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Original Article

Development and evaluation of a new methodology for Soft Tissue Artifact compensation in the lower limb

AUTHORS

Bhriku K. Lahkar^{1,*}, Pierre-Yves Rohan¹, Ayman Assi^{1,2}, Helene Pillet¹, Xavier Bonnet¹, Patricia Thoreux^{1,3}, Wafa Skalli¹

AFFILIATIONS

¹ *Institut de Biomécanique Humaine Georges Charpak, Arts et Métiers Sciences et Technologies, Paris, France*

² *Laboratory of Biomechanics and Medical Imaging, Faculty of Medicine, University of Saint-Joseph, Beirut, Lebanon*

³ *Université Sorbonne Paris Nord, Bobigny, France*

*Corresponding author: Bhriku K. Lahkar

Institut de Biomécanique Humaine Georges Charpak, Arts et Métiers ParisTech, 151 Boulevard de l'Hôpital, 75013 Paris, France

E-mail address: bhriku_kumar.lahkar@ensam.eu

28 **Abstract**

29 Skin Marker (SM) based motion capture is the most widespread technique used for motion
30 analysis. Yet, the accuracy is often hindered by Soft Tissue Artifact (STA). This is a major
31 issue in clinical gait analysis where kinematic results are used for decision-making. It also
32 has a considerable influence on the results of rigid body and Finite Element (FE)
33 musculoskeletal models that rely on SM-based kinematics to estimate muscle, contact and
34 ligament forces. Current techniques designed to compensate for STA, in particular multi-
35 body optimization methods, assume anatomical simplifications to define joint constraints.
36 These methods, however, cannot adapt to subjects' bone morphology, particularly for patients
37 with joint lesions, nor easily can account for subject- and location-dependent STA. In this
38 perspective, we propose to develop a conceptual FE based model of the lower limb for STA
39 compensation and evaluate it for 66 healthy subjects under level walking motor task.

40 Both hip and knee joint kinematics were analyzed, considering both rotational and
41 translational joint motion. Results showed that STA caused underestimation of the hip joint
42 kinematics (up to 2.2°) for all rotational DoF, and overestimation of knee joint kinematics (up
43 to 12°) except in flexion/extension. Joint kinematics, in particular the knee joint, appeared to
44 be sensitive to soft tissue stiffness parameters (rotational and translational mean difference up
45 to 1.5° and 3.4 mm). Analysis of the results using alternative joint representations highlighted
46 the versatility of the proposed modeling approach. This work paves the way for using
47 personalized models to compensate for STA in healthy subjects and different activities.

48 **Keywords**

49 Soft Tissue Artifact, *In vivo* joint kinematics, Model personalization, Hip and knee joint,
50 Finite Element Analysis

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53 **1. Introduction**

54 Accurate assessment of *in vivo* kinematics is essential for providing insights into normal joint
55 functionality (Akbarshahi et al., 2010) and investigation of lower limb joint pathology
56 (Andriacchi and Alexander, 2000). Skin Marker (SM) based motion capture is the most
57 widespread technique used for estimating skeletal kinematics of the lower limb. However, the
58 accuracy of such technique is affected by the relative movement of soft tissues with respect to
59 the underlying bone; a bias commonly referred to as Soft Tissue Artifact (STA). If not
60 compensated for, STA can lead average kinematic errors up to 16 mm in translation and 13°
61 in rotation for the knee joint (Benoit et al., 2006). Such errors may significantly influence the
62 assessment of pathology or the treatment effects in clinical gait analysis (Seffinger and
63 Hruby, 2007).

64 Different methods have been proposed in the literature to reduce the effect of STA on
65 bone pose estimation (e.g., single-body optimization (Chèze et al., 1995), double anatomical
66 landmark calibration (Cappello et al., 1997), point cluster technique (Andriacchi et al., 1998),
67 and Multi-body Optimisation (MBO) (Lu and O'Connor, 1999)). Amongst these, MBO,
68 which relies on a predefined kinematic model with specific joint constraints, is increasingly
69 used. Initially, simple kinematic constraints such as hinge or spherical joints were
70 considered to represent hip and knee articulation (Charlton et al., 2004; Leardini et al., 2017;
71 Lu and O'Connor, 1999; Reinbolt et al., 2005). Later, anatomical joint constraints (parallel
72 mechanism, coupling curves, ligament length variation, and elastic joint) were introduced,
73 providing encouraging 3D kinematics as they allowed joint displacements (Bergamini et al.,
74 2011; Duprey et al., 2010; Gasparutto et al., 2015; Richard et al., 2016). However, regardless
75 of the joint constraints imposed, generic (unpersonalized) model-derived kinematics were
76 shown inaccurate (knee kinematic error up to 17° and 8 mm) as these models could not adapt
77 to patient-specific geometry, particularly in pathological conditions (Clément et al., 2017).

78 On the other hand, personalization of model geometry based on medical images was shown
79 promising in improving joint kinematics accuracy (Assi et al., 2016; Clément et al., 2015;
80 Nardini et al., 2020).

81 Joint simplification has indirect consequences on the predictive accuracy of both rigid
82 body musculoskeletal (MSK) models, and Finite Element (FE) based MSK models. Studies
83 that used FE-MSK models to predict local joint mechanics using *in vivo* joint kinematics (Shu
84 et al., 2018; Xu et al., 2016) assumed the knee joint as 1 DoF. Such assumption might result
85 in propagation of uncertainties on the predicted kinematics and would affect the joint reaction
86 as well as muscle and ligament forces.

