Subject-specific body segment parameter estimation using 3D photogrammetry with multiple cameras

Inertial properties of body segments, such as mass, centre of mass or moments of inertia, are important parameters when studying movements of the human body. These quantities are, however, not directly measurable. Current approaches include using regression models which have limited accuracy; geometric models with lengthy measuring procedures; or acquiring and post-processing MRI scans of participants. We propose a geometric methodology based on 3D photogrammetry using multiple cameras to provide subject-specific body segment parameters while minimizing the interaction time with the participants. A low-cost body scanner was built using multiple cameras and 3D point cloud data generated using structure from motion photogrammetric reconstruction algorithms. The point cloud was manually separated into body segments and convex hulling applied to each segment to produce the required geometric outlines. The accuracy of the method can be adjusted by choosing the number of subdivisions of the body segments. The body segment parameters of six participants (four male and two female) are presented using the proposed method. The multi-camera photogrammetric approach is expected to be particularly suited for studies including populations for which regression models are not available in literature and where other geometric techniques or MRI scanning are not applicable due to time or ethical constraints.

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4 1. Introduction

5 Inertial body segment parameters (BSP), such as mass, centre of mass (CoM) or moment of 6 inertia, are used in motion analysis in research as well as in clinical settings. Accurate values are 7 essential for techniques such as inverse dynamic analysis to allow the calculation of joint torques based on measured segmental accelerations (Winter, 1979). It is, however, not straightforward to 8 9 measure these quantities from subjects directly. One approach is to use mathematical models of 10 the body segments and rely on anthropometric measurements to determine the dimensions of the modelled segments. This type of methods requires a multitude of anthropometric measurements 11 of the participants and is limited by the accuracy of the mathematical model of the body 12 13 segments. The first mathematical model suggested by Hanavan in 1964 represented 15 body 14 segments as cylinders and spheres and required 25 anthropometric measurements (Hanavan, 15 1964). More detailed models presented by Hatze or Yeadon required a total of 95 or 242 measurements respectively rendering these methods inefficient for studies with a large number of 16 17 participants because of the time and discomfort for the participant to acquire all the measurements needed (Hatze, 1980; Yeadon, 1990). Other types of approaches rely on X-ray or 18 19 MRI based tomography to extract subject-specific BSP from participants. Unlike other methods, 20 CT or MRI scans provide information about internal structures such as tissue composition which 21 should improve the reconstruction accuracy (Martin et al., 1989; Mungiole & Martin, 1990; 22 Pearsall, Reid & Livingston, 1996; Bauer et al., 2007). These approaches are, however, also 23 difficult to implement in large-scale studies due to cost and ethical constraints. Alternatively, it is 24 possible to approximate inertial BSP by adjusting previously reported average values or using 25 regression models that require only a very few subject-specific measurements (commonly subject height and weight). Such average values and regression models were derived from 26 27 cadavers or participants in a number of famous studies, such as the ones by Clauser, Dempster or 28 Zatsiorsky (via de Leva) (Dempster, 1955; Clauser, McConville & Young, 1969; McConville, Clauser & Churchill, 1980; Leva, 1996). The reliability of such regression models is, however, 29 30 rather low and the models are only applicable to a population similar to the one used to derive 31 the regression equations.

Recently, other methods have been explored to obtain volumetric data of body segments that, in combination with body density assumptions, can provide subject-specific inertial BSP. Sheets et al. used a laser to scan the body surface of participants and morphing a generic model, which

