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4 **Title:** Modeling of the Thigh: a 3D deformable approach considering
5 muscle interactions

6

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11 **Deadlines:**

12 - End of March 2016: first draft submission

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17

18 **Notes:**

19 Authors involved in parts 2 and 3 (chapters 6 to 25) are asked to limit their text to 7 500 words
20 (approximately 15 pages). There should be a maximum of 6/7 figures and/or tables per chapter with
21 2 to 3 figures that can be printed in color.

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Abstracts (135 words). *“should be 100-150 words”*

A 3D deformable model of the thigh’s musculoskeletal system has been developed. An adjucted multi-body rigid model was used to compute the bone kinematics and the muscle activation levels which drive a Finite Element model of the musculoskeletal system integrating contractile muscles. A thermomechanical beam model for muscle fibers was embedded in the passive muscle matrix meshed with hexahedrons with a hyperelastic quasi incompressible material law. Contacts between surrounding anatomical structures, transmitting radial forces, were modeled to investigate the effect of the interactions between muscles on the production of muscle forces. Two steps of the gait cycle were simulated. This study demonstrated that integrating such a deformable model could update the musculo-tendon forces traditionally computed from a multi-body rigid model that fails to take into consideration 3D fiber distribution as well as muscle tightening effect.

Keyword: thigh, 3D deformable contractile muscles, interactions between muscles, radial forces, multi-body rigid model, Hill-type model, gait

1 **Words:** 7353.

2 Modeling of the Thigh: a 3D deformable approach considering muscle 3 interactions.

4

5 Introduction

6 Muscles contribute to the control of the bones kinematics and the joint stiffness, the stability during
7 movements, and the propulsion for the movement task. The muscles transmit the force generated by
8 the sarcomeres to bones through the myotendinous junction, but also transmit the force to
9 surrounding anatomical structures by the connective tissue. Therefore, the relative position of
10 muscles with respect to the surrounding anatomical structures could be of major importance on the
11 muscle force production (Yucesoy, 2010). So dealing with muscles as isolated functional units
12 producing force according to force-length/force-velocity relationships could lead to misestimate their
13 actions. If the literature based on experiments and case-study models have largely investigate the
14 myofascial force transmission as a shear loading (Purslow, 2010), Siebert et al. (2014) reported
15 recently that the muscle force production could decrease due to a transverse compression: this
16 opens up new understandings on the muscle force transmission to surrounding anatomical
17 structures through radial loading. This appears as essential in the muscular architecture of the thigh.
18 The musculoskeletal thigh system is made of the largest skeletal muscles of the whole body. The
19 powerful thigh muscles, between the pelvis and the knee, contribute to the locomotion, the stability
20 maintenance, and the support of the full body weight. Due to the large muscle loads applied, the hip
21 joint, tibiofemoral and patellofemoral are ones of the most degenerative joints (Kim and James,
22 2012; Zhang and Jordan, 2010). If the lower limb is still one of the most investigated part in the body
23 through musculoskeletal modeling, the complexity of muscular architecture of the thigh system often
24 yielded to simplify its numerical representation.

25 Investigating musculoskeletal functions and disease requires a framework integrating a relevant soft
26 tissue modeling as well as a physiological contractile ability within muscle models for relevant stress
27 field and pressure estimation. However developing such a framework is still challenging, by gathering
28 together mechanical, physiological and numerical issues under the constraints of allowing for future
29 personalization (Fernandez et al., 2006). So the few authors investigating it chose a multi-body rigid
30 model with reasonable computation cost for the computation of the boundary conditions
31 (kinematics and musculo-tendon forces), to drive a deformable joint finite element model, most of
32 the time, in quasi-static simulations (Adouni and Shirazi-Adl, 2013). However, driving a deformable

1 musculoskeletal model require to deal with large deformation of soft tissue, contact issues (implying
2 heavy computational costs), and, in dynamic simulations, kinematically incompatible boundary
3 conditions between the multi-body rigid model and the deformable musculoskeletal one, which
4 limits the level of details of the latter. Moreover, modeling the whole musculoskeletal thigh system
5 with a solid representation remains mainly based on a continuum mesh of the detailed anatomical
6 structures, with passive muscle modeling as hyperelastic materials using first order Ogden law (Al-
7 Dirini et al., 2016). In computer graphics applications (Lee, 2011), the simplest approach consists in a
8 pure geometric deformation (i.e., without any material law) of the muscle surface according to the
9 kinematics of bones, also called skinning (Dicko et al., 2013; Dong et al., 2002; Kohout et al., 2013;
10 Murai et al., 2016). For more realistic approaches Li et al. (2013), Fan et al. (2014) and Sachdeva et al.
11 (2015) proposed an Eulerian-on-Lagrangian (EOL) approach for skin, muscle and tendon models
12 respectively. The Eulerian discretization allows efficient handling of large incompressible
13 deformations, collision detection between many contacting muscles on a single and uniform grid per
14 bone. However, the authors use a pure isotropic elastic material law without taking into account
15 fibers, and reported numerical issues leading to unrealistic soft tissue deformations at the boundary
16 of the grids. Authors from computer graphics were also ones of the firsts to drive complex
17 musculoskeletal deformable models with bone kinematics and muscle activation levels computed
18 from inverse dynamics analysis using an adjusted multi-body rigid model (Lee, 2011). The most
19 tremendous framework was initiated by Teran et al. (2005). The authors constructed a model of the
20 musculoskeletal system of the upper limb from the Visible Human dataset (with more than thirty
21 muscles). To assure numerical convergence for a reduced computational cost, the authors developed
22 a finite volume approach (which was a variation of the linear tetrahedral finite element formulation)
23 corrected to deal with wrapped elements. Based on this work, Lee et al. (2009) increased the level of
24 details of the musculoskeletal deformable model of the upper body, with 68 bones and 814 muscle
25 actuators based on the linearized Hill-type law for the adjusted multi-body rigid model. Finally, Si et
26 al. (2014) used this framework to simulate the whole human body swimming (with a model made of
27 103 rigid bones and 823 muscles). In all the three above mentioned studies, the muscle tissue was
28 represented as a transverse isotropic material law: a Mooney-Rivlin rubber-like model for the passive
29 part plus a deviatoric stretch in the along-fiber direction based on the activation level. The fiber
30 direction inside each tetrahedron resulted from a B-spline interpolation. To avoid contact elements
31 and speed up the simulation, a coarse tetrahedral mesh embedded the detailed musculoskeletal
32 model. This embedding framework, called the "simulation mesh" by the authors, was initiated and
33 theorized for general soft body deformation in Sifakis et al. (2007): the "simulation mesh" was an
34 adaptively coarsen embedding tetrahedral mesh in which the detailed skin surface (triangulated
35 mesh for rendering and collision handling), the fatty tissue, the muscles and the bones were

1 embedded. In the Teran/Lee approach, a network of spring constraints linked the deformable
2 embedding mesh to the rigid articulated skeleton. Nodes of the embedding part close to the surface
3 nodes of the embedded muscles were links for the muscle surface model, leading to a frictionless
4 contact representation between muscles and avoiding the spurious separation of muscles. Another
5 published framework proposed to use springs to model the interfaces between the biceps brachii
6 muscle and its neighbors to conserve reasonable computation time over surface contact elements
7 (Böl et al., 2011a, 2011b). The authors proposed a finite element model of the musculoskeletal upper
8 arm to simulate isometric contraction in five arm positions. No bone kinematics was taken into
9 account, the reconstruction of the geometry of the muscles being segmented from MRI at each
10 position. The passive response of the muscle was based on a hyperelastic incompressible transversal
11 isotropic constitutive law, with an along-fiber stretch contribution coming from the activation by
12 electromechanical coupling. The interface spring stiffness were identified to fit experimental surface
13 displacements on one point.