87 In light of the aforementioned contexts, reliable estimation of skeletal kinematics with
88 SM-based motion data is still a major challenge (Richard et al., 2017). Furthermore,
89 extensive time and complexity associated with customization of models to subjects' geometry
90 prohibit large sample size. In that context, methods for 3D reconstruction of bony segments
91 from medical imaging modalities, particularly biplanar X-ray imaging, are promising in
92 research and clinical routine (Chaibi et al., 2012). Also, there is a need for adaptable
93 modeling approaches that can account for subject-, task- and location-dependent STA.

94 In a previous study, a conceptual FE model was proposed for STA compensation
95 (Skalli et al., 2018). The model consists of bone segments (pelvis, femur and tibia), skin
96 markers, virtual markers, connecting elements between skin markers and corresponding
97 bones, and joint models for the hip and the knee joint. The potential advantage of the
98 proposed model is its versatility with regards to soft tissue stiffness personalization and
99 alternative joint model representation. The objective of the current study was to develop the
100 conceptual model for the lower limb and to implement it on healthy volunteers considering
101 subject-specific models.

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103

104 **2. Materials and methods**

105 First, the conceptual model is presented. Then implementation of the model is
106 illustrated within an IRB approved (CEHDF285) study. Finally, the consistency and
107 versatility of the model were investigated through sensitivity of various parameters, including
108 the joints representation.

109 *2.1 Conceptual FE model of the lower limb*

110 The rationale underlying the conceptual model is twofold: First, the spring connecting
111 the virtual marker and the skin marker is a simple way of modeling globally and grossly the
112 soft tissue deformation, while being able to adjust the spring stiffness to differentiate both
113 between anatomic regions (such as the pelvis, thigh and shank), and between populations of
114 different skin types (tight or loose). Second, considering virtual markers just beneath the skin
115 markers allows easy post-processing of the results to estimate the corrected position of skin
116 markers. These corrected marker positions are analogous to the model determined markers in
117 standard MBO approaches. Such post-processing helps to use the classical gait analysis
118 software.

119 The conceptual model of the lower limb consists of bone segments, nodes
120 representing skin markers and virtual markers, joint elements, and elements that connect the
121 skin markers to the corresponding bones. Bone segments are represented by a set of high
122 stiffness (quasi-rigid) beams. The joints between the segments are represented by rigid links,
123 allowing free rotations at the joint and controlled displacements. The connection between a
124 skin marker and the corresponding virtual marker is represented by a linear spring, where all
125 the soft tissue deformation is reported. The connection between the virtual marker and the
126 corresponding bone segment is established with high stiffness beams (Fig. 1).

127 The proposed model requires only the measured optoelectronic skin marker locations
128 as imposed boundary conditions, without any need for additional force nor any further
129 optimization process. The output of the Finite Element Analysis of the model is the
130 mechanical response resulting in bone motion and the virtual marker positions. From the
131 virtual marker positions, corrected marker positions are obtained.

132 Skin markers are denoted by S , differentiating between those of the pelvis (SP), femur
133 (SF) and tibia (ST). The number of markers for the pelvis (NMP), femur (NMF) and tibia
134 (NMT) is variable and depends on the protocol being considered. Each skin marker is
135 therefore referred to using the corresponding subscript : SP_i , SF_i and ST_i respectively for the
136 pelvis (SP_1 to SP_{NMP}), femur (SF_1 to SF_{NMF}) and tibia (ST_1 to ST_{NMT}). Using the same
137 convention, virtual markers are denoted SC (SCP_i , SCF_i and SCT_i for the pelvis, femur and
138 tibia respectively), and the bone points are denoted as B (differentiating between those of the
139 pelvis (BP), femur (BF) and tibia (BT)). These bone points are different bone landmarks,
140 automatically annotated in the 3D models (Chaibi et al., 2012), which serve as nodes for the
141 FE model. As illustrated further in Fig. 2(a), the pelvis bone was represented by 6 nodes
142 (BP_1 to BP_6), the femur by 7 nodes (BF_1 to BF_7), and the tibia by 6 nodes
143 (BT_1 to BT_6)(refer supplementary material for details). Beam elements with elastic modulus
144 (E) of 12 GPa (Choi et al., 1990) were used to connect the nodes for each bone segment. Hip
145 and knee joints are denoted by HJ and KJ respectively.

146 **Modeling of the skin marker-bone connection:**

147 Each pelvis skin marker (SP_1 to SP_4) was linked to the pelvis bone by a combination
148 of spring element that connects the skin marker to the corresponding virtual marker
149 (SCP_1 to SCP_4), and a beam element that connects the virtual marker to the bone. The springs
150 were assigned with stiffness (k) values in the range 5 kN/m to 65 kN/m (Dumas and
151 Jacquelin, 2017; Gittoes et al., 2006; McLean et al., 2004), whereas the beams were

152 considered highly stiff and assigned the same elastic modulus as that of the bones. The same
153 combination of elements was used to connect the skin markers to the femur and tibia bone
154 segments.

155

156 **Modeling of the joints:**

157 As a first option, hip and knee joints were represented each by a rigid link allowing
158 free rotation while controlling the relative displacements (through the length of the link). For
159 the hip joint, the rigid link connected the acetabulum center and femur head center. For the
160 knee joint, the rigid link was defined in the line joining the centroid of the two femoral
161 condyle centers to the centroid of the two tibial plateau centers (Fig. 2(b)) (refer
162 supplementary material for details).

