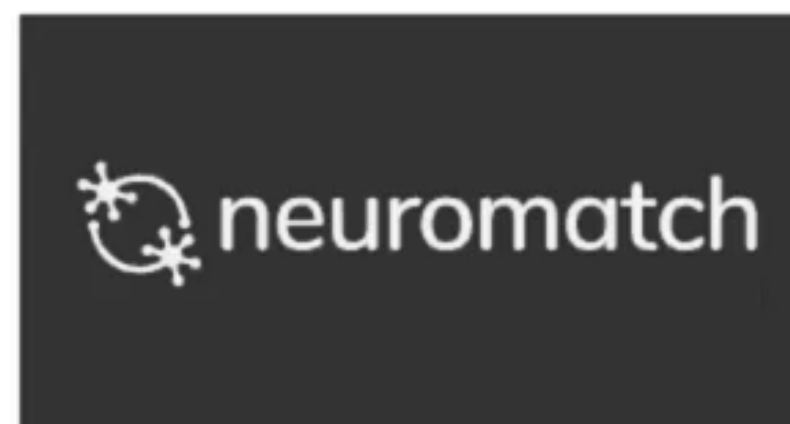




Fast and scalable learning of generative models for chaotic dynamical systems and neural data

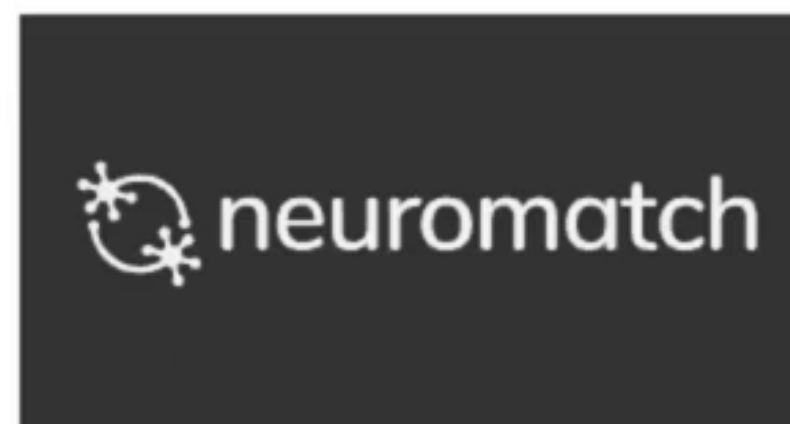
Leonard Bereska, Po-Chen Kuo, Manuel Brenner, Daniel Durstewitz
Central Institute of Mental Health Mannheim, University of Heidelberg
28.10.2020, Neuromatch 3.0





Fast and scalable learning of generative models for chaotic dynamical systems and neural data

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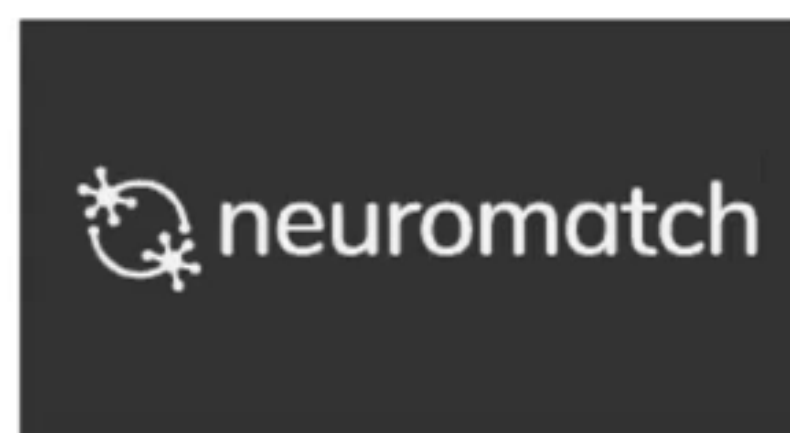




Fast and scalable learning of generative models for chaotic dynamical systems and neural data

Vollbildmodus Ctrl+L

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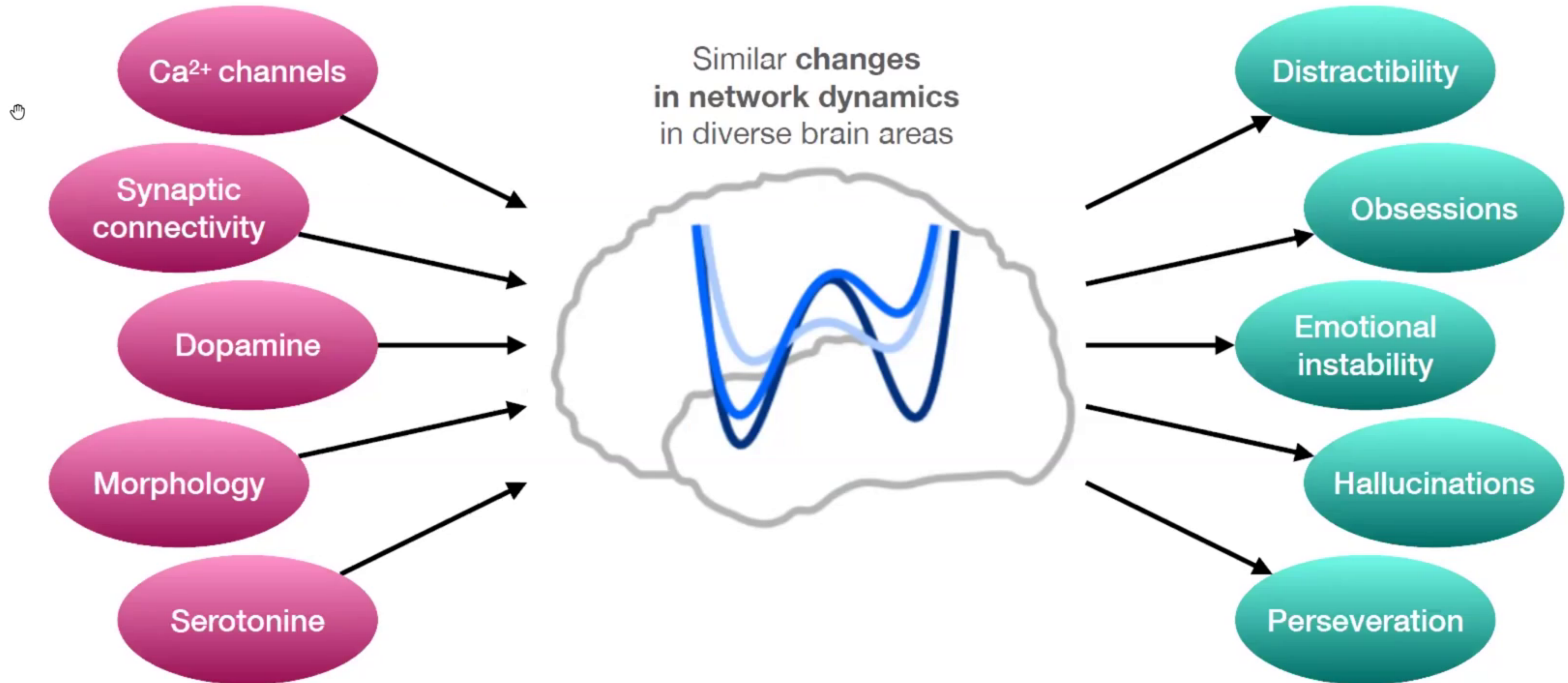




Motivation

Diverse biophysical and structural **causes**

Diverse **changes** in cognitive and emotional **experience**





Inferring Generative Models from Data

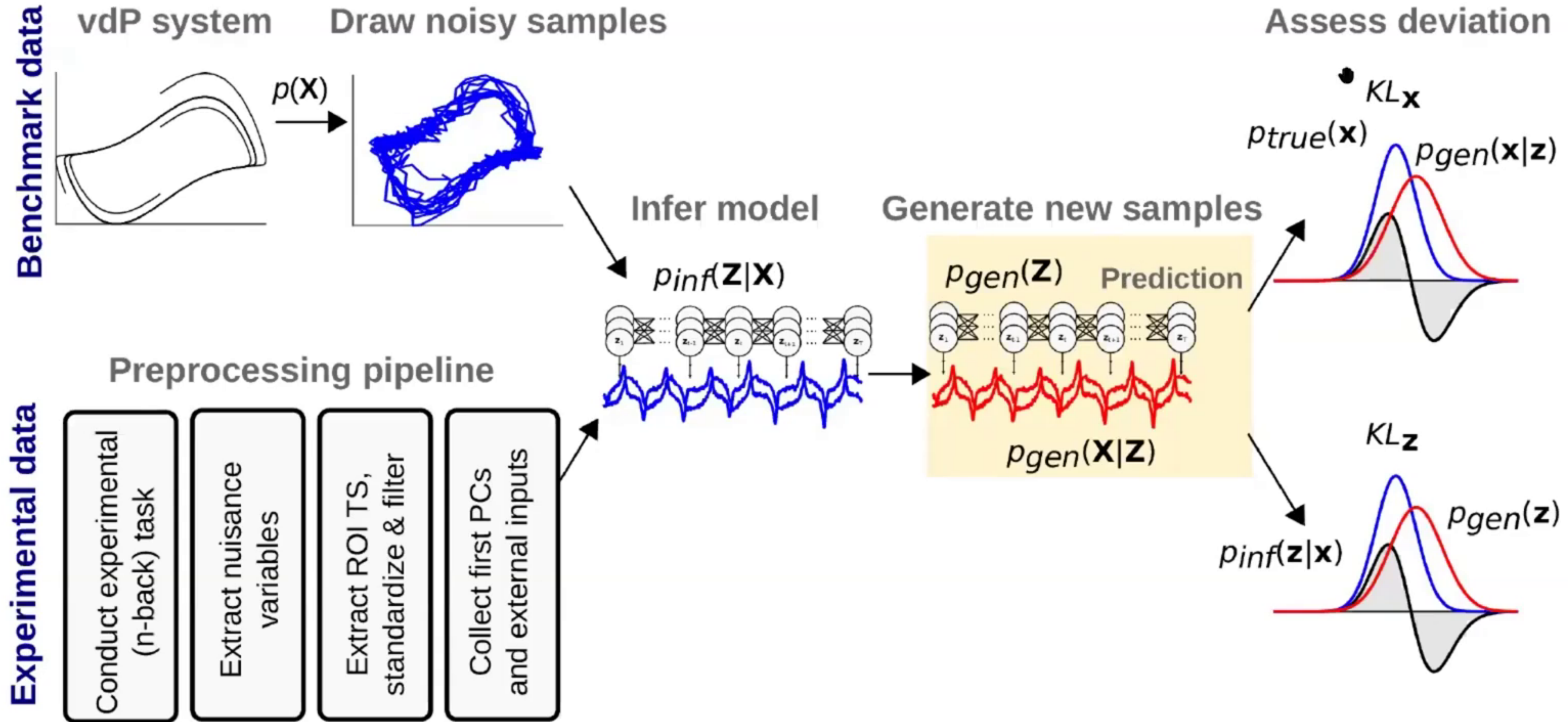
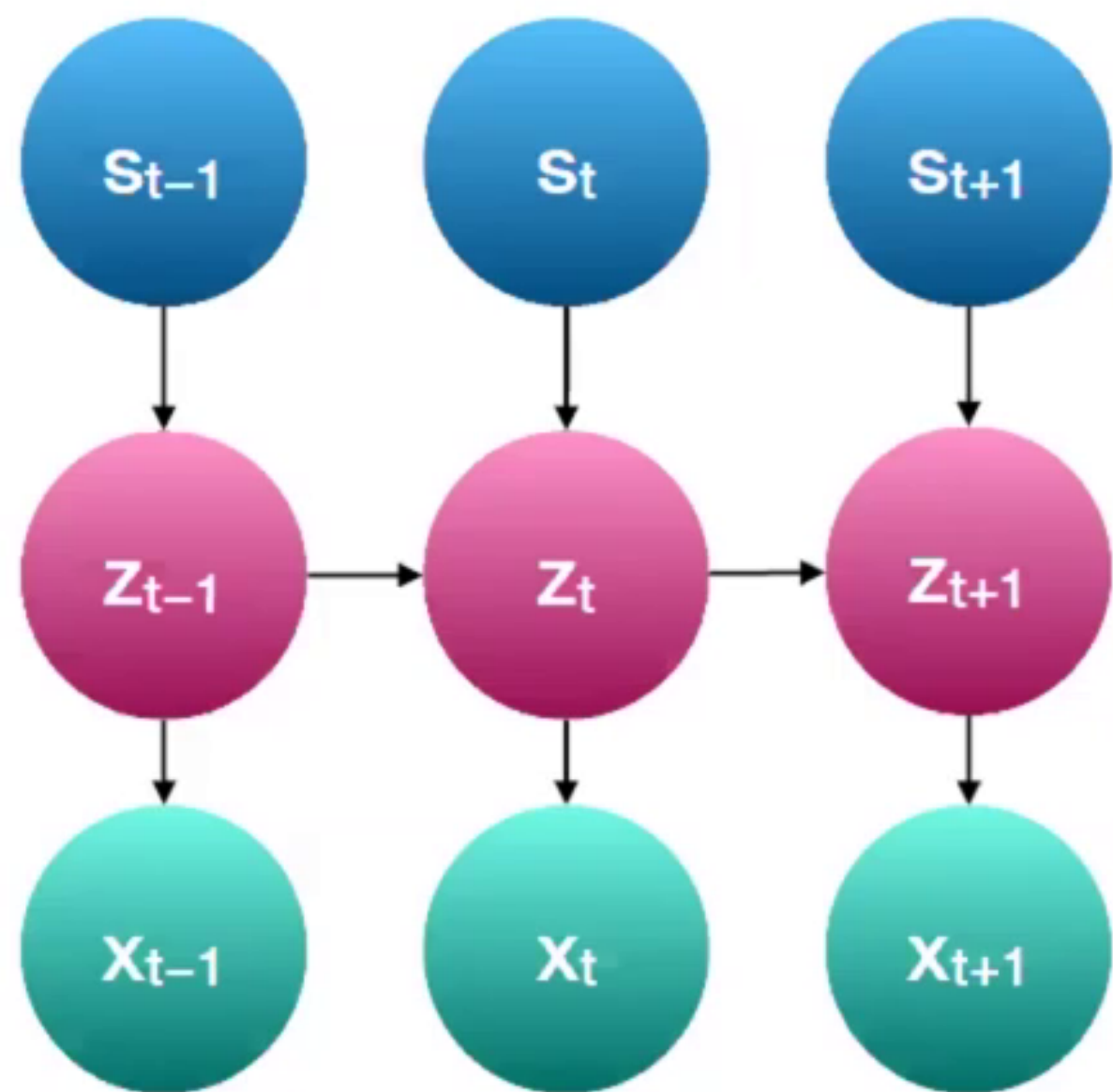


Figure from: Koppe, G., Toutounji, H., Kirsch, P., Lis, S., & Durstewitz, D. (2019). Identifying nonlinear dynamical systems via generative recurrent neural networks with applications to fMRI. PLoS Computational Biology, 15.



