Predictive Collective Variable Discovery

with Deep Bayesian Models in Atomistic Systems.

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Predictive Collective Variable Discovery

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Which set of collective variables (CVs) captures resilient and parsimonious features in atomistic systems?



Figure 1: Alanine Dipeptide. Exemplary collective variables.

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Figure 1: Alanine Dipeptide. Exemplary collective variables.

• Relevant coordinates are highly clustered around a set of *lower dimensional collective variables*.

[Banon]

 Vast combinatoric possibilities for *choosing* CVs. [Chakraborty et al. (2018)]

Questions we address

- How to identify good CVs?
- What are good CVs?
- How to quantify "good"?
- Are identified CVs physically interpretable?

Motivation

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- How to quantify "good"?
- Are identified CVs physically interpretable?

Probabilistic predictions

- How to use utilize identified CVs for *predictive* purposes?
- How to quantify predictive *uncertainty*?

Methodologies for identifying collective variables at our disposal:

- 1. Intuition-, path-, state-based coordinates.
- 2. Linear dimensionality reduction methods.
 - Principal component analysis (PCA) [F.R.S. (1901); Hotelling (1933)].
 - Multidimensional scaling (MDS) [Troyer and Cohen].

Methodologies for identifying collective variables at our disposal:

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 - Principal component analysis (PCA) [F.R.S. (1901); Hotelling (1933)].
 - Multidimensional scaling (MDS) [Troyer and Cohen].
- 3. Nonlinear dimensionality reduction methods.
 - Isometric feature map (Isomap) [Tenenbaum et al. (2000)].
 - Sketch map [Ceriotti et al. (2011)] .
 - Diffusion maps (and locally scaled diffusion maps) [Coifman et al. (2005); Nadler et al. (2006)] .
 - Kinetic map [Noé and Clementi (2015)] and Commute maps [Noé et al. (2016)].

Physical intuition does not suffice for identifying good collective variables [Rohrdanz et al. (2013)].

Problem definition - Equilibrium statistical mechanics

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tomistic model
$$p(\mathbf{x}) \propto e^{-\beta U(\mathbf{x})}$$

• $\mathbf{x} \in \mathcal{M}$: atomistic coordinates

A

- U(x): atomistic potential
- Observables: $\mathbb{E}_{p(\mathbf{x})}[a] = \int a(\mathbf{x}) p(\mathbf{x}) d\mathbf{x}$

CV representation

$$z = \mathcal{R}(x), \quad \text{dim}(z) << \text{dim}(x)$$

- z: (reduced) collective variables.
- *R*: operator (mapping onto CV space).

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Introduce $p(\mathbf{z})$ and $p(\mathbf{x}|\mathbf{z})$, thus the joint $p(\mathbf{x}, \mathbf{z})$:

Methodology - Probabilistic generative model





Data $\mathbf{x}^{(i)} \sim p(\mathbf{x})$

Methodology - Probabilistic generative model

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Data $\mathbf{x}^{(i)} \sim p(\mathbf{x})$

Proposed generative model [Jordan (1999)]

adopted from Predictive Coarse-Graining [Schöberl, Zabars, and Koutsourelakis (2017)] [Felsberger and Koutsourelakis, Wed.]

$$\underbrace{p(\mathbf{z})}_{CV.} \xrightarrow{p(\mathbf{x}|\mathbf{z})} \overline{p}(\mathbf{x}) = \int p(\mathbf{x}|\mathbf{z}) \ p(\mathbf{z}) \ d\mathbf{z}$$

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• CVs z as latent generator for atomistic coordinates x.

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- CVs z as latent generator for atomistic coordinates x.
- Probabilistic map from CVs z to the atomistic coordinates x, via p(x|z).

