Crafting Hierarchical Strand-based Hair Geometry with Frequency-decomposed Representative Guide Curves

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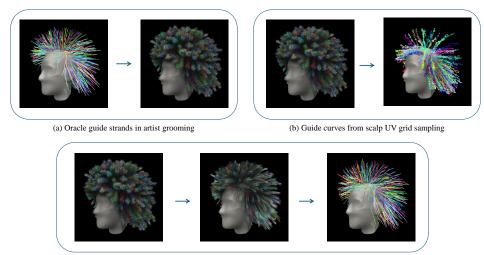
Abstract

We introduce a hierarchical generative representation for strand hair geometry that progresses from coarse, low-pass filtered guide strands to densely populated hair strands rich in high-frequency details. Unlike traditional methods that sample guide strands from a regular 2D grid on a UV scalp map, our model utilizes k-medoids clustering centers from low-pass filtered dense strands as guide curves, which more accurately retain the hairstyle's inherent characteristics. We employ the discrete cosine transform (DCT) to separate low-frequency structural curves from high-frequency curliness and noise, avoiding the Gibbs' oscillation issues associated with the standard Fourier transform in open curves. Our variational autoencoder-based model, with its permutation-equivariant architecture, facilitates flexible, off-the-grid guide strand geometry modeling and enables the completion of dense strands in any quantity and density, drawing on principles from implicit neural representations. Empirical evaluations confirm the capacity of the model to generate convincing guide curves and dense strands, complete with nuanced high-frequency details.

1 Introduction

The quest for realistic virtual humans is a cornerstone of modern computer graphics, with high-quality 3D hair strand generation being one of its most intricate challenges. Our work focuses on the generation of strand hair with machine learning algorithms, which alleviates the labor-intensive process inherent in crafting digital hairstyles—a task that requires meticulous attention to detail and consumes a disproportionate amount of time and artistic resources. In addition, it strives to establish a generative prior, a foundational blueprint that not only streamlines the creation process but also enhances the reconstruction capabilities essential for virtual human applications.

Hair exhibits a wide spectrum of morphological details, from the subtlest of low-frequency principle directions and waves to the complexity of high-frequency curls, as described by the physics of elastic rods and Kirchhoff's theories [4, 22] on the dynamics of twisting filaments. These high-frequency elements often include curly helical structures [3, 15] as well as extraneous noise that can detract from accurately modeling the hair's principal growth direction. To counteract this, we advocate for a frequency decomposition approach to extract the principal direction as a low-frequency signal. This technique is vital in distilling the essence of hair's natural trajectory, ensuring that the core path of growth is clearly defined and free from the visual clutter of high-frequency noise. Such a focus on low-frequency signals is not only important but also highly effective in generating hair models that are true to the natural flow and inherent physical properties of hair, providing a robust basis for more nuanced and detailed simulations.



(c) Guide curves from frequency decomposition and representative subset extraction (ours)

Figure 1: (a) The oracle guide curves from a grooming project in Blender [8] for principle hair growing directions. (b) Existing works simply sample guide curves from dense strands regularly on a UV grid of the scalp, which may contain unnecessary signals of high-frequency noise for neighboring dense strands. (c) We model guide curves by frequency decomposition and k-medoids clustering for a representative set of smooth guide curves, which are utilized in training our hierarchical hair generation model.

The complexity of hair data has led to the adoption of sparse guide strands as a foundational step in many hair reconstruction and generation methods, reflecting the guide hair-based authoring used in strand-based hair grooming in Fig. 1(a). These existing methods [32, 40, 46, 41], however, often simplify the process by downsampling from a regular grid on a 2D UV map of the scalp, as in Fig. 1(b), with notable drawbacks. Regular grid sampling can miss intricate hairstyle details and may produce guide strands that are not as concise or representative, burdened by high-frequency details and noise irrelevant to adjacent strands. In response to these challenges, our proposed method introduces low-pass filtering alongside the extraction of representative curve sets for guide curves, as illustrated in Fig. 1(c). This approach aligns more closely with contemporary computer graphics tools designed to craft artistically groomed hairstyles. By refining the guide curves this way, we aim to capture the essential form and structure of hairstyles more effectively, paving the way for more accurate and visually pleasing hair modeling.

To encapsulate our methodology and the contributions of our work:

- Our approach involves constructing a hierarchical generative model for hair strands that begins with the formation of coarse guide curves through low-pass filtering and culminates in the generation of dense hair strands that incorporate intricate high-frequency details.
- We utilize the discrete cosine transform (DCT) to craft the coarse and low-pass filtered guide curves. This technique effectively distinguishes the fundamental shape of the hair from the more complex aspects of curliness and noise, thereby circumventing the Gibbs' oscillation problem that often plagues the standard Fourier transform when applied to open curves.
- The guide curves are derived using k-medoids clustering centers from the dense hair data. This novel method proves to be superior in preserving the original hairstyle's features compared to the conventional practice of sampling from a regular 2D grid on a UV scalp map.
- We have developed a permutation-equivariant architecture that facilitates the learning hierarchical
 generation of a set of strands in low and high frequencies. This architecture is designed to support
 our flexible guide curve modeling approach, which does not rely on a fixed grid. The transition
 from sparse to densely populated strands is inspired by implicit neural representation, capable of
 handling an unlimited number of fine strands.

