Neuromechanical Coupling is Reflected in the Spatial Organization of the Spinal Motoneuron Pools

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Abstract

Recent advances in neuroprosthetics have shown that activating the surviving spinal circuitry below the level of spinal damage can produce complex movements and restore function. This is because the spinal cord circuitry plays an important role in the control of complex movements by integrating descending execution-related signals with feedback sensory signals. To generate efficient movements through muscle contractions, the nervous system embeds the anatomical and dynamical properties of the controlled body parts, a concept termed neuromechanical tuning. The motoneurons innervating the skeletal muscles of the body form complex spatial structures that span multiple segments. The goal of this study is to determine if these structures can support neuromechanical tuning. We performed a comparative analysis of the spatial organization of the motoneuron pools and the functional organization of the muscles they innervate. We developed a three-dimensional model of the spatial organization of motoneuron pools innervating upper limb muscles in the macaque and quantified their relative distances. We then utilized a musculoskeletal model of the macaque upper extremity to characterize the functional relationship between the muscles innervated by the motoneuron pools. We found that the distances between motoneuron pools mimicked the agonistic or antagonistic actions of muscles they innervate. This shows that the spatial organization of the motoneuron pools embeds the functional musculoskeletal anatomy evolved to support the large repertoire of upper limb movements in primates. This spinal anatomy may play an important role in the neuromechanical tuning and simplify the sensorimotor control of the arm.

Introduction

Epidural electrical stimulation is the most promising technique to restore functional movement after spinal cord injury [1–8]. Models of the spinal cord anatomy play a crucial part in surgical planning and tuning of the neuroprosthesis for the needs of the individual [7]. Here we developed a model of the anatomical arrangement of spinal motoneuron (MN) pools of the upper limb and related it to the function of the muscles that MN pools innervate.

The spinal cord drives the musculoskeletal system by rapidly integrating ongoing execution-related (efferent) signals with proprioceptive (afferent) signals to enable motor function in

presence of uncertain environmental conditions. Preserving and taking advantage of
proprioceptive signals is crucial for the optimal augmentation of voluntary movement after spinal
cord injury using neuroprostheses [3]. The afferent signals are somatotopically organized carrying
information from specific body parts to separate spinal cord segments [9,10]. We also know that the
primary motor cortex, one of the main generators of the efferent signals for the motor control of
the upper limb, is somatotopically organized [11–14]. This organization effectively embeds the
anatomical and dynamical properties of the controlled body parts in the neural substrate, a
concept referred to as internal models or, specific to the embedding of the musculoskeletal
system, neuromechanical tuning [15–18]. The neuromechanical tuning serves to simplify the
processing of the proprioceptive signals for the generation of efficient efferent commands. The
question remains as to how much the spinal cord contributes to the neuromechanical tuning.
Answering this question will help develop new methods to tune the neuroprosthetics for maximal
effectiveness.
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Comparative analysis between the MN pool anatomy and the function of muscles they innervate
can help understand how the location of MN pools help integrate the afferent and efferent neural
commands. The general understanding of the anatomical organization of MN pools innervating
the human leg and some arm muscles comes from postmortem studies [19–21]. There is also detailed
anatomical data of the distribution of MN pools innervating muscles of the lower limbs in cats
obtained with retrograde staining [22], which has been comprehensively analyzed [23]. We also know
that the rostrocaudal distributions of these MN pools are similar to those in humans [21,24].
However, the detailed anatomical data about the distribution of MN pools innervating the upper
limb muscles in humans is sparse and lacks identification of their targets [19]. In this study we used
the detailed anatomical organization of MN pools innervating the upper limb muscles of
macaques obtained with retrograde staining [25] to create a spatial model of the MN distributions
in the spinal cord ventral horn. We then took advantage of the detailed information about the
organization of the musculoskeletal system developed using advanced imaging and modeling
techniques [16,26,27]. Our goal is to determine if there is an anatomical substrate in the spinal cord
that can support neuromechanical tuning. We will achieve this goal through the comparative
analysis of the spatial organization of the MN pools and the functional organization of the
muscles they innervate.
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73 **Materials & Methods** All data processing and analyses were done in Matlab (Mathworks,
74 Natick, MA).
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Model of the Spatial Organization of Motoneuron Pools
Locations of motoneuronal cell bodies innervating macaque forelimb muscles were acquired
from a study utilizing retrograde transport of horseradish peroxidase in 10 *M. mulatta* and 4 *M.*
fascicularis monkeys [25]. Images of spinal cord sections from the rostral and caudal parts of the
5th cervical segment (C5) through the 1st thoracic segment (T1) were scanned, digitized using