35 contained joint location information, to the scanned surface (Sheets, Corazza & Andriacchi, 36 2010). Bonnechere et al used a Kinect sensor to estimate body segment lengths but not their 37 volumetric data required to estimate inertial properties (Bonnechère et al., 2014). Clarkson 38 evaluated the Kinect sensor as a surface scanner using a mannequin, but found the scanning 39 resolution to be quite low (Clarkson et al., 2012). Another approach to gain surface data is to use 40 photogrammetry. In 1978, Jensen proposed the use of stereophotogrammetry to estimate BSP 41 parameters (Jensen, 1978). In his model, the human body was divided into elliptical disks with a 42 thickness of 20 mm and the radii of the elliptical disks were estimated using images from the front and side. The drawback of this approach lies in the simplifying assumptions of representing 43 body segments as the elliptical disks. It is, however, possible to reconstruct the surface of a 3D 44 45 object from multiple uncalibrated 2D images taken from different positions without requiring any assumptions to the geometry of the body. This principle is referred to as "structure from 46 47 motion" and was initially used for producing 3D models of static objects and landscapes. Perhaps the most striking example to date is the "Building Rome in a Day" project which used 48 images from the Flikr web site (http://www.flickr.com) to generate a 3D model of the whole city 49 50 (Agarwal et al., 2009). The reconstruction of a 3D surface from multiple cameras is two-stage 51 process. In stage one, the position, orientation and the parameters of the camera optics are estimated. This is achieved by the bundle adjustment algorithm (Triggs et al., 2000) that 52 53 minimizes the error between the re-projected feature points using estimated camera pose and parameters with the actual feature points in the images. In theory, feature points could be chosen 54 55 manually but this would be cumbersome and not very accurate. Instead, Scale Invariant Feature 56 Transform (SIFT) algorithms are employed which automate this process by identifying possible common points between multiple images (Lowe, 1999). Stage two uses the calibrated views to 57 produce a dense point cloud model of the 3D object. There are a number of possible approaches 58 59 to achieve this (for review see (Seitz et al., 2006)) but probably the most widespread current 60 approach is patch-based multi-view stereo reconstruction (Furukawa & Ponce, 2010). This 61 photogrammetric approach has gained wide acceptance for producing 3D models in areas such 62 as archaeology (McCarthy, 2014) and palaeontology (Falkingham, 2012), and is even used for markerless motion capture (Sellers & Hirasaki, 2014). The aim of this paper is to investigate 63 64 whether an approach based on structure form motion photogrammetric reconstruction can provide person-specific body segment parameters and to identify the strength and weaknesses of 65 66 such an approach.

67 2. Methods

68 Photogrammetry relies on obtaining multiple photographs taken from different locations. These 69 photographs can be taken with any suitable device and for objects that do not move, the most 70 cost effective option is to take 50+ photographs with a single camera that is moved around the 71 object. This has the additional advantage that a single intrinsic calibration can be used since the 72 camera optics can be considered identical for multiple images. However for subjects that can 73 move, all the photographs must be taken simultaneously so that the subject is in exactly the same 74 position for all the images. Simultaneous photographs can be achieved in several different ways including multiple still cameras with synchronised remote controls, multiple USB web cameras, 75 76 or multiple networked cameras. There is probably little to choose between these methods but 77 initial experimentation found that network/IP cameras provided a cost effective solution that 78 scaled well. The camera resolution should be as high as reasonably possible since higher 79 resolution images provide more information for the feature extraction algorithms and higher point density in the eventual reconstruction. This means that low resolution cameras such as low 80 81 cost web cameras and standard resolution video cameras may not be suitable.

82 2.1. 3D body scanner design

Photogrammetric reconstruction can work well with as few as 4 cameras (Sellers & Hirasaki, 83 2014) but more cameras are necessary to provide a relatively gap free reconstruction. We used a 84 85 fixed dummy and a single camera moved around the subject every 5° and compared reconstructions using 72, 36, 24, 18, 12 and 9 images (see Fig. 1A). Acceptable reconstruction 86 87 were found with 18 or more cameras although using larger numbers of cameras certainly 88 improved the reconstruction quality. The network camera was implemented using Raspberry Pi 89 (RPi) modules, type A, each equipped with an 8GB SD card and a Pi camera 90 (http://www.raspberrypi.org). These modules run the Linux operating system and provide a 91 flexible and cost-effective 5 megapixel network camera platform. 18 cameras were attached to a 92 4.8 m diameter frame on top of which the RPis were mounted pointing towards the central area 93 on the floor. Angling the camera view downwards allowed the pattern on the floor to be seen by 94 each camera which greatly aided the camera calibration algorithm which relies on shared 95 features seen in multiple fields of view. Each RPi module was provided with a USB WiFi 96 receiver (Dynamode WL-700-RX) and power was provided using the standard RPi power 97 adapter plugged into a multi-socket attached to each support pole. Four 500 W Halogen

- 98 floodlights were mounted to provide additional lighting to increase the image quality. A
- 99 schematic of the RPi scanner is shown in Fig. 1B.

