# Multi-hypotheses Conditioned Point Cloud Diffusion for 3D Human Reconstruction from Occluded Images

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### predicts the *pixel-aligned* 3D shapes of humans *robustly* from occluded images.



Input image

Segmented images

3D reconstruction as point cloud

# Motivation

3D human reconstruction from 2D images play a significant role in metaverse.

Social interaction (human-object, human-human) make it more challenging due to occlusions.

## **3D Human Reconstruction**

## Parametric body models (SMPL/SMPL-X)

well regularized with human body priors.

#### robust to occlusion.

lack geometric details like clothing and hair.



### Implicit-function-based models

conditioned on SMPL estimation. predict the pixel-aligned 3D shapes of humans. sensitive to occlusion, cannot inpaint the invisible regions.

## MHCDIFF: Multi-hypotheses Conditioned Point Cloud Diffusion



## **MHCDIFF: Multi-hypotheses Conditioned Point Cloud Diffusion**

#### **4.2 Local features from SMPL**

signed distance and normal obtained from the closest surface of SMPL mesh independent of global pose.

generalize well in diverse SMPL estimation due to occlusion.

 $X_t^{SMPL} = [\gamma(d(X_t|S)), \mathbf{n}(X_t|S)]$ 

#### **4.3 Multi-hypotheses condition**

effectively captures the distribution of multiple plausible SMPL meshes. robust to the noise of each SMPL estimation due to the occlusion.

$$X_t^{SMPL} = [rac{1}{s} \sum_{i=1}^s \gamma(o(X_t|S_i)), \gamma(d(X_t|S_i)))$$

 $(S_{\overline{i}})), \mathbf{n}(X_t|S_{\overline{i}})]$ 



## MHCDIFF: Multi-hypotheses Conditioned Point Cloud Diffusion

### 4.4 Conditioned point cloud diffusion model

capture the global consistent features and generate the invisible parts. correct the misaligned SMPL estimation during the denoising process.

$$\mathcal{F}_{ heta}(\cdot): \mathbb{R}^{(3+c+4L+3)N} {
ightarrow} \mathbb{R}^{3N}$$



#### **Quantitative evaluation**

MHCDIFF outperforms prior implicit-function-based methods and SMPL/SMPL-X estimation methods on occluded images, and shows comparable performance on full-body images.

#### **Randomly masked CAPE**

	Methods	Chamfer Distance (cm)	Point-to-Surface (cm)		Methods	single	occluded single	two natural-inter	two closely-inter	three
Α	PaMIR [106]	12.912	12.619	A	PaMIR [106]	0.690	2.349	5.154	3.752	4.714
	ICON [92]	2.896	2.789		ICON [92]	0.555	0.549	0.563	0.786	0.669
	ICON (PIXIE estimation)	3.329	3.212		SIFU [104]	0.644	3.335	4.796	3.503	3.264
	SIFU [104]	14.397	14.087		HiLo [96]	0.606	2.808	4.139	3.346	4.398
	HiLo [96]	13.711	13.405	В	PIXIE (SMPL-X) [15]	0.868	0.813	0.755	0.951	0.809
В	PIXIE (SMPL-X) [15]	2.705	2.662		ProPose (SMPL) [14]	0.675	0.567	0.574	0.766	0.688
	ProPose (SMPL) [14]	2.370	2.307	Ours	MHCDIFF	0.591	0.491	0.536	0.703	0.673
Ours	MHCDIFF	1.872	1.810	0 410		<u>,</u>				<u></u>

#### MultiHuman

### **Qualitative evaluation (CAPE)**



#### **Qualitative evaluation (MultiHuman)**





#### **Qualitative evaluation (Hi4D)**



#### **Qualitative evaluation (in-the-wild)**



### **Qualitative evaluation (in-the-wild)**





Input image

Segmented images

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Ours



## Conclusion

### Contributions

Multi-hypotheses conditioning mechanism effectively captures the distribution of multiple plausible SMPL meshes.

Point cloud diffusion model captures the global consistent features and inpaints the invisible parts.

## Limitations

Inference is slow due to iterative denoising procedures. Point cloud may not be directly usable in real-world applications.

# Thank you!

## Website: <a href="https://donghwankim0101.github.io/projects/mhcdiff">https://donghwankim0101.github.io/projects/mhcdiff</a> Code: <a href="https://github.com/DonghwanKIM0101/MHCDIFF">https://github.com/DonghwanKIM0101/MHCDIFF</a>

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