1.1 Related Work

Recent learning algorithms for strand hair focuses on capture and reconstruction from single [33, 42, 45] or multiple views [23, 32, 40], often relying on intermediate representations of volumetric occupancies [33, 44, 42, 40] or orientation fields from image gradient cues [33, 44, 40, 45, 42].

Our work focuses on building a generative model directly on human hairstyles in the form of a set of strands, which has been an established representation in industry [2],[10], [38], owing to its compatibility with physics-based applications. Consequently, strands have emerged as a favored representation in numerous computer vision, machine learning, and graphics projects. The most related works to our approach are recent methods of GroomGen [46] and HAAR [41], both relying on off-the-shelf VAE codec representations of strands [32] mapped on the domain of discretized UV map, following a coarse-to-fine pipeline in accordance with different UV grid resolutions. In contrast to these approaches, we extract guide strands at flexible locations on the scalp that are more loyal to the original hairstyle. We also utilize frequency decomposition by the DCT to achieve a compact representation without requiring training an off-the-shelf codec and optimization can be end-to-end. In addition, the generated grooming can sample any number of dense strands from arbitrary location without being restricted by the UV grid resolution.

2 Extracting the Hierarchy from Hair Strand Data

A human's hair can be seen as a set of strands $\mathcal{H}=\{l_i\}_{i=1}^N$, where each strand is a 3D curve $l(t):[0,1]\mapsto\mathbb{R}^3$. In practice, strand l is usually available as a polyline with discretely sampled control points in a sequence of n 3D points: $l=[l(0),l(1),\ldots,l(n-1)]\in\mathbb{R}^{n\times 3}$ sampled from the continuous curve. The roots of the curves are attached to the head scalp $\mathcal{M}: l(0)\in\mathcal{M}$. In this work we consider all the hairstyles are aligned to the same human head scalp geometry.

We aim at a hierarchical modeling of human hairstyles due to the high complexity of the human hair data. Inspired by the guide strand-based modeling in artist strand hair creation, we propose to smoothen the strands with frequency decomposition and extract representative coarse guide curves.

2.1 Frequency decomposition of strands as open curves

Human hair exhibits a spectrum of morphologies, ranging from straightness to inherent curvature of planar waves or 3D helices from a confluence of the strand's physical attributes, such as mass, length, and tensile elasticity [22]. For undulating and helical strands, the predominant axis of growth can be conceptualized as a smooth, low-frequency trajectory, whereas the detailed twists and noise are considered high-frequency features. Artists creating hairstyles typically start by making smooth, basic guide strands and then add detailed waves, curls, and slight randomness [2, 8]. Notwithstanding, contemporary learning-based algorithms for guide strand modeling have largely overlooked this intrinsic frequency paradigm.

In our approach, we separate hair strands into low- and high-frequency components, aligning with traditional artist-driven hair modeling techniques that utilize smooth guide strands. Initially, we model the smoothed, low-frequency base of the strand, then we enhance it with high-frequency waves, curls, and localized noise. This method not only adheres to the spectral bias principle [28] but also streamlines the learning process.

The discrete Fourier transform (DFT) is a widely applied method for frequency decomposition. However, the DFT assumes strong periodicity of the signal, thus suffers from the Gibbs' phenomenon [12] with significant oscillations when processing on open curves whose start points and endpoints do not coincide with each other [11, 9], as depicted in Fig. 2.

Therefore, we look into another Fourier-related transform, the discrete cosine transform (DCT) [1], related to the DFT but with only the real part of cosine components in the basis. DCT is widely employed in image, video, and audio signal compression. [30] The DCT has a strong energy compaction property and more flexible boundary conditions, facilitating encoding open curves with fewer harmonics. Formally, we apply the DCT to transfer the hair curve l(t) with $t=1,\ldots,n$ into frequency domain $\tilde{l}(\tau)$ by

$$\mathrm{DCT}(l(t)): \tilde{l}_X(\tau) = (\frac{2}{n})^{\frac{1}{2}} \sum_{t=0}^{n-1} w(t) \, l_X(t) \, \cos{[\frac{\pi}{2n}(2t+1)\tau]}, \quad \text{where } w(t) = \begin{cases} \frac{1}{\sqrt{2}} & \text{if } \tau = 0, \\ 1 & \text{otherwise.} \end{cases}$$

And the corresponding inverse DCT is the inverse function $iDCT = DCT^{-1}$. l_X and \tilde{l}_X denote the signal component on the X-axis coordinate in spatial and frequency spaces respectively. The same process also applies to Y- and Z-axes.

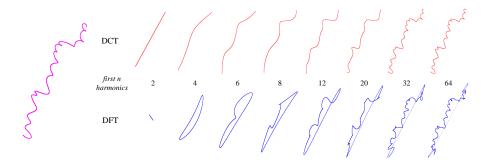


Figure 2: Comparing frequency-based smoothing of an exemplar curly strand. Smoothing from the DFT biases towards closed curves and suffers from Gibbs' oscillations [12] for hair strands as open curves. In contrast, The DCT has a strong energy compaction property and more flexible boundary conditions, facilitating encoding open curves with fewer harmonics.