Adobe Illustrator (Adobe Inc) and imported into Matlab. Metadata from these images
representing cell body locations in the three orthogonal planes (mediolateral, dorsoventral, and
rostrocaudal axes) and the segmental outlines of white and gray matter were used to generate a
three-dimensional (3D) model, similar to our previous work (Yakovenko et al., 2002). One
representative set of transverse grey matter outlines (*M. fascicularis* subject 79-1 from Jenny and
Inukai [25]) was selected as the standard in our model and used for the spatial normalization of
both outlines and MN coordinates along the dorsoventral and mediolateral axes. The published
data for the anatomical size of spinal segments in monkeys [28] was used to scale the rostrocaudal
coordinates of labeled MNs.

The digitization of MNs from published bitmap images of spinal cord section was challenged by
image resolution, color representation, and the superposition of cells in the plane of each slice
and the resolution. To mitigate this, additional data from the published histograms of the quantity
of MNs in each section was utilized [25]. The MN locations obtained from transverse sections
were resampled to match the numbers reported in the histograms for each spinal cord section.
Briefly, if the dataset extracted from transverse sections included higher number of MNs than
reported in the histogram for the corresponding pool in a given section due to mislabeling of
irregular shapes in pixelated images, then MN indices for the given MN pool and section were
randomly selected for removal. Conversely, the reduction in the number of digitized MNs was
also possible due to the potential omission of some images of the transverse sections in the
manuscript figures. This was compensated by adding missing MNs in accordance with the spatial
distribution of the given MN pool and section. The mean position and standard deviation within
transverse slices and the range across spinal segments were held constant for each MN pool.
Symmetry was assumed between left and right sides of the spinal cord when combing data from
different animals to generate a bilateral model (Fig. 1).

To analyze the anatomical organization of MN pool locations, first the centers of MN pools were
calculated by averaging the coordinates of all MNs comprising them. Then Euclidean distances
were then calculated between each MN pool center. Unfortunately, the rostrocaudal distribution
in our sample of MN pools was limited, with most centers of MN pools co-localizing between
T8 and C1 (Fig. 1). Therefore, Euclidian distances were calculated not only in 3D, but also in
2D, with the rostrocaudal distribution excluded to control for the potential noise it introduced in
our analysis.

Musculoskeletal Models of Upper Extremity
We used open-source simulation software, OpenSim (version 4.1, Stanford University, Stanford,
CA, USA), to model the musculoskeletal anatomy of both the macaque and human right upper
limb. The macaque musculoskeletal model developed by Chan and Moran [29] was modified to
add forearm muscles innervated by the MN pools included in the model described above. The
following eight forelimb muscles have been added or modified based on anatomical data
(Berringer, 1968): *adductor pollicis*, *dorsal interossei*, *extensor digitorum*, *extensor pollicis*

longus, *flexor digitorum profundus*, *flexor digitorum superficialis*, *lumbricals*, and *ventral*

interossei (Fig. 2A). Muscles with multiple heads and/or points of insertion were modeled as

separate musculotendon actuators. For example, *biceps brachii* was modeled as two muscles,

biceps long head and *biceps short head*. Similarly, the *extensor digitorum* was modeled as 4

muscles, sharing a similar origin, but inserting on each corresponding distal phalange of digits 2-

5 (index through pinkie). Abbreviations of muscle names used in this study can be found in

Table 1. To simulate more accurately the length changes of the additional muscles in different

postures and during movement, we extended the model to include the carpometacarpal (CMC),

metacarpophalangeal (MCP), and interphalangeal (IP) joints of the digits, increasing the total

number of degrees of freedom (DOFs) in the macaque model to 27. The carpometacarpal joint of

the thumb and the metacarpophalangeal joints of digits 2-4 were modeled with 2 DOFs,

representing flexion/extension and abduction/adduction around the x- and z-axes, respectively.

