TRAJEVAE: Controllable Human Motion Generation from Trajectories

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Abstract

The creation of plausible and controllable 3D human motion animations is a long-standing problem that requires a manual intervention of skilled artists. Current machine learning approaches can semi-automate the process, however, they are limited in a significant way: they can handle only a single trajectory of the expected motion that precludes fine-grained control over the output. To mitigate that issue, we reformulate the problem of future pose prediction into pose completion in space and time where multiple trajectories are represented as poses with missing joints. We show that such a framework can generalize to other neural networks designed for future pose prediction. Once trained in this framework, a model is capable of predicting sequences from any number of trajectories. We propose a novel transformer-like architecture, TRAJEVAE, that builds on this idea and provides a versatile framework for 3D human animation. We demonstrate that TRAJEVAE offers better accuracy than the trajectory-based reference approaches and methods that base their predictions on past poses. We also show that it can predict reasonable future poses even if provided only with an initial pose.

1. Introduction

Creating realistic human animation is one of the key components in robotics, game, and movie industries. Typically when working on an animation, the animator starts with defining a character’s skeleton. Parts of this skeleton are manually created and defined to be in a specific relation such that each joint can influence the position of other joints. Together, these parts form a kinematic chain. While these relations are helpful to maintain skeleton movement constraints, they are not sufficient to create realistic animation. In fact, the generation of such animations requires manual key-framing of the joint positions throughout the sequence. That becomes quickly an unfeasible task for complex motions.

The procedure can be aided with recent advances in machine learning. The automation of character animation is a long-standing problem with multiple solutions proposed, including those based on neural networks and probabilistic models \cite{16,27,28,55,56,61}. The main goal of these methods is to generate sequences of joint positions given some conditioning information, \textit{i.e.} control signal. This control signal can be any partial future information, for example, the direction of the movement, speed, type of an action be-
ing performed, coordinates of a particular joint, or any combination of the above [11, 17, 24, 26, 51, 65, 71]. However, these methods can handle only simple motions of a body, such as walking, running, or side-stepping, and cannot infer fine-grained motions of each of the joints. In many cases, this formulation becomes impractical, if we wanted to simulate a crowd where characters perform vivid actions such as jumping, bending, or hand waving.

To solve this issue, we propose a data-driven approach to train a model to handle a variable number of pieces of information, trajectories, for human motion generation. A single trajectory refers to a particular skeleton joint, e.g. elbow, and specifies where that joint should be located in each time step. We can choose whether we want to specify trajectories of a few joints and leave the rest to be generated by the model or generate a more specific motion by adding more trajectories. This formulation introduces an unprecedentedly flexible framework that encompasses all previous approaches while offering adjustable control over the generated motion.

We formulate the problem of predicting future poses from trajectories as a pose completion problem to achieve our goal. We trace the inspiration for that formulation to the evolutionary cognitive skills of humans who are able to hallucinate [32] the rest of a human body out of a few markers that exhibit a human motion. While a similar formulation was firstly used in [25], it was applied to a significantly different task of predicting future motions from past frames. Similarly, [35] predicts missing poses in a sequence where only some of pose frames are given. However, we notice that without introducing a structured bias into the training of these methods, they fail at generating realistic poses even if we provide several trajectories. Moreover, they are deterministic by design and cannot generate multiple, diverse motions.

Since we cast our problem as structured pose completion, we leverage recent advancements in stochastic tensor completion [75] and show the application of that paradigm on a novel motion generation model which we call RAJEVAE. Thanks to our formulation, we achieve a desirable property that the accuracy of generated motions increases when more information is provided at the input. At the same time, our model can predict future poses even if no trajectory is given. In industrial applications, our method can generate full-body animations for automatically-tracked joints while naturally handling missing information if some of the joints are not seen by the model. RAJEVAE outperforms trajectory-based baselines and methods based on several past full body frames in terms of accuracy. We additionally show that our formulation can be adapted to existing methods targeting the defined task, thus improving their results.

We summarize our contributions as follows:

• a simple and general training paradigm that enables controllable generation of future poses from a variable number of input information pieces,

• TRAJEVAE — the first generative model that predicts diverse poses from any number of input trajectories,

• empirical study showing that our formulation can be successfully applied to existing methods for generating motion from a single trajectory to improve their results and enable them to use multiple trajectories.

