# Self-Supervised Learning of Wrist-Worn Daily Living Accelerometer Data Improves the Automated Detection of Gait in Older Adults

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### **Supplementary Information**

#### Ablation Studies

The supplementary tables present evaluations of various model configurations aimed at optimizing ElderNet, reporting performance using the F1 score and standard deviation across three seeds. Given the limited number of seeds, p-values were not reported as they would not provide meaningful statistical insights.

Supplementary	Table S1.	The effect	of using	the UK	Biobank	pre-trained	model.
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	Trained from	Pre-trained		
Model	scratch	model		
ResNet-V2	77.15 (0.30)	82.59 (0.89)		

Performance is reported as the F1 score (standard deviation between different seeds). The pre-trained UK Biobank model was a ResNet-V2 with the MTL approach. We employed the same architecture and MTL approach to train the SSL model from scratch using the MAP data.

**Supplementary Table S2**. The effect of customizing the SSL model for older adults using the MAP data.

Model's head	MAP	Without MAP
FC (without non-linearity)	84.67 (0.44)	82.55 (0.08)
FC (with non-linearity)	84.74 (0.51)	83.21 (0.45)
U-Net	83.02 (0.86)	83.90 (0.08)

Performance is reported as the F1 score (standard deviation between different seeds). In this table, we employed the combined model (i.e., pretrained UK Biobank model + additional model's head) with the optimal SSL configuration, termed ElderNet. FC: Fully-connected.

## Supplementary Table S3. The effect of dense labeling.

		Dense
Model	Window Labels	Labeling
ElderNet	84.74 (0.51)	81.99 (0.41)

Performance reported as the F1 score (standard deviation between different seeds). The model employed for both labeling approaches is identical. The key distinction lies in the final

layer of the model: in the first approach, it projects the output for each window, while in the dense labeling approach, it projects the output for each sample.

Sequence	Accuracy	Specificity	Recall	Precision	F1 score
length					
<30	95.99	97.54	80.29	76.28	78.23
seconds					
>30	99.20	100.00	81.25	100.00	89.66
seconds					

Supplementary Table S4. Performance stratified by sequence length.

# Supplementary Table S5. ElderNet setting.

Method	Optimizer	Learning-	Batch	Epochs	Learning-rate Scheduler
		rate	size		
MTL	Adam	1e-4	6000	40	Linear scaling with a 5-epoch warm-up
SimCLR	Adam	1e-4	6000	40	Cosine Decay with a 5-epoch warm-up

Each batch contains 1500 windows from 4 unique participants. The windows were sampled in proportion to their STD, as described in Yuan et al.<sup>38</sup>. For the MTL, we employed linear scaling for the learning rate to align with the pre-trained UK Biobank model, which utilized MTL and linear scaling. For SimCLR, we utilized the cosine decay scheduler, as outlined in the original paper of this method<sup>32</sup>.



**Supplementary Figure S1**. Precision-Recall curves for different configurations of normalizations. Normalization was implemented per subject, making each axis of its acceleration signal zero-mean and a standard deviation of 1. AUC: Area under the curve.



**Supplementary Figure S2**. a) On the left, it illustrates the estimated daily walking durations using a model that utilized dense labeling in the fine-tuning phase and provided a per-sample output. b) On the right, the figure represents the estimated walking duration using a model that outputs per-window prediction