100 RPi cameras can record either still images or movie files. For this application we needed to 101 trigger all the cameras to record a single image at the same instant. This was achieved using the 102 open source "Compound Pi" application (http://compoundpi.readthedocs.org), which uses the 103 UDP broadcast protocol to control multiple cameras synchronously from a single server. Once 104 the individual images have been recorded, the application provides an interface to download all 105 the images obtained to the server in a straightforward manner. Since UDP broadcast is a one-to-106 many protocol, all the clients will receive the same network packet at the same time and the 107 timing consistency for the images will be of the order of milliseconds which is adequate for a 108 human subject who is trying to stand still. Higher precision synchronisation can be achieved 109 using a separate synchronisation trigger but this was unnecessary in this application.

110 2.2. Data acquisition

111 Full body scans using the RPi setup were obtained from six voluntary participants. Additionally,

their body weight and height was measured (Table 1). The male visible human was used as an

113 additional data set for validation (National Library of Medicine's Visual Human Project (Spitzer

et al., 1996)). The experimental protocol (reference number 13310) was approved by the

115 University of Manchester ethics panel. In accordance with the experimental protocol, written

116 consent was obtained from all participants.

117 The reconstruction algorithms rely on finding matching points across multiple images so they do

not work well on images that contain no textural variation. We therefore experimented with

using different types of clothing in the scanner, such as sports clothing, leisure clothing, and a

- 120 black motion capture suit equipped with Velcro strips to aid feature detection. Clothing was
- 121 either body-tight or tightened using Velcro strips if they were loose since loose clothing would
- 122 lead to an overestimation of the body volume. The participants stood in the centre of the RPi
- setup with their hands lifted above their head (see Fig. 2) and the 18 images were then acquired.

124 2.3. Data processing

- 125 The 3D point cloud reconstruction was initially done using open source application VisualSFM
- 126 (http://ccwu.me/vsfm/) which performed adequately, but we then switched to using Agisoft
- 127 PhotoSc http://www.agisoft.com) which proved to be much easier to install and use. The
- 128 program runs identically on Windows, Mac or Linux. The full 3D reconstruction with 18 images
- 129 took an average of 30 minutes using an 8 core 3GHz Xeon Macl with 12GB RAM. The actual
- 130 time taken was variable depending on the image file size and the reconstruction parameters. The
- 131 output of the Agisoft PhotoScan is an unscaled 3D point cloud of the participants and
- 132 surrounding scenery (see Fig. 2), which requires further post-processing to calculate BSP values.
- 133 First, the point cloud was scaled and oriented using CloudDigitizer (Sellers & Hirasaki, 2014),
- the oriented point clouds were then divided into anatomical segments using Geomagic
- 135 (http://geomagic.com), and the convex hulls computed in Matlab®
- 136 (http://www.mathworks.com). The reference points for the body segmentation are listed in the
- 137 supporting information Table S1. The body segments were all oriented into the standard
- 138 anatomical pose before the volume, centre of mass and inertial tensor were calculated based on
- the hull shape and segment density using a custom function implemented in Matlab® (see
- 140 supporting information). The choice of body density is an interesting issue. Different tissues
- 141 within segments have different densities and tissue composition is moderately variable between
- 142 individuals. Indeed variations in density are commonly used to estimate body fat percentage
- 143 (Siri, 1961; Brožek et al., 1963). MRI and CT based techniques can allow individual tissue
- 144 identification and can compensate for this but surface volumetric techniques need to use an
- 145 appropriate mean value. Segment specific densities are available (e.g. (Winter, 1979)) but the
- 146 quoted trunk density is after subtraction of the lung volume. For a surface scan model, we need
- 147 to use a lower value trunk density that incorporates the volume taken up by the air within the
- 148 lungs. Therefore for the purpose of this paper a trunk density value of 940 kg/m³ was chosen,
- 149 while a uniform density of 1000 kg/m³ was assumed for all other body segments (Weinbach,
- 150 1938; Pearsall, Reid & Ross, 1994). The body mass calculated from the volume was never
- exactly the same as the recorded body mass so the density values were adjusted pro-rata to
- 152 produce a consistent value for total mass.