Interphalangeal joints were modeled as a single DOF representing flexion/extension about the x-

axes.

A detailed model of the human arm was used to assess musculoskeletal differences between

macaque and human musculoskeletal anatomy. The original model developed by Saul, et al.

(2015) was previously expanded to include 23 DOFs and 52 musculotendinous actuators

representing all major joints and 33 major muscles of the human arm (Gritsenko, et al. 2016)

(Fig. 2B). The moment arms and maximal muscle forces were validated against published

measurements [30]. Only the DOFs spanned by muscles innervated by the MN pools modeled

above were included in the comparative analysis of muscle lengths. These DOFs were elbow

flexion/extension, hand pronation/supination, wrist joint flexion/extension and

abduction/adduction, CMC joint flexion/extension and abduction/adduction, MCP joints 1-5

flexion/extension, and all IP joints 1-5 flexion/extension.

To analyze the functional organization of the musculoskeletal systems, musculotendon lengths

were calculated for each forelimb posture across the physiological range of motion within the

model. Then a correlation matrix of muscle lengths across all postures was then computed as in

Gritsenko et al. [16]. Positive correlations indicate a synergistic action of muscles, while negative

correlations indicate antagonistic action. The correlations between muscle pairs whose MN pools

were located closer vs. further apart were compared as described in the Statistical Analysis

section.

<u>Statistical Analysis</u>

The distances between MN pool centers were regressed against the correlation coefficients (R)

between the lengths changes of muscles innervated by the corresponding MN pools. Nearby MN

pools are more likely to share the descending inputs and afferent feedback, which makes them

more likely to innervate synergistic muscles. Therefore, short distances that fell below the

median of the distance distribution were analyzed separately from long distances that fell above

the median. Thus, four regressions were done using 1) short 3D distances; 2) long 3D distances;

3) short 2D distances in the transverse plane; and 4) short 2D distances in the transverse plane.
Familywise error for conducting the four tests was addressed using Bonferroni adjustment [31],
alpha was set to 0.0125.
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Results

<u>Motoneuron clustering in the macaque spinal cord</u>
The MN pools in the cervical enlargement of macaques were distributed along multiple segments
(Fig. 1) similar to those in the lumbar enlargement of cats [23] and humans [19,32]. MNs innervating
proximal muscles in the macaque, such as the *biceps* and *triceps*, together with *flexor carpi*
radialis and *extensor carpi radialis* were located in the rostral part of the cervical enlargement,
while the rest of the MN pools were located in the caudal cervical and thoracic segments (Fig.
1B). In humans, the rostro-caudal distribution of MN pools has been estimated from the
distribution of ventral roots across spinal segments and the muscles they innervate (Kendall and
McCreary, 1983). The rostrocaudal distributions of human and macaque MN pools overlapped
for all muscles (Fig. 3A). Although the MN pools innervating digit and some wrist muscles
appeared to be more rostrally located and span more segments in humans compared to macaques.
This may be due to the fewer number of spinal segments in humans compared to macaques [33],
which may allow for more separation between MN pools in the rostrocaudal dimension in
macaques. The distribution of MN pools in the transverse plane in macaque and human spinal
cords also overlapped (Fig. 3B). In humans, the distribution of the MN pools in the transverse
plane was estimated using postmortem staining by cresyl violet, as summarized in [19]. The human
MN pools in the lateral division of the ventral grey horn overlapped with those in the macaque
(Fig. 3B). Although it was not possible to identify which muscles the human MNs innervated,
the overlap with the macaque data suggests that the human MN pool labeled as 5 may innervate
biceps and the human MN pool labeled as 7 may innervate triceps (Fig. 3B). Overall, these data
show that there is a notable overlap between MN pools of humans and macaques.
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To quantify the anatomical organization of spinal MN pools we calculated the Euclidian
distances between the centers of MN pools. The MN pools innervating *biceps* and *extensor carpi*
radialis were the furthest apart from the rest of the MN pools, while being a relatively short
distance from each other (Fig. 4A, dark red cluster in the top left corner). Moreover, several
other MN pools innervating synergistic muscles were co-localized, such as the MN pool of *flexor*
digitorum superficialis was a relatively short distance apart from that of *flexor digitorum*
profundus. The MN pools innervating two long thumb muscles (*extensor* and *abductor pollicis*
longus) were co-localized. The MN pools innervating two other thumb muscles (*adductor*
pollicis and *abductor pollicis brevis*) were also co-localized with those innervating intrinsic hand
muscles (*dorsal interosseous* and *lumbricalis*) (Fig. 4A, dark red cluster in the bottom right
corner). The close anatomical arrangement of MN pools remained the same when the
rostrocaudal distribution was excluded from the distance calculations (Fig. 4B). This anatomical

