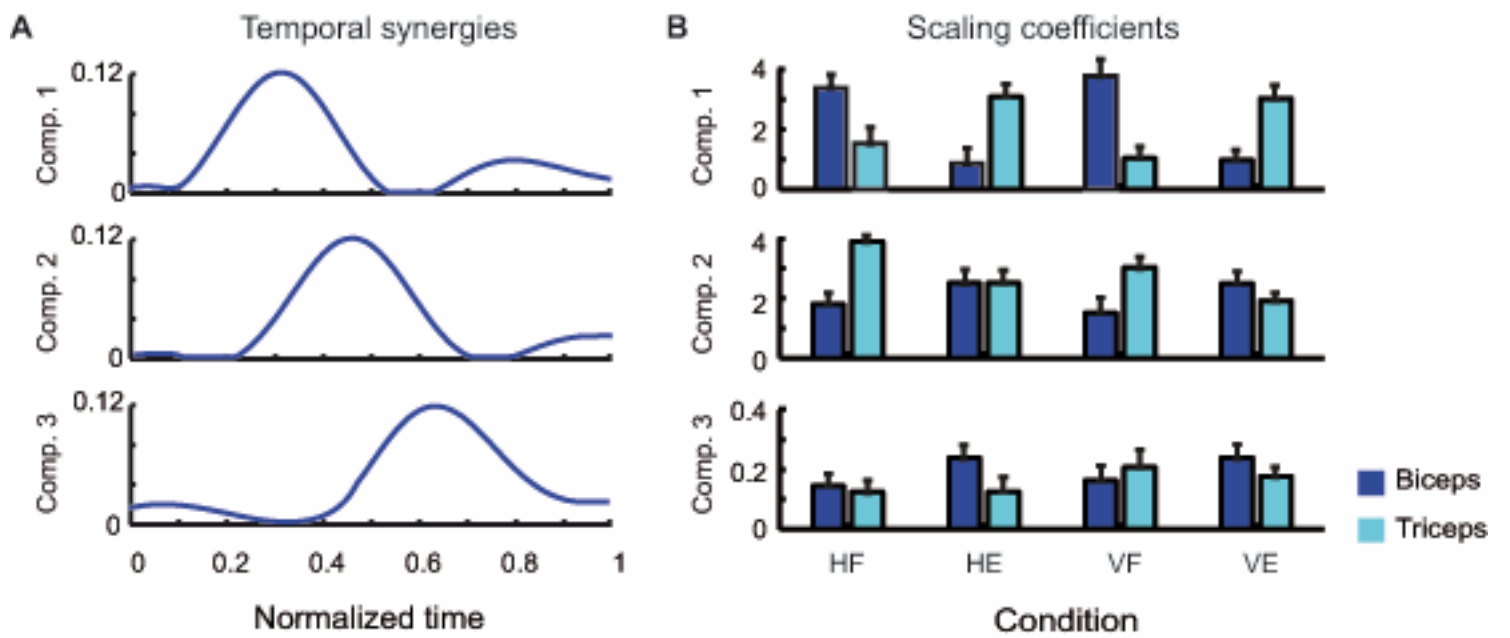


Figure 5.TIF

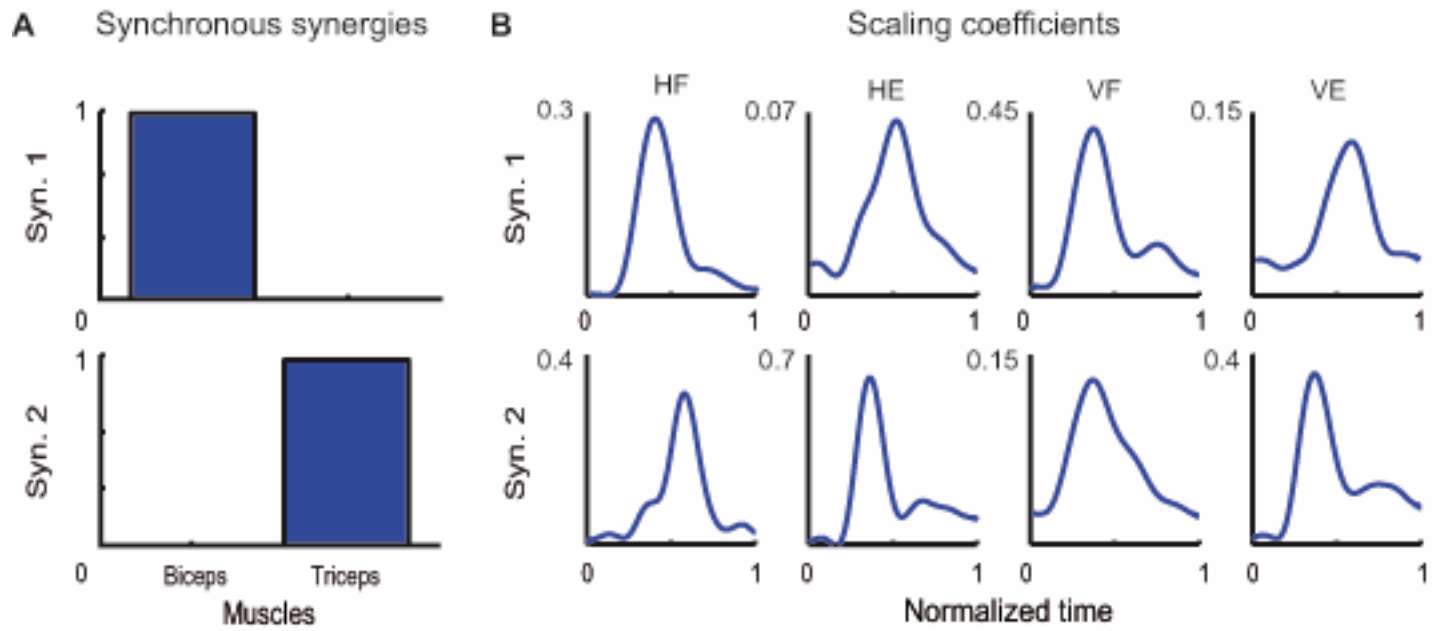