Piece-wise Linear Recurrent Neural Network



Latent Model $z_t \in \mathbb{R}^M$

$$z_t = \underset{\text{Diagonal}}{A} z_{t-1} + \underset{\text{Off-diagonal}}{W} \phi(z_{t-1}) + h_0 + C s_t + \epsilon_t, \quad \epsilon_t \sim \mathcal{N}(\mathbf{0}, \Sigma)$$

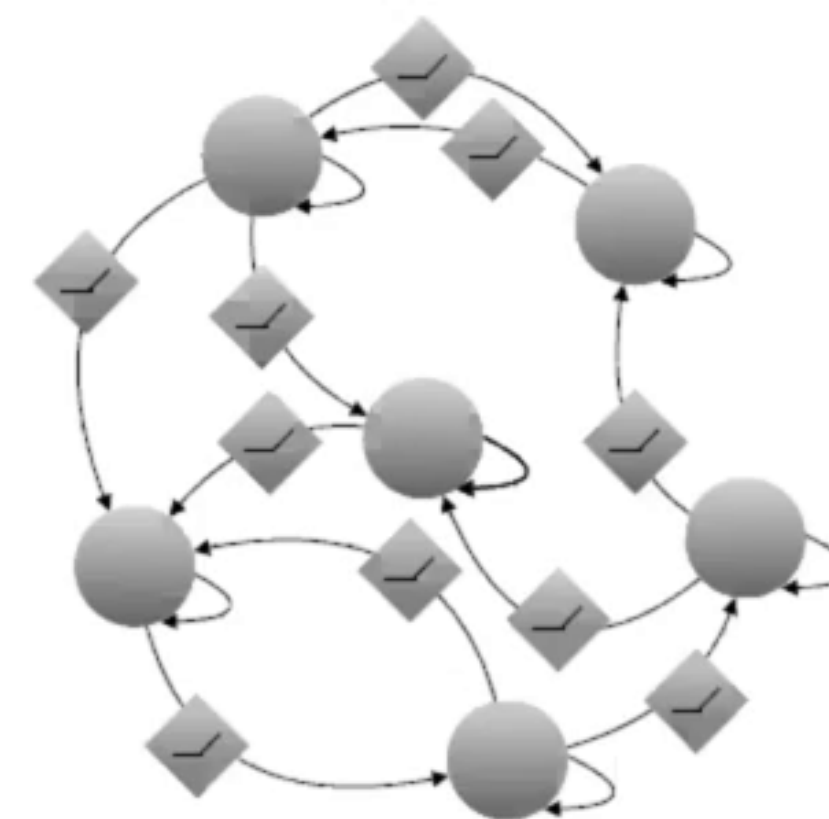
Observation Model

$$x_t = B z_t + \eta_t, \quad \eta_t \sim \mathcal{N}(\mathbf{0}, \Gamma)$$

Complete Likelihood

$$\begin{aligned} \log p_{\theta}(\mathbf{x}, \mathbf{z}) = & -\frac{1}{2} (z_1 - \mu_0 - s_1)^T \Sigma^{-1} (z_1 - \mu_0 - s_1) \\ & -\frac{1}{2} \sum_{t=2}^T (z_t - A z_{t-1} - W \phi(z_{t-1}) - s_t)^T \Sigma^{-1} (z_t - A z_{t-1} - W \phi(z_{t-1}) - s_t) \\ & -\frac{1}{2} \sum_{t=1}^T (x_t - B z_t)^T \Gamma^{-1} (x_t - B z_t) \end{aligned}$$

Activation Function



$$\phi(z_{t-1}) = \max(0, z_{t-1})$$

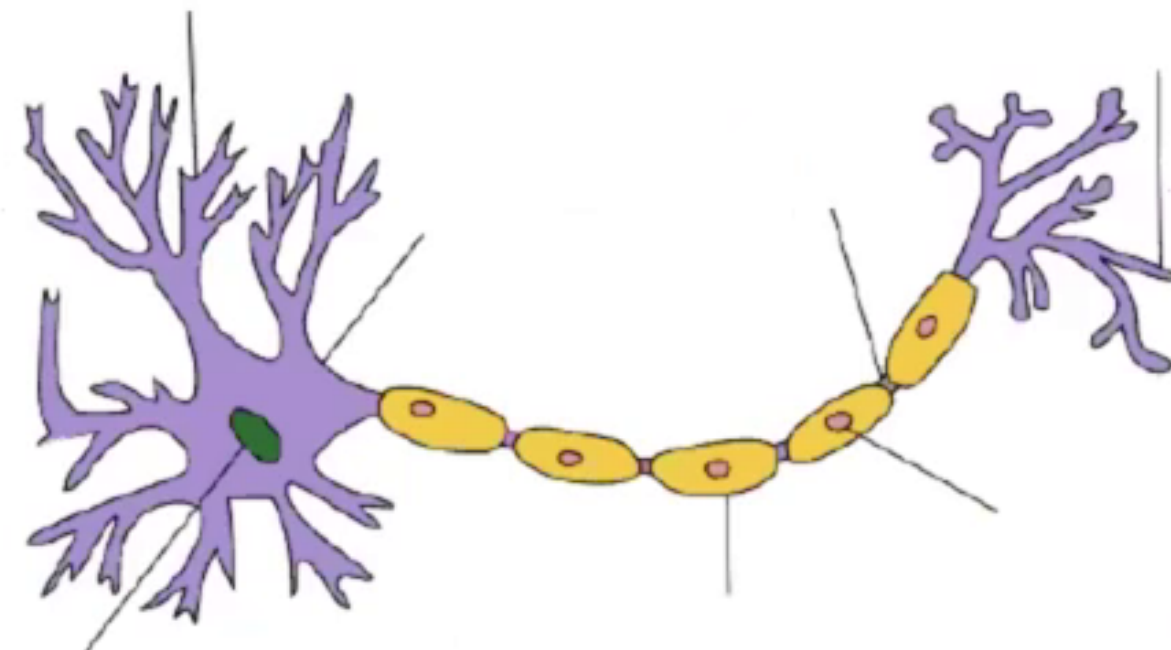
¹ For **fMRI observation model** refer to: Koppe, G., Toutounji, H., Kirsch, P., Lis, S., & Durstewitz, D. (2019). Identifying nonlinear dynamical systems via generative recurrent neural networks with applications to fMRI. PLoS Computational Biology, 15.



Increasing Computational Capacity

- $z_t = Az_{t-1} + W\phi(z_{t-1}) + h_0 + Cs_t + \epsilon_t, \epsilon_t \sim \mathcal{N}(\mathbf{0}, \Sigma)$

- Can we reduce the dimensionality of latent space?
- Retain piece-wise linear form
- Neurophysiological analogy: dendritic computation or neuronal diversity

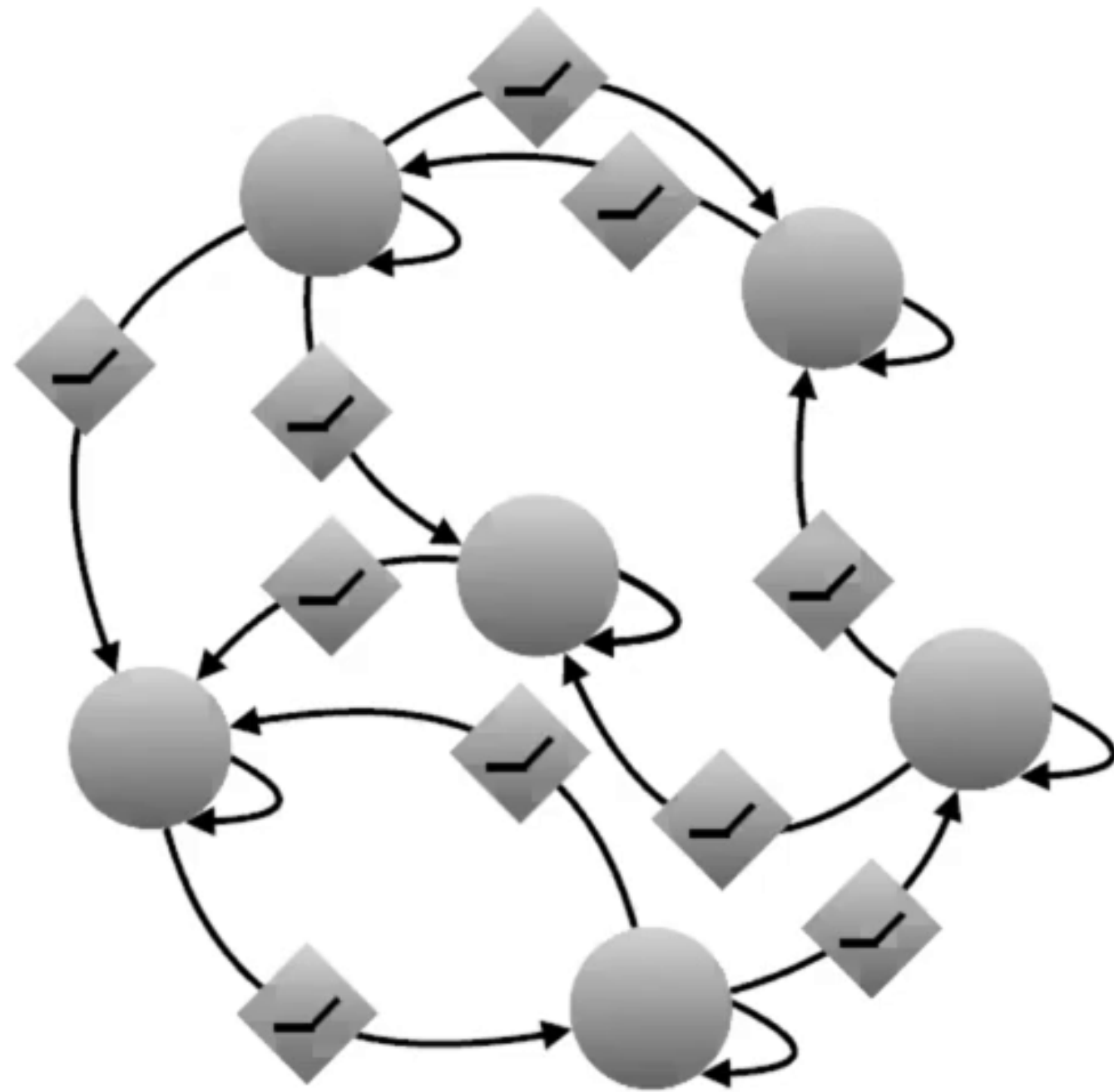




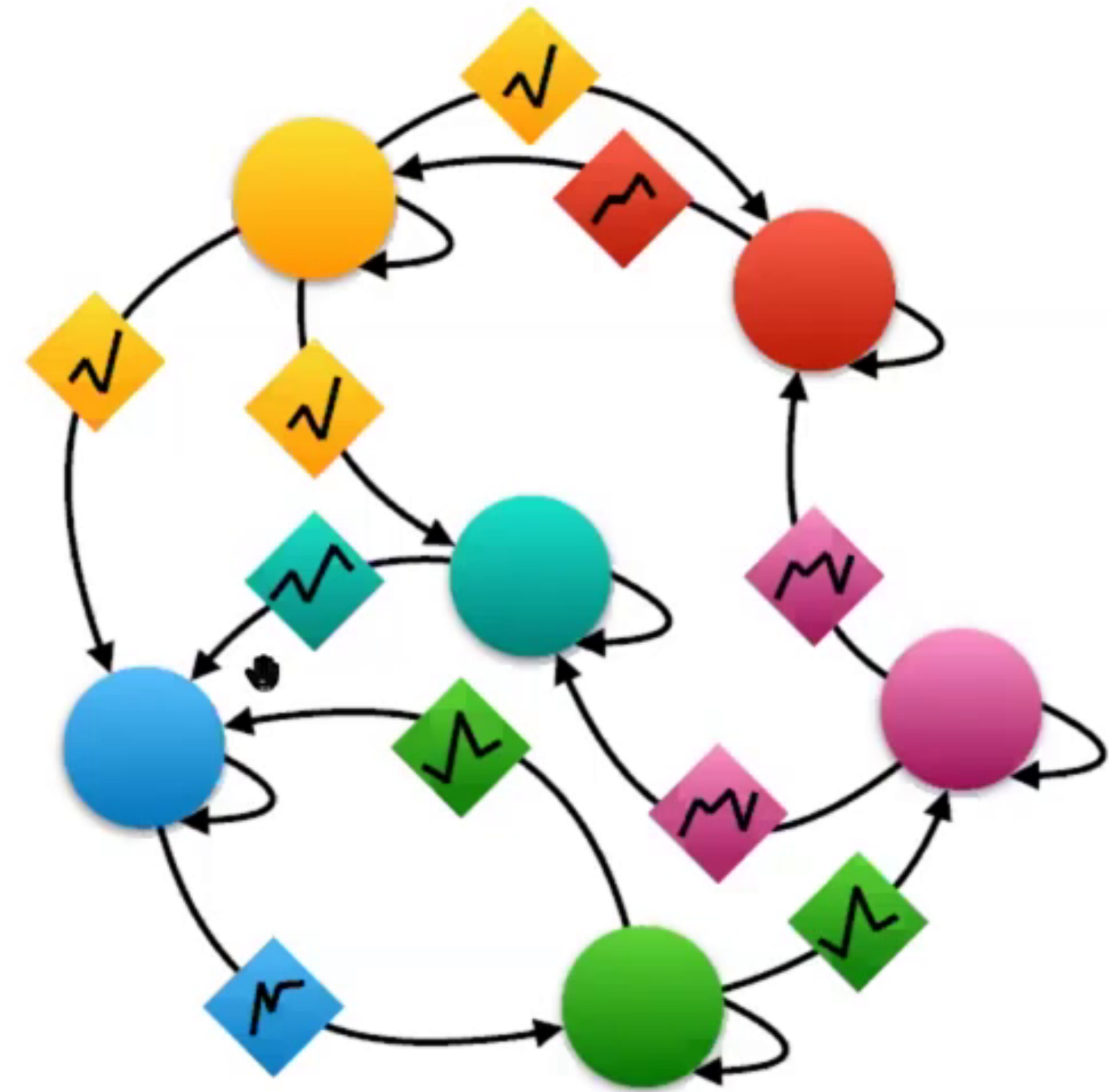
Basis Expansion

New activation function in latent model

$$\mathbf{z}_t = \mathbf{A}\mathbf{z}_{t-1} + \mathbf{W}\phi(\mathbf{z}_{t-1}) + \mathbf{h}_0 + \mathbf{C}\mathbf{s}_t + \boldsymbol{\epsilon}_t, \quad \boldsymbol{\epsilon}_t \sim \mathcal{N}(\mathbf{0}, \boldsymbol{\Sigma})$$



$$\phi(\mathbf{z}_{t-1}) = \max(0, \mathbf{z}_{t-1})$$



$$\phi(\mathbf{z}_{t-1}) = \sum_{b=1}^B \alpha_b \max(0, \mathbf{z}_{t-1} - \mathbf{h}_b)$$