163 *2.2. Model implementation*

164 *2.2.1. Data Acquisition*

165 66 healthy volunteers were included (age range: 18-60 years; weight: 71.3 ± 15 Kg;
166 height: 170 ± 10 cm) in this study. The only exclusion criterion was previous record of
167 orthopedic surgery of the lower limbs.

168 Quantitative Movement Analysis was performed on an optoelectronic analysis system
169 comprising 7 video-cameras (Vicon Motion System Ltd., Oxford Metrics, UK). The
170 optoelectronic markers were positioned following the Plug-in Gait[®] method (Davis et al.,
171 1991), and participants were asked to perform level walking at self-selected speed (Fig. 3(a)).
172 Biplanar radiographs were then acquired using the EOS system (EOS Imaging, France). 3D
173 digital models of bones were obtained using a 3D reconstruction algorithm validated by
174 previous studies (Fig. 3(b)) (Chaibi et al., 2012). The location of skin markers was also
175 computed from biplanar X-Rays.

176 *2.2.2. Subject-specific FE model development and simulation*

177 From the 3D digital models of the bones, subject-specific anatomical landmarks were
178 automatically identified, resulting in nodal coordinates of each bone, as represented in Fig.
179 3(c). The distance between the skin and the corresponding virtual marker was arbitrarily
180 chosen as 1 mm (i.e., spring length). A rigid link represented the hip and knee joints. The
181 choice of the rigid link (L_k) for the knee joint was motivated by the *in vitro* kinematic
182 response obtained in a previous study on 12 cadaveric specimens showing that overall knee
183 translations were nearly 20 mm (Rochcongar et al., 2016). For the hip (L_h), the joint length
184 was fixed to 1 mm based on an unpublished data on the hip joint translation quantified using
185 biplanar X-rays. For simplicity, stiffness parameter of the springs was kept constant across all
186 segments and assigned 50 kN/m.

187 The measured skin marker displacements at each frame from the motion capture were
188 introduced to the model as a prescribed boundary condition. A solution was computed at each
189 frame using commercial FE package ANSYS®, with a default Newton-Raphson algorithm,
190 an implicit scheme widely used in numerical procedures for Partial Differential Equations
191 (Bathe, 1996).

192 2.2.3. Kinematic computation

193 The positions of the resulting bone segments and virtual markers were used to define
194 corrected markers (CF_i) with the same consideration as those of the virtual markers, i.e., rigid
195 links with the bone segment (Fig. 4). The positions of the corrected markers were used to
196 compute STA Compensated (STAC) joint kinematics. Hip and knee joint rotational
197 kinematics were expressed in the pelvis and femur anatomical reference frames (EOS-based)
198 respectively, and with Cardan sequence $YX'Z''$. Hip joint translation was defined as the
199 relative displacement between points A_c' and F_c' expressed in the pelvis anatomical reference
200 frame. Similarly, knee joint translation was defined as a relative displacement between points
201 C_c' and T_c' expressed in the femur anatomical reference frame. Anatomical reference frames

202 were defined as described in (Schlatterer et al., 2009) for the femur and tibia, and in (Dubois,
203 2014) for the pelvis (Fig. 4). A customized Matlab (MathWorks, Massachusetts, United
204 States) routine was used for both SM-based and STAC kinematic processing. In each case,
205 before and after STA compensation, joint kinematics were obtained using an internal
206 procedure implemented in our previous studies (Azmy et al., 2010; Pillet et al., 2016;
207 Rochcongar et al., 2016). Briefly, skin marker coordinate systems were registered on the
208 bone anatomic reference frames to get the joint kinematics.

209 Joint kinematics (mean \pm 1SD) for the hip and knee joint were plotted for all DoFs over
210 time normalized gait cycle. The difference in range of motions (dROM) was also computed
211 between SM-based and STAC kinematics.

212 *2.3. Illustration of versatility*

213 *2.3.1. Sensitivity of spring stiffness and joint length*

214 Two different stiffness values for the springs were implemented (5 kN/m and 65
215 kN/m) to investigate the influence of stiffness parameters on joint kinematics.

216 Furthermore, two different knee joint lengths ($L_k = 21$ mm and 31 mm) were
217 arbitrarily considered to investigate the impact of joint lengths on estimated kinematics.
218 Implemented knee joint lengths were based on the minimum and maximum value found in
219 the population. In this case, spring stiffness was kept constant with a value of 65 kN/m.

220 Differences between the two groups with different spring stiffness values and then
221 joint lengths were analyzed with a Student's *t*-test or Wilcoxon sign-rank test depending on
222 the outcomes of the Shapiro-Wilk test of normality, using a customized Matlab routine. For
223 all the tests, the significance level was set 0.05 *a priori*.

224 *2.3.2. Alternative joint representations*

225 Two other alternative joint models were considered to illustrate the versatility of the
226 lower limb FE model.

227 These joint models were:

228 *Parallel Mechanism*: The centers of the medial and lateral condyles and corresponding tibial
229 plateaus were considered to model the knee joint with two rigid links approximating the
230 femur-tibia contact behavior. The hip joint model was left unaltered (single-link model).

231 *Spherical joint*: Spherical joint model at the hip and knee joint was considered. The joint
232 constraint location was placed on the femur head center for the hip joint and the mid-point of
233 the two femoral condyles for the knee joint. Such consideration was similar, as reported in
234 previous studies (Sauret et al., 2016).