Figure 6.TIF

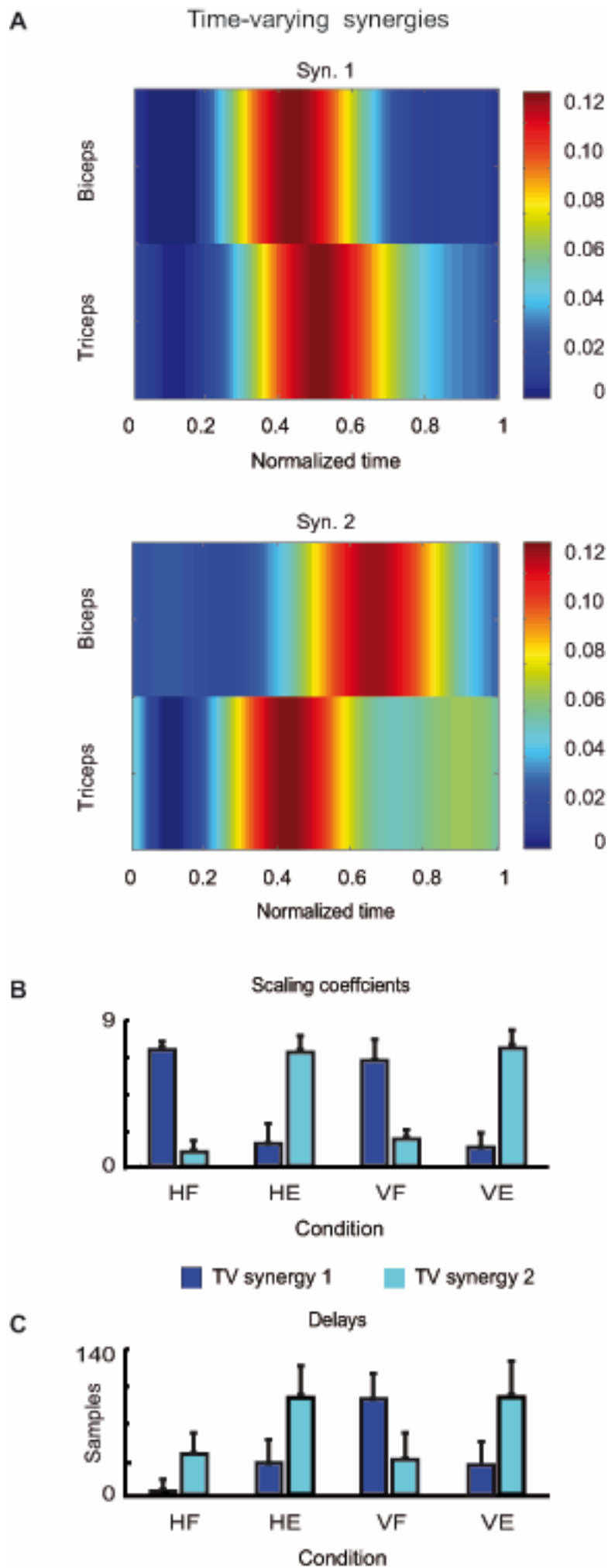


Figure 7.TIF