Manuel Brenner

Variational Inference

$$\mathbf{x} = \{\mathbf{x}_t \mid t = 1 \dots T\}$$

$$\mathbf{z} = \{\mathbf{z}_t \mid t = 1 \dots T\}$$

Variational Lower Bound

$$p(\mathbf{x}) \geq \mathcal{L}(\theta, \phi, \mathbf{x}) = \mathbb{E}_{q_\phi(\mathbf{z}|\mathbf{x})}[\log p_\theta(\mathbf{x}, \mathbf{z})] + \mathbb{E}_{q_\phi(\mathbf{z}|\mathbf{x})}[\log q_\phi(\mathbf{z}|\mathbf{x})]$$



Complete Likelihood

$$\log p_\theta(\mathbf{x}, \mathbf{z}) = -\frac{1}{2}(\mathbf{z}_1 - \boldsymbol{\mu}_0 - \mathbf{s}_1)^T \boldsymbol{\Sigma}^{-1}(\mathbf{z}_1 - \boldsymbol{\mu}_0 - \mathbf{s}_1)$$

$$-\frac{1}{2} \sum_{t=2}^T (\mathbf{z}_t - \mathbf{A}\mathbf{z}_{t-1} - \mathbf{W}\phi(\mathbf{z}_{t-1}) - \mathbf{s}_t)^T \boldsymbol{\Sigma}^{-1}(\mathbf{z}_t - \mathbf{A}\mathbf{z}_{t-1} - \mathbf{W}\phi(\mathbf{z}_{t-1}) - \mathbf{s}_t)$$

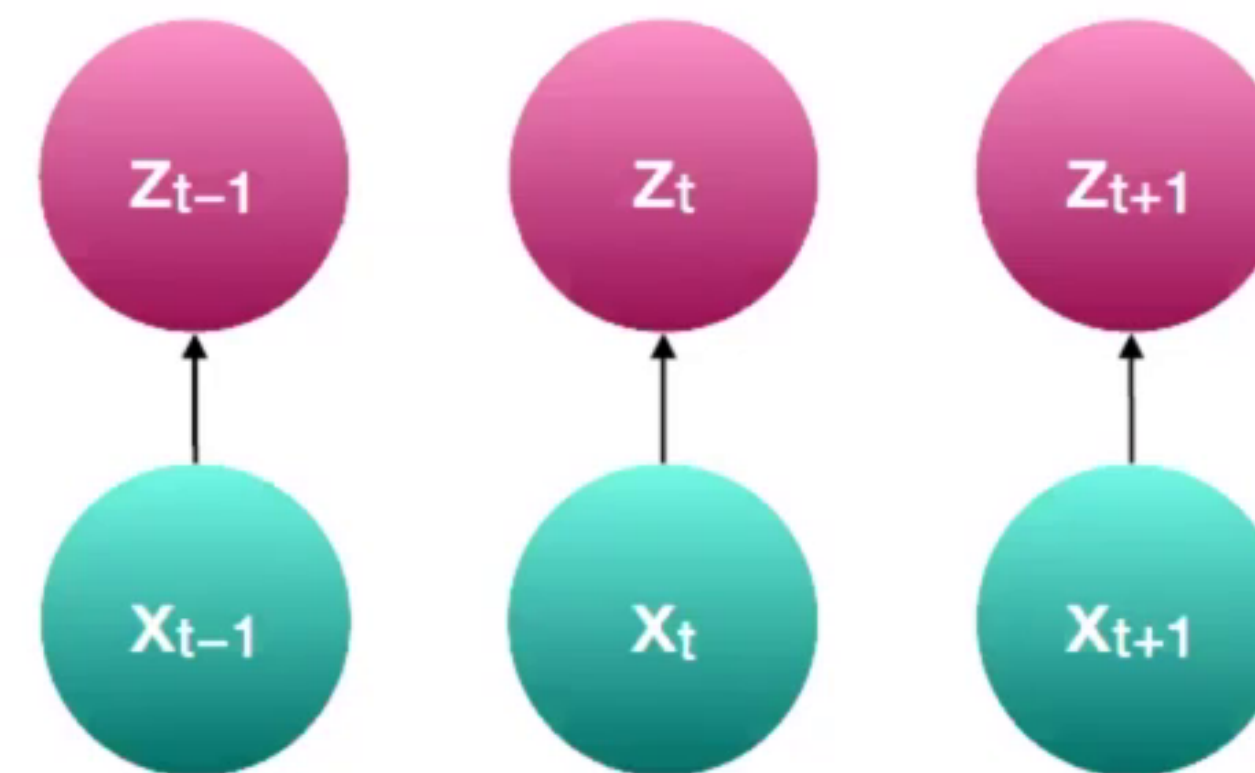
$$-\frac{1}{2} \sum_{t=1}^T (\mathbf{x}_t - \mathbf{B}\mathbf{z}_t)^T \boldsymbol{\Gamma}^{-1}(\mathbf{x}_t - \mathbf{B}\mathbf{z}_t)$$

Mean-Field Approximation

$$q_\phi(\mathbf{z}|\mathbf{x}) = \prod_{t=1}^T q_\phi(\mathbf{z}_t|\mathbf{x}_t) = \prod_{t=1}^T \mathcal{N}(\boldsymbol{\mu}_\phi(\mathbf{x}_t), \boldsymbol{\sigma}_\phi(\mathbf{x}_t)^2)$$

$$\boldsymbol{\mu}_\phi(\mathbf{x}_t) = \text{NN}_\mu(\mathbf{x}_t)$$

$$\boldsymbol{\sigma}_\phi(\mathbf{x}_t) = \text{NN}_\sigma(\mathbf{x}_t)$$

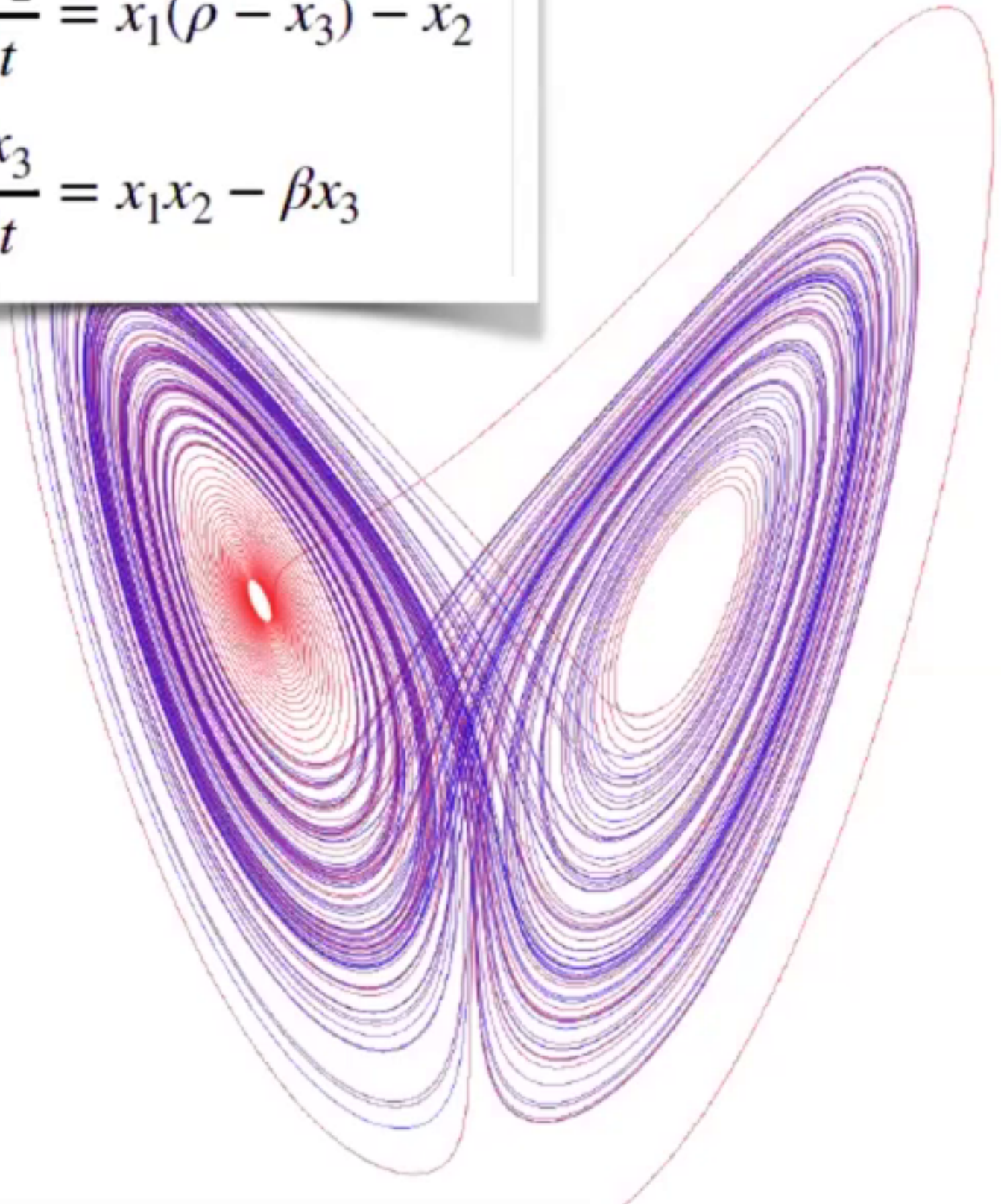




Benchmark Dynamical Systems

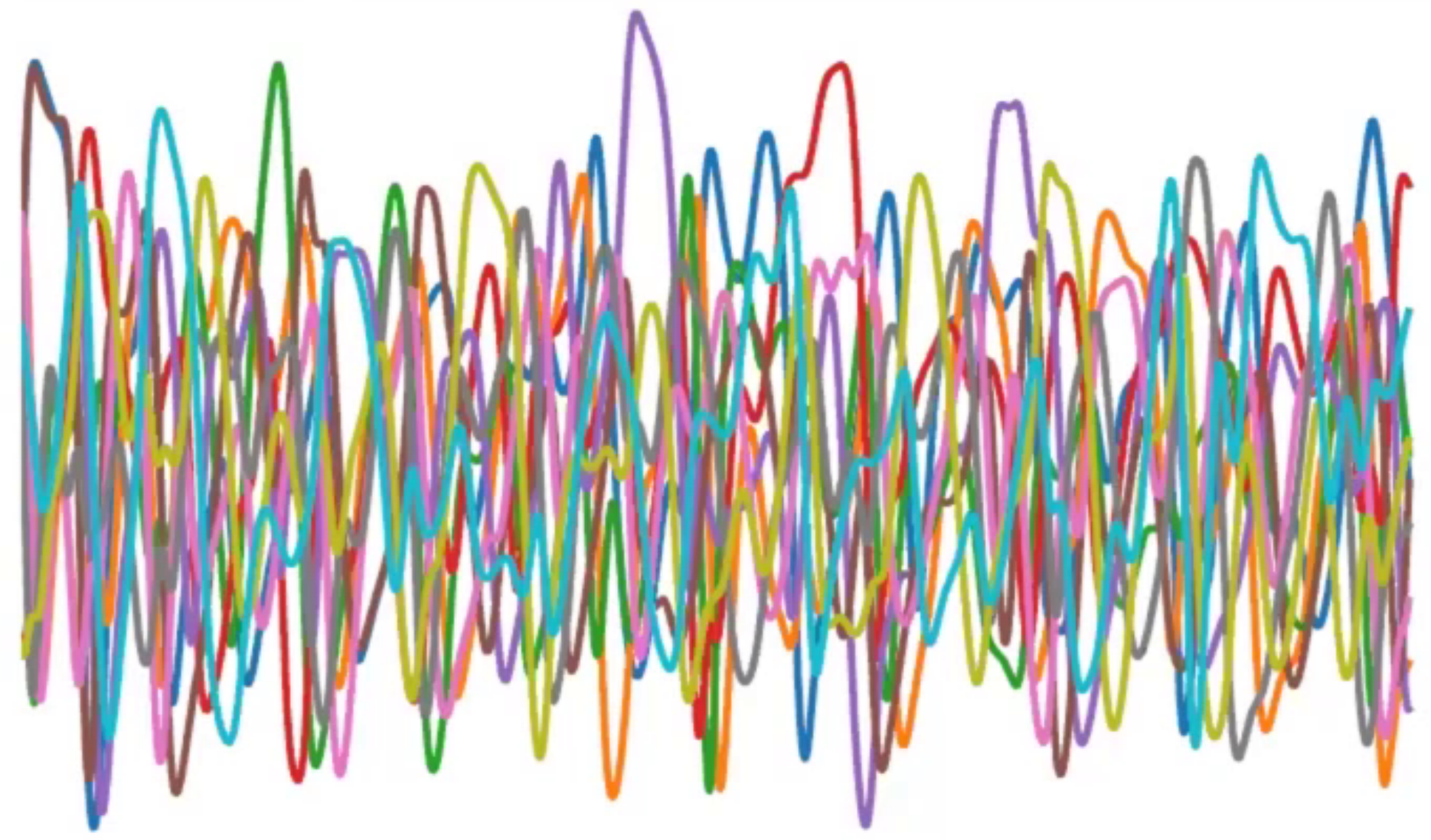
Lorenz Attractor

$$\begin{aligned}\frac{dx_1}{dt} &= \sigma(x_2 - x_1) \\ \frac{dx_2}{dt} &= x_1(\rho - x_3) - x_2 \\ \frac{dx_3}{dt} &= x_1x_2 - \beta x_3\end{aligned}$$