235 Differences between alternative joint models were analyzed with a Student's *t*-test or
236 Wilcoxon sign-rank test depending on the outcomes of the Shapiro-Wilk test of normality,
237 using a customized Matlab routine. For all the tests, the significance level was set 0.05 *a*
238 *priori*. Spring stiffness value of 65 kN/m was assigned for all the joint models.

239 *2.4. Model comparison with multi-body optimization*

240 As no reference kinematics (artifact-free motion) were available, the FE model results
241 were compared to a standard MBO method with spherical joint modeling for both the hip and
242 knee joints (Lu and O'Connor, 1999). The joint constraints and locations incorporated in the
243 MBO were in accordance with the FE model. To compare the kinematic results of the
244 subject-specific FE models with MBO, the same anatomical reference frames were defined
245 for the MBO bone segments.

246 Differences between the two methods were analyzed with a Student's *t*-test or
247 Wilcoxon sign-rank test depending on the outcomes of the Shapiro-Wilk test of normality,
248 using a customized Matlab routine. For all the tests, the significance level was set 0.05 *a*
249 *priori*.

250 **3. Results**

251 Each FE model with 5 DoF joints required less than 45 sec of run time on a single
252 processor desktop PC to simulate a complete gait cycle of approximately 200–300 frames.
253 All results are synthesized in Table 1.

254 *3.1. Joint kinematics*

255 Both rotational and translational kinematics estimated with skin marker measurements
256 and FE model embedding the 5 DoF joint model are illustrated in Figs. 5 and 6 for the hip
257 and knee joints respectively. The joint kinematics are plotted over time-normalized gait cycle.

258 Overall, for the hip joint, STAC and SM-based kinematics exhibited qualitatively
259 similar pattern. However, differences in range of motion (dROM) varied across all DoFs,
260 with a maximum value of -2.2° for Abduction/Adduction (Abd/Add) followed by -1.6° and
261 -0.3° for Flexion/Extension (Flex/Ext) and Internal/External (Int/Ext) rotation respectively.
262 Maximum joint displacement up to 1 mm was observed for STAC kinematics while showing
263 up to 41.5 mm for SM-based kinematics in Posterior/Anterior (Post/Ant) direction. In the
264 Medial/Lateral (Med/Lat) and Inferior/Superior (Inf/Sup) direction, joint displacement
265 exhibited less than 1 mm, whereas SM-based kinematics showed up to 28 mm.

266 For the knee joint, maximum dROM value was observed for Int/Ext (12.5°) followed
267 by Flex/Ext (-6.3°) and Abd/Add (1.5°) rotation respectively. A maximum of 20 mm of
268 joint displacement was noted in (Post/Ant) direction, while remaining DoFs showed up to 9
269 mm (Med/Lat) and 3 mm (Inf/Sup) for STAC kinematics. These results showed up to 30.5
270 mm, 12.5 mm, and 21 mm respectively, for SM-based kinematics.

271 *3.2. Sensitivity study*

272 *3.2.1. Spring stiffness parameter and joint length*

273 With two different values of spring stiffness parameters (5 kN/m and 65 kN/m), no
274 statistical significance in dROM was noted for the hip joint kinematics. As for the knee joint,
275 different spring stiffness revealed significant dROM for all DoFs except Flex/Ext.

276 With different knee joint lengths (21 mm and 31 mm), hip translational kinematics
277 displayed significant variability for Lat/Med motions (by 2 mm), while showing less than 1
278 mm change for remaining DoFs. As for knee translational kinematics, significant dROM was
279 observed for the Post/Ant and Inf/Sup motions.

280 *3.2.2. Influence of alternative joint models on kinematics*

281 Different joint representations displayed varying kinematic changes across all DoFs
282 for the hip and knee joints. Among hip joint kinematic results, significant dROM was
283 observed only for the Lat/Med (up to 4 mm) and Post/Ant motion (up to 2.7 mm). Similarly,
284 knee joint kinematics significantly varied up to 3.4° in dROM for Int/Ext rotation when joint
285 model was altered, along with Post/Ant motion (up to 7.6 mm), for the single link Vs parallel
286 mechanism.

287 *3.3. FE Model comparison with MBO*

288 Statistically significant differences between MBO-based and FE-based STA
289 compensation for the hip and knee joints were found ($p < 0.05$). Those differences, however,
290 were always in the range of 0.7° to 2°.

291 **3. Discussion**

292 Soft Tissue Artifact compensation is essential for accurate estimation of *in vivo* joint
293 kinematics in both research and clinical routine; however, personalization and versatility of
294 current model-based methods still represent a challenge. The purpose of this study was to
295 develop and evaluate a conceptual FE model of the lower limb for STA compensation. The
296 proposed method was evaluated on a population of 66 subjects. This model is
297 computationally fast (less than 45 sec run time), and its main advantage is versatility allowing
298 a wide range of parameters and joint representations to be considered.