14 For applications related to medical engineering and research in biomechanics Fernandez et al. (2005)
15 developed a framework based on cubic Hermite elements, to investigate the patella kinematics and
16 pressure on it, driven by a flexion of the tibia, with deformable models of the quadriceps muscles
17 (rectus femoris and the vastus group). The quadriceps were modeled without contacts between
18 anatomical structures, the vastus muscles being fixed on the femur. The contact modeling was
19 restricted to the interactions between the patella, the femur and the tendons from the patella. The
20 muscle passive matrix was modelled with an orthotropic law with volume conservation. The fiber law
21 was issued from cardiac activation tissue law: the fiber-based active contraction law guaranteed a
22 local contraction field in the fiber field direction. However the activation level was arbitrary chosen
23 only to assure the numerical stability. Behr (2006) proposed a solid model of the musculoskeletal of
24 the lower limb for car crash applications. The hexahedron passive muscle mesh was associated to a
25 viscoelastic anisotropic quasi-incompressible material law, with various elastic modulus depending
26 on the fiber direction. Beams embedded in the solid mesh were chosen to model the fibers, and the
27 activation level was represent as a thermal load. The contacts between muscles were represented,
28 and the muscle activation was set through a thermomechanical coupling. However the lower limb
29 was fixed in sitting posture and the activation levels were set arbitrarily. Blemker reported a model of
30 the hip muscles and chiefs of the quadriceps (Blemker et al., 2005; Blemker and Delp, 2006). The
31 authors proposed to model the muscle as a quasi-incompressible, fiber-reinforced composite with
32 transversely isotropic material law to integer along-fiber shear strain, cross-fiber shear strain, and
33 volume strain. The activation resulted in a modification of the along-fiber stretch contribution within
34 the material law. If the authors proposed a framework to take into account a physiological complex

1 fiber field within the muscle material law, few muscles were finally modeled and the knee flexion
2 was simulated using bone kinematics and a homogenous 20% activation level.

3 In this context, the present study presents an efficient approach for modelling the human thigh with
4 3D deformable contractile muscles driven by a set of muscle activations. The specific aim is to
5 investigate the effect of the interaction between muscles, leading to radial loading, on the
6 production of muscle forces. To succeed in modeling such a complex anatomical system, an adjucted
7 ulti-body rigid model was used to compute the bone kinematics and the muscle activation levels.
8 These inputs drove a 3D deformable model of the musculoskeletal thigh system, with a robust
9 contractile model based on a thermomechanical coupling: the fibers were modeled as beams with
10 thermomechanical properties to represent the activation level as a variation of temperature, and
11 then these fibers were associated to the passive matrix meshed with hexahedrons with a
12 hyperelastic quasi incompressible law. Such an approach should guarantee to simulate a part of the
13 gait cycle in a reasonable computational cost, taking into account the contacts between surrounding
14 anatomical structures transmitting radial forces.

15

16 [Background: anatomical and functional description of the organs](#)

17 The osteoarticular system of the thigh is made of the femur, articulated with the pelvis through the
18 hip joint and with the shank through the knee joint (tibiofemoral and patellofemoral joints).

19 Muscles of the thigh are clustered in three compartments, surrounding by the Fascia Lata (cf. Figure
20 2.1): the anterior compartment (including ctineus, sartorius, and the quadriceps with four chiefs:
21 rectus femoris, vastus lateralis, medialis and intermedialis), the hamstring muscles of the posterior
22 compartment (semimembranosus, semitendinosus, and the long and short heads of biceps femoris),
23 the medial compartment muscles (adductor magnus, longus and brevis nd gracilis). At the hip level,
24 the gluteal group is made of the gluteal muscles (minimus, medius, and maximus) and the Tensor
25 Fascia Latae muscle.

26

[Insert Figure 2.1 here]

27

[Insert Figure 2.2 here]

28 As reported in figure 2.2, several muscles are involved in the thigh musculoskeletal system, leading to
29 complex interactions during their contractions. Apart for the well-known agonist-antagonist
30 synergies, mechanical connections through collagenous linkages also contribute to transmit forces
31 intermuscularly through epimuscular myofascial force transmission (Yucesoy et al., 2010). Moreover,

1 the intramuscular extracellular matrix (made of the endomysium, perimysium and epimysium), but
2 also the muscular septi, the aponeuroses and the joint capsules (depending of the fascia definition
3 (Schleip et al., 2012)) are surrounding by the fascia. The fascia are any connective tissue linking all
4 the anatomical structures (organ, muscles etc.) leading to a continuum throughout the human body
5 (Findley and Shalwala, 2013). The connective tissue results in myofascial force transmission between
6 adjacent muscles or group of muscles, and between muscular and non-muscular tissues. Fascia
7 allowing the sliding between anatomical structures, the network of connective tissue should transmit
8 shear loading between muscles. Fascia form an extensive network especially throughout the limb,
9 with deep fascia defining a functional organization groups of muscles (i.e., the muscle compartments
10 described above).

11

12 Material and Methods

13 1. Multi-body rigid modelling of the lower limb

14 The multi-body rigid model of the musculoskeletal system of the lower limb, used in adjunction to the
15 deformable model, was developed previously (Moissenet et al., 2014). Five segments (pelvis, thigh,
16 patella, shank and foot) were considered as rigid bodies, and articulated by four joints (ankle,
17 tibiofemoral, patellofemoral and hip). The hip joint was modelled as spherical. The tibiofemoral joint
18 was modelled as an equivalent mechanism made of two sphere-on-plane contacts for the medial and
19 lateral compartments and three isometric ligaments (anterior cruciate, posterior cruciate and medial
20 collateral ligaments). The patellofemoral joint was modelled as an equivalent mechanism made of a
21 hinge joint between the patella and the femur and an isometric patellar tendon. The ankle joint was
22 also considered as an equivalent mechanism, with a spherical talo-crural lateral contact and two
23 isometric ligaments (between the tibia and the calcaneus, and between the fibula and the
24 calcaneus).

25 The muscles were represented by 43 lines of actions (Delp et al., 1990), with via-points from the
26 origins to the insertions to specify the muscle path. Some muscle geometries were divided into one
27 to three parts to take into account extended insertion sites. The muscle lever arms were computed
28 for each action line, yielded to a muscular redundancy. The multi-body rigid model of the
29 musculoskeletal system of the lower limb is illustrated here after (cf. Figure 2.3).

30

[Insert Figure 2.3 here]

31 First, the segments positions, velocities and accelerations resulted from a typical multi-body
32 kinematic optimization (under rigid body and joint-related kinematic constraints), to minimize the

1 squared distances between experimental and model-determined skin markers positions (Duprey et
2 al., 2010). Second, the muscular redundancy problem was solved by inverse dynamics and static
3 optimization, minimizing simultaneously the sum of the squared musculo-tendon forces contact
4 forces, ligament forces and bones forces (Moissenet et al., 2014).

5 This model of the lower limb has been previously evaluated according to the « Grand Challenge
6 Competition to predict in vivo knee loads » dataset, previously described in details (Fregly et al.,
7 2012). A part of the second dataset was reused in the present study (Moissenet et al., 2014): the
8 experimental data recorded for one subject (male, 83 year-old, 67 kg, 1.72 m) walking at a self-
9 selected speed. Marker trajectories and ground reaction forces for one gait cycle were extracted
10 from this dataset and used in this study. The multi-body rigid model was scaled to the subject
11 anthropometry to fit the segments length and the bones kinematics and musculo-tendon forces were
12 computed at every frame of the gait cycle. These information were used to drive the deformable
13 model of the musculoskeletal system of the lower limb. For that, from the musculo-tendon forces
14 and the musculo-tendon dynamics, the muscle activation levels were computed using a Hill-type
15 model (Chèze et al., 2015).