arrangement suggests that the MN pools innervating agonistic muscles that perform a single
function, such as spreading fingers apart in the case of the last cluster, are located closer together
than the MN pools innervating antagonistic muscles that perform distinct functions. We have
previously conducted a rigorous analysis of the agonistic and antagonistic relationships between
muscles using a model of musculoskeletal anatomy of the human arm [16]. Here we have included
a musculoskeletal model of the macaque upper limb and compared the functional relationships
between muscles to the anatomical relationships between MN pools that innervate those muscles.

To evaluate the functional relationship between muscles we correlated changes of muscle lengths
across the whole range of possible upper limb postures [16]. The logic here is that muscle length
changes that are positively correlated indicate agonistic action, while muscle length changes that
are negatively correlated indicate antagonistic action. For example, during wrist extension
extensor carpi ulnaris and *extensor digitorum communis* would shorten and their lengths would
be positively correlated indicating their agonistic action, while *flexor digitorum profundus* and
flexor digitorum superficialis would lengthen and their lengths would be negatively correlated
with that of the former two muscles indicating their antagonistic action. Indeed, all muscles but
dorsal interosseous and *lumbricalis* had strong agonistic or antagonistic relationships with at
least one other muscle based on the length comparisons using the macaque musculoskeletal
model (Fig. 4C). This shows that there are strong functional relationships between most muscles
of the upper limb that are innervated by the MN pools included in the spinal cord model.
Moreover, the functional relationships between all major muscles of the upper limb of the
macaque were very similar to those observed in human arm (Fig. 5). This shows that the
functional relationships are preserved across species with similar anatomy.

Next, we compared the functional relationship between muscles with the spatial relationship
between MN pools innervating these muscles. We examined separately the 3D Euclidian
distances between MN pool centers that fell below and above the median of the distribution (Fig.
6A). There was no significant linear regression between the relative locations of nearby MN
pools and the functional relationships of the corresponding muscles (Fig. 6B). However, when
the Euclidian distances were calculated only in the transverse plane, i.e., excluding the
rostrocaudal distribution (Fig. 6C), a significant linear regression was revealed between the
relative locations of nearby MN pools and the functional relationships of the corresponding
muscles (Fig. 6D). No significant regressions were found for MN pools that were located further
apart from each other (3D locations: $R^2 = 0.127$, $p = 0.16$; 2D locations: $R^2 = 0.071$, $p = 0.57$).
This shows that the spatial distribution of the MN pools in the transverse plane mimics the
functional relationships between the muscles they innervate.

Discussion

Here, we have developed the first spatial model of cervico-thoracic MN distributions in the
macaque. Using this model, we have shown that the MN distributions in the macaque spinal cord

are broadly similar to those in human spinal cord (Fig. 3). Using the musculoskeletal models of
the macaque and human upper limbs we have also shown that the muscle anatomy of both
species serves similar agonistic and antagonistic actions (Fig. 5). We then demonstrated
quantitatively that the spatial distribution of MN pools co-localized in the transverse plane
mimics the functional relationships between the muscles these MNs innervate (Fig. 4, 6). This
suggests that the spatial organization of the MN pools embeds the functional musculoskeletal
anatomy evolved to support the large repertoire of human arm movements [16].