299 Qualitatively similar kinematic patterns were observed between SM-based and FE-
300 based STA compensated (STAC) results for both the hip and knee joints, with differences in

301 range ROM across all DoFs. SM-based kinematics were comparable to the literature
302 (D'Isidoro et al., 2020; Fiorentino et al., 2017). Results obtained showed that overall
303 rotational ROM was underestimated by SM-based results up to 2.2° for the hip joint, thus
304 confirming similar observations reported in studies that compared SM-based ROM to dual
305 fluoroscopic measurements (Fiorentino et al., 2020). For the knee joint, SM-based ROM was
306 smaller by 6.3° for the Flex/Ext, whereas other DoFs revealed up to 12° higher as compared
307 to STAC kinematics. For translational kinematics of the knee joint, SM-based results were
308 higher as compared to STAC kinematics. STAC knee kinematics were comparable to studies
309 that reported either bone-pin-based or fluoroscopy-based kinematics (Benoit et al., 2006;
310 Kozanek et al., 2009; Myers et al., 2012). Nevertheless, we observed overall higher dROM
311 values between SM-based and STAC as compared to the studies that reported in the range
312 4.4° – 5.3° for rotational kinematics, and up to 13 mm for translational kinematics (Benoit et
313 al., 2006; Leardini et al., 2005). These discrepancies may arise from the experimental
314 protocol, such as number of markers, cluster configuration and location.

315 Sensitivity study showed that joint kinematics (particularly the knee joint) were
316 sensitive to spring stiffness exhibiting dROM value up to 1.5° for the rotational kinematics
317 and up to 3.5 mm for translational kinematics. Different joint representations revealed that
318 alternative joint models have considerable influence on the estimated kinematics, particularly
319 knee Int/Ext rotation (up to 3.4°) and translations (up to 7.6 mm), establishing similar
320 remarks as reported in the literature (Duprey et al., 2010; Richard et al., 2017).

321 Joint kinematics computed using the proposed FE-based STA compensation model
322 were able to consider the joint simplifications used in a standard MBO method in the
323 literature, producing consistent results. However, the proposed approach aims to overcome
324 the limitations of the MBO method underlined by (Clément et al., 2017), who showed that
325 simplified joint constraints (kinematic or anatomical) were inadequate for clinical

326 applications. Indeed, such simplifications considered in MBO methods may be sufficient for
327 many movement analysis applications; however, this is not the case for pathological cases.
328 For example, in a previously unpublished study using EOS imaging, the translations at the
329 hip joint (distance between the femur head and acetabulum center) during a change of posture
330 from standing to sitting was quantified varying up to 5 mm in pathological population. In
331 such cases, it is apparent that standard joint representation, such as a spherical joint, is not
332 relevant. The main advantage of the new procedure is its versatility. Indeed, it can be
333 modified to incorporate more or less detailed joint representations to improve joint mechanics
334 estimation. It could also evolve into subject-specific modeling for clinical applications. For
335 example, the spring stiffness could be personalized based on quantitative soft tissue
336 deformation that can be assessed using medical images (Südhoff et al., 2007). Such avenues
337 are currently under investigation.

338 This study has some limitations. First, there was no reference kinematics to compare
339 the results to. Therefore, the joint kinematics exhibited by the FE models were compared to
340 those computed with the MBO method. Nevertheless, as we cannot consider MBO as a fully
341 reliable solution for STA compensation (Richard et al., 2017), such comparison is only for
342 assessing the qualitative performance of the FE model. Second, STA parameters
343 implemented in the model were arbitrary as there is a lack of data in the literature.
344 Personalization of such parameters is, however, essential to encompass different range of
345 subjects (young, adult, patients with CP and OA etc.). Third, joint representation in this
346 model is still simplified, which could be insufficient for investigating local joint mechanics
347 for healthy or pathological joints (Adouni et al., 2012; Lenhart et al., 2015; Shu et al., 2018;
348 Valente et al., 2014). Nevertheless, as a preliminary step, the current contribution only
349 focused on exploring and facilitating personalization of the parameters that are important for
350 STA compensation. Moreover, even with simplified joint representation, the model could

351 limit the effect of STA in joint kinematics. Fourth, although the proposed approach may give
352 the impression that it complicates the process of STA compensation in gait analysis, the
353 perspectives are numerous, as already highlighted. Fifth, no external forces nor inertial/mass
354 forces were imposed on the model. The only boundary conditions were the external skin
355 markers displacements. Considering the inertial/mass forces would be necessary when
356 dynamic phenomena are essential to take into account, for example, in sports biomechanics.
357 Finally, the study was based on a single motor task, i.e., level walking. Therefore, the results
358 may vary with other motor tasks (hopping, cutting, stand-to-sit) and hence the interpretations.

359 The proposed approach may serve in two major fields of applications: First, in gait
360 analysis for research, where classical scaling techniques are used to obtain subject-specific
361 geometry instead of image-based model personalization. Compensating for STA with such
362 method is possible with an approximated model geometry, while being able to differentiate
363 soft tissue stiffness parameters between different sub-groups. To be noted that as such scaling
364 techniques often consider gross anthropometry of the subject and disregards distinctive
365 features of the joint, therefore can be considered “not actually personalized” (Nardini et al.,
366 2020; Smale et al., 2019). Second field of application can be clinical gait analysis, where
367 image-based model personalization could capture anatomical details of the joint structures.

368 In conclusion, as a first study, we presented a conceptual FE model of the lower limb
369 for STA compensation and evaluated it in a population of 66 subjects with varying
370 morphologies. The model appeared to be satisfactory in compensating for STA and versatile,
371 facilitating parameters necessary for model personalization. The methodology developed and
372 evaluated in this study may improve the accuracy of kinematic predictions, which is
373 instrumental for MSK models as well as making clinical decisions. In the current
374 contribution, the human model used for the computations consists only of the lower limbs
375 (pelvis, femur and tibia). However, the same approach can be considered for the whole body,

376 which could be particularly interesting for the shoulder joint (Duprey et al., 2017). Future
377 work could focus on further model evaluation based on *in vivo* data, such as dual
378 fluoroscopy.