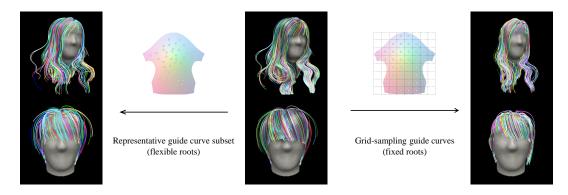


Figure 3: Comparison of guide curves extracted by k-medoids and grid-sampling from reference hairstyles in the middle, both with an equal count of curves. It is evident that grid-sampling can omit crucial hairstyle details. In contrast, the representative guide curve subset derived from our -medoids method more accurately captures the essence of the original hairstyle.

Due to the even symmetry of the cosine function, the DCT is equivalent to the DFT operating on the symmetrically extended signal sequence of twice the length, and the sine components are canceled out. As a result, the periodicity extension of the signal no longer introduces strong discontinuity, which facilitates modeling open curves, as corroborated by Fig. 6. We refer to the literature for details [30, 9].

2.2 Extracting representative coarse guide curves by k-medoids clustering

Learning-based approaches to strand-based hair modeling typically adopt a coarse-to-fine strategy, necessitating the generation of a set of coarse guide strands due to the computational demands of processing dense strands. This concept aligns with the traditional artist-driven hair creation workflow. Conventional methods predominantly employ a strategy of sampling guide strands by extracting those emanating from scalp locations situated at the grid centers of an unwrapped UV map of the scalp manifold. However, this regular grid sampling approach is suboptimal, often failing to capture certain strand clusters critical for detailing the hairstyle, as illustrated in Fig. 3. In contrast, we introduce a methodology for extracting a sparse subset of k off-the-grid strands from the comprehensive dense hair strand set $\mathcal{H}^L = \{l^L\}$, with l^L denoting low-pass filtered strand curves from the DCT as pridominent growing directions, resulting in a representative curve set \mathcal{G}_k^* that more accurately mirrors the characteristics of the dense strands.

Definition 1 (Representative coarse guide curves). Given a set of dense low-pass filtered hair curves \mathcal{H}^L , the representative coarse curve set \mathcal{G}_k^* of cardinality k is the subset of \mathcal{H}^L which has the minimum possible bidirectional chamfer distance from \mathcal{H}^L . Formally, $\mathcal{G}_k^* = \arg\min_{\mathcal{G}_k \subset \mathcal{H}^L: |\mathcal{G}_k| = k} \operatorname{chamfer}(\mathcal{H}^L, \mathcal{G}_k)$

Our investigation reveals that the k-medoids clustering method [16] delivers an optimal approach for identifying the representative strand subset. k-medoids is a classical partitioning technique that organizes data points into clusters with the aim of minimizing the distance between the points within a cluster and a specific point within the same cluster, known as the medoid, which serves as the cluster's nucleus. Unlike the popular k-means algorithm which allows the cluster's centroid to be a virtual average of the data points, k-medoids selects actual data points as the medoids, enhancing the interpretability of the cluster centers. Formally, the objective of k-medoids is to partition the set \mathcal{H}^L into a collection of disjoint clusters $\mathbb{S} = \{\mathcal{S}_1, \dots, \mathcal{S}_k\}$, each with its corresponding medoids $\mathcal{U} = \{u_1, \dots, u_k\}$ in such a way that the aggregate of dissimilarities to all elements within each cluster is minimized, as defined by the following objective:

$$\min_{\mathbb{S}} \sum_{i=1}^{k} \sum_{l^L \in \mathcal{S}_i} d(l^L, u_i), \quad \text{where } u_i = \underset{l^L \in \mathcal{S}_i}{\arg\min} \sum_{l'^L \in \mathcal{S}_i} d(l^L, l'^L), \tag{2}$$

and $d(\cdot, \cdot)$ is a dissimilarity measurement. This objective leads to the following observation:

Theorem 1. The medoid set $\mathcal{U} = \{u_1, \dots, u_k\}$ from the k-medoids clustering of \mathcal{H} is the representative curve set \mathcal{G}_k^* , if aggregated squared Euclidean distance $d(l, l') = \frac{1}{n} \|l - l'\|_2^2 := \frac{1}{n} \sum_{t=0}^{n-1} \|l(t) - l'(t)\|_2^2$ for two individual curves is used as the divergence function for k-medoids.

Proof. Since each element is closest to its own cluster center, the objective of k-medoids in eq. 2 can be reformulated as finding the set of medoids $\mathcal{U} = \{u_1, \dots, u_k\} \subset \mathcal{H}$ such that $\mathcal{U} = \arg\min_{\mathcal{G}_k} \sum_{i=1}^k \sum_{l^L:d(l^L,u_i)=\min_{u_j\in\mathcal{G}_k} d(l^L,u_j)} d(l,u_i)$. So $\mathcal{U} = \arg\min_{\mathcal{G}_k} \sum_{l^L\in\mathcal{H}} \min_{u_i\in\mathcal{G}_k} d(l^L,u_i) = \arg\min_{\mathcal{G}_k} \sum_{l\in\mathcal{H}} \min_{u_i\in\mathcal{G}_k} \|l^L-u_i\|_2^2$. On the other hand, $\mathcal{G}_k^* = \arg\min_{\mathcal{G}_k} \operatorname{chamfer}(\mathcal{H},\mathcal{G}_k) = \arg\min_{\mathcal{G}_k} \frac{1}{|\mathcal{H}|} \sum_{l^L\in\mathcal{H}} \min_{u_i\in\mathcal{G}_k} \|l^L-u_i\|_2^2 + \frac{1}{|\mathcal{G}_k|} \sum_{u_i\in\mathcal{G}_k} \min_{l^L\in\mathcal{H}} \|l^L-u_i\|_2^2$ in which the second term is zero because $\mathcal{G}_k \subset \mathcal{H}$. Therefore $\mathcal{G}_k^* = \arg\min_{\mathcal{G}_k} \sum_{l^L\in\mathcal{H}} \min_{u_i\in\mathcal{G}_k} \|l^L-u_i\|_2^2 = \mathcal{U}$.