Methodology - Probabilistic generative model

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$$\underbrace{p(\mathbf{z})}_{\text{CV space.}} \xrightarrow{p(\mathbf{x}|\mathbf{z})} \overline{p}(\mathbf{x}) = \int p(\mathbf{x}|\mathbf{z}) \ p(\mathbf{z}) \ d\mathbf{z}$$

- CVs z as latent generator for atomistic coordinates x.
- Probabilistic map from CVs z to the atomistic coordinates x, via p(x|z).
- Posterior distribution p(z|x) = p(x|z)p(z)/p(x)/p(x) represents code/dictionary for mapping atomistic coordinates x into its lower dimensional CV embedding z. (intractable, e.g. use approximate inference thus approximate posterior with q(z|x)).



Methodology - Learning



The marginal log-likelihood of the data set $\mathbf{X} = {\mathbf{x}^{(i)}}_{i=1}^{N}$:

$$\log p(\mathbf{x}^{(i)}, \cdots, \mathbf{x}^{(N)}) = \sum_{i=1}^{N} \log p_{\boldsymbol{\theta}}(\mathbf{x}^{(i)}),$$

with $\log p_{\theta}(x^{(i)})$ for one datapoint $x^{(i)}$:

Lower bound the log-likelihood

[Dempster et al. (1977)]

$$\begin{split} \log p_{\theta}(\mathbf{x}^{(i)}) &= \log \int p_{\theta}(\mathbf{x}^{(i)} | \mathbf{z}^{(i)}) p_{\theta}(\mathbf{z}^{(i)}) \, d\mathbf{z} \\ &= \log \int q_{\phi}(\mathbf{z}^{(i)} | \mathbf{x}^{(i)}) \frac{p_{\theta}(\mathbf{x}^{(i)} | \mathbf{z}^{(i)}) p_{\theta}(\mathbf{z}^{(i)})}{q_{\phi}(\mathbf{z}^{(i)} | \mathbf{x}^{(i)})} \, d\mathbf{z} \\ &\geq \int q_{\phi}(\mathbf{z}^{(i)} | \mathbf{x}^{(i)}) \log \frac{p_{\theta}(\mathbf{x}^{(i)} | \mathbf{z}^{(i)}) p_{\theta}(\mathbf{z}^{(i)})}{q_{\phi}(\mathbf{z}^{(i)} | \mathbf{x}^{(i)})} \, d\mathbf{z} = \mathcal{L}(\theta, \phi; \mathbf{x}^{(i)}) \end{split}$$

The log-likelihood is decomposed into: [Beal and Ghahramani (2006); Kingma

and Welling (2013); Rezende et al. (2014)]

$$\log p_{\theta}(\mathbf{x}^{(i)}) = \underbrace{\mathcal{L}(\theta, \phi; \mathbf{x}^{(i)})}_{\text{variational lower bound, since } D_{KL} > 0} + \underbrace{D_{KL}(q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})||p_{\theta}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)}))}_{\geq 0}$$

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$$\log p_{\theta}(\mathbf{x}^{(i)}) \ge \mathcal{L}(\theta, \phi; \mathbf{x}^{(i)}) = -\underbrace{D_{KL}(q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})||p_{\theta}(\mathbf{z}))}_{\text{Regularize } \phi, \text{ such that } q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})} + \underbrace{\mathbb{E}_{q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})}[\log p_{\theta}(\mathbf{x}^{(i)}|\mathbf{z}^{(i)})]}_{\text{Expected neg. reconstruction error.}}$$

with the approximate posterior $q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})$, e.g. a distribution of the exponential family.

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with the approximate posterior $q_{\phi}(\mathbf{z}^{(i)}|\mathbf{x}^{(i)})$, e.g. a distribution of the exponential family.

Discovery of CVs as approximate Bayesian inference, i.e. identify the code/dictionary $q_{\phi}(z|x)$.

• MLE estimate

 $\max_{\boldsymbol{\phi},\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta},\boldsymbol{\phi};\mathbf{X})$

• MAP estimate

$$\max_{\boldsymbol{\phi},\boldsymbol{\theta}} \mathcal{L}(\boldsymbol{\theta},\boldsymbol{\phi};\boldsymbol{X}) + \underbrace{\log p(\boldsymbol{\theta})}_{log-prior}$$

• MLE estimate

$$\max_{\phi,\theta} \mathcal{L}(\theta,\phi;\mathsf{X})$$

MAP estimate

$$\max_{\phi,\theta} \mathcal{L}(\theta,\phi;\mathsf{X}) + \underbrace{\log p(\theta)}_{log-prior}$$

• Approximate Bayesian posterior of decoding parameters θ , $p(\theta|\mathbf{X})$, with Laplace approximation.