16

17 2. Deformable modelling of the thigh

18 *Geometry and Meshing*

19 The detail geometry of the musculoskeletal system of the right thigh was issued from the manual
20 segmentation of the Visible Human Project® (Laine) imagery dataset. The anatomical structures were
21 manually segmented and cleaned before meshing. The five bones (pelvis, femur, patella, tibia, and
22 fibula) were represented by the external wall of the cortical bones, to be considered as rigid bodies.
23 The five ligaments (lateral [LL], medial collateral [ML], anterior cruciate [ACL], posterior cruciate
24 [PCL]) and the patellar ligament [PL] were modeled with non-linear spring elements. The quadriceps
25 tendon was meshed with tetrahedrons due to its thin shape.

26 The muscle matrix was meshed with hexahedral elements, oriented close to the line of action of each
27 muscle. The solid meshing approach was based on a block template iteratively deformed to fit the
28 muscle geometry, with a constraint on the quality of elements (scaled jacobian metrics upper than
29 0.4). The muscle fibers were modeled by a network of beam elements, in the principal orientation of
30 each muscle (i.e. its line of action), and merged with the edge of hexahedrons (cf. Figure 2.4). The
31 tetrahedral formulation was not considered, despite of meshing facility, to guaranty both a smooth
32 field of deformation for a limited number of elements, and high quality elements supporting the

1 surface contacts for an effective transmission of loads even with large deformation. If recent complex
2 methods could provide a full hexahedral mesh of the muscles, the deformation of a block to mesh
3 muscles provided a simple space decomposition of the solid mesh into external layers and a deep
4 volume with contraction capability, with a coherent numbering of nodes appropriate for the edge
5 detection. In addition, the orientation of the solid mesh according to the line of action of muscles
6 allowed to use beams for the fibers directly connected to the nodes of hexahedrons.

7 **[Insert Figure 2.4 here]**

8 17 muscles were considered to represent the musculoskeletal system of the thigh, with the three
9 compartments of the thigh and the gluteal one for the hip. Quadrangle shell elements were
10 extracted from the boundary of the solid part of the muscle to represent the aponeuroses of the
11 muscle. For the muscle insertion and origin, the muscle mesh was attached to bones by using quasi-
12 rigid elements, located at the anatomical insertions surfaces. The Tensor Fascia Latae muscle was not
13 introduced in the model: after segmentation and meshing, the extremities of this muscle were far
14 from the original ones in the image dataset to ensure quality of elements, leading to very long quasi-
15 rigid elements modeling the insertion and, thus, non-physiologic lever arms. The minimal initial gap
16 between anatomical structures was set to 5 mm. The whole deformable model of the lower limb
17 consisted to 160.000 elements and 135.000 nodes (cf. Figure 2.5).

18 **[Insert Figure 2.5 here]**

19

20 *Mechanical properties*

21 The material properties of the ligaments and tendons were considered as elastic and isotropic, with
22 Young's modulus of 0.15 MPa and a Poisson's ratio of 0.495. A hyperelastic law (Mooney-Rivlin, A = 8
23 MPa, B = 2 MPa, Poisson ration 0.490) for the quadriceps tendon. The passive and active properties
24 of the muscles were represented, and a thermomechanical coupling approach was used to introduce
25 the activation capability. A hyperelastic law (Mooney-Rivlin, A = 3 kPa, B = 0.75 kPa) was assigned to
26 the quadrangle elements and to the hexahedral elements of the muscle matrix (Gras et al. 2011). A
27 linear thermo-mechanical law ($E = 1 \text{ MPa}$, $\text{ALPHA} = 0,01 \text{ mm}^3 \cdot \text{K}^{-1}$) was assigned to the beam
28 network embedded into the solid mesh. The activation of the fibers was driven by the temperature
29 imposed to the beams, yielding to a reduction in length as for the contraction task. A constant
30 activation field was used for all the fibers of each muscle. For each muscle, the behavior was fitted on
31 the Hill-type model (Zajac, 1988), by optimizing first the equivalent modulus of elasticity of the
32 Mooney-Rivlin law for the passive behavior from the initial values above, and then the temperature

1 field for the global (i.e. passive and active) behavior for a given level of activation as reported by
2 Stelletta et al. 2013 (cf. Figure 2.6). In the present study, only the active component was identified
3 due to small elongations of the muscles for the gait cycle steps considered. The CC Hill-type model
4 was chosen according to Siebert's recommendation between the two widely used Hill-type models, to
5 better represent the nonlinear contraction dynamics in musculoskeletal modeling (Siebert et al.,
6 2008).

7 *Boundary conditions and simulation*

8 Two main steps of the gait cycle, the Heel Strike (HS) and the Loading Response (LR) steps, were
9 simulated thanks to an explicit analysis with the LS-DYNA software (LSTC, Livermore, CA). The initial
10 position of the lower limb model corresponded to the HS configuration as defined by the multi-body
11 rigid model (i.e., multi-body kinematic optimization). First, the HS activation levels computed from
12 the multi-body rigid model (i.e., static optimization) were applied to the fiber muscles of the
13 deformable model according to a ramp function. From the previous configuration, the LR kinematics
14 of the segments (corresponding to a 15° knee flexion) and the LR activation levels, both computed
15 from the multi-body rigid model, were applied to the rigid bones and the fibers muscles of the
16 deformable model, also through a ramp function. The kinematics and the activation levels were
17 maintained to stabilize the mechanical behavior of the explicit integration scheme. A slight mass
18 scaling was added to increase the time step from 10^{-6} to 10^{-5} s.

19 To investigate the effect of the interaction between muscles, two configurations were evaluated for
20 each gait cycle step. The first configuration corresponded to the simulation based on the previous
21 model; the second configuration consisted in adding contact surface elements between anatomical
22 structures, to take into account the interaction between muscles and between muscles and bones.
23 The perfect sliding contact model guarantee to transmit loads normal to the boundary only,
24 according to the anatomical statement that "under the deep fascia, the muscles are free to slide
25 because of their epimysium" (Stecco et al., 2011, 2008).

26 *Post-processing*

27 At the end of the Heel Strike (HS) and the Loading Response (LR) steps, the longitudinal force inside
28 each thigh's muscle was computed. All the elements of the muscle mesh intersected by the cross
29 section plane contributed to the longitudinal force: the net forces obtained in the beam, the solid
30 and the shell elements intersected by the cross section were projected onto the normal of the plane.
31 Thus, the addition of the projected total forces combine both the active (from the beam net forces)
32 and passive contribution (from the solid and shell net forces). For comparison purpose, the musculo-

1 tendon force vectors computed from the multi-body rigid model were sum up when the muscle was
2 modelled by multiple lines of actions.

3

4 Results

5 The multi-body rigid model of the musculoskeletal system of the lower limb was used to compute
6 from experimental data the segment kinematics and the activation levels in all thigh muscles, for all
7 the frames of the gait cycle. For both the Heel Strike (HS) and the Loading Response (LR) steps, some
8 muscles presented a level of activation equal to zero. Using this information, the finite element
9 simulations of the two gait cycle steps were successfully achieved in 10 minutes per configuration on
10 a HP Z600 workstation (Quad Core Intel Xeon E5620 2.4Hz). Passive and active forces inside each
11 muscle were computed from the finite element simulations. The total forces were reported in Table
12 2.6.

