LEVERAGING ANTHROPOMETRIC MEASUREMENTS TO IMPROVE HUMAN MESH ESTIMATION AND ENSURE CONSISTENT BODY SHAPES SUPPLEMENTARY MATERIAL

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A ANTHROPOMETRIC MEASUREMENTS

015 The selection of the anthropometric measurements is mainly adopted from AnthroNet (Picetti et al., 016 2023). In total, 36 measurements are selected, which can be divided into 23 lengths and 13 cir-017 cumferences. All measurements are taken based on the standard SMPL-X T-pose. The reference 018 landmarks are chosen by matching the vertices on the default mesh with the landmarks defined by 019 the anthropometric survey of the U.S. army personnel (Gordon et al., 2014). A visualization of the landmarks can be found in Figure 1 and 2a. The lengths are calculated by computing the Euclidean 020 distance between two landmarks or the difference along the coordinate axis pointing upwards for 021 certain heights. The lenghts are visualized in Figure 2b and 3. Table 1 lists the enclosing landmarks 022 for each length. To measure the circumferences, we adopt the code from (Bojanic, 2023). For each 023 measurement, a plane is created, the intersection between the mesh and the plane are extracted and the convex hull of the result is calculated. During this process, the mesh is restricted to the body 025 part to be measured. A visualization of the circumferences can be found in Figure 4 and a list of the 026 landmarks and the normal vectors spanning the plane in Table 2. 027





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56	Idx	Length Name	From	То
57	1	Shoulder width	Left shoulder tip (left acromion)	Right shoulder tip
58	2	Back torso height	Cervicale	Back belly button
59	3	Front torso height	Suprasternale (top of the breast bone)	Belly button
50	4	Head	Head top	Cervicale
1	5	Midline neck	Chin	Suprasternale
2	6	Lateral neck	Center between the ears	Cervicale
-	7	Height	Head top	Center between heels
	8/9	Hand right/left	Center between middle and ring finger	Stylion rotated downwards
	10/11	Arm right/left	Acromion	Wrist
	12/13	Forearm right/left	Elbow	Stylion rotated downwards
	14/13	Calf right/left	Vice point at the femur (frochanterion)	Anklo
	10/17	Foot width right/left	Small too	Big toe
	20/21	Heel to ball right/left	Heel	Ball
	22/23	Heel to toe right/left	Heel	Big toe
			litter	Dig toe
2				head
л			e a	
4				
5		i e	lateral neck	
6		Cervicale		midline neck
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5		1		
				torse from front
		button	torso from back	
		Dutton .		V
		14		height
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		(a) Visualization of a	subset of (b) Visualization of used langth	- he with a standard
		(a) visualization of a the used landmarks	T-nose SMPL -X mesh	ns with a stanuaru
		uie useu fanumarks.	1-pose Sivii L-A mesii.	
		Figure 2. Sidevi	ew visualizations of landmarks (a) and	lengths (b)
		1 15010 2. SIdevi	and and an	ionguis (v).
	Table '	2. Definitions of circum	ferences by landmarks and the normal	vector spanning the plane
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Idx	Circumference	Normal Vector	Position
1	Waist	Up	Belly button
2	Chest	Up	Nipple
3	Hip	Up	Pubic bone
4	Head	Up	Head temple
5	Neck	Spine to head	Adam's apple
6/7	Upper Arm	Shoulder to elbow	Center of the bicep
8/9	Forearm	Elbow to wrist	Widest point of the forearm
10/11	Thigh	Up	Center of the thigh
12/13	Calf	Up	Widest point of the calf



Table 3: GT data analysis for MPI-INF-3DHP (Mehta et al., 2017). Bone length analysis based on the 3D joint locations (left) and on SMPL-X annotations by NeuralAnnot (right). Standard deviation σ , relative standard deviation $\frac{\sigma}{avg}$ and relative range $\frac{\max - \min}{avg}$ of anthropometric measurements are reported. Standard deviations are given in cm, except for the β parameters. The values are averaged between left and right body parts and between all persons in each dataset. The β parameter standard deviation is averaged over all β parameters.

3E) joint a	nnotations	5	SMPL-X annotations				
Measure	σ	r. σ	r. range	Measure	σ	r. σ	r. range	
head	0.19	1.03%	2.08%	head	0.21	0.75%	4.87%	
hip width	0.22	0.89%	1.80%	hip circ.	1.16	1.16%	9.13 %	
forearm	0.21	0.87%	1.77%	forearm	0.45	1.80%	9.75%	
upper arm	0.29	0.90%	1.82%	arm	0.83	1.59%	8.19%	
lower leg	0.60	1.49%	3.06%	lower leg	1.05	2.56%	11.54%	
thigh	3.83	7.91 %	41.90%	thigh	0.77	2.02%	9.47%	
-				height	2.76	1.56%	8.24%	
				β param.	0.18			

Table 4: GT data analysis for Human3.6m (Mehta et al., 2017): Analysis of SMPL-X annotations by NeuralAnnot. Standard deviation σ , relative standard deviation $\frac{\sigma}{avg}$ and relative range $\frac{\max - \min}{avg}$ of anthropometric measurements are reported. Standard deviations are given in cm, except for the β parameters. The values are averaged between left and right body parts and between all persons in each dataset. The β parameter standard deviation is averaged over all β parameters.