153
$$s = \frac{m_{Participant}}{\sum m_{SegmHull,i}}$$
 (1)

154 The factor *s* effectively scales the body densities and is thus also applied the moments and 155 products of inertia obtained from the convex hull segments.

156 3. Results

157 Six participants were scanned using the RPi photogrammetry setup and their point cloud 158 segmented. In order to be able to calculate the inertial properties, the point cloud needs to be 159 converted into a closed surface mesh. To calculate the volume of an arbitrary shape defined by a 160 surface mesh, the mesh needs to be well defined, i.e., it should be two-manifold, contain no holes 161 in the mesh, and have coherent face orientations. The processing of converting a point cloud to a 162 well defined mesh is known as hulling and there are several possible methods available. The simplest is the minimum convex hull where the minimum volume convex shape is derived 163 mathematically from the point cloud. This approach has the advantage of being extremely quick 164 165 and easy to perfor not it is very tolerant of point clouds that may contain holes where the 166 reconstruction algorithm has partially failed. However it will always overestimate the volume 167 unless the shape is convex. There are also a number of concave hulling approaches. Some are 168 mathematically defined such as AlphaShapes (Edelsbrunner & Mücke, 1994) and Ball Pivoting 169 (Bernardini et al., 1999) and require additional parameters defining the maximum level of 170 permitted convexity. Others are proprietary and can require considerable manual intervention such as the built in hole-filling algorithms in Geomagic. This latter group provides the highest 171 172 quality reconstructions but at the expense of considerable operator time. For this paper we 173 concentrated on convex hulls under the assumption that the level of concavity in individual body 174 segments was likely to be relatively small. The relative segment mass of all participants are 175 reported in Fig. 3 (the segmented convex hulls are shown in Fig. S1 in the supporting 176 information). Figure 3 also displays average values from literature. As the participants were 177 imaged wearing shoes, the foot volume is overestimated significantly, which is why its relative 178 mass is systematically higher than the values reported in literature. It is possible to adjust the 179 value using a foot-specific scaling factor that accounts for this overestimation although of course 180 if the subsequent use of the BSP parameters is in experiments with participants wearing shoes then the shoe mass becomes an important part of the segment. The moments of inertia are shown 181 in in Fig. 4 together with average values from literature. Geometrop methods also allow us to 182 183 calculate the products of inertia which are otherwise simply assumed to be zero. The average products of inertia are depicted in Fig. 5 (absolute values shown only, signed values reported in 184 185 the supporting information Table S2-S4). Some segments, e.g. the thigh or trunk, have products 186 of inertia that are of a similar order of magnitude as their moments of inertia, which is indicative

187 of a noticeable difference between the inertial principal axes and the anatomical principal axes. 188 The majority of the products of inertia are however significantly smaller than the moments of 189 inertia (of the same segment) by one to two orders of magnitude. Figure 6 contains the relative 190 centre of mass in the longitudinal segment direction, i.e. along the z-axis with the exception of 191 the foot whose longitudinal axis corresponds to the x-axis (see Fig. 2). Figure 7 shows the shift 192 of CoM from the longitudinal axis in the transverse plane (x-y plane). The CoM values in 193 literature assume a zero shift from the principal anatomical (longitudinal) axis. The shift values 194 we found with our geometric method are generally unequal to zero, but they have be to viewed 195 with caution as the placement of the reference anatomical axis itself has uncertainties associated 196 with it. The numerical values presented in Fig. 3-7 and the segment lengths are reported in the 197 supporting information (Tables S2-S13)