13

[Insert Table 2.6 here]

14 The two configurations of the deformable model, without and with contacts between anatomical
15 entities, provided muscle forces always consistent with the multi-body rigid modelling of the lower
16 limb, for the Heel Strike (HS) step. The mean [MAX] absolute difference on the total muscle forces
17 was around 4.3N [20N] without contact, and 2.7N [10.3N] with contact. From a general view, there
18 was no difference regarding the two configurations. Indeed for this gait cycle step, the bones were
19 fixed and the only inputs were the muscle activation levels from the multi-body rigid model. So the
20 contraction of the muscles was simulated isometric, without an evolution of the muscle cross
21 section. In addition, in the absence of any variation of the muscle conformation due to the
22 kinematics, the interaction between the muscles was limited: no geometric penetration was noted in
23 the absence of contact element. The main differences with the muscle force patterns from the multi-
24 body rigid model was due to a few modification of the muscle lever arms. As the origins and
25 insertions were fixed, only the contraction tended to reduce the curvature of the muscles, resulting
26 in a slight modification of the lines of actions and so of the produced muscle forces.

27

[Insert Figure 2.7 here]

28 For the Loading Response (LR) step, the two configurations of the deformable model yielded to
29 muscle forces in the same order of magnitude than from the multi-body rigid modelling of the lower
30 limb. The mean [MAX] absolute difference on the total muscle forces was slightly higher than for the
31 HS step, around 13.0N [26.9N] without contact, and 17.5N [112.0N] with contact. For this step of the

1 gait cycle, the muscle deformable models presented large penetrations in the absence of contact (cf.
2 Figure 2.7): the bone kinematics contributed to modify the interaction between structures in
3 addition to a higher level of contraction.

4 Some muscles did not produced any forces according to the multi-body rigid model. In the
5 deformable model, the solid passive part of the muscles had a material behavior yielding to a
6 production of forces even for a negative strain field. However, the behavior of the muscle
7 deformable model was optimized to fit the Hill-type model in traction, so the passive contribution
8 was limited in compression. In addition, the muscles without activation being mainly in the medial
9 and posterior compartments of the thigh, they moved away during the flexion of the knee. In the
10 absence of the skin or any fascia clustering muscles in coherent groups, the contact status was not
11 active for these muscles.

12

13 Discussion and Perspectives

14 This study reports a framework for modeling the musculoskeletal system of the human thigh,
15 combining an adjucted multi-body rigid model to compute bone kinematics and muscle activation
16 levels from experiments, and a driven solid deformable muscle models with contractile ability. The
17 specific aim was to investigate the effect of the interaction between muscles, leading to radial
18 loading, on the production of muscle forces. The kinematics and muscle activation levels were
19 computed from experimental data by using a model of the lower limb which was previously validated
20 against in vivo joint contact forces (Moissenet et al., 2014). The contractile ability in the deformable
21 solid musculoskeletal system was modelled by a thermomechanical coupling on muscle fiber beams
22 embedded in the muscle passive solid hexahedral meshes. The global behavior of each of the 17
23 muscles was fitted to represent a Hill-type model. Interactions and so radial loading between
24 anatomical structures was considered through surface-to-surface contact elements.

25

26 Modelling choices

27 Several authors modelled a set of deformable muscles with a hyperelastic quasi-incompressible law
28 model for the passive part (often used for rubbers), with a contribution resulting from a deviatoric
29 stretch in the along-fiber direction based on the activation level (Blemker and Delp, 2006; Böl et al.,
30 2011b; Lee et al., 2009; Sifakis, 2007; Si et al., 2014; Teran et al., 2005). However, the Teran/Lee
31 approach did not include neither any anisotropic shear behavior relative to the fiber axis nor a
32 realistic muscle constitutive model: the activation parameter was limited to produce a bulk length

1 from the contraction along the muscle fiber directions. As pointed by (Sifakis, 2007), this simplified
2 approach had to be improved to investigate a more physiological behavior of the muscles. This was
3 proposed by Blemker et al. (2005): the authors reported a muscle material model integrating an
4 anisotropic material law with along-fiber stretch, with volume control and cross-fiber shear. This full
5 material model was reused later, for example by Sanchez et al. (2014). For the fiber field, Blemker et
6 al. (Blemker and Delp, 2006, 2005) deformed a complex template from a generic cube to the muscle
7 surface (as Webb et al. (2014) and Berranen et al. (2014) later). Defining the fibers' force
8 contribution as a contribution to the total strain energy function provided a simple solution to take
9 into account the fiber field within the muscle matrix. Conversely, our choice was to dissociate the
10 active and passive contributions of the fiber and the matrix. Few authors reported such a discrete
11 approach to model the muscle through a truss/beam network embedded in a passive hexahedron
12 mesh. To our knowledge, Behr (2004) was the first published work reporting such a reinforced
13 approach. Its numerical simplicity and so robustness allowed to model 16 muscles for the lower limb,
14 under one activation level for on position of the limb. Böl and Reese (2008) improved this approach
15 in initiating the Sartorius muscle mechanical response (i.e. the fiber truss stretch) by motoneuron
16 discharges, to vary the muscle force in modulating the rate of impulses driving the muscle fibers.
17 Note that these authors decomposed the hexahedral elements in tetrahedral ones. Roux et al. (2016)
18 reported a reinforced discrete formulation evaluated on a passive generic muscle, even if such an
19 approach should not be able to take into account shear loading within the muscle tissue. Using an
20 activation being computed from an adjusted multi-body rigid model to drive a deformable muscle
21 make sense only if the deformable muscle provide a global behavior compatible with a Hill-type
22 model (as we made by optimization). Using a 1D Hill-type element as fiber elements within a solid
23 matrix (Hedenstierna et al., 2009), instead of a thermomechanical relationship, would not have
24 prevented to make such an optimization process in order to ensure a consistent global muscle
25 behavior. In addition, the initial length being different for each beam modeling the fiber field, this 1D
26 Hill-type element approach required to define one material law per fiber, which is quite tedious
27 (Howley, 2014). Berranen et al. (2014, 2012) proposed to embed Hill-type contractile 1D elements
28 within the tetrahedral mesh of the matrix. They reused a force function for their contractile element
29 previously reported by Böl and Reese (2008): they introduced the maximum isometric force of the
30 whole muscle and the fiber density in the Hill-type elements to assure a coherence with the classical
31 Hill-type model as used in multi-body rigid models, even for bipennate or pennate organization,
32 without additional optimization. The link between the fibers and the passive tissue was obtained by
33 considering the fiber forces due to contraction as external forces applied to the embedding passive
34 muscle mesh. Another solution would have been to use constraint equations to link non-coincident
35 nodes between fibers and solid elements. However, this could lead to numerical issues in large

1 deformation. So Berranen solved the well-known numerical issues in large rotation by filtering out
2 local rotational modes, as for co-rotational FEM. Finally some authors provided more realistic
3 approaches to model the muscle, until the integration of cellular models; but these formulations,
4 only investigated on models representing one isolated muscle, are recognized as presenting too high
5 computational complexity for an integration in a complete musculoskeletal model (Kockova and
6 Cimrman, 2009; Röhrle et al., 2012; Wisdom et al., 2015; Zöllner et al., 2012).