3 Learning to Generate Hierarchical Strand Hair

3.1 Processing hair data into hierarchy

In our approach to hair data abstraction, we extract the low-frequency structure disentangled from high-frequency details for each strand as the principle growing directions, followed by selecting a representative coarse subset of guide strands from the low-pass filtered dense hair data.

For each strand l, we map the root point on the scalp manifold $l(0) \in \mathcal{M}$ to its UV coordinate on the UV mapping of the scalp as $r \in \mathcal{M}_{\mathrm{UV}} \subset \mathbb{R}^2$. We translate all strands to centralize the root points at the origin of the 3D space, and achieve the low-pass filtered curve with the first \tilde{n}^L components in the DCT. We apply the DCT on the derivative $\dot{l}(t) = l(t+1) - l(t)$ instead of the original sequence of coordinates to ensure the same root position. To condense the signal, the reconstructed curve l_i^L in the spatial domain is resampled to a reduced resolution of $2 \cdot \tilde{n}^L$ by fitting a cubic spline, adhering to the Nyquist sampling theorem [35] for precise discretization. For high-frequency details, we maintain the signal in the form of DCT coefficients, denoted as $\tilde{l}^H \in \mathbb{R}^{(\tilde{n}^H - \tilde{n}^L) \times 3}$ A cut-off frequency \tilde{n}^H is strategically selected to ensure compactness and exclusion of signal components exceeding \tilde{n}^H as high-order noise. Practically, we set $\tilde{n}^L = 8$ and $\tilde{n}^H = 40$. Next, we employ k-medoids to identify the indices of the representative coarse guides for each hairstyle, maintaining a consistent k = 512.

This way, we represent each strand as a tuple (r, l^L, \tilde{l}^H) , where r is the root point's UV coordinate, l^H is the low-pass filtered strand curve, and \tilde{l}^H encapsulates the high-frequency details as DCT harmonics. Strands within the representative coarse guide curve set are denoted as (r^*, l^{L*}) while high-frequency signals are excluded from guide curves. The learning process for hair generation is structured hierarchically: it commences with the generation of the guide curve set $\{(r^*, l^{L*})\}$, progresses to the densification of strands $\{(r, l^L)\}$, and culminates with the integration of high-frequency details $\{\tilde{l}^H\}$ when necessary, for hairstyles with heavy curliness or non-smoothness.

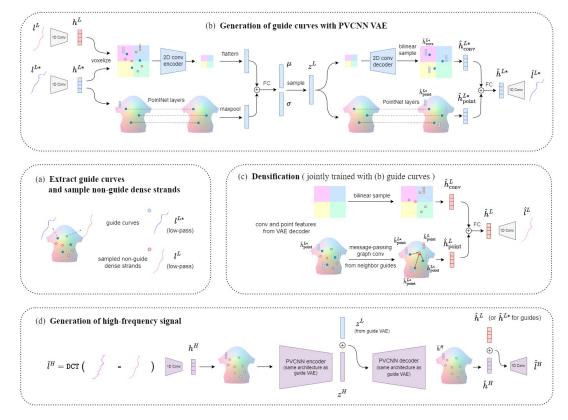


Figure 4: Generation pipeline overview: (a) During training, a small batch of dense strands and extracted guide curves are sampled from the UV map of the scalp to optimize the networks. (b) Guide curves are generated using a PVCNN VAE that incorporates 2D convolution and PointNet layers. (c) Dense strands are generated by aggregating features from the convolution grid via bilinear sampling and from neighboring guide curves through graph convolution, with the densification module being jointly trained with the guide curve model. (d) High-frequency signals are refined using another PVCNN VAE, conditioned on the global and local latent features derived from the network responsible for generating the low-pass filtered principal strand signal.

3.2 Generation pipeline

The generative process of our model is depicted in Fig. 4. We employ variational autoencoders (VAE) [18, 31] to capture the distribution of hierarchical hair structures. The choice of VAE is motivated by its straightforwardness and proven effectiveness and efficiency in optimization and sampling, while other categories of generative models are left to future exploration. The objective of a VAE is to learn a generative model $p_{\theta}(x,z) = p_{\theta}(z)p_{\theta}(x|z)$ for data x and latent variables z. Since the true posterior is intractable, we approximate it using the latent encoder model $q_{\phi}(z|x)$ and optimize the variational lower bound (ELBO) on the marginal likelihood p(x): $\mathcal{L}_{\text{VAE}} = \mathbb{E}_{q_{\phi}(z|x)}[\log p_{\theta}(x|z)] - \text{KL}(q_{\phi}(z|x)||p_{\theta}(z))$, where the second term is the Kullback-Leibler divergence, quantifying the difference between the approximate and true posteriors.