Figure 2: Laplace approximation: $p(\theta|X) \approx \mathcal{N}(\mu, S)$

· $p(\boldsymbol{\theta}|\mathbf{X}) \approx \mathcal{N}(\boldsymbol{\mu}, \mathbf{S})$

$$\cdot \; oldsymbol{\mu} = oldsymbol{ heta}_{ extsf{MAP}}$$

•
$$S^{-1} = -\frac{\partial^2 \mathcal{L}(\theta, \phi; X)}{\partial \theta_k \partial \theta_l} - \frac{\partial^2 \log p(\theta)}{\partial \theta_k \partial \theta_l}$$

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Numerical Illustration - Alanine Dipeptide





- 22 atoms $\rightarrow \text{dim}(\mathbf{x}) = 66$.
- Implicit solvent.
- No pre-processing of data the data: x⁽ⁱ⁾ are the cartesian coordinates of all atoms in the system.



Auto-Encoding Variational Bayes

According [Kingma and Welling (2013)].

$$\bar{p}(\mathbf{x}|\boldsymbol{\theta}) = \int p(\mathbf{x}|\mathbf{z},\boldsymbol{\theta}) p(\mathbf{z}|\boldsymbol{\theta}) d\mathbf{z}$$

Mapping $z \rightarrow x$ (decoder):

$$p(\mathbf{x}|\mathbf{z}, \boldsymbol{\theta}) = \mathcal{N}(\boldsymbol{\mu}(\mathbf{z}; \boldsymbol{\theta}), \ \boldsymbol{\sigma}^2 \boldsymbol{I})$$

With,

- $\cdot \mu(\mathsf{z}; \theta)$ output of fully connected decoding neural network.
- σ^2 independet of z due to [Mattei and Frellsen (2018)].



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With,

- $\cdot \mu(z; \theta)$ output of fully connected decoding neural network.
- σ^2 independet of z due to [Mattei and Frellsen (2018)].
- Proposed model does not pre-assume any physical insight.
- No a priori assumption or data treatment needed.



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Auto-Encoding Variational Bayes

$$\bar{p}(\mathbf{x}|\boldsymbol{\theta}) = \int p(\mathbf{x}|\mathbf{z},\boldsymbol{\theta}) p(\mathbf{z}|\boldsymbol{\theta}) d\mathbf{z}$$

We assume CVs are normal distributed according,

$$p(z) = \mathcal{N}(0, I)$$

- Here: $\dim(z) = 2$.
- Explore assigned meaning given $\dim(z) = 2$.



Auto-Encoding Variational Bayes

The approximate posterior is of the following form,

$$q(\mathbf{z}|\mathbf{x}, \boldsymbol{\phi}) = \mathcal{N}(\boldsymbol{\mu}(\mathbf{x}; \boldsymbol{\phi}), \boldsymbol{\sigma}(\mathbf{x})^2 \boldsymbol{I}).$$

With,

- $\mu(\mathbf{x}; \phi)$ and $\sigma^2(\mathbf{x}; \phi)$ described by a fully connected encoding network.
- · Last layers separate into $\mu(\mathbf{x}; \phi)$ and $\sigma^2(\mathbf{x}; \phi)$.

Numerical Illustration - Model





Figure 3: Schematic draft of the Bayesian variational autoencoder employed [Kingma and Welling (2013); Rezende et al. (2014)].



Training with N = 1500 datapoints, a batch size of M = 64 for 1500 epochs, dim(z) = 2

Figure 5: Lower bound.

Figure 4: CVs assigned to training data (latent representation of training data). Discovered CVs separate conformational modes well.



Prediction for
$$\{z|z_1 = [-4, 4], z_2 = 0\}$$
.