379 **Conflict of Interest**

380 None

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1 **Figure Captions**

2

3 **Figure 1.** Schematic illustration of the conceptual lower limb FE model. Detailed illustration shown
4 only for the femur segment. BP_i , BF_i and BT_i denote pelvis, femur and tibia bone nodes. SF_i and SCF_i
5 are the skin and virtual markers respectively of the for the femur segment. HJ and KJ denote the hip
6 and knee joint respectively.

7 **Figure 2.** (a) Detailed representation of the lower limb FE model with generic anatomical bony
8 landmarks. Anatomical landmarks for the pelvis (BP_1 to BP_6): right antero-superior iliac spine, right
9 postero-superior iliac spine, left antero-superior iliac spine, left postero-superior iliac spine, and right
10 and left acetabulum centers. For the femur (BF_1 to BF_7): femur head center, greater trochanter, two
11 diaphyseal points, medial and lateral condyle centers and center of the two condyles. For the tibia
12 (BT_1 to BT_6): center of the two plateaus, two diaphyseal points, medial and lateral malleoli and center
13 of the two malleoli. (b) joint modeling of the hip (L_H) and knee joint (L_K) with rigid links allowing
14 free rotation and controlled relative displacement.

15 **Figure 3.** Schematic illustration of FE model personalization (a) the locations of the skin markers
16 throughout gait cycle obtained from motion capture and (b) 3D digital models of the pelvis, femur
17 and tibia built from two orthogonal radiographs (c) anatomical landmarks were identified from the 3D
18 digital models resulting the nodal coordinates of each bone.

19 **Figure 4.** Illustration of corrected markers (CF_i, CP_i, CT_i) and anatomical reference frames
20 ($R_{fem}, R_{pel}, R_{tib}$) for the (A) femur, (B) pelvis and (C) tibia respectively. Corrected markers are
21 obtained from the virtual marker in a direction orthogonal to the bone segment and 1mm away from
22 virtual marker. Anatomical reference frames for the femur and tibia were defined as described in
23 (Schlatterer et al., 2009) and for pelvis (Dubois, 2014).

24 **Figure 5.** Hip joint kinematics during gait presented as Mean \pm 1SD. Mean values for skin marker-
25 based (green) and FE model predicted results (blue) are shown as solid lines, while standard deviation
26 in lighter shades. Differences in ROM (dROM) between SM-based and STAC results are depicted as
27 insets for all DoFs

28 **Figure 6.** Knee joint kinematics during gait presented as Mean \pm 1SD. Mean values for skin marker-
29 based (green) and FE model predicted results (blue) are shown as solid lines, while standard deviation
30 in lighter shades. Differences in ROM (dROM) between SM-based and STAC results are depicted as
31 insets for all DoFs.

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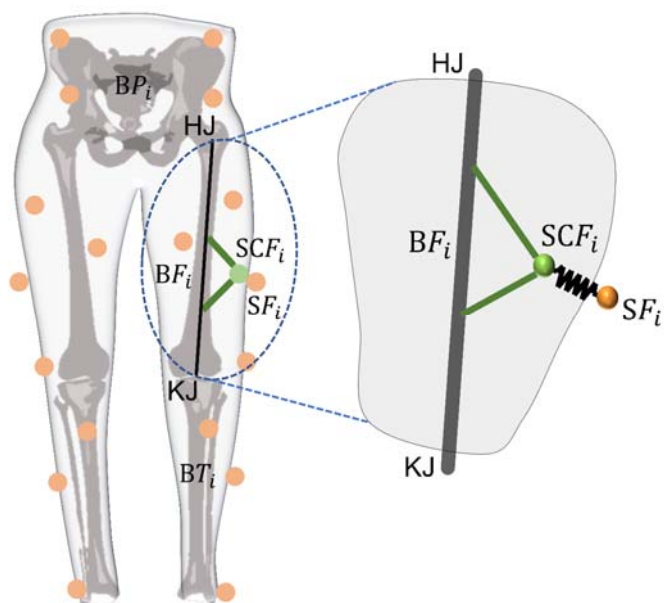
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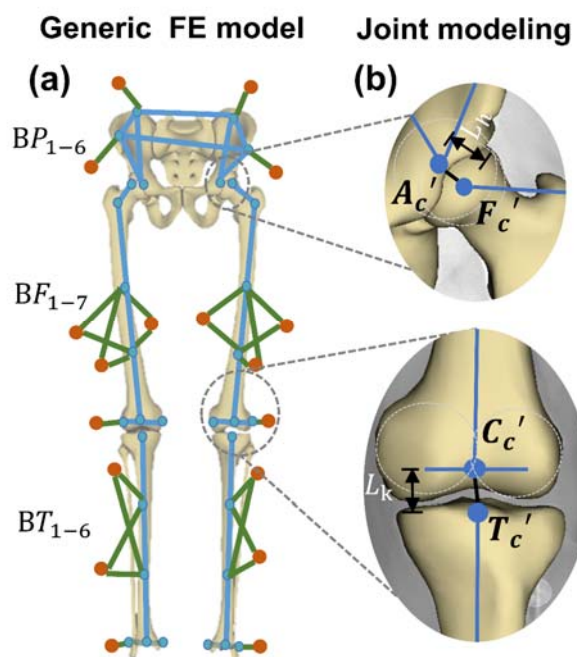
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41 **Figure 1.**



