



Diffusion Models for Unified Scanpath Prediction Giuseppe Cartella¹, Vittorio Cuculo¹, Alessandro D'Amelio²,

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Modeling Human Gaze Behavior with

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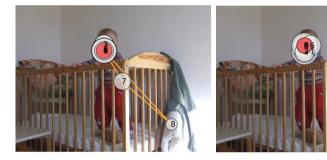
1. The Variability of Human Visual Exploration

Human Visual Exploration is inherently variable, even for the same image/task.



- The decision of where to look next at any given moment is neither entirely deterministic nor completely random [1].
- The stochasticity of gaze allocation enables the observer to remain responsive to new signals and promotes a flexible shift of attention.

The Failure of SOTA Models













Most existing approaches generate averaged behaviors, failing to capture the variability of human visual exploration.

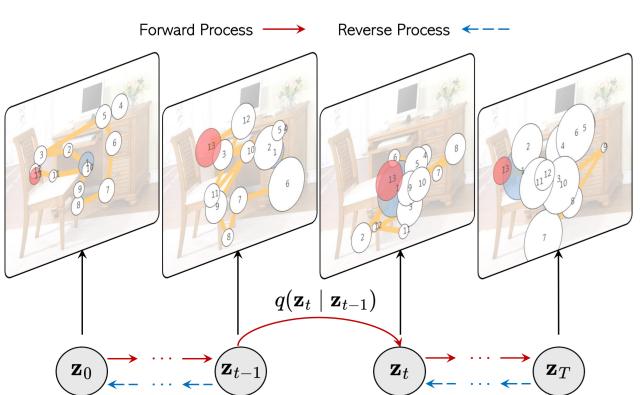
However, achieving good performance in terms of MultiMatch (MM), ScanMatch (SM) or Sequence Score (SS) DOES NOT mean that the model is able to consider the inter-subject variability.

2. Diffusion Models are a Natural Fit

Our Idea

Leverage stochastic sampling to model uncertainty in gaze allocation.

