# UNIVERSITY of WASHINGTON

Most 2D-3D human pose estimation models are learning-based and are susceptible to domain gaps, leading to low generality across new datasets. However, optimization methods suffer from low performance but can function case by case. Our method takes advantage of both by proposing an optimization algorithm that applies a 3D prior diffusion model to predict accurate 3D poses based on 2D keypoints. Additionally, our method shows great zero-shot and crossevaluation capacities across 3DPW, H36m and 3DHP.

### Algorithm 1 ZeDO pipeline

**Require:** Initial 3D pose  $P_{init}$ , Target 2D pose  $p_{2d}$ , 2D pose confidence scores  $C_{2d}$ , Camera intrinsic K, Diffusion timestamp t, Pre-trained diffusion model  $\theta_q(P, t)$  $R_0, T_0 \leftarrow \arg\min_{R_0, T_0} \|K(R_0 P_{init} + T_0) - p_{2d}\|_2$ // Initial Pose Optimization  $P_0 \leftarrow R_0 P_{init}$ // Iterative Optimization and Denoising  $r \leftarrow K^{-1}p_{2d}$  $\hat{r} \leftarrow \frac{r}{\|r\|_2}$ for  $i \leftarrow 0$  to n - 1 do if i < warmup then  $T_i \leftarrow T_0$ else  $T_i \leftarrow \arg\min_{T_i} \|C_{2d}(K(P_i + T_i) - p_{2d})\|_2$ end if *II Project 3D keypoints to rays*  $\widetilde{P}_i \leftarrow \left( (P_i + T_i) \cdot \hat{r} \right) \hat{r} - T_i$  $P_{i+1} \leftarrow \theta_g(\widetilde{P}_i, t(i))$ end for return  $P_n$ 



We apply diffusion model as your 3D Human Pose prior generator.

## **Back to Optimization: Diffusion-based Zero-Shot 3D Human Pose Estimation**

Zhongyu Jiang<sup>1</sup>, Zhuoran Zhou<sup>1</sup>, Lei Li<sup>2</sup>, Wenhao Chai<sup>1</sup>, Cheng-Yen Yang<sup>1</sup>, Jenq-Neng Hwang<sup>1</sup> <sup>1</sup>University of Washington, <sup>2</sup>University of Copenhagen

Algorithm Outline :

- Set KNN-cluster as the initial pose. Run
- optimization for appropriate R and T.
- Compute camera rays with camera parameters 2. and 2D pose.
- Move 3D pose to the rays to minimize projection 3. error. Update the new T unless it is during warmup.
- Adjust 3D pose by the prior diffusion model.
- Repeat steps 2-5 1000 times.





Our method's pipeline consists of four components illustrated in the algorithm outline. The optimization block and truncated diffusion model repeat 1000 iterations to achieve an accurate 3D prediction,  $P_{1000}$ .



### Experiment

Methods	CE	Opt	PA-MPJPE $\downarrow$	$ MPJPE\downarrow $
Kolotouros et al. [24]			59.2	96.9
Kocabas <i>et al</i> . [22]			51.9	82.9
Kocabas et al. [23]			46.4	74.7
Li <i>et al</i> . [25]			<u>45.0</u>	<u>74.1</u>
Ma <i>et al</i> . [32]			41.3	67.5
Li et al. [25]	✓		50.9	82.0
Kocabas et al. [22]	$\checkmark$		56.5	93.5
Kocabas <i>et al</i> . [23]	$\checkmark$		50.9	82.0
Gong <i>et al</i> . [11]	$\checkmark$		58.5	94.1
Gholami <i>et al</i> . [9]	$\checkmark$		46.5	<u>81.2</u>
Chai <i>et al</i> . [5]	$\checkmark$		55.3	87.7
Song <i>et al</i> . [47]	$\checkmark$		55.9	-
Choutas <i>et al</i> . [6]	$\checkmark$		52.2	-
<b>ZeDO</b> $(S = 1, J = 17)$	$\checkmark$	$\checkmark$	40.3	<b>69.7</b>
<b>ZeDO</b> $(S = 1, J = 14)$	$\checkmark$	$\checkmark$	<u>43.1</u>	76.6

Dataset	Diff Model	RO WU			S = 1		S = 50		
			WU	RA	GT	$\mathbf{MPJPE}\downarrow$	$\textbf{PA-MPJPE} \downarrow$	$\mathbf{MPJPE}\downarrow$	$\textbf{PA-MPJPE} \downarrow$
H36M	H36M					75.0	52.7	53.4	42.7
H36M	H36M	$\checkmark$				77.2	53.7	52.7	42.4
H36M	H36M	$\checkmark$	$\checkmark$			<b>65.7</b> (9.3 ↓)	<b>49.0</b> (3.7 ↓)	<b>51.4</b> (2.0 ↓)	<b>42.1</b> (0.6 ↓)
H36M	H36M	$\checkmark$	$\checkmark$	$\checkmark$		69.5	51.4	52.9	42.5
H36M	H36M	✓	$\checkmark$		$\checkmark$	50.1	35.8	37.0	27.5
3DHP	H36M				✓	148.3	88.8	93.4	59.0
3DHP	H36M	$\checkmark$	$\checkmark$		$\checkmark$	113.8	74.1	80.1	56.0
3DHP	H36M	<ul><li>✓</li></ul>	$\checkmark$	$\checkmark$	$\checkmark$	<b>99.9</b> (48.4 ↓)	<b>67.9</b> (20.9 ↓)	<b>69.9</b> (23.5 ↓)	<b>49.0</b> (10.0 ↓)
3DHP	3DHP	✓	$\checkmark$		$\checkmark$	86.5	55.9	55.2	38.6





Minimize Reprojection Error

#### Ablation Study on H36M and 3DHP



Results on 3DPW