Generation of representative coarse guide curve set with PVCNN network The guide VAE for the generation of the representative guide curve set $\{(r^*, l^{L*})\}$ with a fixed cardinality k is detailed in Fig. 4(b). The guide curves $\{(r^*, l^{L*})\}$ are conceptualized as a collection of data points, with the sequence l^{L*} transformed to a feature vector $h^{L*} \in \mathbb{R}^{d_{h^{L*}}}$ using a 1D convolutional encoder embedded on the domain of UV coordinates $r^* \in [0,1]^2$. Our network architecture draws inspiration from the Point-Voxel CNN (PVCNN) [19], leveraging its demonstrated efficiency and effectiveness in the domain of point set learning.

The architecture, akin to Point-Voxel CNN (PVCNN), utilizes a dual-branch approach to process point features: a PointNet branch and a convolution branch. Within the encoder, the point branch consists of a shared multilayer perceptron (MLP) that operates on all points, employing max pooling

for aggregation as per the PointNet design [27]. The convolution branch, on the other hand, begins by partitioning the UV space $[0,1]^2$ into a 2D grid of $W\times H$ resolution. This step is analogous to the "voxelization" in PVCNN but is executed in 2D. Each grid cell is assigned an averaged feature vector derived from all data points whose root coordinates fall within it, yielding a feature tensor of dimensions $\mathbb{R}^{W\times H\times d_{h^{L*}}}$. This tensor is then processed by successive downsampling 2D convolution layers to distill a global feature. to enhance the feature representation in the encoder's convolution branch, we incorporate features from the sampled dense strands. These additional features are not utilized in the PointNet branch to avoid increasing the memory footprint, whereas the convolution branch feature is constrained to a fixed resolution. The mean and standard deviation for reparameterizing the VAE latent code z is the aggregated features from both branches.

The decoder's architecture mirrors that of the encoder, except that the point feature is bilinearly sampled from the feature grid at the end of the convolution branch, following the hybrid grid-implicit representations [26, 5]. The interpolated point feature is then reconstructed into the low-passed curve sequence l^{L*} .

Sampling root positions from generated density map The proposed PVCNN architecture learns generation of the curve sequence $p(\{l^{L*}\}|\{r^*\})$ given root positions $p(\{r^*\})$. Our objective extends to mastering the joint distribution $p(\{r^*,l^{L*}\})$, which we approach by decomposing $p(\{r^*\},\{l^{L*}\})$ into $p(\{r^*\}) \cdot p(\{l^{L*}\}|\{r^*\})$ as optimizing the root coordinates and curve sequence jointly presents significant empirical challenges. While the PVCNN network is utilized for $p(\{l^{L*}\}|\{r^*\})$, learning $p(\{r^*\})$ could potentially be achieved with a distinct generative network, such as SetVAE [17]. Nevertheless, we employ a trick to generate a $W \times H$ grid map of root densities at the convolution branch decoder, achieved by tallying the roots within each grid cell. During inference, $p(\{r^*\})$ is derived from this grid density map, before the sequence signal $\{l^{L*}\}|\{r^*\}$ is generated.

Densification To accommodate our innovative off-the-grid guide curve design, we introduce a densification module inspired by the concept of implicit neural representations [24, 21, 7, 39] This approach enables the generation of an arbitrary number of fine strands, independent of the scalp UV map's resolution, by leveraging hybrid grid [26, 5] and graph [6] representations. Given a randomly sampled query root position $r_i \in \mathcal{M}_{\text{UV}}$, our goal is to derive fine strand features that align with the guide curve features within the VAE decoder. Features from the convolution branch $\hat{h}_{\text{conv},j}^{L*}$ are bilinearly sampled from the convolution branch, while for the PointNet branch, the query aggregates pointwise features $\hat{h}_{\text{point},j}^{L*}$ from the query's neighboring guide curve locations $r_j^* \in \mathcal{N}_{r_i}$, employing a graph-hybrid implicit representation strategy [6] This method considers guide curves as anchor points and propagates signals to arbitrary query locations through message-passing graph convolution [13]: $\hat{h}_{\text{point},i}^L = \frac{1}{|\mathcal{N}_{r_i}|} \sum_j w_j (\text{MLP}(r_i, r_j^* - r_i, \hat{h}_{\text{point},j}^{L*}) + \hat{h}_{\text{point},j}^{L*})$ with a two-layer MLP, and $w_j = \frac{1}{\|r_j^* - r_i\|_2} / \sum_{j'} \frac{1}{\|r_{j'}^* - r_i\|_2}$ is a decaying weight. The resulted features are passed to the same 1D convolution decoder in the guide curve VAE model.

It is important to note that the generation of representative guide curves and sampled non-guide strands is trained jointly. This pipeline is designed to be invariant to the sampling density, allowing for the generation of any desired number of dense strands during inference.

Refinement of high-frequency details The model for learning the high-frequency components $\{\tilde{l}^H\}$ is a conditioned VAE with the same PVCNN architecture as the model that generates guide curves, except that the model is conditioned by concatenating the global latent from the guide model and the local dense strand feature. This way, we can sample different forms of high-frequency details of noise and curly structures from the same generated low-frequency components of strands.