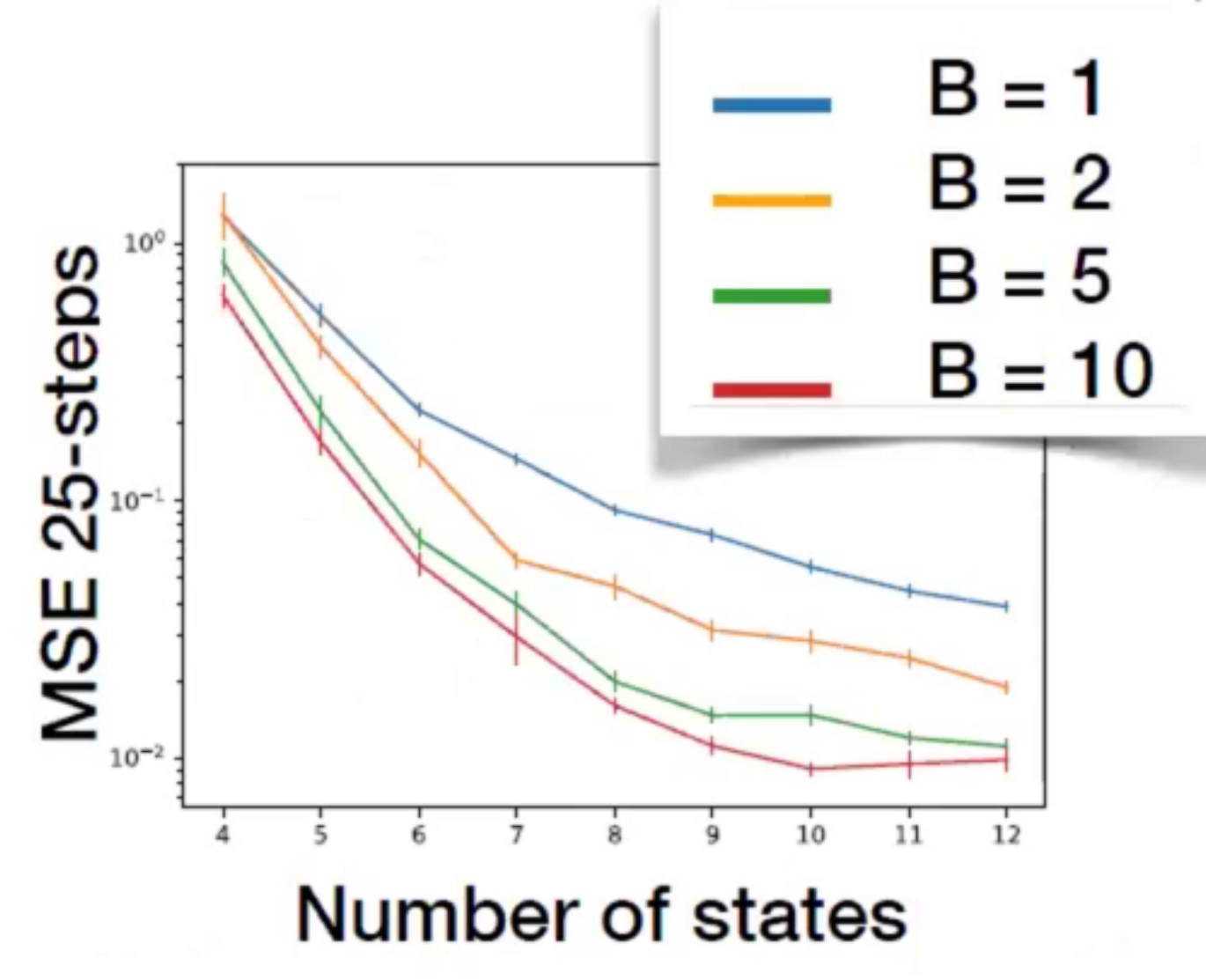
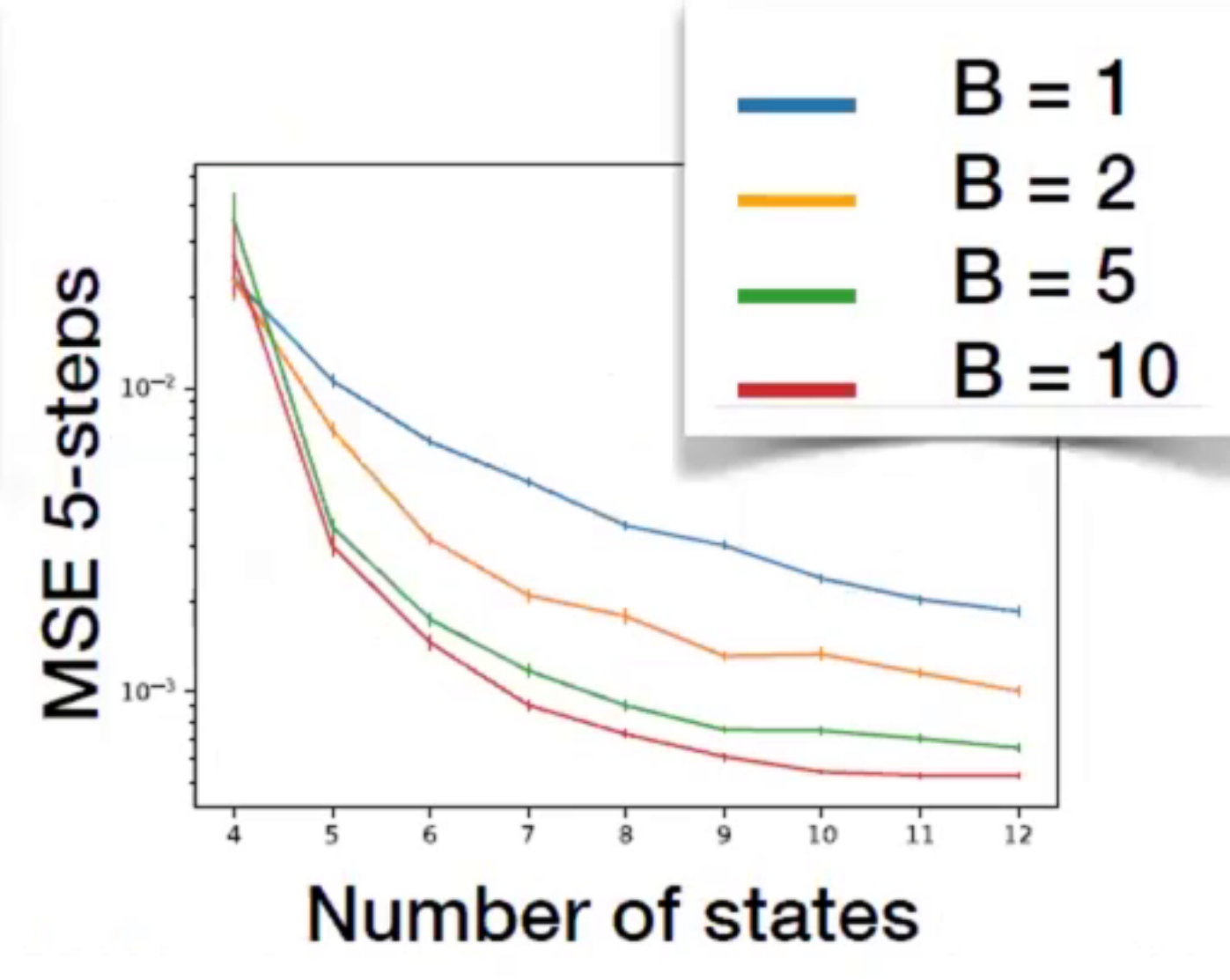
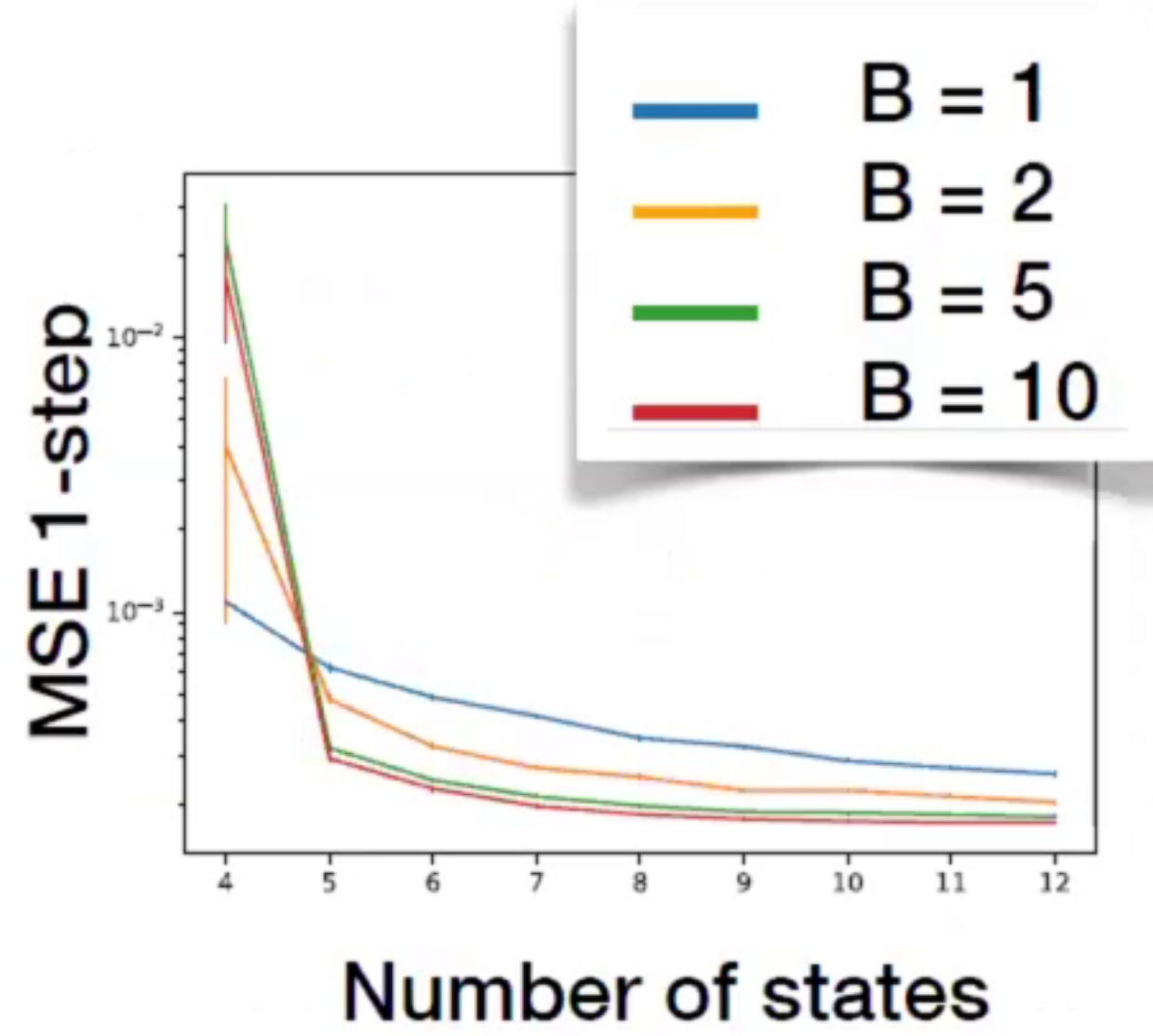
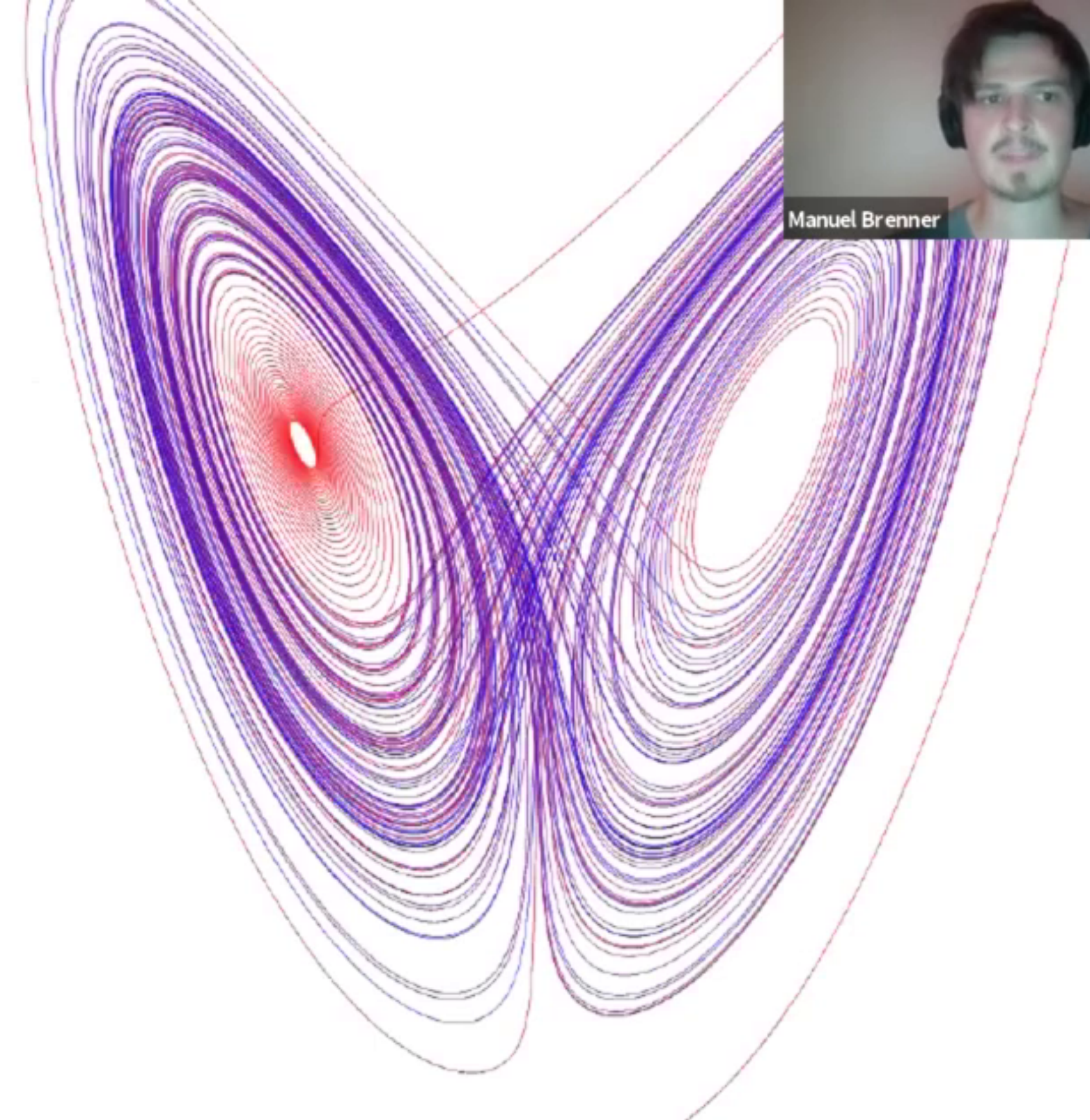
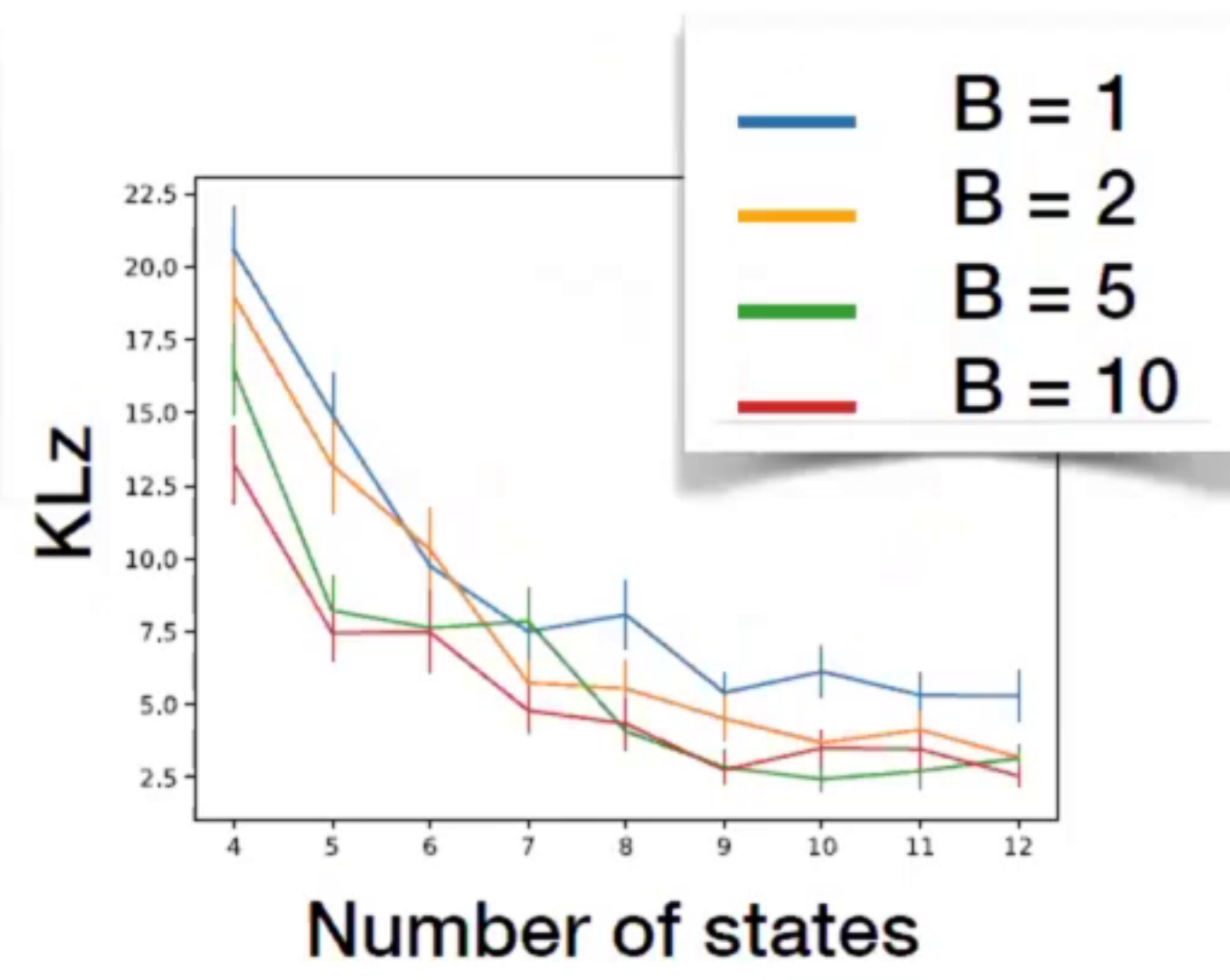
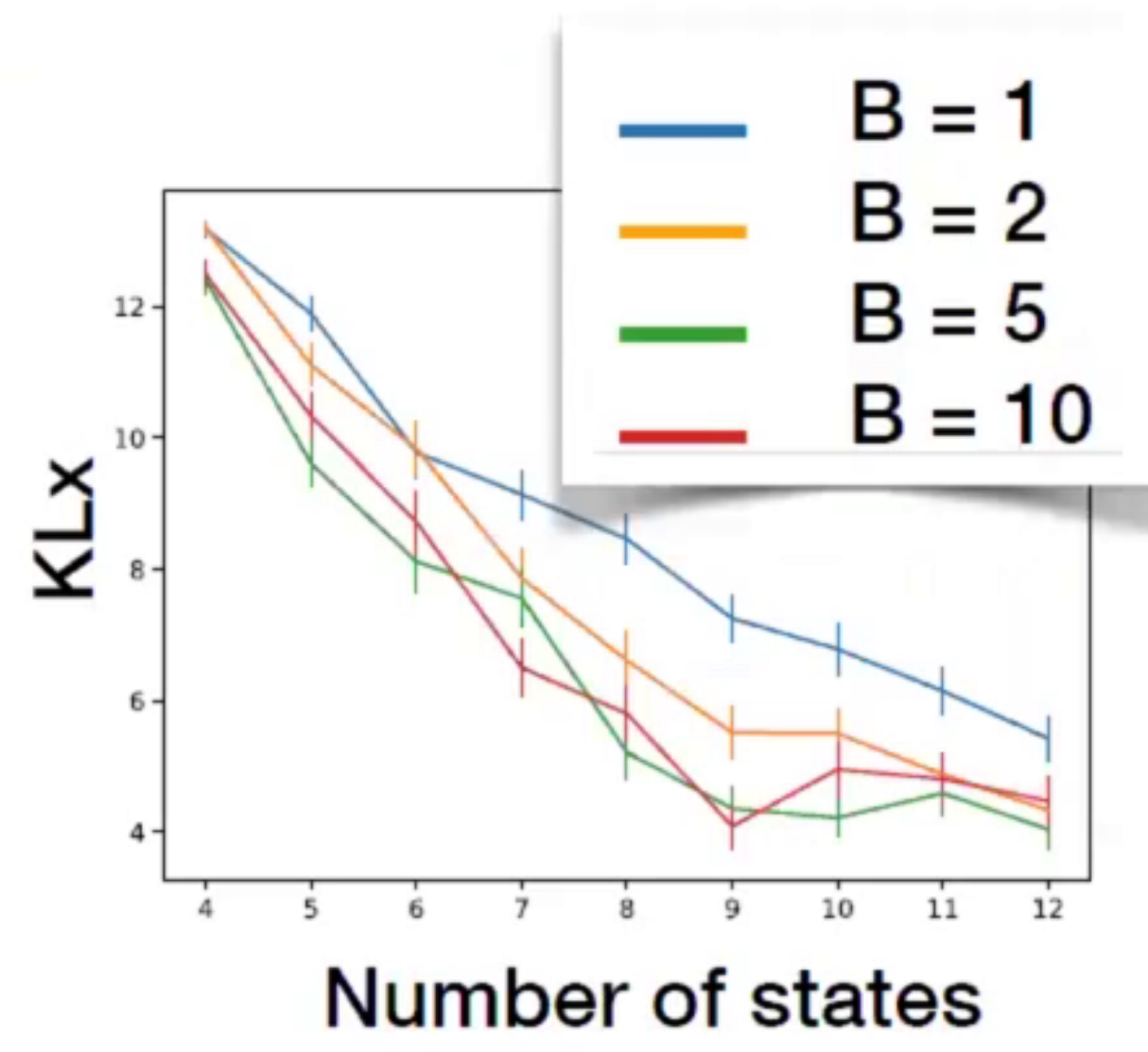