7 We choose to model the muscle as an isotropic passive mesh reinforced by fiber beams. Hodgson et
8 al. (2012) reported, by using a 2D numerical model, that the material properties of the passive
9 muscle matrix could significantly influence the deformation field within a contracting muscle as well
10 as the force produced, despite of various pennation angles. Thus they suggested to use an
11 anisotropic passive matrix. As reported by Böhl et al. (2014) on porcine biceps femoris muscles, the
12 passive muscle tissue with the fibers embedded in the extracellular matrix presents an anisotropic
13 response perpendicularly to the fiber direction, limiting the relevance of decomposing the passive
14 muscle in an isotropic part relating to the extracellular matrix and an anisotropic part related to
15 fibers.

16

17 [Interactions between muscles](#)

18 The results of the present study demonstrate an effect of taking into account the interactions
19 between anatomical structures on the forces produced by muscle forces computed from classical
20 multi-body rigid modelling (i.e., inverse dynamics and static optimization) based on experimental
21 data. Both with and without contacts between anatomical entities, the deformable model provided
22 muscle forces consistent with the multi-body rigid models of the lower limb. In detail, for both the
23 two gait cycle steps, some muscles provided exactly the same total forces without and with contacts,
24 while some muscles produced slightly more forces when introducing contacts (the deepest vastus
25 medialis and the superficial Sartorius in HS, the deeper semimembranosus and the superficial gluteal
26 medius in LR). Basically, this study pointed out the limitations of multi-body rigid modelling and the
27 pertinence of finite element modelling to investigate the biomechanics of muscle contraction. The
28 limitations of multi-body rigid models are well-known: muscles are represented by their lines of
29 action between origins and insertions, sometimes with intermediate via-points (more rarely with
30 warping surfaces) for keeping consistent moment arms (especially for the muscles crossing the hip).
31 Such a simplified approach assumes mass-less muscle, the segment mass being embedded to a fixed
32 point of the bone. The physiological 3D fiber distribution as well as muscle tightening effects are
33 ignored. The inability to represent the transmission of radial forces is less often stated but of
34 paramount importance. Several authors reported a force transmission effect as a “transverse

1 loading” or a “lateral force transmission”, which consists in the transmission of forces by shear
2 stresses in the intramuscular connective tissue or in the extracellular matrix of skeletal muscle
3 (Purslow, 2010). This kind of force transmission corresponds to a tangential loading with respect to
4 the muscle surface. Others loadings are the axial loading (i.e. in the direction perpendicular to the
5 cross section plane of the slender muscle structure, also called “longitudinal force” by some authors)
6 and the radial loading (i.e. normal to the surface of the muscle, the one currently investigated in this
7 study). Authors pointed, mainly on animal models (Huijing et al., 1998; Maas et al., 2001; Yucesoy et
8 al., 2010), that the connecting tissue between muscles influenced the longitudinal muscle force, due
9 to myofascial force transmission from surrounding muscles (Monti et al., 1999). Within isolated
10 intact mammalian muscles, Ramaswamy et al. (2011) reported that the axial and the tangential force
11 transmission were equal in rats and mice, for different force levels. More recently Zhang and Gao
12 (2014) concluded on the impaired tangential force transmission in old rats.

13 The effect of the *in vivo* radial loading (i.e. due to muscle compression by surrounding muscles,
14 connecting tissue, and bones) on the muscle force have not been investigated to our knowledge. By
15 considering any muscle almost as isochoric (the muscle contains about 80% of water (Van Loocke et
16 al., 2008)), the surrounding muscles should radially load any deeper muscle, leading to an increase of
17 the muscle force because the internal muscle pressure is related to the muscle force under isometric
18 conditions (Davis et al., 2003; Winters et al., 2009). But Siebert et al. (2016) reported that increasing
19 the internal pressure lead to decrease the longitudinal muscle force. Thus, in our study the deepest
20 gluteus minimus yielded to a total force drastically lower by taking into account contacts between
21 muscles, at LR only. The gluteus minimus was caught between the pelvis and the other gluteal
22 muscles, displaying high contraction. Siebert et al. (2014) reported, from isometric experiments on
23 rat, that increasing the transversal (i.e. radial) muscle loading resulted in a decrease in the muscle
24 force. Van Loocke et al. (2006) pointed that only a transversely isotropic hyperelastic model was able
25 to capture the passive muscle in compression; and Van Loocke et al. (2008) reported a significant
26 stiffening effect with compression rate and fiber orientation for passive muscle in compression
27 (approximately 50% of total stress at $0.5 \text{ \%strain.s}^{-1}$).

28

29 Limitations

30 *Fiber field representation*

31 The fibers were aligned with the curved direction between the origin and insertion of each muscle, to
32 stay as close as possible to the adjucted multi-body rigid model with Hill-type muscle model. To
33 improve our model, the physiological 3D fiber field could be extracted from imaging (Kwah et al.,
34 2013; Levin et al., 2011; Schenk et al., 2013; Zhukov and Barr, 2003) with a preference for DTI over

1 ultrasound (Bolsterlee et al., 2015; Damon et al., 2011) even with 3D position tracker system, based
2 on a template fitted to the muscle geometry either from anatomical observation (Blemker and Delp,
3 2005) or from digitized *in vitro* muscle specimens (Lee et al., 2015; Sanchez et al., 2014), or
4 computed from theoretical approaches (Choi et al., 2015; Inouye et al., 2015; Wong and Kuhl, 2014).
5 If more physiological curvature and length of the fascicles could explain non uniform strain inside the
6 muscle volume especially around the aponeurosis (Azizi and Deslauriers, 2014; Blemker et al., 2005)
7 with greater intramuscular pressure in deeper regions of muscles (Rana et al., 2013), the total force
8 produce by the muscle should stay unchanged. Moreover, our aim was not to compute the strain
9 field within muscles (which is also influenced by the chosen activation function (Stelletta et al., 2013)
10 or the aponeurosis morphology (Rehorn and Blemker, 2010)), but the total force produced by the
11 muscles. In addition to the 3D architecture, Siebert et al. (2015) pointed also the necessity to use
12 specific force-length and force-velocity relationships according to the considered muscles. In our
13 quasi-static approach, thermomechanical parameters were optimized to identify a behavior coherent
14 with the normalized force-length relationship: the global behavior of each muscle was in agreement
15 with the Hill-type model classically used in multi-body rigid model, with distinct force-length
16 relationship considered for all the thigh muscles (Delp et al., 1990; Klein Horsman et al., 2007).

17

18 *Aponeuroses, Fascia and tendon representation*

19 Neither the external fascia nor the skin are taken into account in our model, so the muscle packaging
20 effect is *a priori* insufficiently represented. Turrina et al. (2013) reported that “endomysium,
21 perimysium, epimysium and deep fasciae have not just a role of containment, (...), but are
22 fundamental elements for the transmission of muscular force”. The deep fascia define compartments
23 separating functional groups of muscles, which are generally named according to the actions of their
24 contained muscles: flexors, extensors, adductors, etc. (Benjamin, 2009). Two studies (Eng et al.,
25 2014; Pancheri et al., 2014) pointed that the biaxial material properties of Fascia Lata could be
26 related to the ability of the longitudinal Fascia Lata to transmit force, while allowing larger strains in
27 the transverse direction to allow expansion in case of muscle contraction. This comment regarding
28 the biaxial material properties is the same for the aponeuroses, which exhibit an anisotropic 3D
29 structure, with a transverse stiffness four times lower than the longitudinal one (Azizi and Roberts,
30 2009). In our model, the aponeuroses of the muscles were simplified to a shell approach with an
31 isotropic nonlinear material law. A more realistic material model could contribute to homogeneous
32 fascicle length change during contraction (Muramatsu et al., 2002). In addition, our muscle geometry
33 (and so the aponeurosis morphology) was simplified during the meshing process to guarantee a high
34 element quality: Rehorn and Blemker (2010) and Fiorentino and Blemker (2014) reported that a
35 simplified aponeurosis model could drastically decrease the along-fiber stretch across all the muscle

1 tissue, the peak of the along-fiber stretch within the muscle being related to geometry of the
2 aponeuroses.