SMPL-X annotations							
Measure	$\mid \sigma$	r. σ	r. range				
head	0.41	1.51%	10.28%				
hip circ.	1.24	1.19%	8.90%				
forearm	0.83	3.30%	27.93%				
arm	0.77	2.58%	22.88%				
lower leg	0.43	1.18%	12.20%				
thigh	0.66	1.27%	9.43%				
height	3.40	2.06%	15.66%				
β param.	0.20						

C EVALUATING A2B MODELS

We measure two types of errors to evaluate the performance of our A2B models. The first type (β error) shows the error if we take the GT β parameters, derive anthropometric measurements (B2A), input them into the A2B models and evaluate the MSE of the predicted β parameters. The second type (A error) calculates B2A from the predicted β parameters and evaluates the mean difference between the GT and predicted anthropometric measurements (all 36) in mm. These evaluations are a kind of cycle consistency evaluation for A2B and B2A. Figure 5 provides a visualization of the evaluation scheme. The part that is also included in the training is highlighted.





216 **KEYPOINT SELECTION FOR FIT3D** D

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We use the fit3D (Fieraru et al., 2021) dataset for our evaluations, since this is the only sports dataset 219 with public SMPL-X annotations. We evaluate on the SMPL-X joints, since these are trivial to 220 obtain from SMPL-X meshes and there is no regressor available for the fit3D annotated 3D joints. SMPL-X has 144 defined joints. Since our focus is mainly on the body and not on the hands and face, we remove most of these joints. In the end, we select a subset of 37 SMPL-X joints: pelvis, 222 left hip, right hip, spine1, left knee, right knee, spine2, left ankle, right ankle, spine3, left foot, right 223 foot, neck, left collar, right collar, head, left shoulder, right shoulder, left elbow, right elbow, left 224 wrist, right wrist, left index, left thumb, right index, right thumb, left big toe, left small toe, left heel, 225 right big toe, right small toe, right heel, right eye, left eye, right ear, left ear, nose. 226

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GENERATION OF PSEUDO GT ANTHROPOMETRIC MEASUREMENTS E

230 As we do not have access to the athletes of ASPset and fit3d to obtain real anthropometric measurements, we need an alternative to kind-of simulate this process. For ASPset, as a first step, we 231 run IK on the GT 3D joint locations. We obtain the necessary anthropometric parameters from the 232 generated meshes with B2A. Then, we use the median values of these measurements as the GT an-233 thropometric values. We call these parameters pseudo GT throughout this paper, since this is not 234 directly the GT, but obtained from IK executed on the GT 3D joint locations and the B2A computa-235 tion from the created meshes. These parameters are used in this paper to generate the pseudo GT β 236 parameters by A2B prediction. 237

We do not have access to the athletes of the fit3D dataset either. Therefore, we need some kind of GT 238 data to mimic measurements. Obviously, there is no GT available for the official test set evaluation 239 on the evaluation server. We therefore split the official training dataset into a training, validation, 240 and test set for our evaluations. Details can be found in the main paper. With this selection, we 241 have GT shape parameters available. We do not use these directly, but apply B2A and use the 242 median measurements over time in order to mimic the measuring process and obtain a single set of 243 anthropometric measurements per person (which is not the case for the provided GT parameters, see 244 Section 3 in the main paper). In real applications, this step is omitted because the anthropometric 245 parameters can be measured directly from the athletes before starting the recording.

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F FINETUNING HME MODELS WITH PSEUDO GT MESHES

249 Finetuning existing HME models on pure 3D joints datasets is not possible, since they need mesh 250 annotations for training. However, with IK, we can generate pseudo GT meshes. We exemplary 251 test a finetuning of SMPLer-X on ASPset with this approach. Experiments show that using their 252 finetuning script with 1.6M iterations leads to worse results than the results without finetuning. 253 Therefore, we reduce the number of iterations with early stopping and achieve better results with finetuning only for 32K iterations. 254

255 The results shown in Table 5 prove that finetuning on IK generated meshes can lead to a significant 256 improvement of the scores. Replacing the β parameters of the finetuned results with the A2B β 257 parameters boosts the performance even more. These are the best results achieved with any existing 258 HME model throughout this study. 259