2. Related Works

Deterministic motion prediction In recent years, multiple methods were proposed for predicting a single future motion based on a corresponding past sequence of poses [2,3,10,12,18,20–22,40,41,45–47,51,53,60] or video frames with missing poses [16, 30, 33, 47, 52, 72, 73]. Cai et al. [10] and Aksan et al. [2] use a transformer-like architecture to achieve this goal. Mao et al. [45, 46] extend the pose representation by performing Discrete Cosine Transform (DCT) [1] on joint coordinates. They additionally applied Graph Convolutional Networks (GCN) to incorporate the spatial information relationships between joints. Lebailly et al. [39] adapted inception modules [59] to handle different temporal resolutions of the data. Kaufmann et al. [35] uses a U-Net architecture to complete missing poses in the input tensor. Similar to our approach, Ruiz et al. [25] treats the motion prediction as a pose completion problem. However, these consider only randomly incomplete data while we train our method explicitly to leverage a variable number of available joint trajectories. While being successful, these methods are limited to predicting a single future pose sequence.

Stochastic motion prediction To model the distribution of possible motions, recent works [4, 23, 25, 51, 35, 38, 43, 63, 67, 70] leverage advances for generative modeling and build upon models such as Generative Adversarial Networks (GANs) [19], Variational Autoencoders (VAEs) [37] or normalizing flows [54]. These models enable sampling multiple future pose sequences and to accomplish this, they are often built as conditional models (CGANs [19] and CVAEs [37]). Barsoum et al. [6] use Wasserstein GAN [5] in a sequence-to-sequence framework. To make poses more realistic, they regularize bone lengths and deviations between poses in consecutive frames. Walker et al. [63] apply VAE for the same goal and uses predicted poses to generate structurally consistent images. Zhang et al. [75] consider the motion generation given an action label to performed by the generated character. The authors also consider a transformer architecture to be suitable for time sequence modeling. While having high generation accuracy, these methods do not provide fine-grained control over generated outputs.
Stochastic prediction from trajectories  Existing methods [11, 15, 24, 26, 51–53, 57, 71] methods handle a single trajectory often represented as a target pelvis coordinate projected onto the floor’s plane, or as a tuple of velocities for each axis. Pavllo et al. [53] encode a control signal in the form of the desired trajectory. Hunter et al. [24] input to the model a t control signal and past control signals to predict t + 1 pose. Holden et al. [26] split the motion into phases and model them with a phase-conditioned neural network. Yet these approaches are designed for a single trajectory and cannot generate motions like jumping jacks or hand waving where multiple trajectories need to be specified.

Image completion  As our method draws inspiration for image completion literature, we briefly summarize recent advanced in that field.

Image inpainting [7] is a well-known problem in computer vision. We posit that several of the already proposed approaches [9, 44, 49, 66, 68] can be successfully applied for pose completion where only a part of the pose is given. This way, we can leverage principles of these methods to improve general results. Liu et al. [44] defines a partial convolution where the image is convolved only over pixels that are available in the input. Yu et al. [68] incorporates generative adversarial networks [19] and an attention mechanism to improve overall results. Zheng et al. [75] define a probabilistic model in the VAE framework that allowed the authors to generate diverse and realistic image completions.

3. Method

We introduce a novel paradigm of trajectory representation that enables fine-grained motion control with an arbitrary number of joint trajectories. We show that generating full-body poses from trajectories can be treated as a pose completion problem. Then, we introduce TRAIEVAE that builds on the paradigm and allows us to sample multiple, diverse poses which follow the conditioning trajectories. Throughout the whole paper, we parametrize trajectories and poses in 3D global coordinates. We show the overview of our model in Fig. 2.

3.1. Handling multiple trajectories

We formulate the problem of predicting poses that follow a particular trajectory as a pose completion problem. We denote a trajectory \( Y = \{y_1, \ldots, y_T\} \) of length \( T \) as vectors with \( k \leq J \) known joint positions from a corresponding pose sequence \( X = \{x_1, \ldots, x_T\} \), where \( x_t, y_t \in \mathbb{R}^J \). The trajectories of the unknown \( J - k \) joints in \( Y \) are set to 0. The goal of the pose completion task is to predict \( X \) given \( Y \).