3 In our model, the tendons were represented as an elastic beam network with a global behavior
4 stiffer than muscle tissue converging towards a stiffness matrix (Calvo et al., 2010): introducing the
5 tendon material law should not pose additional complexity, the nonlinear stress–strain relationship
6 being similar to aponeuroses (Scott and Loeb, 1995).

7
8

9 *History of activation and muscle force enhancement*

10 We considered the muscle activation for two steps in the gait cycle, by simulating the Heel Strike (HS)
11 step followed by the Loading Response (LR) one. Simulations were performed with an explicit
12 scheme but represented quasi-static conditions which are consistent with the activation levels
13 computed from the adjusted multi-body rigid model. In the literature, a muscle force enhancement
14 (Herzog et al., 2008) has been noted both from in vitro experiments and from in vivo ones on large
15 human muscles under physiological conditions (Abbott and Aubert, 1952; Hahn et al., 2007; Seiberl
16 et al., 2010). Herzog (2004) pointed that the force history dependence could raise up to 50% of the
17 corresponding isometric reference force, through the active and the passive structures (Herzog and
18 Leonard, 2005). However, for example the muscle force enhancement is not related to a history-
19 based variation of fascicle lengths and pennation angles (Tilp et al., 2011). So modeling the force
20 history dependence is difficult due to various hypothesis (mainly out of mechanical parameters) to
21 explain this phenomenon (Campbell and Campbell, 2011), leading to more complex continuum-
22 mechanical model of skeletal muscle including actin–myosin interactions and actin–titin interactions
23 (Heidlauf et al., 2016). Due to the quasi-static approach, no history dependence of the forces
24 produced by the muscles in isometric state was taken into account in the model, as for the Hill-type
25 model. So the history of the boundary conditions applied to the model was the simulation of the
26 muscle contraction, followed by the introducing the bone kinematics to facilitate the numerical
27 convergence.

28

29 *Conclusion and perspectives*

30 In our approach, the skeletal muscles were modelled as a fiber reinforced continuum, allowing the
31 force transmission between fibers through the passive matrix's solid elements, particularly by shear
32 loading. Regarding the force transmission between muscles, only the radial loading was considered in
33 this study by using sliding contacts between anatomical structures. The tangential loading between
34 structures was also not possible because the connecting tissue was not represented. For a given
35 activation level, the muscle stiffness (activated or not) could increase due to the radial forces

1 transmitted from the surrounding muscles. The present results could extend the concept of
2 “continuity of tissues” between muscles to the radial loading.

3 No definitive conclusion regarding the quantification of the interaction effects on the production of
4 muscle forces from given activation levels would be taken from these results. For a given activation
5 level, the deep muscles could produce higher forces by taking into account the ability to transfer
6 loads between muscles, their whole surfaces being loaded. Moreover, the presence of contacts on
7 the superficial muscles modified their lines of action during the movement, and therefore altered
8 their forces. The main conclusion is that integrating a deformable model with interactions between
9 anatomical structures could correct the musculo-tendon forces traditionally computed from a multi-
10 body rigid model. As a matter of fact, the force transmission limited only to the tendons is a clear
11 limitation of multi-body rigid modelling that is rarely stated in the literature. An extended study
12 should be pursue on others steps of the gait cycle, even if the two chosen ones were so because of
13 their complementarity in terms of boundary conditions. The effect of the interactions between
14 muscles could depend on the bone kinematics, the activation levels of surrounding muscles, the
15 spatial curvature of the fiber network, and the initial conformation of the muscles.

16 In the present study, the fasciae, the skin and the connecting tissue were not modeled. An initial gap
17 between all anatomical structures was assessed to guaranty non initial penetration which
18 compromises any simulation. Only the surface muscles were able to radially interact with deep
19 muscles if their lines of action changed, either because of their contraction (if they were activated) or
20 their traction (resulting from the bone kinematics, here the knee flexion). Grouping muscles in
21 compartment through fasciae should be assessed in the future, to keep the coherence of anatomical
22 structures and thus render a possible muscle packaging effect. However, such an approach induces
23 to mesh the muscles in a different way, to drastically respect the initial geometry and guarantee no
24 initial penetration. Presently, the initial geometry of the muscle were modified to guarantee high
25 quality hexahedron elements. Further Investigation on meshing methods should be required to
26 preserve the surfaces of muscles, and modeling the tendons as deformable structures, which could
27 modify the global stiffness of the muscle-tendon complex and thus the produced muscle force. In
28 addition, the next meshing step to improve the modeling of the muscles should be focused on
29 dissociate the active truss/fibers from the solid passive muscle mesh, to consider a more realistic
30 fiber field embedded in the muscle matrix mesh as in concrete modeling.

31 The final improvement step should be focused on evaluating the coherence of the model. Regarding
32 the modeling of each muscle, one could find the inspiration in Sturmat et al. (2012) and then Böl et
33 al. (2013) who reported presently one of the most comprehensive approaches to quantify the muscle

1 behavior on animal models. Regarding groups of muscles in their surrounding structures, few
2 quantitative information are easily measurable from in vivo experiments. Ultrasound elastography
3 for studying in vivo muscle stiffness appears presently as the state-of-the-art non-invasive (and
4 simplest) method, with a high potential to investigate muscle properties, despite an underlined
5 theory built on strong hypothesis (Brandenburg et al., 2014; Eby et al., 2013). The strain field within
6 human muscles during in vivo contraction is also measurable using magnetic resonance imaging
7 (Huijing et al., 2011), cine PC MRI (Pappas et al., 2002; Sinha et al., 2015; Zhong et al., 2008) or by
8 combining diffusion and strain tensor MRI (Englund et al., 2011). The advantage of finite element
9 modelling, giving access to strain in the muscle tissue, appears evident for such validations.
10 Moreover, the deformable model of the thigh was driven by the bone kinematics and activation
11 levels corresponding to experimental dataset taken from the « Grand Challenge Competition to
12 predict in vivo knee loads ». This experimental dataset include the tibiofemoral contact forces
13 measured by an instrumented prosthesis, usable to validate the present deformable model as it was
14 done for the multi-body rigid model. For this purpose, the boundary conditions in terms of bone
15 kinematics shall be further released until joint equilibrium is obtained, which is often considered as a
16 great numerical challenge.

17

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23

24 [Conflict of interest statement](#)

25 There are no conflict of interest for any of the authors.

26

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46

1

2 [Figure and table captions.](#)

3

4 **Figure 2.1:** Origins and insertions of the thigh's muscles

5 **Figure 2.2:** Cross sections of the thigh's muscles, according to Figure 1

6 **Figure 2.3:** multi-body rigid model of the lower limb, with the equivalent mechanisms for the joints
7 (left) and the muscle lines of action (right). From Moissenet et al. (2014).

8 **Figure 2.4:** meshing the muscles: (1) generic block of hexahedron with four parameters to define the
9 number of elements per dimension, (2) generic block fitted on the reconstructed boundary of the
10 muscle - the edges in red were reinforced with the thermomechanical beams used for the fiber
11 modelling, (3) 3D solid mesh of the muscle, (4) quasi-rigid elements used to link the muscle to the
12 bone – insertions on the muscle in red, on the bone in green.