To estimate the effect of the convex hull approximation on the mass estimation versus the 198 199 original body segment shape, the volumes of a high resolution 3D body scan and of their convex 200 hull approximation were calculated and compared. A detailed surface mesh was obtained from 201 the National Library of Medicine's Visual Human Project (Spitzer et al., 1996) by isosurfacing 202 the optical slices using the VTK toolkit (http://www.vtk.org) and cleaning up the resultant mesh 203 using Geomagic. The surface mesh of the 3D body scan was separated into body segments and 204 the volume calculated following the same methodology as used for the point cloud data. A 205 convex hull was applied to each body segment and the volume calculated again (see Fig. 8). The 206 volume overestimations for each body segment (averaged between left and right) are shown Fig. 207 9 (column CH). Several body segments showed a large relative volume overestimation (using 208 10% error as a cutoff, although the choice would depend on the required accuracy): foot (26%), shank (31%), hand (47%) and forearm (16%). This is due to the relatively strong curvatures in 209 210 these segments. To minimize the effect, these body segments were subdivided (see Fig. 10) and 211 the convex hulls recalculated. The results of the divided segments are also shown in Fig. 9 212 (column CHD), and the decrease in volume overestimation is apparent. The volume 213 overestimation of the subdivided foot (11%), shank (11%) and forearm (5%) are at a similar 214 level to the other body segments and would probably be acceptable in many cases. The hands 215 show the largest relative mass overestimation still (25%), which is due to its curved position and slightly open fingers. The convex hull error of the hand is, however, expected to be significantly 216 217 smaller if the hand is imaged while being held in a straight position with no gaps between the 218 digits.

219 Figure 11 contains the relative mass estimations of the original surface mesh, the convex hulls 220 with and without subdivision, and the average and regression model values found in literature. 221 With a BMI value of almost 28, the male visible human is not well represented by the average or 222 regression model values found in literature, where the majority of the studies involve relatively 223 athletic people (BMI average of around 24) or obese individuals (BMI over 30). The convex 224 hulls of the subdivided segments (CHD in Fig. 11) give the closest approximation to the original 225 mesh and, with the exception of the hands, are within a relative error of less than 5%. The 226 relative error of the convex hull of the whole segments (CH in Fig. 11) is larger, but still within 227 the range of values found in literature. The moments of inertia are overestimated as well as they 228 are a product of the mass of the segment. Their overestimation follows the same trend as the 229 mass overestimation, i.e. the largest overestimation occurs for the hands, followed by the shanks 230 and feet (see Fig. S2 in supporting information), and the subdivided segments produce more 231 accurate values with an average relative error below 10%.

232 4. Discussion

233 We can see from the results that the proposed methodology is produces values that are very 234 similar to those derived using regression equations. There are no consistent problems although it 235 is clearly important that the hand is held in a suitable flat position but with fingers adducted so 236 that the hulling can provide an accurate volume estimation. We would expect that the 237 photogrammetric process will work as well as any of the published geometrical approaches 238 (Hanavan, 1964; Hatze, 1980; Yeadon, 1990) since it is simply an automated process for 239 achieving the same outcome. The procedure is currently moderately time consuming in total but 240 the interaction time with the participant is extremely short and involves no contact which can be 241 very beneficial for certain experimental protocols or with specific vulnerable participants. Since 242 most of the time is spent post-processing the data, we expect that this post-processing could be 243 streamlined considerably by writing dedicated software rather than the current requirement of 244 passing the data through multiple software packages. The values generated in our sample are relatively close to those generated by using regressing quations but BSP values are highly 245 246 variable between individuals and current regression equations are only suitable for a very limited

range of body shapes. This is particularly the case when we are dealing with non-standard

248 groupings such as children, the elderly or people with particularly high or low BMI values. In

249 general regression equations work well for applicable populations and are probably more

suitable if body mass distribution is not a major focal point of the research, particularly given

that in some cases it can be shown that experimental outcomes are not especially sensitive to the

252 BSP parameters chosen (Yokoi et al., 1998).

However there are a some specific issues with this technique that could to be improved for a more streamlined and potentially more accurate workflow.