To mimic real-life scenarios, at training-time we randomly mask-out some of the joint the input pose sequence. In this way, we reproduct the typical use cases such as occlusions or omissions of the animation artist. At each time step, we sample a matrix \( M \in \{0, 1\}^{T \times 3J} \) that masks the same joints across \( T \) time steps. Therefore, trajectories are obtained as \( Y = X \odot M \) where \( \odot \) is the element-wise multiplication. We motivate the introduced paradigm as follows. Firstly, masking the poses in a principled, structured way introduces a structural bias into the model. The bias aids the model in learning particular distribution of poses. Such a model can outperform previous approaches such as [35] by a significant margin even for a single trajectory. Secondly, thanks to that bias and in contrast to all related works that are limited to a single trajectory, our method allows the user to select how many trajectories are supplied to the network.

As we show in the experiments, this paradigm enables a neural network model to handle a varying number of input trajectories.

3.2. Pose completion with a neural network

To show the applicability of the introduced framework, we design TRAIEVAE — a Conditional Variational Autoencoder (CVAE) [37] with a transformer-like architecture [62, 64], and a learnable prior distribution. The transformer architecture allows us to generate a sequence of poses in
parallel while the learnable prior increases the sampling diversity and plausibility. Our model is an autoencoder with two encoders \( E_{\text{pose}}, E_{\text{traj}} \) and a single decoder \( D \), where most of the blocks are shared. \( E_{\text{pose}} \) produces parameters of the posterior distribution \( q_{\text{pose}} \) from the input poses \( \mathbf{X} \). These parameters are optimized to match the trajectory prior distribution \( p_{\text{traj}} \) parametrized by \( E_{\text{traj}} \). Since we train the model to match distributions \( q_{\text{pose}} \) and \( p_{\text{traj}} \), the decoder \( D \) produces results that are similar for both \( E_{\text{traj}} \) and \( E_{\text{pose}} \). During inference, we do not have access to the ground truth of a complete pose sequence and hence we drop \( E_{\text{pose}} \) that is responsible for encoding poses during training.

**Encoding poses and trajectories** To take advantage of the similarity between representations of poses and trajectories, parameters of \( E_{\text{traj}} \) and \( E_{\text{pose}} \) are shared unless stated otherwise. We firstly encode the 3D coordinates of \( \{ \mathbf{y}_t \}_{t=1}^T \) and \( \{ \mathbf{x}_t \}_{t=1}^T \) with \( E_{\text{traj}} \) or \( E_{\text{pose}} \) to get \( \{ \mathbf{h}_t \}_{t=1}^T \) and \( \{ \mathbf{h}_t \}_{t=1}^T \) respectively. We concatenate them with the initial pose representation \( \mathbf{h}_0 \) obtained from a separate neural network. \( \mathbf{H} = \{ [\mathbf{h}_t; \mathbf{h}_0] \}_{t=1}^T \) for trajectories and \( \mathbf{H} = \{ [\mathbf{h}_t; \mathbf{h}_0] \}_{t=1}^T \) for poses are passed to two self-attention layers [62]. Then, we apply the Discrete Cosine Transform (DCT) for each feature in all vectors independently, and obtain vectors \( \text{DCT}(\mathbf{H}) \in \mathbb{R}^{T \times (|\mathbf{h}|+|\mathbf{h}_0|)} \), \( \text{DCT}(\mathbf{H}) \in \mathbb{R}^{T \times (|\mathbf{h}|+|\mathbf{h}_0|)} \) in the frequency domain. As shown in [74], most the variability of in the pose distribution concentrates in early components of DCT. Hence, sampling from the normal distribution in frequency domain increases diversity of generated poses.

Up to this point, all parameters for processing trajectories \( \mathbf{Y} \) and poses \( \mathbf{X} \) are shared to use the fact that trajectories represent masked future poses. We then split the pipeline into two unshared parts — one for trajectories and one for poses — that are composed of transformer-like encoders that facilitate information sharing between the latent codes. The final multilayer perceptrons produce parameters \( \{ \langle \hat{\mu}_t \rangle_{t=1}^T, \{ \hat{\sigma}_t \}_{t=1}^T \} \) of a normal distribution for trajectories, and \( \{ \langle \tilde{\mu}_t \rangle_{t=1}^T, \{ \tilde{\sigma}_t \}_{t=1}^T \} \) for poses.