Lorenz-96 System

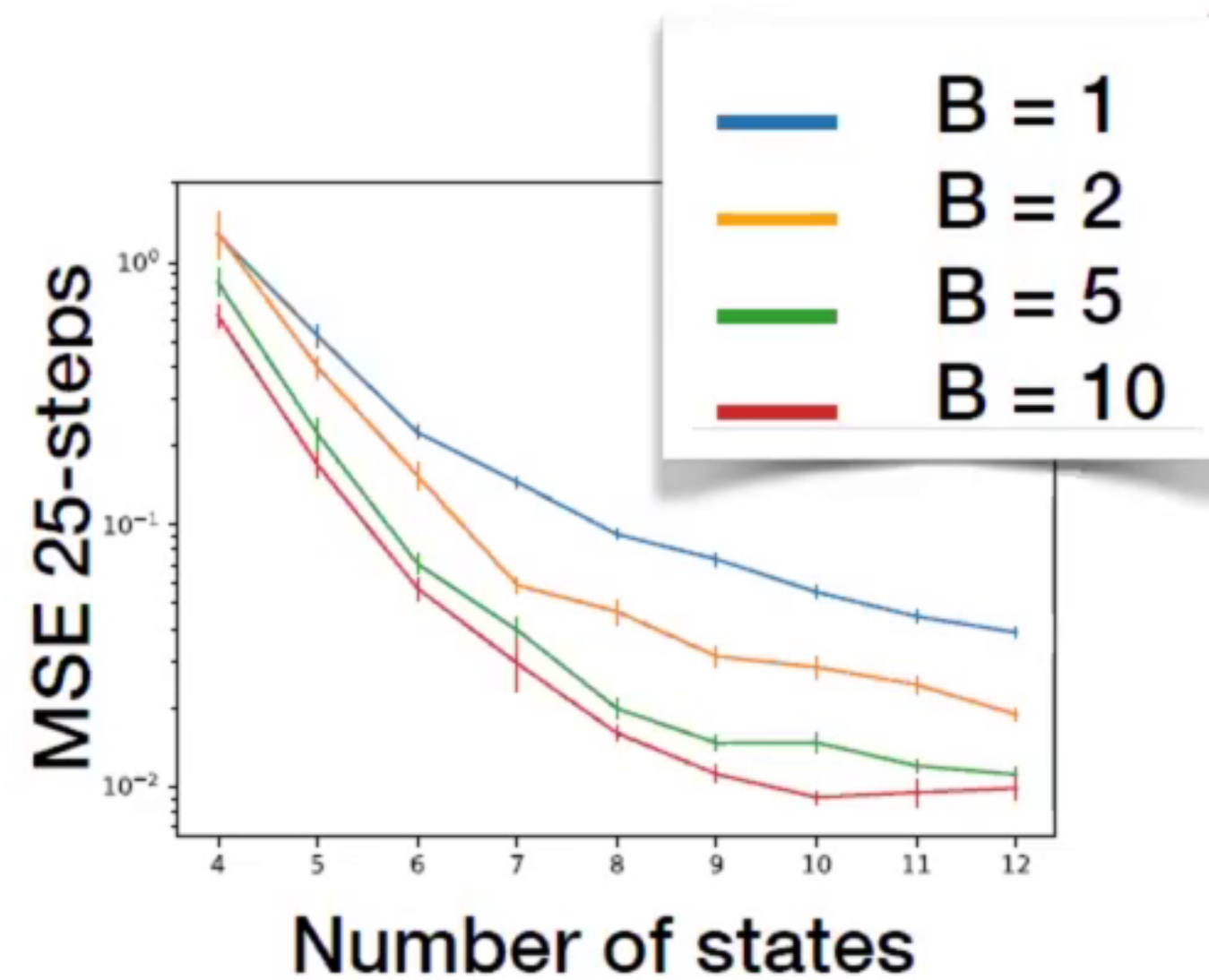
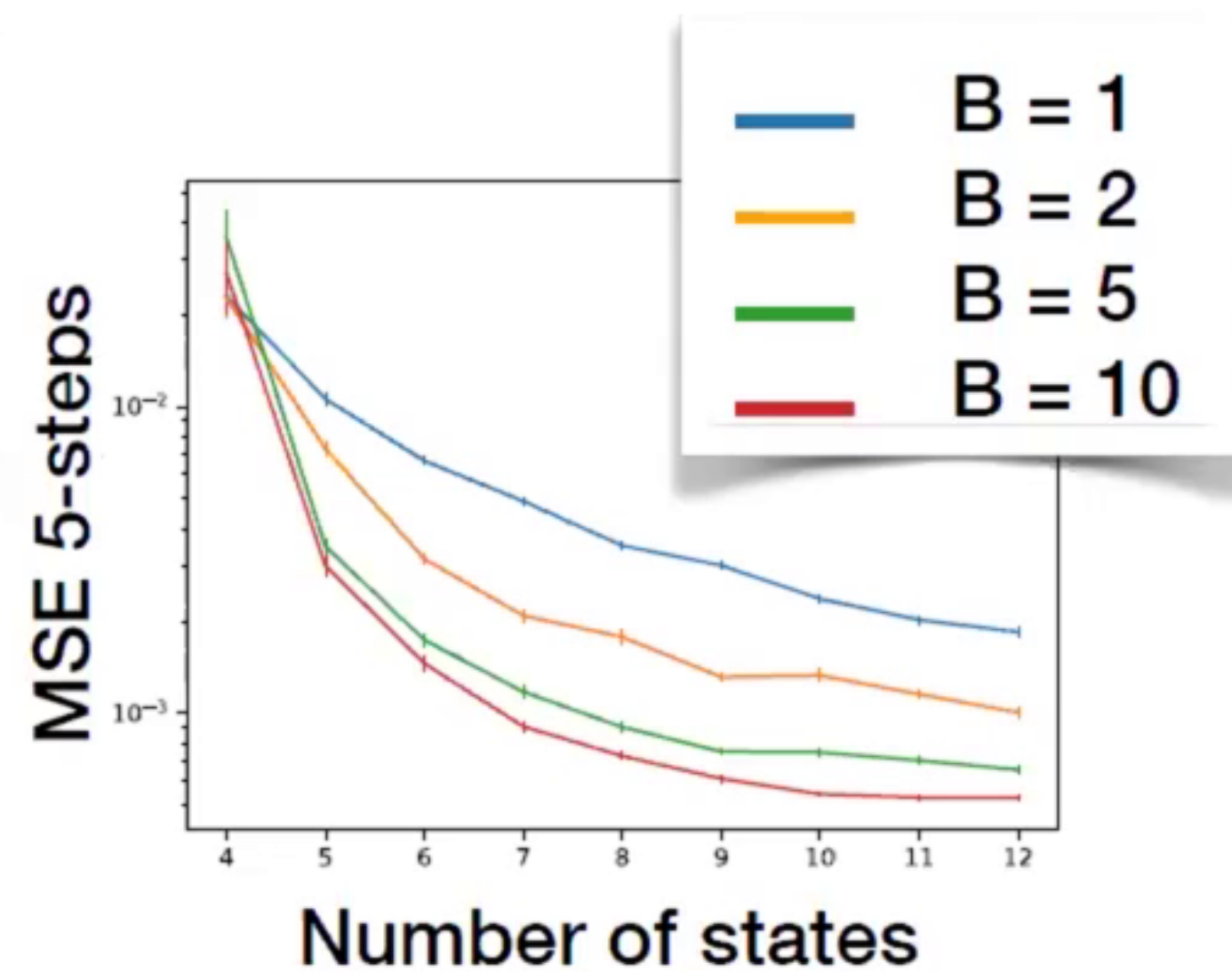
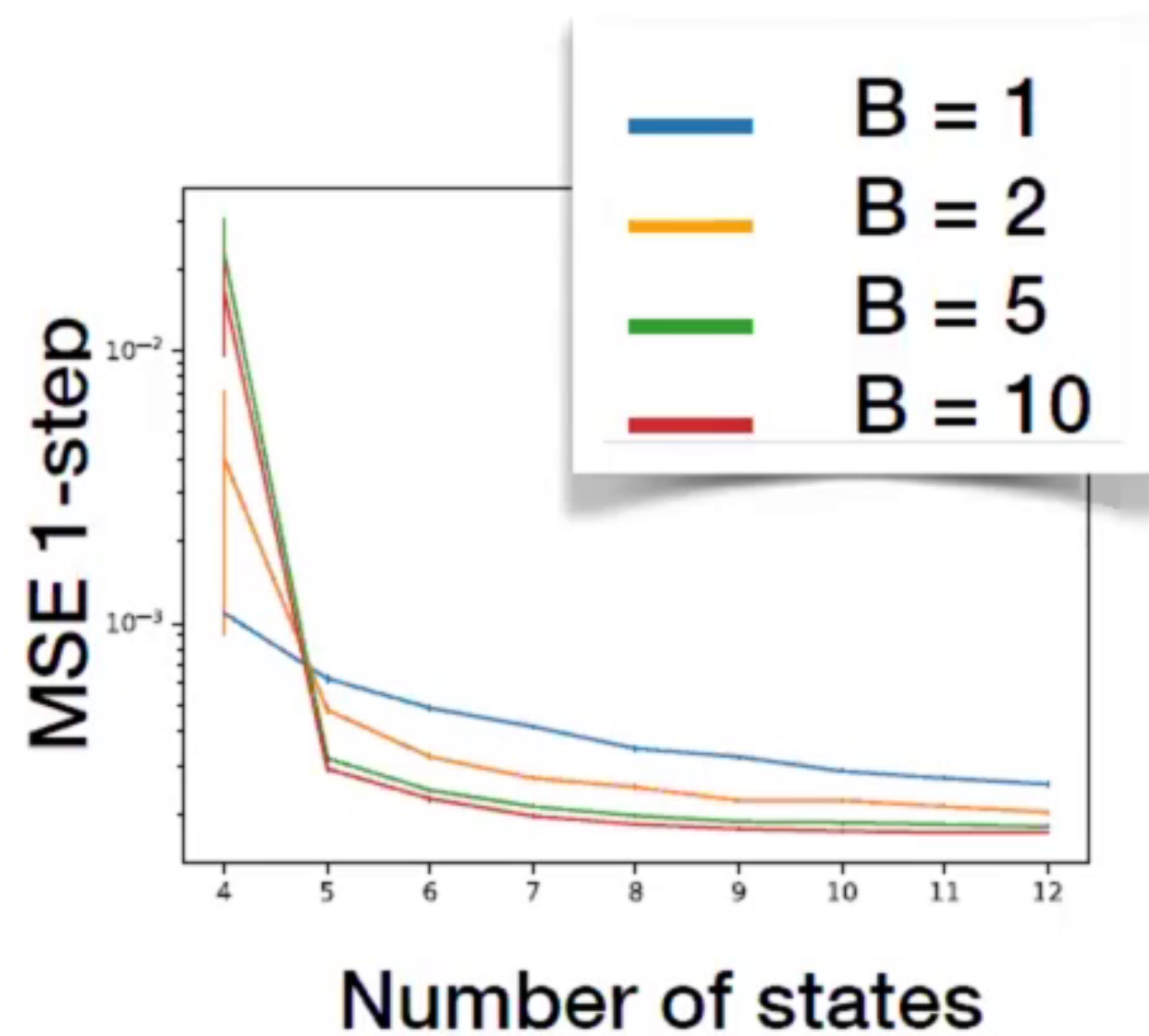
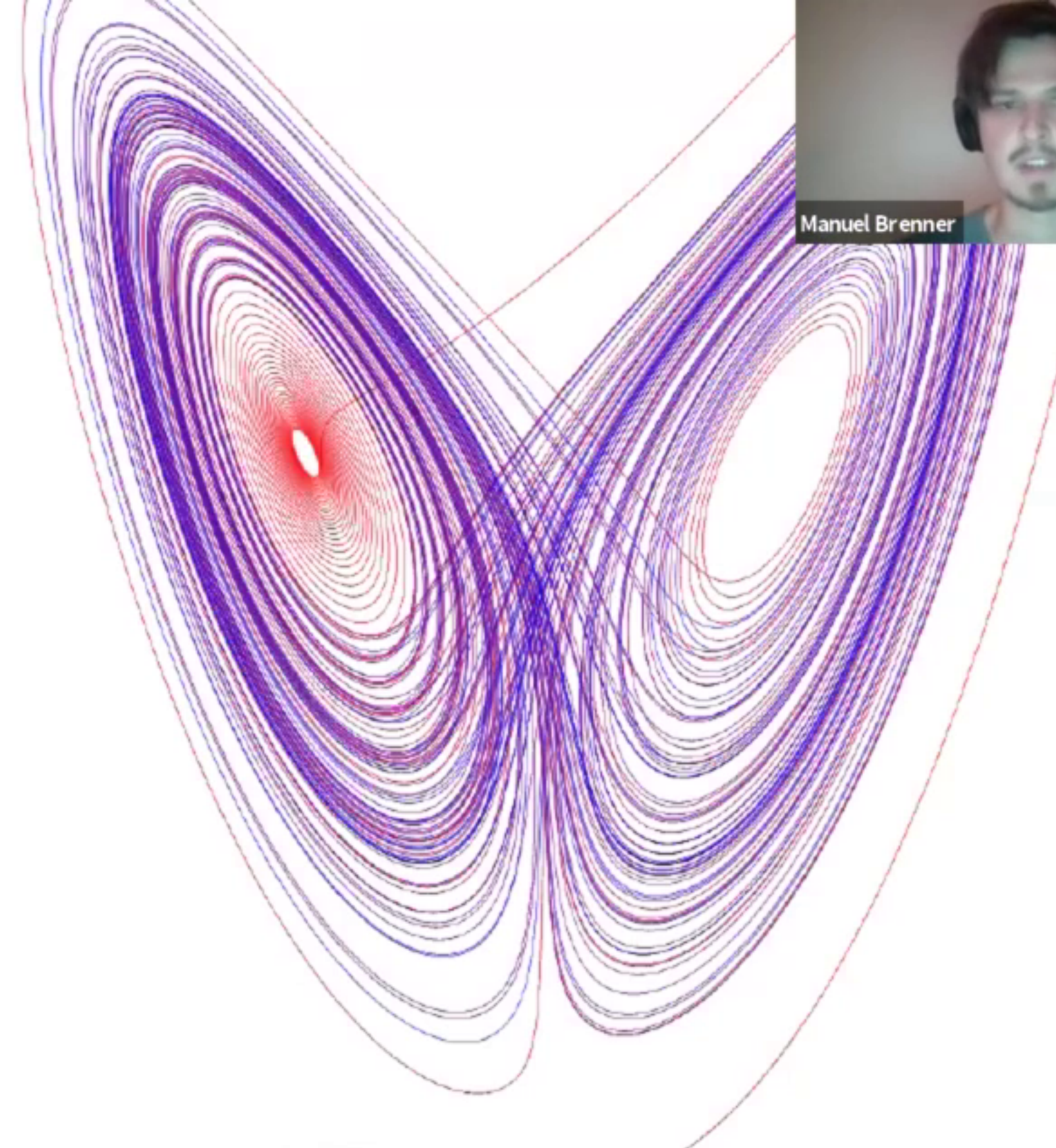