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43 **Figure 2.**

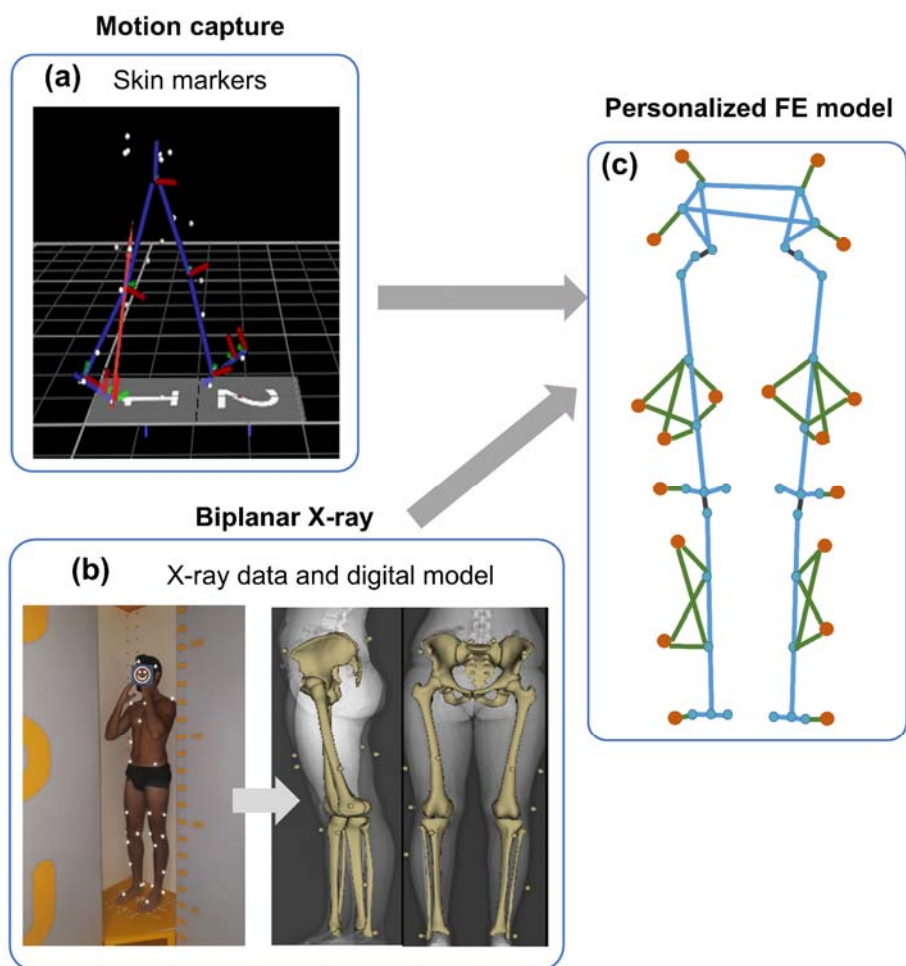


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46 **Figure 3.**

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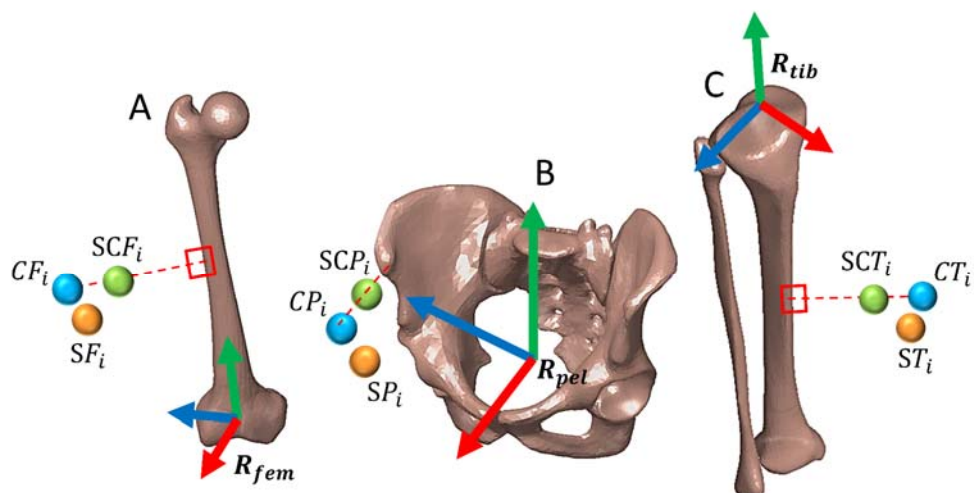
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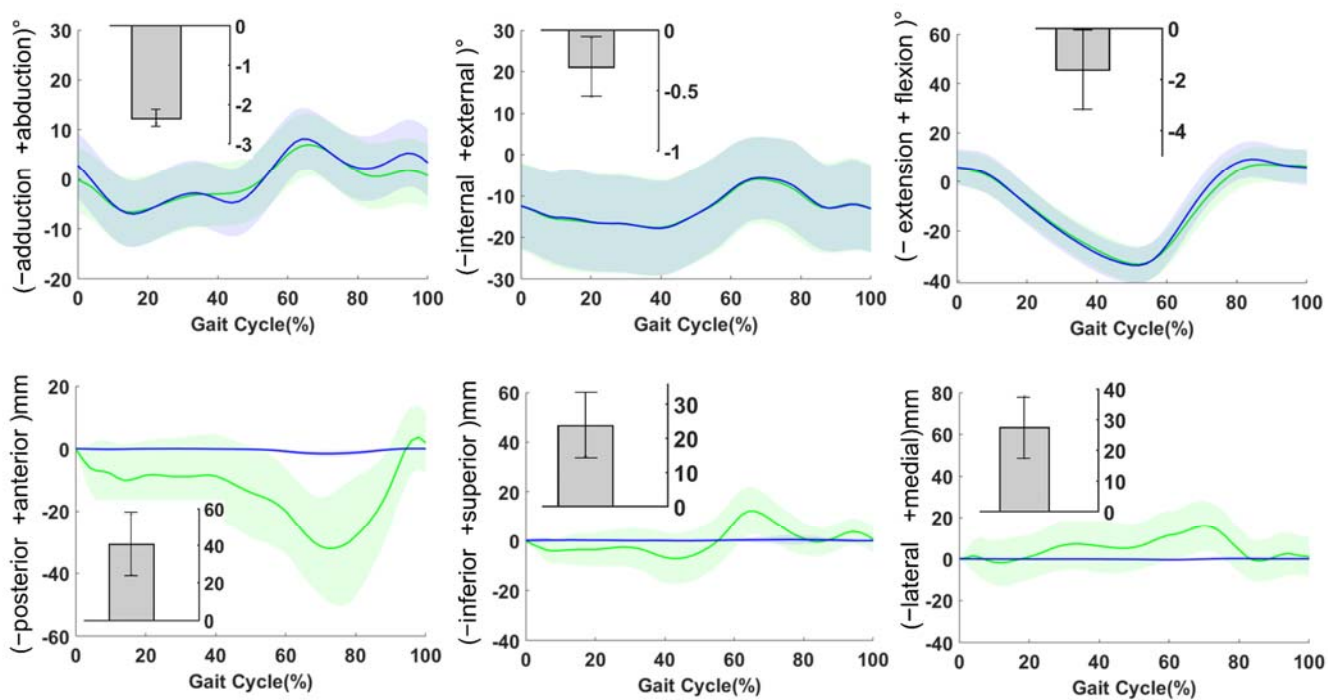