255 Convex hulling of the point cloud is a robust and fast way to produce surface meshes. The fact 256 that it systematically overestimates the volume of concave features can be improved by

subdividing body segments into smaller parts and the decision then becomes what level of

subdivision is appropriate for an acceptable level of accuracy. For example, with only one

259 subdivision of the shank and forearm the relative error of their volume overestimation was

reduced by a factor of three, and the end result was within 10% of the true value which is

261 probably sufficient in most cases, especially given the level of uncertainty in other parameters

such as segment specific density. The adoption of one of the concave hulling techniques is likely

to lead to a similar level of improvement again with a minimum (but not zero) level of additional

work. The level of subdivision required not only depends on the body segment, but also the

265 population studied so it may well be appropriate that the segmentation level is adjusted

266 according to the type of study and its sensitivity to inaccuracies in the BSP (i.e. multiple segment)

267 (subdivisions increase accuracy of volume estimation). In this work, a uniform scaling factor and

constant body density (apart from the trunk) was assumed. It is well known that the density

269 varies among body segments as well as among populations due to different percentages of fat

270 and muscle tissue (Drillis, Contini & Bluestein, 1964; Durnin & Womersley, 1973; Zatsiorsky,

271 2002). Thus, using segment and population specific densities (and scaling factors) may improve

272 (the accuracy of the presented methodology if such values are available or derived. Similarly)

273 (important contributions to segmental mass distribution such as the presence of the lungs within)

274 the torso can be modelled explicitly which may lead to small but important shifts in the centre of

275 mass (Bates et al., 2009).

276 In terms of technology, the current arrangement of using 18 Raspberry Pi cameras is reasonably

277 straightforward and relatively inexpensive. It requires no calibration before use, and the process

278 of moving the subject into the target area is extremely quick. However it does take up a great 279 deal of room in the laboratory and the current software is reliant on clothing contrast for the 280 reconstructions which limits the flexibility of the technique. One area where this could be 281 improved is by projecting a structured light pattern onto the subject so that areas with minimal 282 contrast can be reconstructed accurately (Casey, Hassebrook & Lau, 2008). Our results show that 283 18 cameras is currently the minimum needed for full body reconstruction and a system with 36 284 or more cameras would produce better results. One future use of this technology is clearly the 285 use of such systems and algorithms for complete motion capture (Sellers & Hirasaki, 2014). The 286 limitation currently is that these cameras would need to be closely synchronised and whilst the 287 proposed system is adequate for producing a single still image, it is currently not able to 288 adequately synchronise video. In addition the video resolution is much lower and this makes the 289 reconstruction more difficult. However we predict that markerless, multiple video camera 290 structure from motion systems will become a much more common mainstream tool for 291 experimental motion capture in the near future. Ideally we could imagine that such a system 292 would both do the motion capture and also the body segment parameter reconstruction since 293 much of the computational technology would be shared.

294 Conclusion

295 A methodology based on structure form motion photogrammetric reconstruction has been 296 presented that provides subject-specific body segment parameters. The method relies on the 297 surface depth information extracted from multiple photographs of a participant, taken 298 simultaneously from multiple different view points. The brief interaction time with the participants (taking all required photos simultaneously, and measuring the height and weight 299 300 only) makes this a promising method in studies with vulnerable subjects or where cost or ethical 301 constraints do not allow the use of other imaging methods such as CT or MRI scans. The post-302 processing time is lengthy compared to using regression models or average values from literature

- 303 but not compared to processing MRI or CT data.
- 304 While the results presented in this work were derived using commercial software, such as
- 305 AgiSoft, Geomagic and Matlab[®], we were able to to achieve similar results using open-source
- 306 software only (such as VisualFMS (http://ccwu.me/vsfm/) for calculating 3D point clouds and
- 307 MeshLab (http://meshlab.sourceforge.net/) for point cloud segmentation, hulling and BSP

- 308 calculation). This makes our proposed methodology, in combination with the low hardware
- 309 costs, particularly promising for small-budget projects.