4 Experiments

Data Our method is trained on a dataset comprising 658 synthetic strand hairstyles. Detailed information on the dataset and training procedures is available in the appendix.

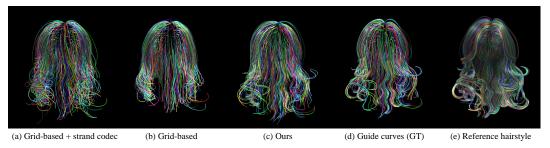
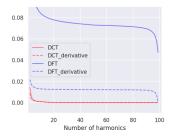
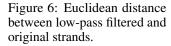


Figure 5: The VAE reconstruction of guide curves, showcasing our method's enhanced performance due to informative guide curves and enriched network representation.





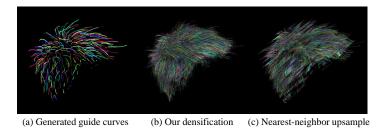


Figure 7: Dense strands from VAE generated guide curves, with our model yielding natural-looking hair, in contrast to the nearest neighbor baseline which displays artifacts.

Frequency decomposition and guide curve extraction Our evaluation focuses on the design choices made in modeling the hierarchical structure of hairstyles. As depicted in Fig. 6, we apply low-pass filtering with a designated cutoff frequency to the strands, either on the original coordinate sequence l(t) or its derivative $\dot{l}(t)$, the latter being the approach we adopt. In both scenarios, the Discrete Cosine Transform (DCT) outperforms the Discrete Fourier Transform (DFT) in reconstructing the signal within the same low-pass frequency band. The derivative form of the signal, being simpler to model, enhances the performance of both DFT and DCT, with DCT showing superior effectiveness. Furthermore, in Table 1, we conduct a comparative analysis of the k-medoids method against grid-based and farthest point sampling techniques for guide curve extraction. The k-medoids approach demonstrably achieves a closer approximation to the dense strand set, as evidenced by the lower bidirectional chamfer distance (CD). These empirical findings align with our theoretical expectations, underscoring the efficacy of our methodological choices in capturing the intricate details of hair structure and texture.

Reconstruction of guide curves with VAE

Our generative hair representation's efficacy is assessed using a set of 30 test examples, as detailed in Table 2 and illustrated in Fig. 5. Our method outperforms traditional grid-based hair representations, which map strands onto a scalp UV grid map and process them with purely 2D convolutional layers. The enhanced performance can be attributed to the rich representation derived from guide curves and our off-the-grid architectural approach. Direct comparison with closely related works such as Groom-Gen [46] and HAAR [41] is challenging due to their unavailability at the time of submission and the fact that they are trained on different datasets. Instead, we benchmark against a conceptually

Table 1: Evaluation on extracted guide curves.

| | Grid-sample | FPS | k-medoids |
|---|-------------|-------|-----------|
| $CD_{(\times 10^{-4})}$ to reference hair | 0.683 | 0.139 | 0.101 |

Table 2: Evaluating reconstruction of guide curves.

| $CD (\times 10^{-4})$ to: | Grid-based | Grid-based + strand codec | Ours |
|---------------------------|------------|------------------------------|-------|
| Input sampled strands | 1.521 | 1.597 | 0.906 |
| GT dense hair | 9.090 | 10.843 | 8.079 |

Table 3: Evaluation of densification on VAE reconstructed guide curves.

| | Nearest-neighbor | Ours |
|---|------------------|-------|
| $CD\ (\times 10^{-4})$ to GT dense hair | 0.311 | 0.209 |

similar baseline: a grid-based representation paired with a pretrained off-the-shelf strand VAE [32]. This alternative, however, falls short in performance. We posit that a strand codec VAE, originally designed as a generative prior for single strand multiview reconstruction, may not be optimal for

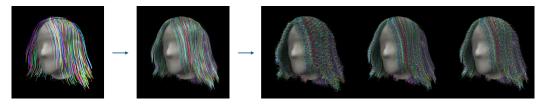


Figure 8: Generating diverse high-frequency details: Our high-frequency conditional VAE adds varied curliness and noise to hairstyles with guide curves and dense strands in low-frequency, demonstrating our model's ability to disentangle low- and high-frequency strand geometry.

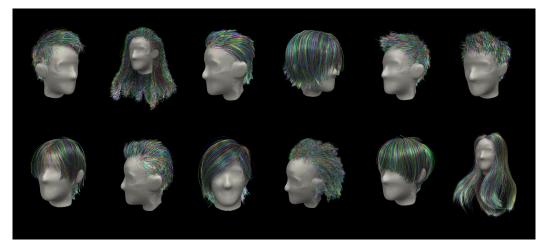


Figure 9: Strand-based hairstyles generated using our approach.

hairstyle-level generative tasks. By jointly optimizing the strand encoder and decoder, our model circumvents the error accumulation inherent in the strand VAE approach, leading to more accurate and realistic hair generation.

Densification Building on the guide reconstruction experiments, we further assess the quality of densified strands against the reference hair, as quantified in Table 3. Our method outperforms the baseline approach, which merely duplicates strands from the closest guide curve. Notably, our model demonstrates the ability to generate high-quality dense strands informed by the generated coarse guide priors, benefiting from the joint training with the guide curve VAE. As illustrated in Fig. 7, our model is capable of refining a set of VAE-generated guide curves into dense strands with fine textural details, in contrast to the nearest neighbor upsampling method, which introduces unnatural artifacts.