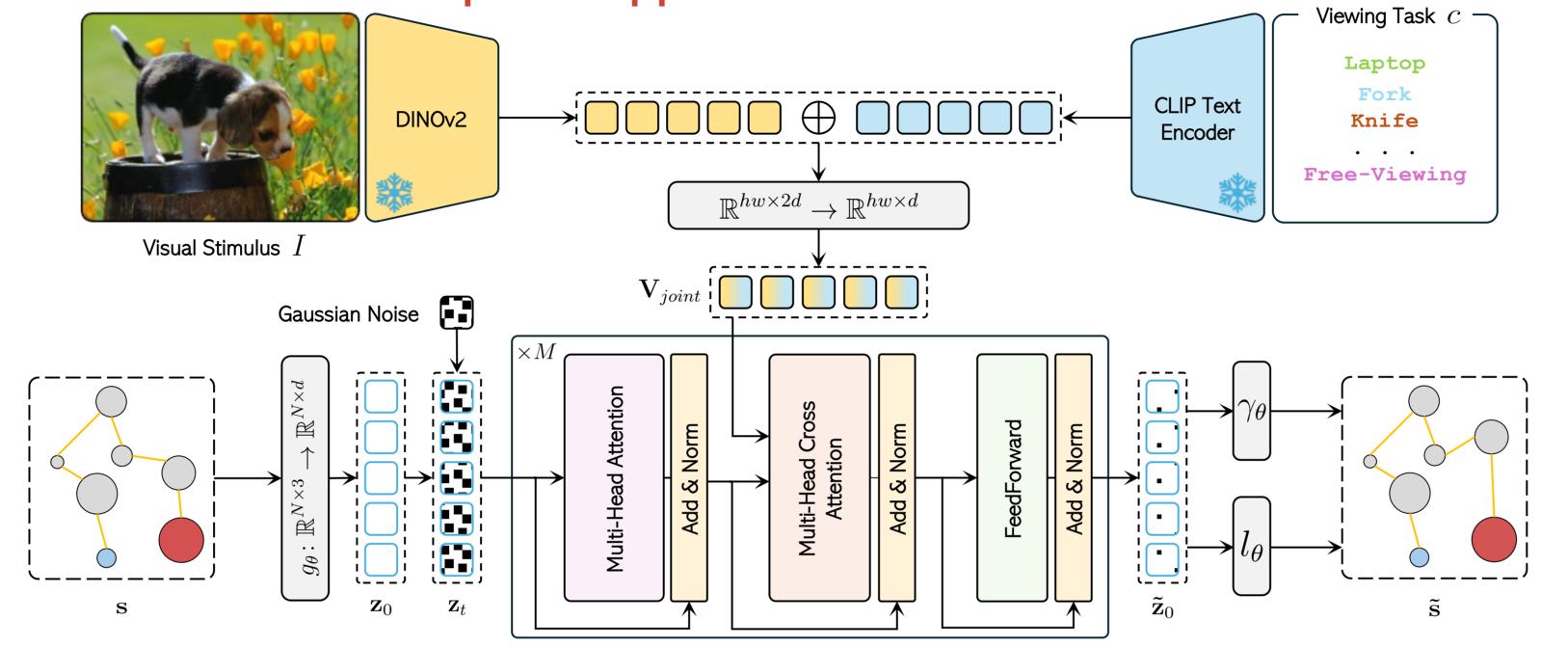
Main Contributions



 $p_{ heta}(\mathbf{z}_{t-1} \mid \mathbf{z}_t, I, c)$

- A diffusion-based architecture for scanpath prediction that reflects the inter-group variability.
- A **novel analysis** on the diversity of the generated scanpaths.
- Introduction of the **Diversity-**Aware Sequence Score.

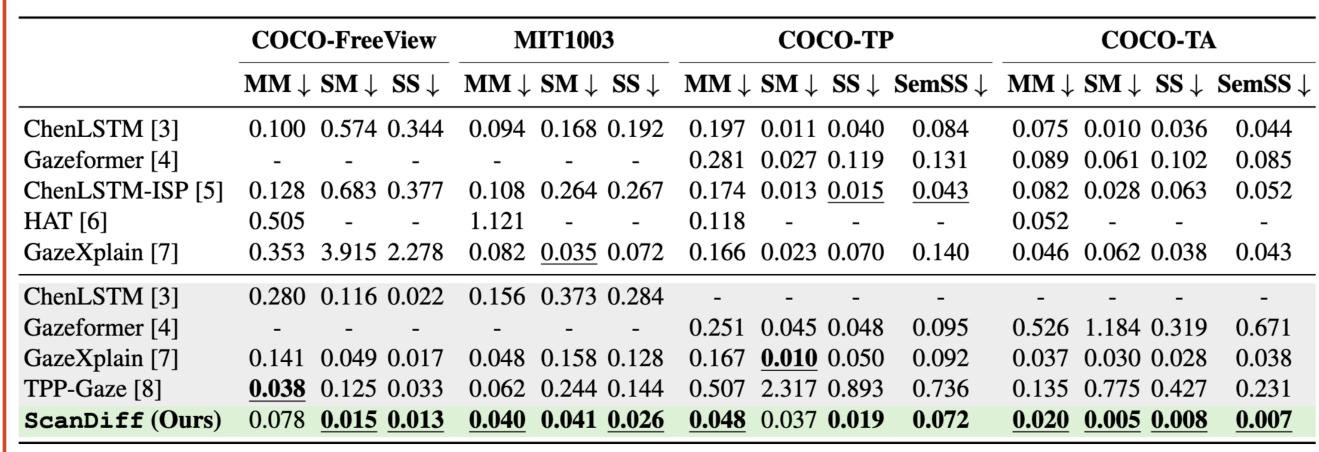




- The scanpath is embedded into the initial uncorrupted
- **Corrupt** the whole embedded sequence z_0 by adding Gaussian noise over T timesteps.
- The **stimulus** is encoded through a **Transformer-based** visual backbone.
- A text encoder embeds the **viewing task**, enabling a unified architecture.
- Visual and textual features are projected in a joint multimodal embedding space.
- A modified Transformer encoder processes the noisy embedded scanpath sequence z_t and the multimodal features serve as conditioning for the denoising process.
- A three layers MLP γ_{θ} reconstructs the original scanpath and a length prediction module l_{θ} predicts the scanpath length.