The functional organization of the spinal cord is modular [34]. This conceptual understanding
emerged from early studies in spinalized animals showing that microstimulation inside the spinal
cord grey matter produces contractions of multiple muscles, which cause converging forces
toward a subset of postures [35]. The idea then emerged that muscles can be controlled in groups
with fewer control signals of varying spatial and temporal patterns that can create a large
repertoire of movements [36]. There is also a parallel line of investigation of the spinal control of
locomotion that reveals the modularity of spinal control circuitry including the central pattern
generator (CPG) [37]. Evidence supports the importance of CPG in human sensorimotor control
[4,38,39]. There is evidence that the different spatiotemporal dynamics of the CPG and its interaction
with the body through afferent feedback can generate different behaviors. This can be seen in
neuromechanical studies with lampreys where varying the spatiotemporal patterns of tonic
excitation to CPG and its propagation along the spinal segments can produce swimming in either
forward or backward direction [40] and change the speed, direction and type of gait [41,42]. Some of
the spatiotemporal dynamics of the CPG during gait has been shown to be reflected in the
anatomical distribution of the MN pools innervating hindlimb muscles of the cat [23], supporting
the idea of the somatotopic "map" of a hindlimb [43]. Our study provides supporting evidence for
the idea of the somatotopic "map" of a forelimb. Our results show that the MN pools innervating
muscles that perform synergistic action are co-localized in the spinal cord. Nearby MN pools are
also likely to share the proprioceptive feedback from the muscles they innervate as has been
shown for cat knee extensor muscles and their afferents [44]. Similarly, nearby MN pools are likely
to receive common descending signals from propriospinal interneurons [45] that can explain
observations of the convergent force fields that are thought to serve as building blocks for more
complex motor control [35,46]. Furthermore, nearby MN pools are likely to receive common
descending signals from the somatotopically organized motor cortex that can explain
observations of the complex movements in different areas of workspace produced as a result of
the stimulation of the primary and premotor cortex [47]. Overall, our results suggest that the spatial
organization of MN pools may play an important role in neuromechanical tuning and simplify
the sensorimotor control of the upper limb.

Recent work has shown that it is possible to restore trunk and leg motor functions using
biomimetic epidural electrical stimulation of the dorsal roots below the lesion caused by the
spinal cord injury [7]. Our results suggest that a similar approach may work for restoring complex

movements of the arm after spinal cord injury. A proof of concept study in macaques have
shown that it is possible to evoke selective muscle responses by stimulating cervical dorsal roots
[48]. Notably, more rostral stimulation in this study evoked responses in *biceps* and *extensor carpi*
radialis, while more caudal stimulation evoked responses in *flexor digitorum superficialis*,
extensor digitorum communis, and *abductor policis brevis*. The rostrocaudal distribution of MN
pools innervating these muscles is similar to that shown by our model (Fig. 3A). These
observations support the validity of our model and indicate potential future uses of it for
designing next generation neuroprosthetics.
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289 **Acknowledgements**
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294 **Tables & figure captions**
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296 **Table 1. Muscle abbreviations used in this study**

Abbreviation	Muscle Name	Abbreviation	Muscle Name
ANC	Anconeus	FCU	Flexor carpi ulnaris
AP	Adductor pollicis	FPL	Flexor pollicis longus
APB	Abductor pollicis brevis	FDP	Flexor digitorum profundus; 2-5 digits
APL	Abductor pollicis longus	FDS	Flexor digitorum superficialis; 2-5 digits
BIC	Biceps; l-long, s-short heads	ISPI	Infraspinatus
BR	Brachialis	LAT	Latissimus dorsi; r-rostral, m-medial, c-caudal heads
BRR	Brachioradialis	LUM	Lumbricalis (lateral)
CORBR	Coracobrachialis	PALL	Palmaris longus
DEL	Deltoid, a-anterior, m-medial, p-posterior heads	PEC	Pectoralis; r-rostral, m-medial, c-caudal heads
DIO	Dorsal interosseous	PQ	Pronator quadratus
DOR	Dorsoepitrochlearis (latissimocondyloideus)	PT	Pronator teres
ECR	Extensor carpi radialis; l-longus, b-brevis	SSCAP	Subscapularis
ECU	Extensor carpi ulnaris	SUP	Supinator

ED	Extensor digitorum; 2-5 digits	SSPI	Supraspinatus
EDM	Extensor digiti minimi	TERMA	Teres major
EDS	Extensor digiti secundi proprius	TERMI	Teres minor
EI	Extensor indicis	TRI	Triceps; lo-long, la-lateral, m-medial heads
EPL	Extensor pollicis longus	VIO	Ventral interosseous
FCR	Flexor carpi radialis		