**Learnable prior** Constraining the latent space to a standard normal distribution \( \mathcal{N}(0, I) \) as in VAEs is too restrictive and impedes the diversity of generated samples significantly. To overcome the problem and to provide a more flexible distribution, we make the prior learnable [13, 75] and define it as \( p_{\text{traj}}(\mathbf{z}_t|\mathbf{y}_1, \ldots, \mathbf{y}_T) \). During training, we match the posterior distribution \( q_{\text{pose}}(\mathbf{z}_t|\mathbf{x}_1, \ldots, \mathbf{x}_T) \) by optimizing the Kullback-Leibler divergence:

\[
-\text{KL}(q_{\text{pose}}(\mathbf{z}_t|\mathbf{x}_1, \ldots, \mathbf{x}_T)||p_{\text{traj}}(\mathbf{z}_t|\mathbf{y}_1, \ldots, \mathbf{y}_T)),
\]

where \( \mathbf{z}_t \sim \mathcal{N}(\mu_t, \sigma_t) \) and \( \mathbf{z}_t \sim \mathcal{N}(\tilde{\mu}_t, \tilde{\sigma}_t) \) are samples from the prior and posterior distributions respectively.

**Decoding poses** We transform latent vectors of the poses \( \{ \mathbf{z}_t \}_{t=1}^T \) during training and trajectory latent vectors \( \{ \tilde{\mathbf{z}}_t \}_{t=1}^T \) during inference into the original time domain \( \{ \mathbf{w}_t \}_{t=1}^T \) with Inverse Discrete Cosine Transform (IDCT) [1]. We additionally encode the initial pose with an MLP to obtain \( \mathbf{w}_0 \) as we found it improves overall results. A set of concatenated vectors \( \{ [\mathbf{w}_t; \mathbf{w}_0] \}_{t=1}^T \) is decoded with a self-attention decoder. The final fully connected layers predict offsets \( \mathbf{y}_t \) of the reconstructed pose \( \tilde{x}_{t-1} \) from the time step \( t-1 \). Finally, the reconstructed pose \( \tilde{x}_t \) in the time step \( t \) is obtained as:

\[
\tilde{x}_t = \sum_{\tau=1}^{t} \tilde{x}_{\tau-1} + \mathbf{o}_\tau, \quad \tilde{x}_0 = x_0
\]

where \( x_0 \) is the initial pose.

Our approach decodes offsets of joints in step \( t \) with respect to the pose \( t-1 \) without the need to access that pose. Therefore, the offsets can be predicted in parallel, and the final poses are obtained by a simple aggregation of offsets and adding them to the initial pose \( x_0 \). This approach contrasts with the current notion of applying fully autoregressive decoders [24, 70, 74] which suffer from slow inference.

**Training** We train our TRAJECTOR to accurately reconstruct poses, while maintaining the posterior distribution close to the prior. We achieve this by optimizing the following objective [37]:

\[
\mathcal{L} = \mathcal{L}_{\text{MSE}} + \mathcal{L}_{\text{KL}},
\]

where \( \mathcal{L}_{\text{MSE}} \) is the reconstruction term expressed as mean squared error:

\[
\mathcal{L}_{\text{MSE}} = \sum_{t=1}^{T} ||\tilde{x}_t - x_t||^2_2,
\]

and \( \mathcal{L}_{\text{KL}} \) keeps the posterior distribution close to the learnable prior by minimizing Kullback-Leibler divergence:

\[
\mathcal{L}_{\text{KL}} = -\sum_{t=1}^{T} \beta \text{KL} (q_{\text{pose}}(\mathbf{z}_t|\mathbf{X}, \mathbf{Y})||p_{\text{traj}}(\mathbf{z}_t|\mathbf{Y})).
\]

**Masking future poses and data augmentation** Providing target poses \( \{ \mathbf{x}_t \}_{t=1}^T \) during training directly leads to overfitting and makes the network unable to match the posterior with the prior. This further degrades the quality of reconstructed poses during inference. To overcome the problem, we mask input poses \( \mathbf{X} \) with the inverse mask \( \mathbf{M} \), that was used to obtain trajectories, as \( \mathbf{X} \odot (1 - \mathbf{M}) \). This way, the pose encoder \( E_{\text{pose}} \) is forced to leverage the information from the prior distribution of trajectories which are structurally complementary to masked future poses.

4. **Experiments**

We evaluate TRAJECTOR in two scenarios. Firstly, we evaluate the performance of our method and of several baselines when we progressively add conditioning trajectories.
in the input. Secondly, we compare our method with recent methods for a stochastic human generation. We also perform an ablation study to validate our design decisions.