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57 **Figure 4.**



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59 **Figure 5.**

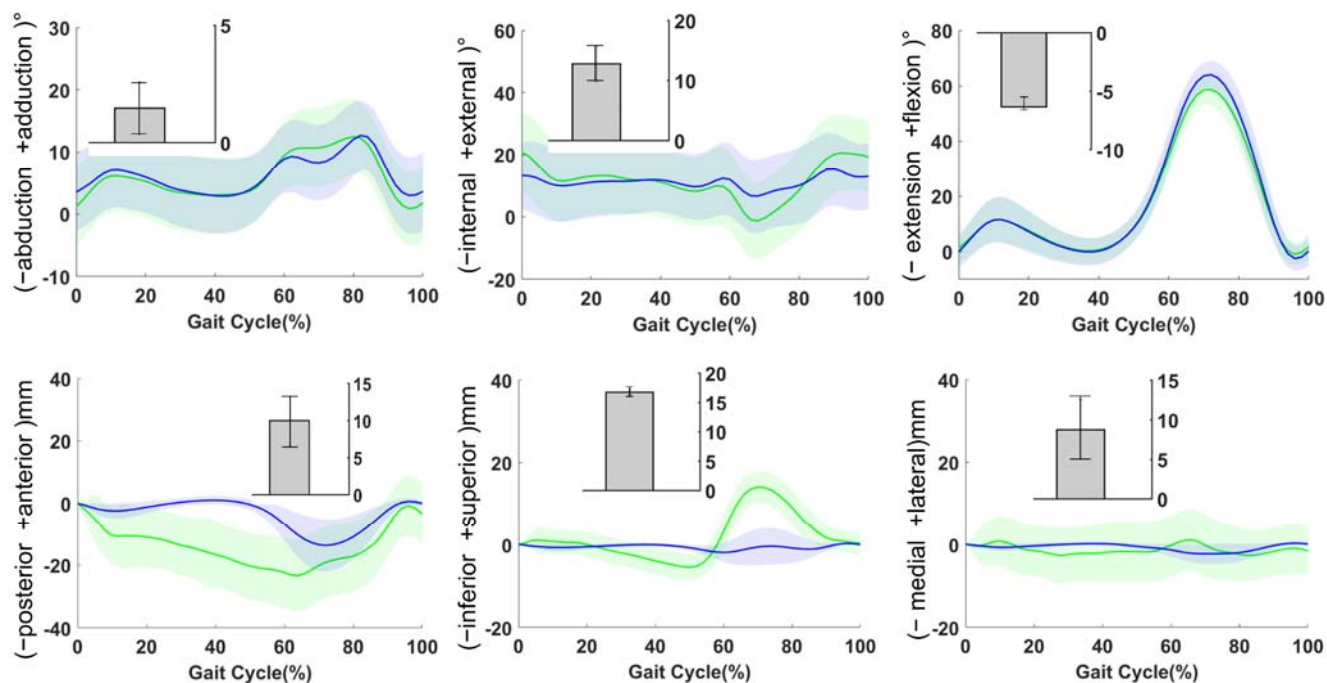


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63 **Figure 6.**



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