Figure 1. Model of motoneuron pool anatomy in a macaque. A. Circles show averaged locations of the MN pools for each muscle in the transverse plane, the shaded areas are standard deviations along the dorsoventral and mediolateral directions calculated across all motoneuron locations in C5 through T1 segments. Grey matter outlines are shown for the caudal portion of the fifth cervical segment (C5C) and the caudal portion of the first thoracic segment (T1C, grey fill). The circle at the axis's origin represents the central canal. **B.** 3D view of the individual motoneuron locations (colored circles) and grey matter outlines for spinal segments C5 - T1. Central canal locations are shown as open circles. Motoneuron pool colors are the same as in A. Insert on the right shows the rostrocaudal distributions of the motoneuron locations with black circles indicating averages.

Figure 2. Musculoskeletal models of a macaque (A) and human (B) upper limbs. Red lines illustrate the origin, insertion, and wrapping geometry of the musculotendinous actuators representing the anatomical arrangement of individual muscles or muscle compartments. For example, two heads of biceps are modelled as two actuators with origin locations on different bones and a common insertion location. All actuators are shown, but not all are labeled. The bones are shown for illustration purposes only, the inertial geometries of limb segments to which the actuators attach are not shown.

Figure 3. Comparison of MN locations between the macaque and human spinal cords. A. Circles show the distribution of MNs along the rostrocaudal direction across spinal segments from the macaque model in Fig 1. Bars show the distribution of corresponding MN pools from a human anatomy textbook [32]. **B.** Circles show MN locations in the transverse plane per spinal segment from the macaque model in Fig 1. Grey matter outlines aligned on the central canal at the origin of the axes are from the macaque model. Black open ovals show the distributions of select MN pools adapted from human tracing studies summarized in [19]. The human distributions are numbered as in the Routal and Pal (1999).

Figure 4. Anatomical relationships of MN pools and the functional relationships between muscles they innervate. Correlation matrix of the relative distances between the centers of MN pools in all dimensions (**A**) and only in the transverse plane with rostrocaudal distribution

excluded (**B**) calculated using the model shown in Fig. 1. Distances were calculated between the centers of MN pools for each pair of MN pools relative to the maximum distance across all pairs. Black indicates short distances, white indicate long distances. **C.** Matrix of Pearson correlation coefficients (R) between muscle length changes evaluated across the full range of motion of the macaque forelimb. Yellow indicates agonistic actions; blue indicates antagonistic actions.

Figure 5. The functional relationships between upper limb muscles of human and macaque. Matrices of Pearson correlation coefficients (R) between muscle length changes evaluated across the full range of motion of the human arm **(A)** and macaque forelimb **(B)** shown in Fig. 2. Yellow indicates agonistic actions; blue indicates antagonistic actions as in Fig. 4C.

Figure 6. Comparative analysis of spatial MN pool anatomy and functional relationships between muscles they innervate. A. Histogram of 3D distances between MN pools. Vertical black line shows the median, only distances below the median were used for the regression in B. **B**. Short 3D distances between MN pools as a function of Pearson correlation coefficients (R) between lengths of muscles those MN pools they innervate. The regression line is not significant as shown by the inserted statistics. **C.** Histogram of 2D distances in the transverse plane between MN pools. Vertical black line shows the median, only distances below the median were used for the regression in D. **D**. Short 2D distances between MN pools as a function of Pearson correlation coefficients (R) between lengths of muscles those MN pools they innervate. The regression line is significant as shown by the inserted statistics.

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A

Pectoralis

Biceps

Brachioradialis

Supinator

Abductor pollicis longus

Extensor digiti minimi

Extensor pollicis longus

B

Deltoid

Triceps

Flexor carpi ulnaris

Extensor carpi ulnaris

Extensor pollicis brevis

Extensor digitorum

A 3D distances between MN pools

B 2D distances between MN pools

Relative Distance

C Mechanical coupling between muscles

R

A Human

B Macaque

A 3D distances between MN pools

C 2D distances between MN pools

B R² = 0.06; p = 0.3580

D R² = 0.29; p = 0.0105