Datasets All our experiments are based on the Human3.6m dataset [29]. It consists of 3.6 million video frames of 11 subjects performing 15 actions. We follow the evaluation protocol used in [70] and hence we use 17-joint poses. The training was done on subjects S1, S5, S6, S7, S8 while subjects S9 and S11 are left for the testing. We test the baselines and our method by predicting 2 seconds of future motion.

We represent human poses and trajectories as a set of joints parametrized as global coordinates. We also normalize each sequence such that the pelvis of the initial pose is located at the (0, 0, 0) coordinate.

Baselines To show the generality of our approach, we define baselines that, thanks to our paradigm, enable generation of high quality human motions for a variable number of trajectories. We compare TrajEVAE with two approaches that are the most similar to ours and also use the principle of recovering the missing joints: MotionGAN [25] and Motion Infilling [35]. We show that using both of them out-of-the-box is not sufficient to produce high quality pose sequences from trajectories. Note also that these models are designed to be deterministic and cannot produce diverse motions.

We also compare to a basic CVAE-RNN based on [70] and an adapted version of MoGlow [24]. MoGlow in its original version supports only walking, running, and stepping motions. The conditioning signal used by the authors is expressed in terms of axis velocities for the pelvis joint. In its basic form, MoGlow can handle only a single trajectory to predict the motion. We provide additional implementation details of these baselines in the supplementary material.

In the second experiment, we follow the evaluation protocol of DLow [70] and use its baselines for the Human3.6M [29] dataset. In contrast to DLow however, our method requires only a single past frame (the initial pose) while the evaluation used by the authors assumed 25 past frames for Human3.6m and it does not allow to control predicted future motions.

Metrics We evaluate the methods using the diversity and accuracy metrics defined in [70]. Average Pairwise Distance (APD) describes the diversity of a set of size $K$ of motions sampled given the same input trajectory. It is expressed as the average $L_2$ distance between all pairs of generated motions.$\frac{1}{K(K-1)} \sum_{i=1}^{K} \sum_{j \neq i} ||\hat{x}_i - \hat{x}_j||_2$. Average Displacement Error (ADE) measures the accuracy of the reconstructed motion and calculates the average $L_2$ distance across all time steps between the ground truth motion and the motion from a generated set of $K$ motions that is the closest to the ground truth $\frac{1}{T} \min_{x \in X} \sum_{t=1}^{T} ||\hat{x}_t - x_t||_2$. Final Displacement Error (FDE) calculates the $L_2$ distance between the pose in the last time step of ground truth motion and the motion from a generated set of $K$ motions that is the closest to the ground truth $\min_{x \in X} ||\hat{x}_T - x_T||_2$. Multi-Modal ADE (MMADE) and Multi-Modal FDE (MMFDE) calculates an average of ADE and FDE respectively between a predicted motion and all samples in a cluster of motion sequences. We group these motions where the $L_2$ distance between their initial poses differs by less than $\epsilon$.

Implementation details At the training time, we obtain masks $M \supset m \in \{0, 1\}^{3J}$ by sampling from the Bernoulli distribution $B(p_m)$ with a probability $p_m$. Then, we replicate the $m$ vector $T$ times to create the structured mask $M \in \{0, 1\}^{T \times 3J}$. We set $p_m = 0.85$ so that the network sees 3 – 4 trajectories on average in the input. We motivate that number to be a sensible trade-off between the accuracy and effort of defining trajectories when using TrajEVAE in practice.

We train TrajEVAE and the corresponding baselines with the Adam optimizer [36] with learning rate set to 0.0001 and multiplied by 0.25 every 80,000 training steps. We set $\beta = 0.01$ in the KL term, the batch size = 64 and we train models for 240,000 steps.

4.1. Qualitative results

Reconstructed sequences To visually examine the proposed method, we generate a set of poses while changing the number of input trajectories. Fig. 3 shows individual frames from selected animation sequences (refer to the supplementary material to see full video clips). When more trajectories are provided, the generated sequence resembles the ground truth more closely. However, even if no trajectory is provided, TrajEVAE generates plausible poses. We achieve this by providing the initial pose to the model during the decoding phase. We note that the initial pose heavily biases the model due to the nature of the dataset.