Created varied high-frequency details in hair Our methodology extends to generating diverse high-frequency details for each hairstyle, building upon the foundation of guide curves and dense strands in low-frequency created by our model. By employing a high-frequency conditional VAE, we introduce a variety of high-frequency details, such as curliness and noise, into the hair strands. This process is vividly illustrated in Fig. 8, which showcases the capability of our approach to effectively disentangle and separately model the components of hair strand signals in both low- and high-frequency domains. This disentanglement allows for the nuanced recreation of hair textures, enhancing the realism and diversity of the generated hairstyles.

Additional results Some outcomes of our hair generation pipeline are showcased in Fig. 9. Ablation experiments are conducted and reported in the appendix.

5 Conclusion

The proposed hierarchical generative representation for strand captures the full spectrum of hair details from low-frequency shapes to dense and fine details. By leveraging k-medoids clustering and the discrete cosine transform, our model surpasses traditional grid representations in preserving the authenticity of hairstyles. The permutation-equivariant architecture of our variational autoencoder ensures the generating any amount of hair strands that are both diverse and realistic, free from

the constraints of grid-based systems. Future work could explore the integration of dynamic hair behaviors and the adaptation of our model to generate a wider variety of hair types, further enhancing the realism and applicability of virtual hair in diverse digital environments.

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A Border Impacts

Advancements in hair generation technology, such as our work, can enrich virtual realism and inclusivity, benefiting industries like gaming and entertainment. However, they also raise concerns about potential misuse in deepfakes and the reinforcement of unrealistic beauty standards. It is essential to balance these innovations with ethical considerations to ensure their positive impact.

B Ablation Studies

We report ablation results to verify the design choices in our approach in Tables 4 and 5.

In Table 4, we assess the VAE reconstruction efficacy with various methodological alternatives. Opting for randomly sampled guides over k-medoids derived ones compromises representativeness, adversely affecting performance. Utilizing frequency space coefficients to represent principal low-pass curves also leads to increased error, justifying our choice to revert low-pass filtered signals to the spatial domain. Our integrated training approach for both strand and hairstyle levels proves superior to employing a pretrained strand VAE codec, as utilized in other studies.

We also explored substituting graph aggregation with the nearest guide's PointNet branch feature for each dense strand's query root position, which negatively impacted the upsampled strands' accuracy.

For the high-frequency model, we benchmarked against using a pretrained strand codec for reconstructing each strand's high-frequency signal. The mean average error in reconstructing high-frequency coefficients suggests that an end-to-end optimization of strand features is more effective than relying on a pretrained codec.

Table 4: Ablation on VAE reconstruction of low-freq. guide curves and densification (CD $\times 10^4 \downarrow$).

| | guide-to-GT guide | guide-to-GT dense | dense-to-GT dense |
|---|-------------------|-------------------|-------------------|
| Ours | 0.906 | 8.079 | 0.209 |
| Randomly sampled strands as guide | 0.997 | 8.220 | 0.213 |
| $l^L ightarrow 	ilde{l}^L$ (frequency space) | 1.152 | 8.304 | 0.226 |
| With off-the-shelf strand codec | 0.966 | 8.117 | 0.216 |
| W/o graph aggregation in densification | 0.913 | 8.102 | 0.214 |

Table 5: Ablation on high-freq. reconstruction (MAE \downarrow).

| | MAE↓ |
|---|-------|
| Ours | 0.137 |
| With off-the-shelf strand codec (freq. space) | 0.145 |

C Implementation Details

Our method is implemented with PyTorch [25]. All experiments are conducted on a single Nvidia RTX A4500 GPU. We use AdamW [20] with learning rate 3×10^{-4} for 100k iterations for both low-frequency and high-frequency models, with an exponential learning rate decay to 3×10^{-6} at the end of training. The batch size is 32.

During training we apply random horizontal flipping and online scale augmentation in a range of [0.95, 1.1]. To prevent optimization of the network from overfitting, in training we oversample 1024 guide curves in data preprocessing and further subsample 512 of them in training, while half of the guide set can still capture the characteristics of the hairstyle. We also sample another batch of 512 non-guide strands in each hairstyle in training to enrich the convolution branch representation. During evaluation we are able to generate arbitrary number of strands but we limit the number to 10000 for visualizations in the paper. The downsampling and upsampling of strands is achieved by fitting cubic splines with corresponding numbers of control points.

We use the visualization tool provided by the code repository in [36] for our work. We use the code from [34] to extract k-medoids.

C.1 Details of the architecture

The 1D convolution encoder and decoder The 1D conv encoder applied on each low-frequency curve $l^L \in \mathbb{R}^{\tilde{n}^L \times 3}$ consists of two downsampling blocks to map the input sequence resolution from 16 to 8 and 8 to 4. Each downsampling block consists of four 1D conv layers with kernel size 3, and every third conv layer of each block applies a stride of 2 for downsampling. In between every two conv layers we add a residual skip connection [14] and if downsampling is involved then the skip connection is replaced by a 1D conv of kernel size 1 with a stride of 2. The dimension of the latent feature is 16. After the sequence length becomes 4 we flatten the sequence data to a vector from 4×16 to 64. Then we attach two fully connected layers to map each layers to map the latent dimension to 32. The 1D conv decoder is in a reversed architecture of the encoder except that the upsampling conv layer is in a depth-to-space design [37].