13 **Figure 2.5:** Deformable model of the lower limb, (1) with a focus on the thigh, (2) on the knee with
14 the modeling of the ligaments, the patella and the tendon, (3) on the vastus medialis with the fiber
15 beams in red embedded in the muscle solid matrix.

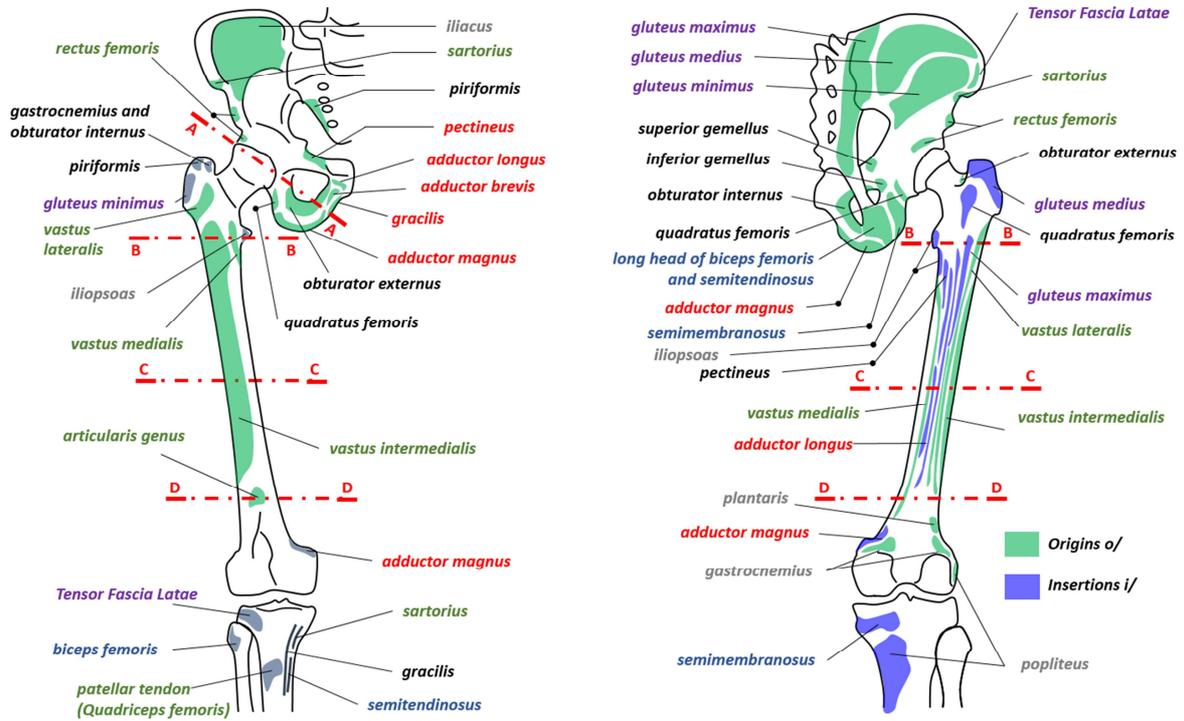
16 **Figure 2.6:** For both the Heel Strike (HS) and the Loading Response (LR) steps, total forces (passive
17 plus active parts) computed for each muscle of the musculoskeletal model, for the multi-body rigid
18 model and the Finite Element (FE) model with the two configurations (without and with contacts
19 between muscles). Note that some muscles did not produced forces and so were did not reported in
20 this table. Abbreviations: gluteus maximus, medius, minimus (G_max, G_med, G_min); adductor
21 longus, brevis, magnus (A_long, A_brev, A_mag); pectineus (Pect); gracilis (Grac); sartorius (Sart);
22 semimembranosus (S_memb); semitendinosus (S_tend); biceps femoris long head, short head
23 (BF_long, BF_short); rectus femoris (RF); vastus medialis, intermedialis, lateralis (V_med, V_int, V_lat)

24 **Figure 2.7:** Geometry of the muscle models after the simulation of the Loading Response step (1);
25 representation of the sliced muscles for the model without contact (2) and with contact (3) two cross
26 sections c1 and c2 (1). Abbreviations were provided with the previous figure.

27

1 **Figure 2.1.**

2 **cf. attached file**



3

4

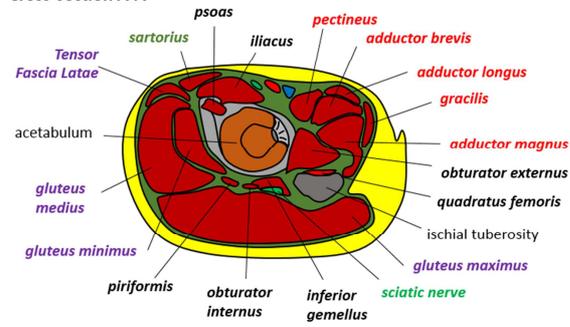
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1 **Figure 2.2.**

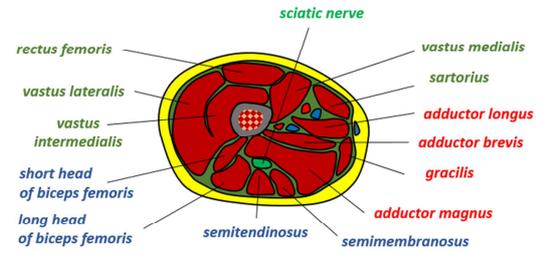
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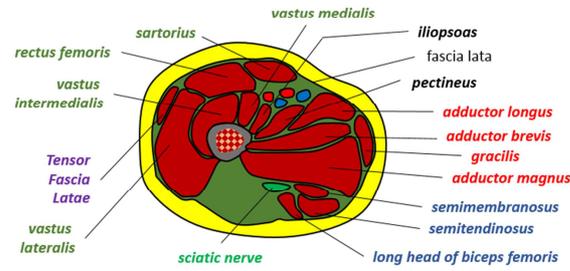
Cross-section A-A