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Figure 1

Body Scanner Design.

A: Point cloud reconstruction with varying number of cameras. B: Schematic representation of the RPi scanner design.



Figure 2

Image processing work flow.

Images from the RPI scanner are converted to 3D point clouds which are then scaled and segmented manually. Subsequently, convex hulling is used to produce a surface mesh around each body segment.



Segment mass (as % of body mass).

P: Average value of all six participants (error bars show standard deviation). Z(m): Male average values reported by Zatsiorsky. Z(f): Female average values reported by Zatsiorsky (Leva, 1996; Zatsiorsky, 2002). D(m): Male average values by Dempster (via Zatsiorsky) (Dempster, 1955; Zatsiorsky, 2002).



Moment of inertia in [10⁴ kg*m²].

P: Average value of all six participants (error bars show standard deviation). Z(m): Male average values reported by Zatsiorsky. Z(f): Female average values reported by Zatsiorsky (Leva, 1996; Zatsiorsky, 2002). The definition of the coordinate system is shown in Fig. 2.



Absolute values of products of inertia in $[10^4 \text{ kg}^*\text{m}^2]$.

The absolute values of Ixy, Ixz and Iyz are shown together with a positive error bar (negative error bar is symmetrical) equal to one standard deviation. The signed values are reported in the supporting information in Tables S2-S4. The Ixy value of the hand is smaller than 10³ kg*m² and is not displayed.



Figure 6

Centre of mass along the longitudinal axis.

P: Average value of all six participants (error bars show standard deviation). Z(m: male, f: female): Average values by Zatsiorsky, adjusted by de Leva . The CoM is given as % of the segment length. The definition of the segments and reference points are given in the supporting information Table S1 - Exceptions: * Foot of participants: Heel and toe end point of participant's shoes instead of foot. ** Forearm and Upper Arm of Z: Elbow reference point is the elbow joint centre instead of the Olecranon (Leva, 1996; Zatsiorsky, 2002).



CoM shift from the anatomical longitudinal axis in the transverse (x-y) plane.

Average values of all six participants are shown (error bars show standard deviation). Due to mirror-symmetry, the y-values of the segments on the left- and right-hand side have opposite signs. To calculate the average, the sign of the segments on the left-hand side was inverted. The CoM is given as % of the segment length. The data of the foot is not included due to the participants wearing shoes.



Figure 8

Visible Human surface mesh.

A: High-resolution surface mesh. B: Convex hull mesh.



Segment volume overestimation of the hulled mesh versus the high-resolution surface mesh of the Visible Human.

Data shown as the relative difference of the hull with respect to the original mesh. CH: Convex hull of body segment. CHD: Convex hull of divided body segments (only segments indicated with an * were subdivided, see Fig. 10).



Subdivision of the body segments with large curvature.

The first row (S) shows the high-resolution surface mesh, the second row (CH) the convex hull of the whole body segment, and the bottom row (CHD) the convex hulls of the subdivided body segments.



Male Visible Human segment mass (as % of body mass) of the high-resolution mesh, convex hull, regression model and average values.

S: High-resolution surface mesh. CH: Convex Hull of whole body segments. CHD: Convex Hull with subdivided body segments (only segments indicated with an * were subdivided as shown in Fig. 10). ZR: Values predicted using Zatsiosrky's linear regression model (using weight and height). Z: Male average values reported by Zatsiorsky. D: Male average values reported by Dempster (Dempster, 1955; Leva, 1996; Zatsiorsky, 2002).



Table 1(on next page)

Participant mass and weight.

P1 – P6: Participants (m: male, f: female). VH: Male Visible Human.

	P1 (m)	P2 (m)	P3 (m)	P4 (m)	P5 (f)	P6 (f)	VH (m)
Mass [kg]	73.4	77.0	88.2	87.8	65.4	55.2	90.3
Height [m]	1.81	1.83	1.85	1.83	1.65	1.58	1.80