Moreover, we experiment with using the SMPLer-X body shape parameters combined with the poses 260 estimated by IK applied to the UU results (see last two rows of Table 5). Using the β parameters 261 from SMPLer-X leads to a slightly better result than the original 3D joint based result (without IK). 262 This evaluation shows that 3D HPE models are better in precisely locating the joints of humans than 263 HME models, but HME models are better in estimating the shape of humans. We also try to use 264 the β parameters of the finetuned variant together with the UU IK poses like before. However, this 265 resulted in a performance drop compared to the body shape parameters from the original SMPLer-X 266 without finetuning. These experiments show that finetuning HME models on pseudo ground truth 267 leads to a better performance regarding the keypoints, but the estimated β parameters have worse quality. This can further be proven by replacing the β parameters from the finetuned SMPLer-X 268 variant with the β parameters from the not finetuned model, which results in a performance gain 269 of over 5 mm compared to the original results from the finetuned version (rows 2 and 4 in Tab.

Table 5: MPJPE results in mm for the test split of ASPset. Results are given for different methods and replaced *beta* parameters with A2B results (columns NN/SVR) or the median of the original β parameters from the model noted in the *measurements* column. SMPLer-X FT stands for the best finetuned variant of SMPLer-X (finetuned with the meshes obtained from IK executed on the GT 3D joints). The *orig* column contains the results without replaced β parameters. We highlight the best result for each model and the best option for the combination of UU IK pose and SMPLer-X β parameters, since this combination outperforms the original UU IK result, too.

-	model	orig.	measurements	NN m	SVR m	NN n	SVR n	median
	SMPLer-X	86.02	SMPLer-X	85.89	85.69	86.03	85.99	86.04
	SMPLer-X FT	79.09	SMPLer-X FT	78.92	78.88	79.44	79.37	79.44
	SMPLer-X FT	-	GT	65.63	65.84	64.71	64.76	-
	SMPLer-X FT	-	SMPLer-X	73.41	73.29	73.65	73.63	73.66
	UU IK	67.54	UU	66.92	66.60	67.25	67.12	67.16
	UU IK	-	SMPLer-X	63.80	63.64	63.81	63.78	63.82
	UU IK	-	SMPLer-X FT	69.46	69.27	69.70	69.63	69.69
	UU IK	-	GT	56.44	56.56	55.14	55.19	-

5). However, our method using the UU IK poses and the A2B body shape parameters with GT anthropometric measurments achieves the overall best results.

We provide a comprehensive summary and visualization of all results on the ASPset dataset in Section G. This includes results of existing HME models, results of our approach, and the finetuning results.

G SUMMARY OF THE RESULTS

We execute a multitude of experiments with different combinations of pose and shape parameters. Figure 7 summarizes the results with their pose and shape origins for ASPset. In general, the poses estimated by IK based on the UU results (red branch in Fig. 7) are more precise than the poses estimated by SMPLer-X (light blue branch in Fig. 7). Further, the body shape parameters from our A2B models with GT anthropometric measurements (green boxes in Fig. 7) achieve the best results for all poses. We provide more qualitative examples comparing SMPler-X with this approach in

Figure 6: Qualitative results of SMPLer-X and our approach for example frames from ASPSet. GT joints and estimated joints are color-coded. Corresponding joints are connected.



Figure 7: Overview of the main results for the ASPset dataset. All results are MPJPE results in mm. Results below *mesh* boxes show the result with the original β parameters. All results after arrows to the right are results with replaced β parameters. The type of the β parameters is noted on the arrow and is color-coded: pseudo GT (green), SMPLer-X (light blue), SMPler-X finetuned (dark blue), UU IK (red).

Figure 6. Without access to the GT, all models benefit slightly from A2B model results with the median anthropometric measurements from B2A of the estimated meshes by the respective model (boxes with same color for all three branches in Fig. 7). Moreover, SMPLer-X A2B body shape parameters perform best when analyzing body shapes without GT access (light blue boxes in Fig. 7). Finetuning SMPLer-X with IK created meshes (dark blue branch in Fig. 7) improves the performance of SMPLer-X, although the quality of the body shape deteriorates. This can be seen as by comparing the shapes from SMPLer-X and finetuned SMPLer-X (dark blue and light blue boxes in Fig. 7) with finetuned and IK poses.

Since fit3D is a larger dataset, finetuning UU works better, which further leads to better IK meshes with an MPJPE of 37.02 mm. Enforcing consistent meshes with GT or IK A2B shape parameters decreases the performance slightly in this case. However, A2B shape parameters achieve slightly better scores than median values. This also holds for OSX and Multi-HMR. Overall, the approach with UU, IK, and A2B body shape parameters achieves an over 33 mm lower MPJPE than any HME model.

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