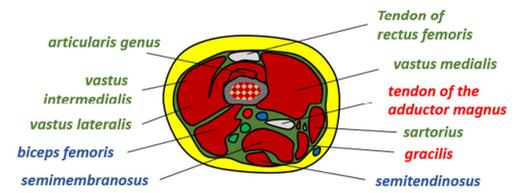
Cross-section C-C



Cross-section B-B



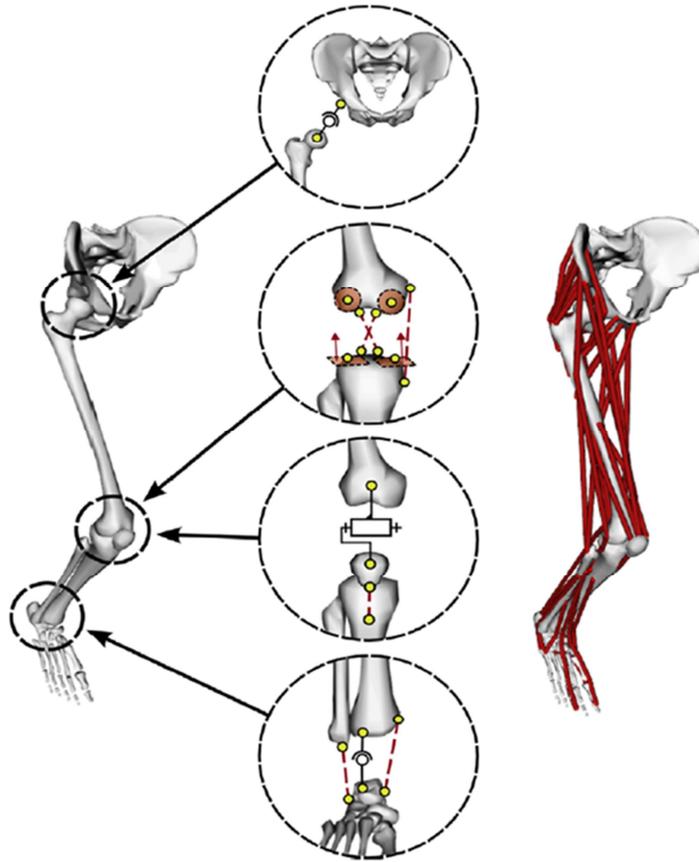
Cross-section D-D



4

5

- 1 **Figure 2.3.**
- 2 **cf. attached file**
- 3

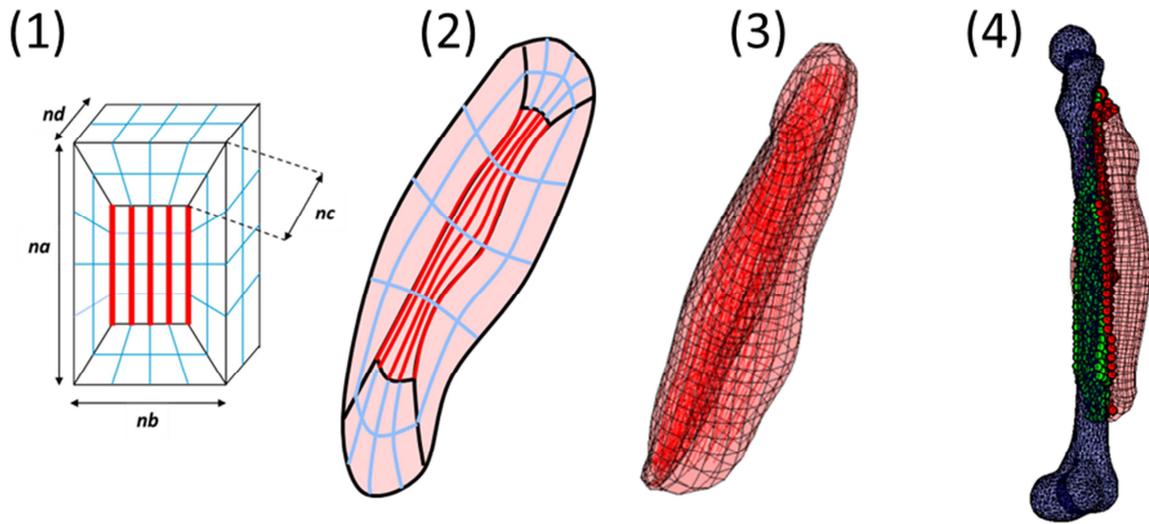


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1 **Figure 2.4.**

2 **cf. attached file**

3



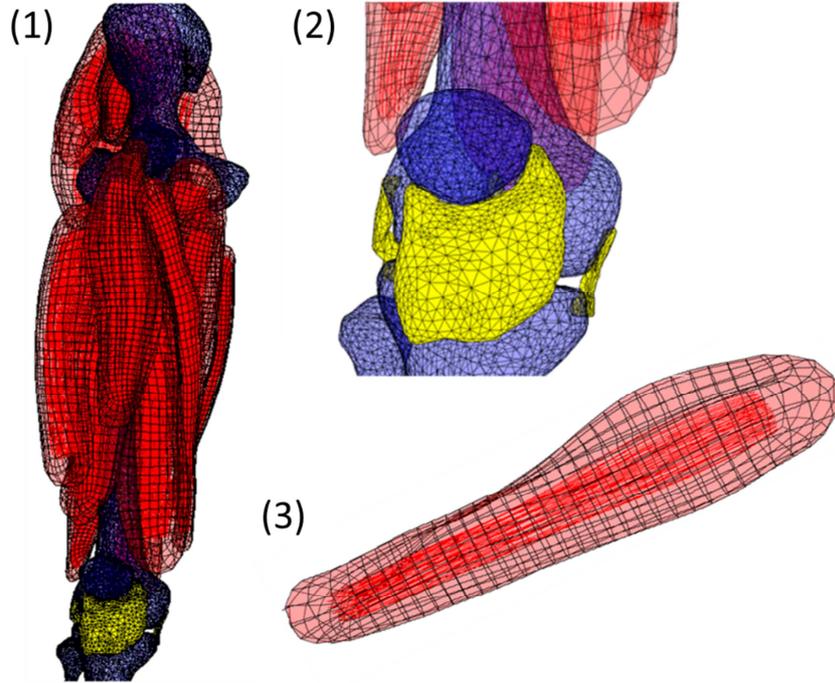
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1 **Figure 2.5.**

2 **cf. attached file**

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1 **Table 2.6.**

2 **cf. attached file**

3

Muscles	Total forces produced by muscles (N)		
	Multi-Body Rigid Model	FE Solid Model	
		Config 1 (w/o contact)	Config 2 (w/ contacts)
V_med	35	16	25
V_int	33	33	30
V_lat	34	34	36
Sart	319	307	324
RF	345	348	344
Pect	44	49	48
A_long	17	18	17
S_tend	187	187	187
BF_short	109	111	111
G_min	17	17	17

4 Heel Strike

Muscles	Total forces produced by muscles (N)		
	Multi-Body Rigid Model	FE Solid Model	
		Config 1 (w/o contact)	Config 2 (w/ contacts)
V_med	127	107	102
V_int	122	138	125
V_lat	122	122	122
Sart	37	31	29
RF	107	112	105
S_memb	62	38	63
G_max	369	368	370
G_med	673	655	677
G_min	559	532	447

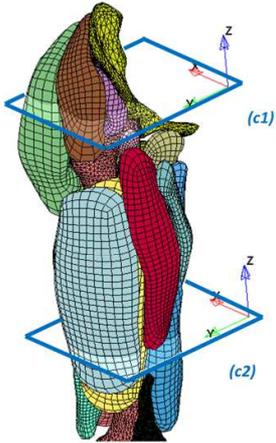
5 Loading Response

1 **Figure 2.7.**

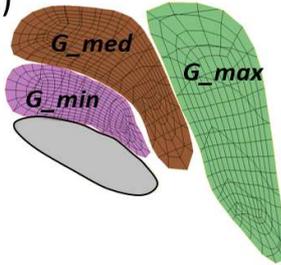
2 **cf. attached file**

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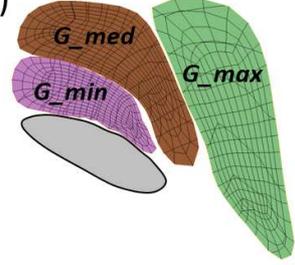
(1)



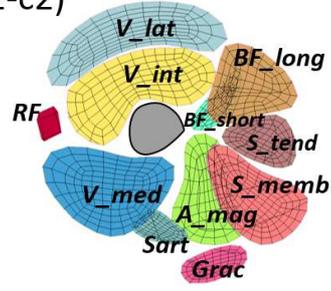
(2-c1)



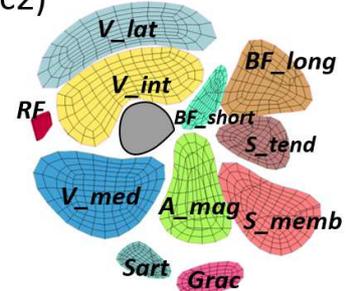
(3-c1)



(2-c2)



(3-c2)



4

5