The **training objective** is defined as the combination of four different components: $\mathcal{L} = \mathcal{L}_{ ext{VLB}} + \mathcal{L}_{rec} + \mathcal{L}_{val} + \mathcal{L}_{T}$

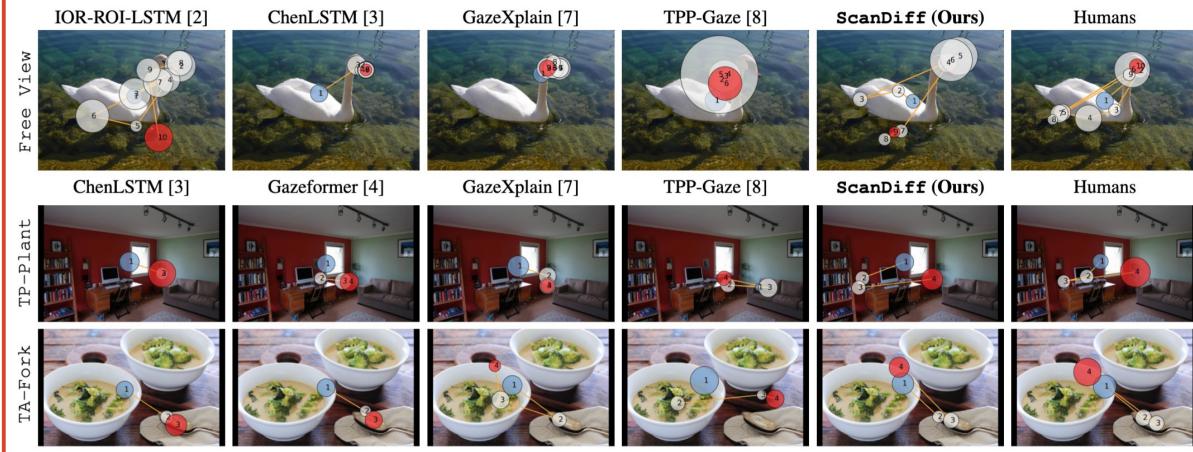
4. Comparison with the State-of-the-Art



We adopt the same evaluation protocol proposed in [7], and use the KL divergence.

ScanDiff achieves SOTA In scanpath prediction generating diverse and plausible gaze trajectories!

5. ScanDiff produces Human-Like Scanpaths

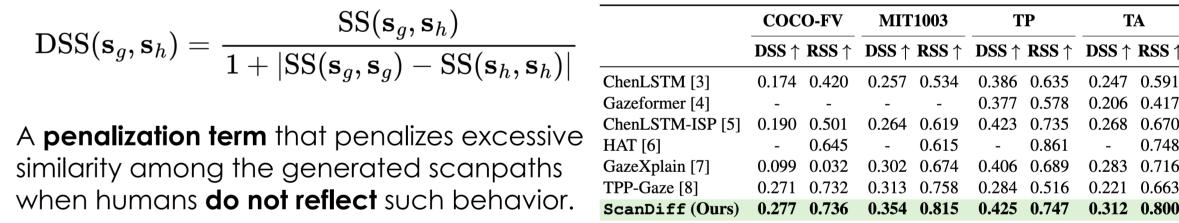


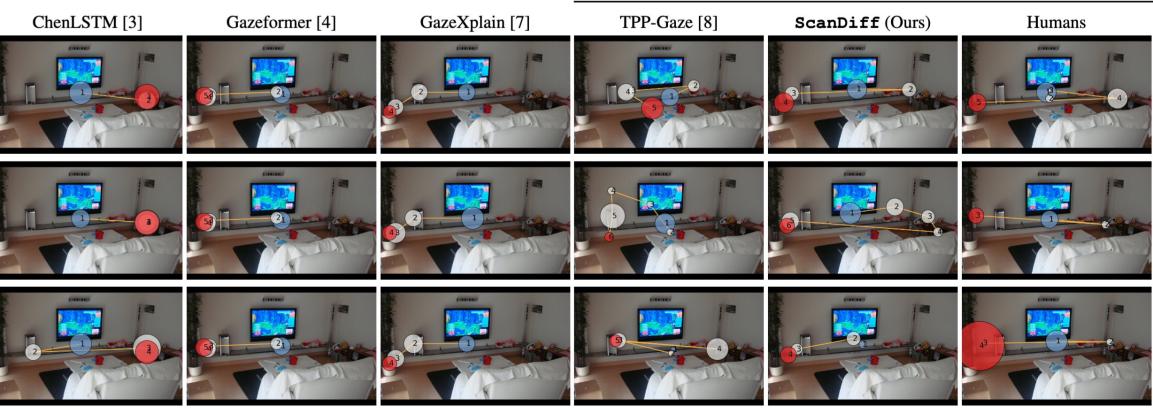
6. Scanpath Variability Analysis

Metrics such as MM, SM, and SS tend to reward models that generate a single representative scanpath (averaged behavior) [9].

This problem stems from the way metrics are computed.

We propose the Diversity-Aware Sequence Score





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9] Kümmerer et al., "State-of-the-art in human scanpath prediction", arXiv 2021