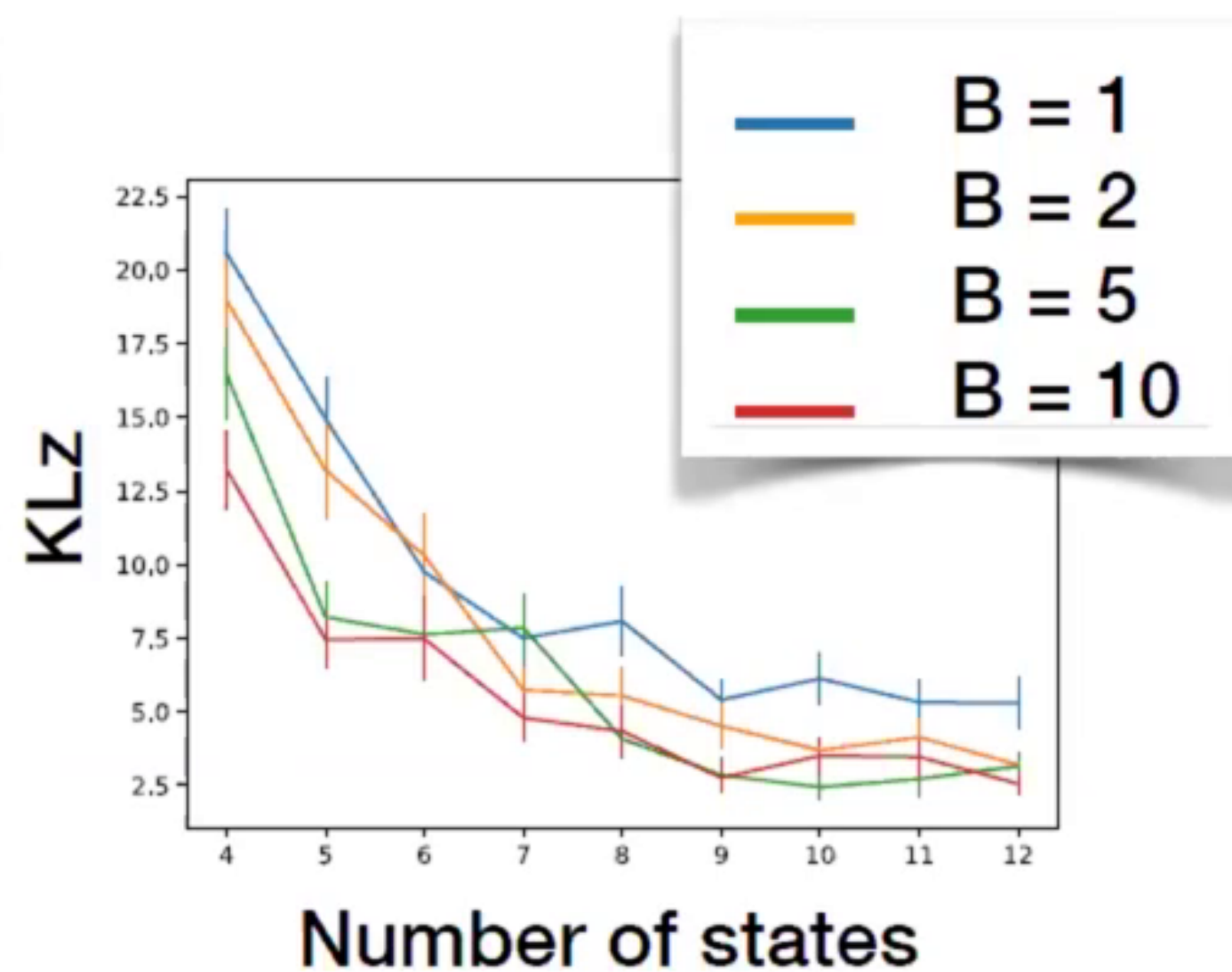
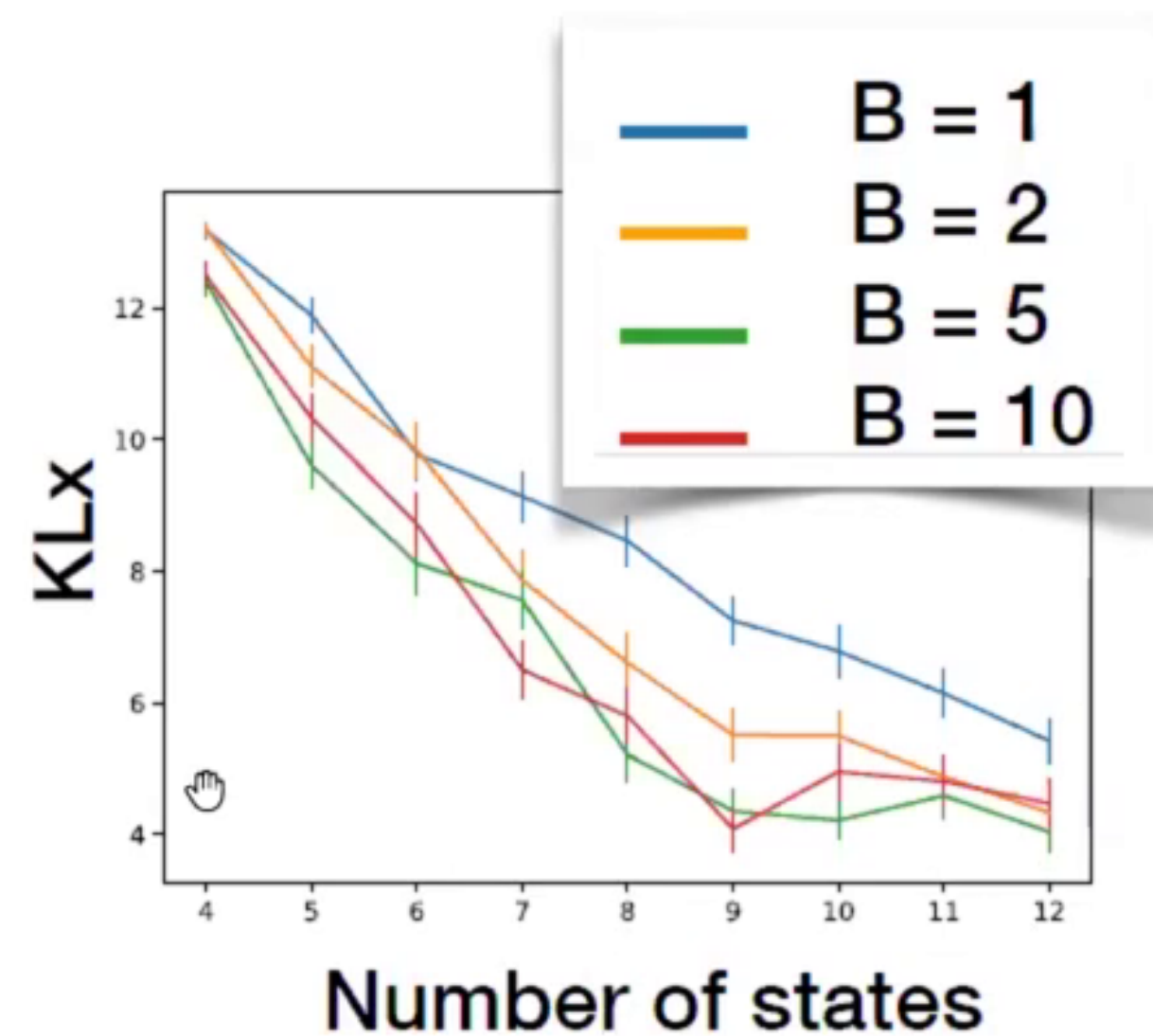
$$\frac{dx_i}{dt} = (x_{i+1} - x_{i-2})x_{i-1} - x_i + F$$