Diversity vs. number of input trajectories Since TrajEVAE allows us to sample latent DCT components from a learnable prior distribution, we can generate multiple, diverse samples for the same set of conditioning trajectories. We show the last frames of such generated samples in Fig. 4. When no trajectories are present, the method generates the most diverse outputs, while retaining the plausibility of poses. As we noticed by examining the generated sequences, such poses can represent waving, bending, or dancing-like motions. When four trajectories are given, generated poses converge towards the ground truth.

Due to the MSE term in Eq. (3), the model is not forced to exactly reproduce the trajectories, and therefore the results plateau when we provide more than ten trajectories. Applying $L_1$ loss instead of $L_2$ mitigates that issue but significantly impedes the diversity of generated samples.

Generalization Finally, we empirically show that, in contrast to other works on controllable human motion gener-
By using TRAJEVAE, we can provide more trajectories in the input to create a realistic pose that follows a particular path. Joints that are not described by a trajectory are completed by our method. Here we show three sequences of motions generated by our method. The joints that have a corresponding input trajectory are depicted as brown spheres. The top row shows the ground truth sequence. Rows below show generated sequences when more trajectories are given, in the order: right foot, left foot, right and left hands. Labels refer to classes in the Human3.6m dataset [29].

TRAJEVAE allows us to use a variable number of trajectories and sample multiple diverse pose sequences. We show in each row input trajectories, end poses for 5 sampled sequences, and the end pose for a sequence decoded from trajectory means \( \{\hat{\mu}\}_t \). The joints that have a specified trajectory are colored in brown.

TRAJEVAE is not limited to only walking, running, or standing, as are other related methods, and can be applied to any motion type. Each row represents a generated sequence for a different motion class given specific trajectories. The joints that have a specified trajectory are colored in brown.

TRAJEVAE can be applied to any set of motions, e.g., dancing, sitting, waving, and others. As we present in Fig. 5, TRAJEVAE generalizes to a variety of different sets of motions that were not handled by previous methods. We additionally show in the supplementary, that given the same trajectory but a different initial pose, our method still generates poses that follow the provided trajectory. It confirms that our approach leverages both the initial pose and trajectories, and can be used to generate animations beyond the ones found in the dataset.
Controlling future motion prediction. We evaluate the quality of generated samples while changing the number of input trajectories. We first compute the metrics with no provided trajectories and then progressively increase their number by adding the following: right foot, left foot, right hand, and left hand. This order was motivated by the variance of coordinates of joints they correspond to, i.e., the joint with highest variance in the dataset was added first. In each case, we sample $K = 50$ poses to calculate metrics. Results are summarized in Table 1. As expected, adding more trajectories decreases the diversity (APD) of samples since the pose is restricted to follow a particular path. At the same time, the accuracy (ADE) of generated samples improves. Our TRAJEVAE obtains the best results in terms of the reconstruction quality in comparison to other methods. The higher diversity of CVAE-RNN is caused by its structure — in contrast to TRAJEVAE, CVAE-RNN encodes the whole sequence into a single latent code instead of multiple components. Therefore, two samples from the prior may have a significantly different structure in the output. However, such an architecture suffers from pose averaging where most of the pose frames are the same in a generated sequence [16,42] which leads to the inferior accuracy of the reconstructed motions. We hypothesize that Motion Infilling [35] obtains high error on the reconstruction due to two issues: it is trained with the $L_1$ reconstruction loss (which also worked detrimentally for TRAJEVAE) and the simple masking scheme where each coordinate for all joints across the sequence is randomly masked out.

In Fig. 6, we show that including more trajectories in the input improves the reconstruction quality. Notice that including too many trajectories plateaus the quality. We conclude that it is caused by the randomness in the latent space, and hence the network is unable to properly encode all the input trajectories to reconstruct poses accurately. Interestingly, the diversity (APD) is the lowest for ten trajectories and slightly increases when we add more trajectories. We argue that this phenomenon may be caused by the order we use when adding trajectories. Due to the low influence of the last seven trajectories on the sequence, they do not affect the quality.

**Comparison with other methods for future motion generation.** We also compare TRAJEVAE with methods that generate future poses from a set of past frames. We evaluate the methods in two scenarios: when sampling $K = 50$ different poses, and when sampling only a single pose. We
The transformer-like architecture already obtains remarkable results. However, these can be improved by using a learnable prior. While DCT reduces the accuracy, we found it to be a necessary component to obtain diverse samples.