The 2D convolution and PointNet branches After each strand sequence is transformed into a vector code, the 2D conv branch encoder aggregates them to a grid of $W \times H = 32 \times 32$ resolution, and downsampled to 2×2 after four downsampling blocks. The downsampling blocks are similar to those in 1D but with 3×3 kernels. And the latent dimension in downsampling blocks are 32, 64, 128, and 256. The $2 \times 2 \times 256$ tensor is then flatterned to a 1024-dim latent vector.

The PointNet branch encoder contains four residual blocks, and each are with two fully connected layers with residual connection. At the beginning of each block we concatenate the roots' coordinates in UV and 3D spaces. At the end of each block we apply a max pooling aggregation and concatenated to the feature of each point cloud, as the input to the next layer, until reaching the end of the fourth block where the max-pooled feature is the global feature from the PointNet branch. The latent dimensions in these blocks are 32,32,64, and 128.

We concatenate the 1024-dim feature from the conv branch and the 128-dim feature, and map them to mean and standard deviation vectors of 1024-dim to reparameterize the latent code of the VAE. The decoder is the reserves of the encoder and similarly the depth-to-space upsampling design [37] in adapted in 2D convs.

In between each two linear or convolution layers throughout the model, we use Swish activation [29] and Group Normalization [43] which divides latent channels into 8 groups.

Densification The two-layer MLP in the message-passing graph convolution has latent dimension of 32 The number of neighboring guide features used in the graph aggregation is 8.

High-frequency VAE The architecture is mostly the same as the guide curve VAE except that some layers need to change the input dimension after the guide VAE latent code is concatenated. In addition, the sequence length of \tilde{l}^H is 32 in the frequency space so there are one more downsampling block in the 1D conv encoder and one more upsampling block in the 1D decoder.

Loss functions The Variational Autoencoder (VAE) framework employs the Evidence Lower Bound (ELBO) which comprises two principal components: the KL divergence term and the reconstruction term. For our low-frequency guide curve model, the reconstruction term is quantified using the mean squared error (MSE) L2 loss on the signal, supplemented by an additional L2 loss on the signal's derivative. Furthermore, a specific term is dedicated to optimizing the reconstruction of the density map, as detailed in Section 3. In the densification process, we introduce a set of losses for the original signal, its derivative, and the density map associated with dense strands. These losses are integrated with a discounting weight of 0.1 to balance their contribution. For the high-frequency model, we incorporate a KL divergence term for the high-frequency latent variable, alongside an L1 loss for the accurate reconstruction of the Discrete Cosine Transform (DCT) harmonics.

Evaluation metric We use aggregated squared Euclidean distance $d(l,l') = \frac{1}{n} \|l - l'\|_2^2 := \frac{1}{n} \sum_{t=0}^{n-1} \|l(t) - l'(t)\|_2^2$ to evaluate distance of two individual strand curves, and the bidirectional chamfer distance based on the individual strand metric to evaluate distance of two hairstyles as sets of strands: $\frac{1}{|\mathcal{H}|} \sum_{l \in \mathcal{H}} \min_{l' \in \mathcal{H}'} d(l,l') + \frac{1}{|\mathcal{H}'|} \sum_{l' \in \mathcal{H}'} \min_{l \in \mathcal{H}} d(l',l)$.

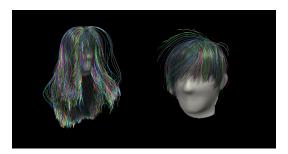


Figure 10: Failure examples.

D Data

The source of synthetic hairstyle data is twofolds. We use 343 synthetic hairstyles that originate from mesh hair cards [15], thus consisting of smooth strands with straight and wavy structures in low frequencies without high-frequency details. In addition, we collected and crafted 26 base particle hair projects using Blender [8] in the form of genuine 3D strands. The 343 hair mesh converted hair and the 26 base strand hair are used for training the low-pass signals in the guide curve VAE and the densification model. For high-frequency signals, the 26 base particle hair are further augmented into 315 examples by operating with the frequency and the magnitude of curliness, and adding random high-frequency noise in Blender. These augmented data are used to train conditional VAE model for high-frequency signal completion, while those from the mesh hair cards are precisely enough to be represented with the first eight DCT harmonics.

E Limitations

The quality of generated hair is constrained by the scarcity of high-quality, strand-based hair datasets. Even with heavy data augmentation as proposed in [46, 41] for local visual features, the variability of the rough geometry of the principal growth directions indicated by the guide curves is limited. Therefore, acquiring a diverse collection of high-quality strand hair examples is crucial for learning a robust generative latent manifold that generates more creative hairstyles.

Another challenge is the efficiency of our proposed method. As it is a geometric deep learning model that handles sets of permutation-equivariant elements, it does not match the runtime efficiency of purely convolutional architectures. For instance, generating guide strands with a batch size of 32 requires 0.7 seconds, whereas a pure 2D convolutional baseline performs inference in just 0.09 seconds. The complete generation process, including densification and high-frequency detail addition, takes about 12 seconds. Despite being less efficient, the generation time remains within a practical range for real-world applications.

Some occasional failure cases include messy isolated flying strands and penetration into head meshes. as in Fig. 10