Results - Lorenz System

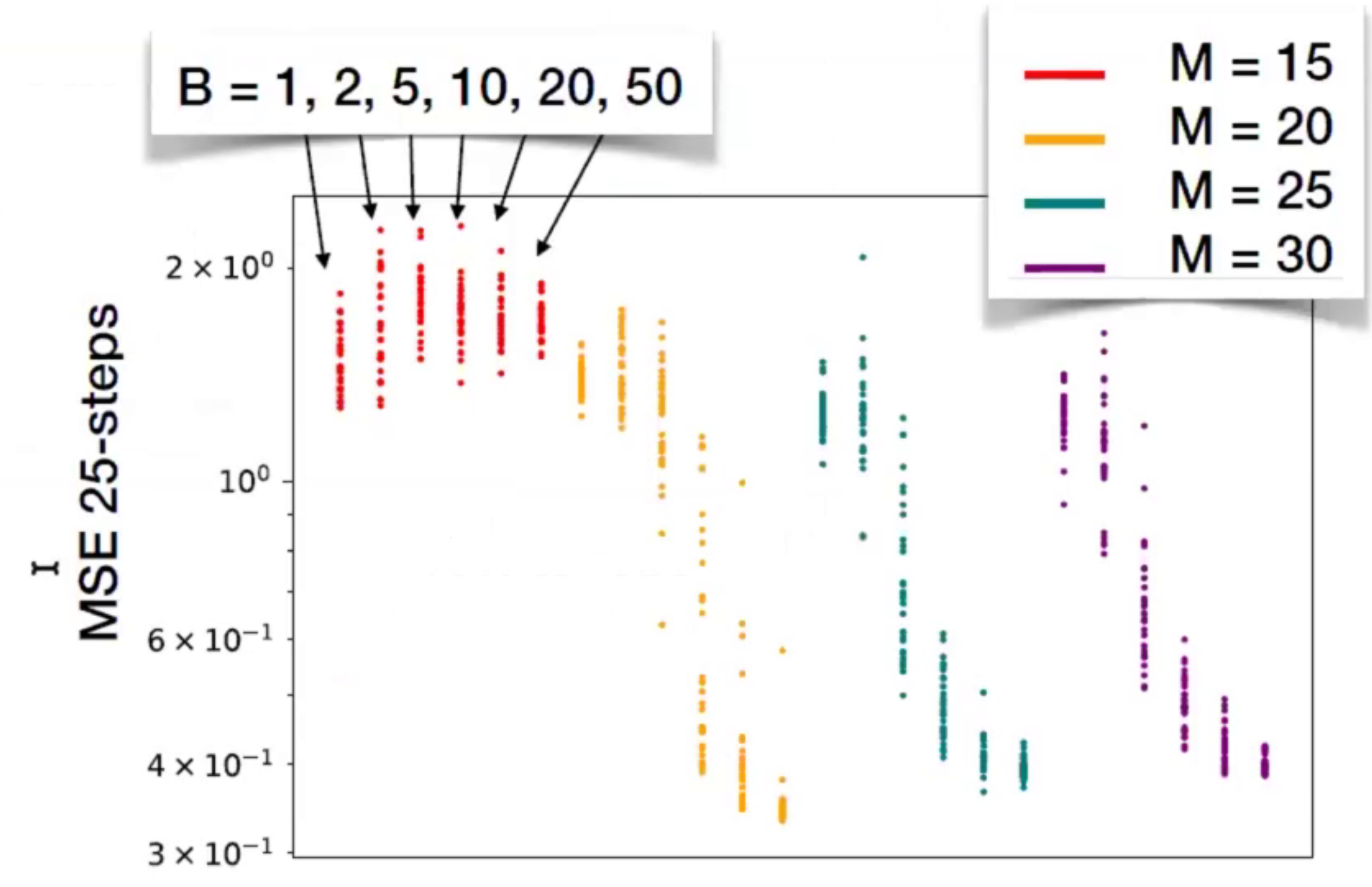
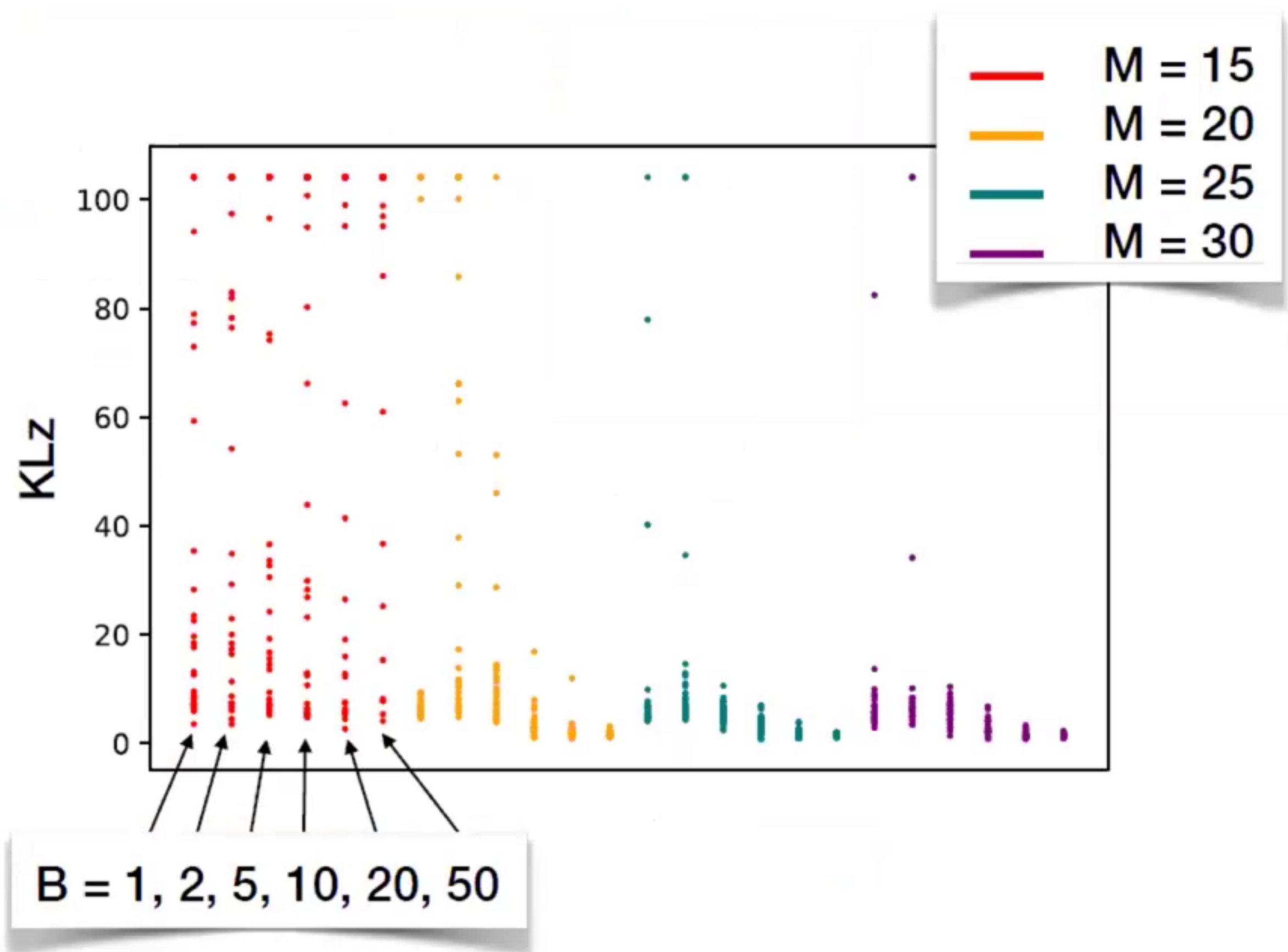


Results - Lorenz System



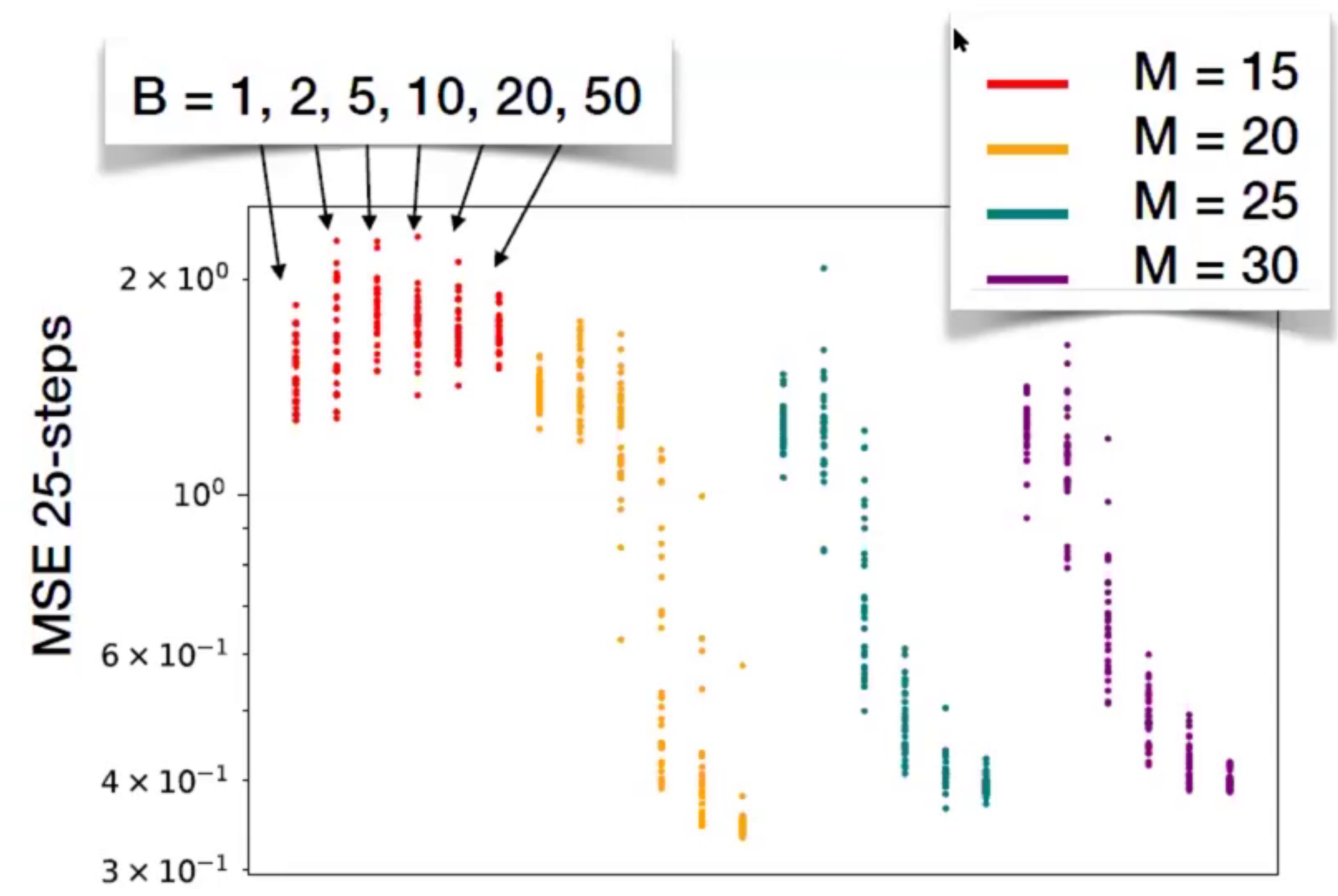
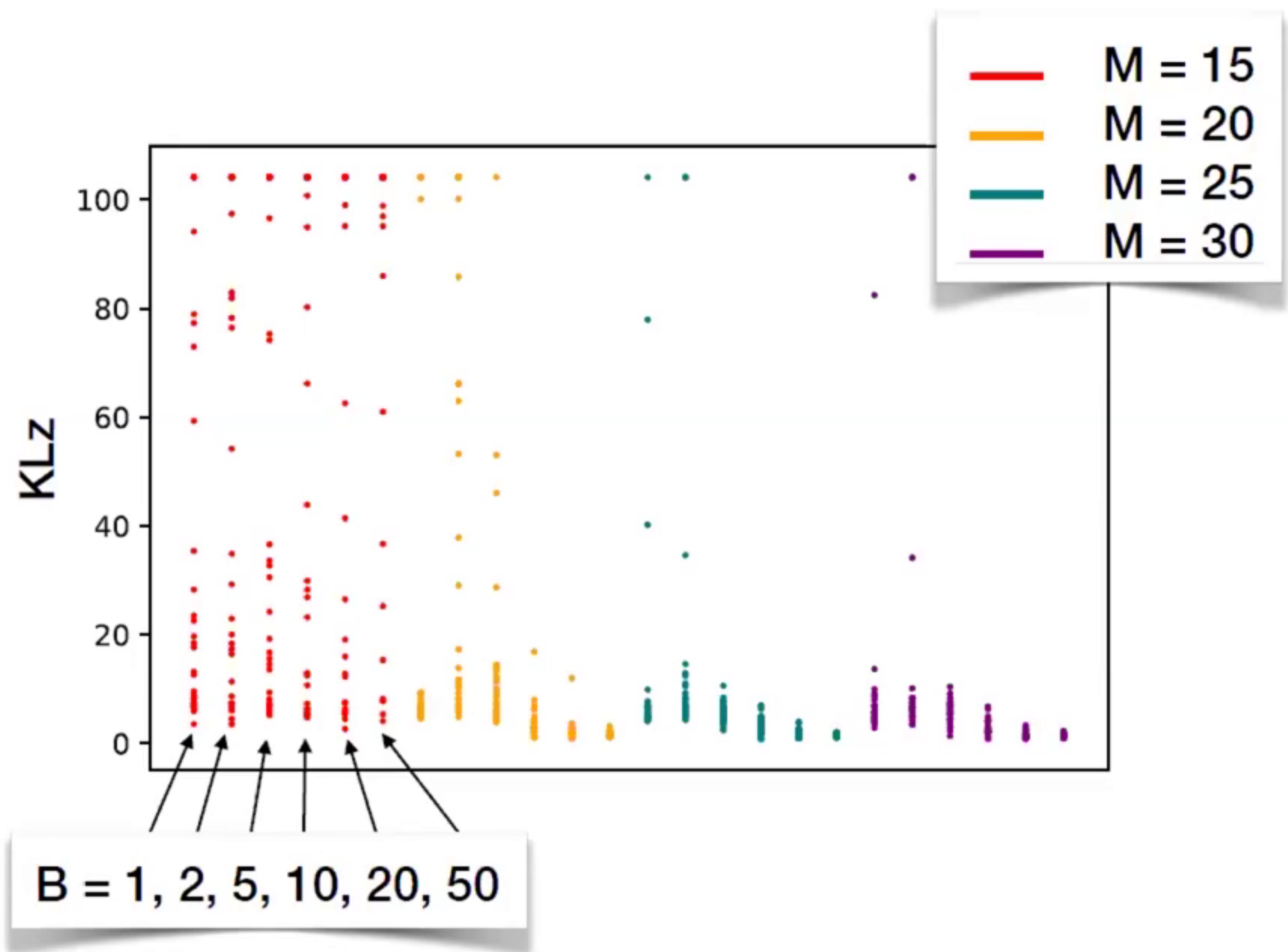