We also identified it beneficial to apply additional regularization technique by masking future poses with the mask $1 - \mathbf{M}$. By masking the poses, the network has to learn to encode more information from poses and stops to rely entirely on the prior distribution.

5. Conclusions and limitations

We introduced the notion of trajectory-conditioned pose generation as a pose completion problem. It allowed us to define TraJEVAE — a method for controllable and stochastic human animation generation. We showed that the paradigm of structured dropping of joints during training, creates a model that can generate realistic poses that follow an arbitrary number of trajectories. Obtained results show the applicability of our method in designing realistic human animations. While our approach trivially generalizes to other data representations, applying it to full-body parametric models, such as SMPL [50, 74], is of high importance.

We identify two limitations of our approach. Firstly, generated poses do not follow the trajectories exactly. While we could resort to a cGAN [19] model as a possible solution for its unprecedented quality of generated samples [34], the application of GANs to structured time series data is still a challenge.

Secondly, when the initial pose is ambiguous about what action it represents, TraJEVAE tends to stretch bones when we input only a single or none trajectories. Applying exponential maps [53] constrains bone lengths but this would limit our method to only work with skeleton structures that have a clearly defined kinematic chain.

6. Ethical concerns

We do not identify immediate abuses of our approach in real world applications. TraJEVAE can be used to re-animate characters, however their skeletons still need to be manually defined. We regard the recent progress in neural radiance field [48] methods and their applications for human reposing [58] as a potential ethical concern.

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Table 3. Quantitative results for the Human3.6M dataset when a single future pose sequence ($K = 1$) is generated. For our method, we assume different scenarios when $k = \{0, 1, 2, 3, 4\}$ trajectories are provided. In this experiment, TraJEVAE decodes predicted means $\hat{\mu}_t$ of trajectories. Best results are in bold.

| Method          | APD ↑ | ADE ↓ | FDE ↓ | MMADE ↓ | MMFDE ↓ |
|-----------------|-------|-------|-------|---------|---------|
| TraJEVAE ($k = 0$) | 1.126 | 1.652 | 2.229 | 0.661   | 0.749   |
| TraJEVAE ($k = 1$) | 1.126 | 1.652 | 2.229 | 0.661   | 0.749   |
| TraJEVAE ($k = 2$) | 1.126 | 1.652 | 2.229 | 0.661   | 0.749   |
| TraJEVAE ($k = 3$) | 1.126 | 1.652 | 2.229 | 0.661   | 0.749   |
| TraJEVAE ($k = 4$) | 1.126 | 1.652 | 2.229 | 0.661   | 0.749   |

Table 4. Influence of design decisions on obtained results for a single given trajectory (a right foot). Best results are in bold. For the ablation study with more trajectories, refer to the supplementary.

| Method                      | APD ↑ | ADE ↓ | FDE ↓ | MMADE ↓ | MMFDE ↓ |
|-----------------------------|-------|-------|-------|---------|---------|
| Base                        | 1.220 | 0.481 | 0.695 | 0.640   | 0.594   |
| + Learnable prior           | 1.749 | 0.448 | 0.622 | 0.618   | 0.753   |
| + DCT                       | 7.014 | 0.487 | 0.637 | 0.602   | 0.703   |
| + Masked future poses       | 6.803 | 0.472 | 0.623 | 0.594   | 0.695   |

Ablation study Finally, we perform an ablation study of our design decisions: making the prior distribution learnable, using the Discrete Cosine Transform in the latent space, and masking future poses with the mask $1 - \mathbf{M}$. We show obtained results in Tab. 4.

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TRAJEVAE: Controllable Human Motion Generation from Trajectories

Supplementary Material

A. Adapting MoGlow

As mentioned in the main text, MoGlow [24] conditions predicted poses on a control signal. This signal represents relative and rotational velocities on the ground plane. Moreover, the authors use the exponential map representation but at the same time claim that MoGlow can be used for any other well-known skeleton representation. We identify the following changes to the original implementation that enabled us to use MoGlow in our framework:

1. We change the exponential map representation to the 3D coordinates of \( J \) joints.
2. We replace the control signal represented as velocities into trajectories defined as poses with some of the joints set to 0. This increases the input signal’s dimensionality from \( 3JT \) to \( 3JT \) for \( T \) time steps. However, it has a negligible effect on the performance.
3. We also removed regularization techniques such as gradient norm clipping and gradient value clipping and disabled data normalization. These techniques deteriorate the learning, and the network does not converge in our scenario.
4. For consistency with other methods, we use the Adam optimizer [36] with the same learning rate regime.