AEVB: Encoded representation of training data: $oldsymbol{\mu}(oldsymbol{x}^{(i)})$





Figure 6: CVs are highly correlated with different conformations.

Figure 7: Mean prediction for x with decoder $p_{\theta}(\mathbf{x}|\mathbf{z})$, given $\{\mathbf{z}|z_1 = [-4, 4], z_2 = 0\}$.



Prediction for $\{z|z_1 = [-4, 4], z_2 = 0\}$.

Figure 6: CVs are highly correlated with different conformations.

Figure 7: Mean prediction for x with decoder $p_{\theta}(\mathbf{x}|\mathbf{z})$, given $\{\mathbf{z}|z_1 = [-4, 4], z_2 = 0\}$.



Identified CVs show high correlation to the known optimal description, the dihedral angles (ϕ, ψ) .



Figure 8: Assessing dihedral angles ϕ given the CVs z.



Figure 9: Assessing dihedral angles ψ given the CVs z.

Numerical Illustration - Physical interpretation?



• $(Z_1, Z_2) \mapsto (\phi, \psi)$



Figure 10: Assessing dihedral angles (ϕ, ψ) given the CVs **z** at $z_2 = 0$.



Figure 11: Assessing dihedral angles (ϕ, ψ) given the CVs **z** at $z_1 = 0$.

Numerical Illustration - Model selection for neural networks

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Motivation

- Network structure highly individual. dim(θ) grows exponentially.
- What are the actually required parameters θ ?
- · Can one search across models?

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Sparsity-enforcing - Hierarchical priors (ARD, [MacKay 1994])

$$p(\boldsymbol{\theta}|\boldsymbol{\tau}) = \prod_{j} p(\boldsymbol{\theta}_{j}|\tau_{j})$$

$$\theta_{j} \sim \mathcal{N}(0, \tau_{j}^{-1})$$

$$\tau_{j} \sim Gamma(a_{0}, b_{0})$$



- Inner EM framework:
 - E-step: Estimate $\langle \tau_j \rangle_{p(\tau_j|\theta_j)} = \frac{a_0+1/2}{b_0+\theta_i^2/2}$
 - M-step: Additive component of the derivative of the log-likelihood: $\frac{\partial}{\partial \tau_{c,j}} = \langle \tau_j \rangle \theta_j$

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Numerical Illustration - Predictive CVs 🥀

- Predictions accounting for uncertainty in θ , with $\theta \sim p(\theta|X)$.
- Metropolis-within-Gibbs sampler [Mattei and Frellsen (2018)] corrects usage of approx. posterior $q_{\phi}(\mathbf{z}|\mathbf{x})$.

Numerical Illustration - Predictive CVs 👫

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10 8 4 2.0 2.1 2.2 2.3 2.4 2.5 2.6 2.7 2.8 Mg/A) meterence MAP 12%-99% quantile

Figure 12: Root-mean-square deviation.

Figure 13: Radius of gyration.



Figure 14: Ramachandran plot

ARD-prior identifies a parsimonious representation: 60% of the θ 's set to zero.

Numerical Illustration - Predictive CVs 👫

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- Metropolis-within-Gibbs sampler [Mattei and Frellsen (2018)] corrects usage of approx. posterior $q_{\phi}(\mathbf{z}|\mathbf{x})$.
- $N = \{526, 1527, 4004\}$



Adequate predictions with a dataset of N = 526. Low data instead of big data.

Summary

Conclusion

- CV discovery as *Bayesian* inference.
- Identified CVs reveal physics based on low data.
- *Physics learned from data*: discovered CVs are attached with physicochemical meaning.
- *Predictive* model obtained for property estimation and uncertainty quantification.

Outlook

- Sparse automatic determination of network structure.
- · VI on network parameters $heta, \phi$; Fully Bayesian approach.
- Utilize identified CVs for enhanced sampling methods (iteratively add data obtained from enhanced sampling).
- Addressing robustness and dependency of network [Mattei and Frellsen (2018)].

Questions?

Thank you!

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