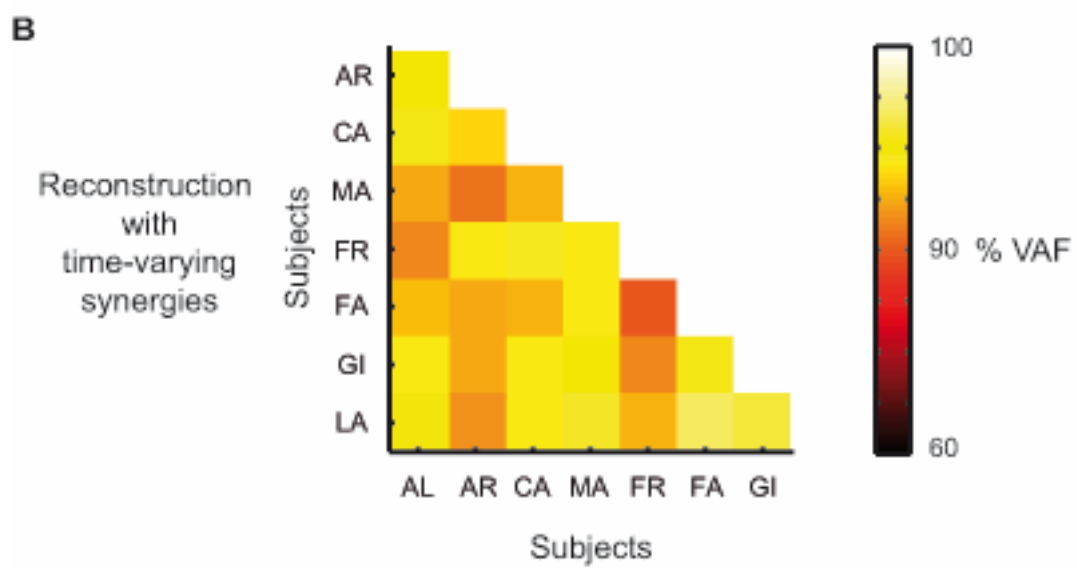
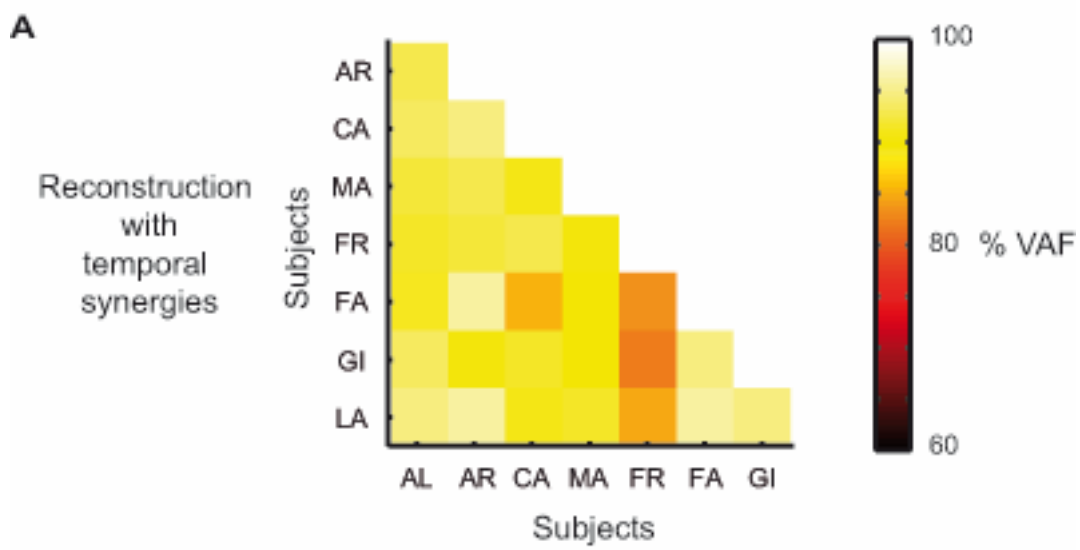


Figure 8.TIF