We left the rest of the implementation unchanged.

B. Implementation details

TRAJEVAE MLPs applied in the input and in the CVAE’s bottleneck output latent codes of size 256. Therefore, vectors \( \mathbf{H} \) and \( \mathbf{H} \) processed by self-attention layers have a dimensionality 512. The initial layer in the decoder \( D \) processes \( \{ [\mathbf{w}_1; \mathbf{w}_0] \} \) and is defined as a function: \( f : \mathbb{R}^{768} \rightarrow \mathbb{R}^{512} \). The final layers outputs vectors of size 3J.

All MLPs responsible for encoding poses and trajectories mentioned in the main text consists of the following structure: \( \text{Linear} \rightarrow \text{Layer Normalization} \rightarrow \text{Leaky ReLU}(\alpha=0.1) \rightarrow \text{Linear} \rightarrow \text{Layer Normalization} \rightarrow \text{Leaky ReLU}(\alpha=0.1) \), where \( \alpha \) is a scale of the negative slope of the function. The initial MLP in the decoder \( D \) has the structure \( \text{Linear} \rightarrow \text{Layer Normalization} \rightarrow \text{Leaky ReLU}(\alpha=0.1) \rightarrow \text{Linear} \).

The CVAE baseline operate with the same dimensionalities as TRAJEVAE. We implement them in the same way as defined in [70]. The model uses GRU network to encode the temporal data. The recurrent decoder also receives a coordinate of the trajectory \( y_t \) in the time step \( t \).

We additionally apply dropout = 0.1 to self-attention layers as described in [62].

C. Same pose, different trajectories

We perform an additional experiment that confirms the generality of our approach. We show results for a scenario when we use different trajectories for the same initial pose. As expected, the generated poses follow different trajectories even though such combinations do not occur in the dataset.

Preparing the data To maintain plausibility that a particular initial pose is physically capable of following a conditioning trajectory, we pair each initial pose \( x_0 \) in the dataset with all trajectories where the distance between \( x_0 \) and coordinates of the trajectory in a time step \( t = 0 \) is below \( \epsilon_0 = 0.01. \) Since obtaining the ground truth sequence \( X \) in such a case is not possible, we assume that the sequence \( X \) corresponding to a given trajectory is a sufficient approximation of the expected sequence. We evaluate TRAJEVAE as previously using APD, ADE, FDE. We omit MMADE and MMFDE for its exponential computational complexity that this scenario creates.

Results We present results in Tab. 5. Even though these trajectories do not come from the same sequence as the initial poses, TRAJEVAE generates a sequence that follows the trajectory. The decrease in accuracy (ADE) between \( k = 2 \) and \( k = 3 \) is caused by adding a trajectory that corresponds to the right hand, while \( k < 2 \) we add only trajectories regarding feet. While feet commonly behave similarly throughout the animation, hands have a significantly different motion from other joints.

The value \( k = 0 \) corresponds to no trajectories, and therefore we omit it in the Tab. 5. Refer to supplementary files to find animations generated for initial poses with different trajectories.

D. Extended ablation study

In experiments, we provide results for an ablation study when only a trajectory for the right foot is provided. We additionally show in Tab. 6 results when we input no trajectories, or progressively add trajectories of the right foot, left foot, right hand, and left hand. The extended results show that our design decisions consistently affect scenarios when we vary the number of the input trajectories.
Table 5. Quantitative results for the Human3.6M dataset when \( K = 50 \) samples are generated for the scenario where we use different trajectories from the whole dataset for the same initial pose. We assume different situations that \( k = \{1, 2, 3, 4\} \) trajectories are provided.

| \( k \) | APD  | ADE  | FDE  |
|-------|------|------|------|
| 1     | 5.373| 0.370| 0.476|
| 2     | 5.400| 0.362| 0.472|
| 3     | 5.096| 0.375| 0.491|
| 4     | 4.175| 0.332| 0.433|

Table 6. Influence of design decisions on obtained results for \( k = \{0, 1, 2, 3, 4\} \) trajectories. These trajectories refer to scenarios when we use no trajectories and then add progressively trajectories for the right foot, left foot, right hand, and left hand. The best results are in bold.