Distributional Autoencoders Know the Score

(NeurIPS 2025)

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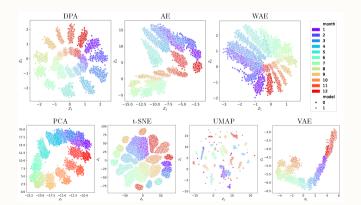


"Nonlinear PCA that learns the data score."

BACKGROUND & MOTIVATION

Unsupervised learning method card

- 1) PCA: linear, ordered components, mean reconstructions only.
- 2) AE: non-linear encodings, no ordering, mean tendency for reconstruction.
- 3) **DPA:** non-linear, ordered components **and** distribution-faithful reconstructions.



INTRODUCTION TO DPA

The goal of the Distributional Principal Autoencoder (DPA) [Shen and Meinshausen, 2024] is distributionally-faithful reconstruction of all data (X) mapped to the same value by the encoder (e):

$$P_{e,x}^* = \operatorname{Law}(X \mid e(X) = e(x)).$$

The encoder-decoder optimization objective is based on the *energy score*:

$$(e^*, d^*) \in \arg\min_{e, d} \ \sum_{k=0}^p \mathbb{E}_X \Big[\mathbb{E}_{Y \sim P_{d, e_{1:k}(X)}} [\|X - Y\|^\beta] \Big] - \tfrac{1}{2} \mathbb{E}_X \Big[\mathbb{E}_{Y, Y'} \underset{\sim}{\text{iid}}_{P_{d, e_{1:k}(X)}} [\|Y - Y'\|^\beta] \Big] \Big]$$

where $P_{d,\,e_{1:k}(X)}$ is the reconstructed distribution using only the first k components of e.

First main result: geometry aligns exactly with the data score (eta=2)

Theorem 1

For $\beta=2$ and under relatively mild assumptions we have, for almost every sample $X\sim P_{\mathrm{data}}$ and encoder level set $\mathcal{L}_{e^*(X)}$, the following balance equation for almost every $y\in\mathcal{L}_{e^*(X)}$:

$$\boxed{ \frac{2 \big(y - c(X) \big)}{\frac{V(X)}{Z(X)} - \| y - c(X) \|^2} \, D_{e^*}^{\top}(y) \; = \; \nabla_y \log P_{\mathrm{data}}(y) \, D_{e^*}^{\top}(y), }$$

where $D_{e^*}(y)$ is the encoder Jacobian at y, whenever the following quantities: the **level-set center-of-mass**:

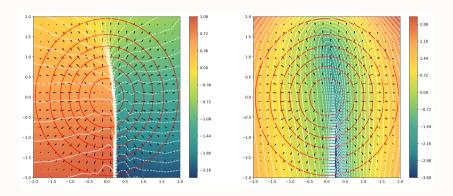
$$c(X) = \frac{1}{Z(X)} \int y \, P_{data}(y) \, \delta(e(y) - e(X)) \, dy, \label{eq:constraint}$$

and the level-set variance:

$$V(X) = \int \|y-c(X)\|^2 \, P_{data}(y) \, \delta(e(y)-e(X)) \, dy$$

are finite, and the **level-set mass** $Z(X) = \int P_{data}(z) \ \delta(e(z) - e(X)) \ dz > 0.$

2D GAUSSIAN INTUITION: TANGENTIAL VS. NORMAL



Rotational symmetry:

One component is tangential (both sides ≈ 0).

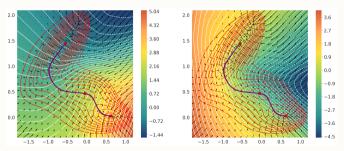
The other is normal (level sets orthogonal to the score); together they recover polar coordinates.

BOLTZMANN DATA: FORCES & MINIMUM FREE-ENERGY PATH (MFEP)

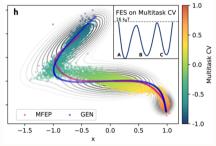
For Boltzmann distributed data, we recover the normal force field:

$$\vec{F}(y) \; D_{e^*}^\top \; = \; 2 \; k_B T \, \frac{y - c(X)}{\frac{V(X)}{Z(X)} - \|y - c(X)\|^2} \; D_{e^*}^\top(y).$$

Encoding of molecular simulation trajectories thus reveals the Minimum free energy path (MFEP):



DPA components trace the MFEP in a single fit.



Existing methods require iteration/supervision [Bonati et al., 2023].

SECOND MAIN RESULT: ENCODING DIMENSIONS BEYOND THE MANIFOLD ARE UNINFORMATIVE

Theorem (Extra dimensions are completely uninformative)

For a manifold that can be approximated in K' dimensions by the encoder, the dimensions $(K'+1,\cdots,p)$ of such optimal encoder obey:

$$P_{d^*,e^*_{1:k}(X)} = P_{d^*,e^*_{1:K'}(X)}, \ \ \text{for} \ k \in [K'+1,\dots,p].$$

Furthermore, these dimensions are conditionally independent of the data X, given the relevant components $(e_1^*,\cdots,e_{K'}^*)$,

$$X \perp \!\!\! \perp e^*_{K'+i}(X) \mid e^*_{1:K'}(X), \qquad \forall i \in [1, \dots, p-K'].$$

or equivalently, they carry no additional information about the data distribution:

$$I\left(X;e^*_{K'+i}(X)\mid e^*_{1:K'}(X)\right)=0, \qquad \forall i\in[1,\dots,p-K'],$$

So WHAT?

Distribution approximation / reconstruction and dimensionality reduction / disentanglement almost always present a trade-off. For example, this is what the β in β -VAE does:

$$\underset{\theta,\phi}{\operatorname{arg\,min}} \ \mathbb{E}_{p_{\mathrm{data}}(x)} \Big[\underbrace{\mathbb{E}_{q_{\phi}(z|x)} [-\log p_{\theta}(x\mid z)]}_{\text{reconstruction}} \ + \underbrace{\beta}_{\text{disentanglement}} \underbrace{\operatorname{KL} \Big(q_{\phi}(z\mid x) \, \big\| \, \prod_{j} p(z_{j}) \Big)}_{\text{disentanglement}} \Big]$$



https://arxiv.org/abs/2502.11583

Thank you!

References

Luigi Bonati, Enrico Trizio, Andrea Rizzi, and Michele Parrinello. A unified framework for machine learning collective variables for enhanced sampling simulations: mlcolvar. The Journal of Chemical Physics, 159(1):014801, July 2023. ISSN 0021-9606, 1089-7690. doi: 10.1063/5.0156343. URL https://doi.org/10.1063/5.0156343.

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