Results - Lorenz-96 System





Results - Lorenz-96 System

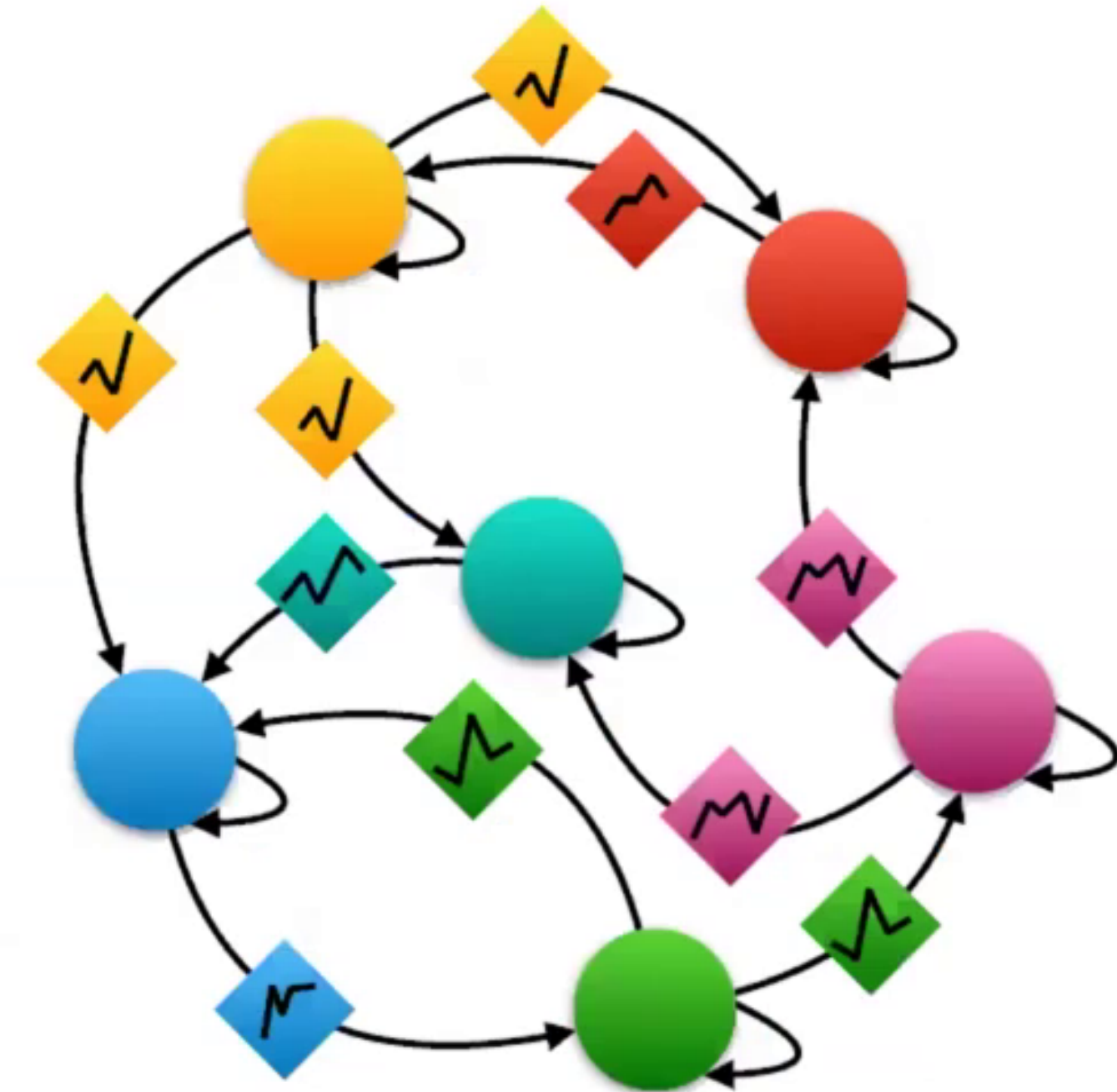


Summary



Basis expansion:

- Improves inference
- Encourages learning of more interesting dynamics
- Reduces dimensionality of the latent space

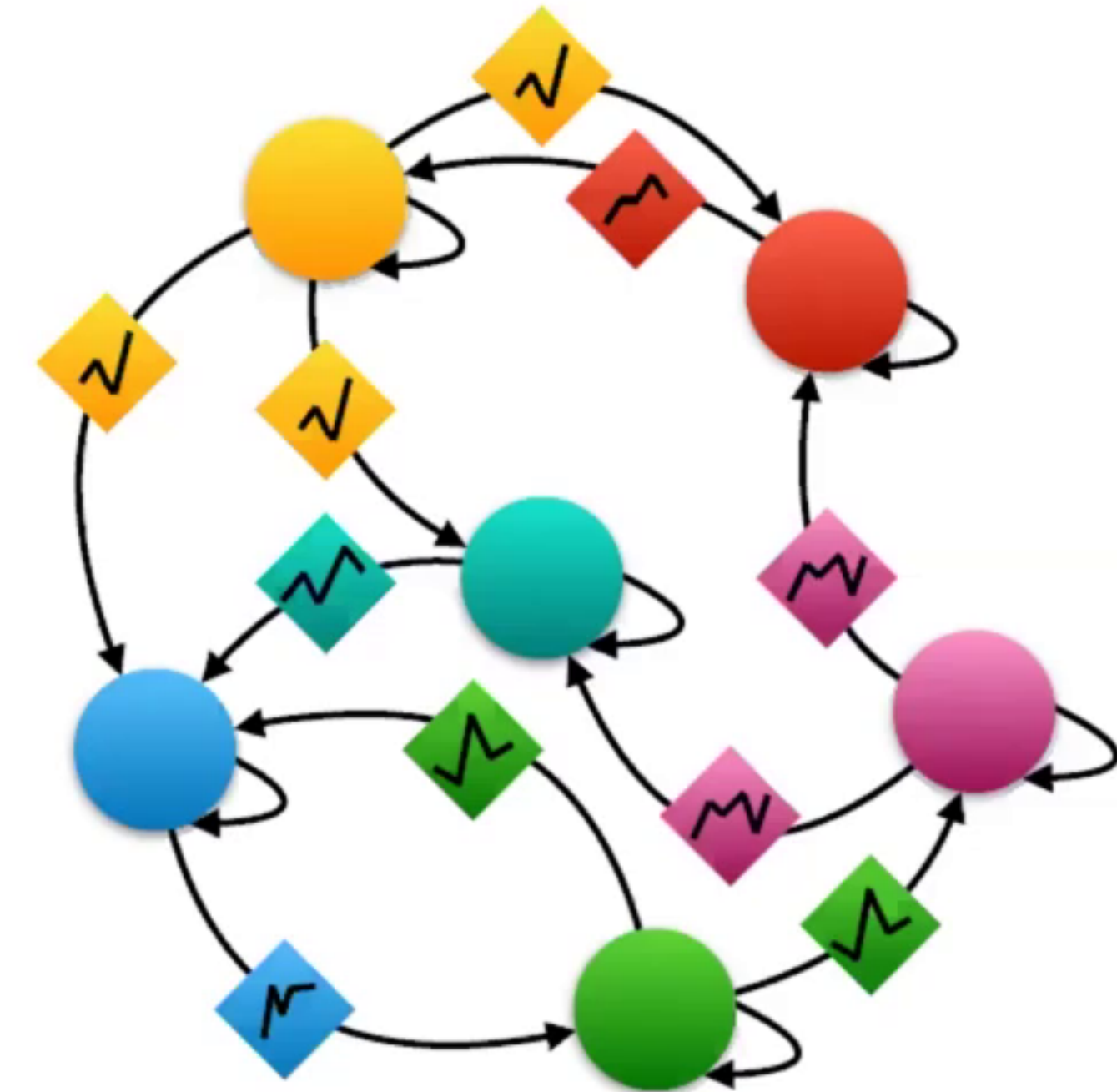


Summary



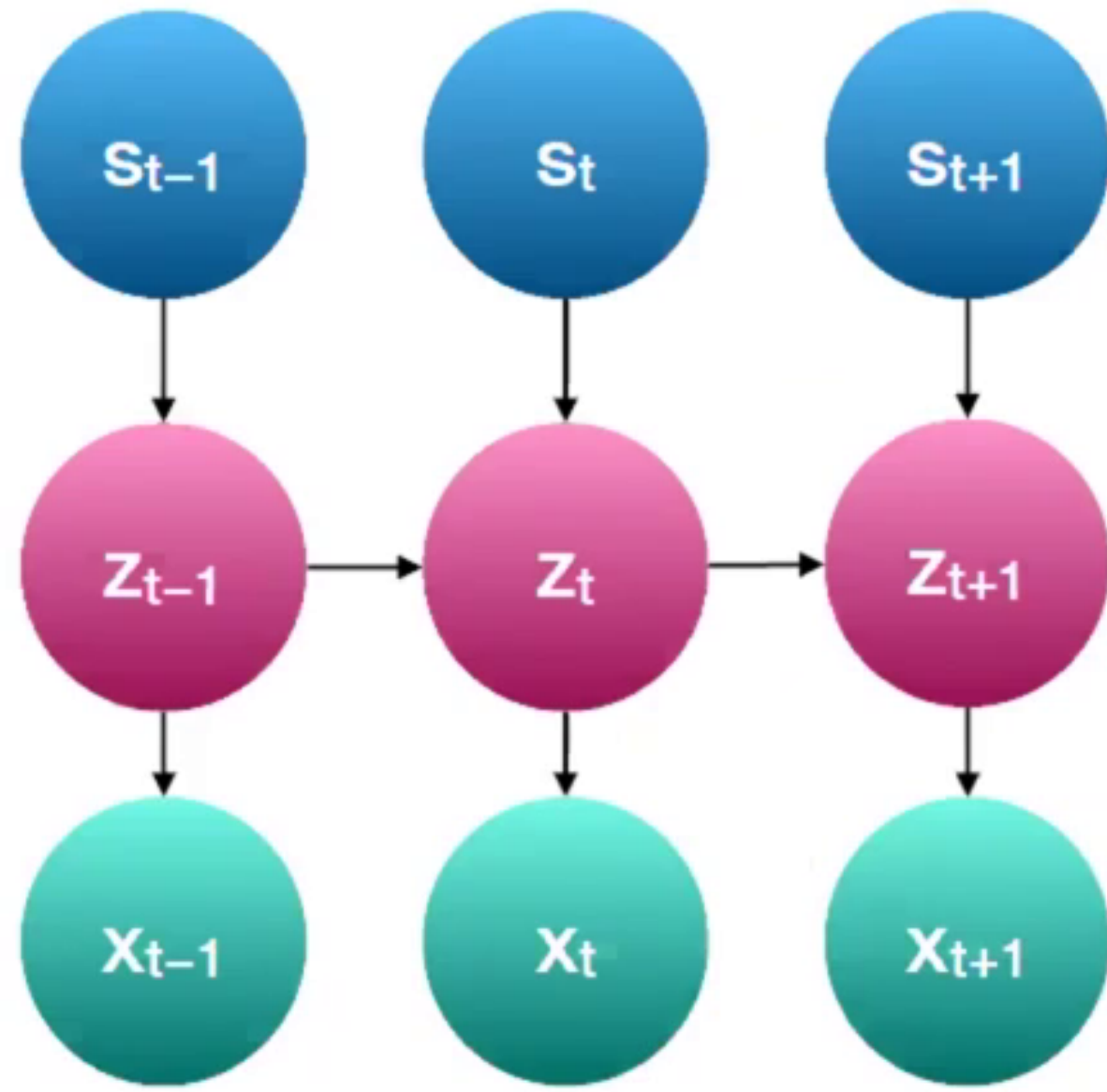
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Piece-wise Linear Recurrent Neural Network



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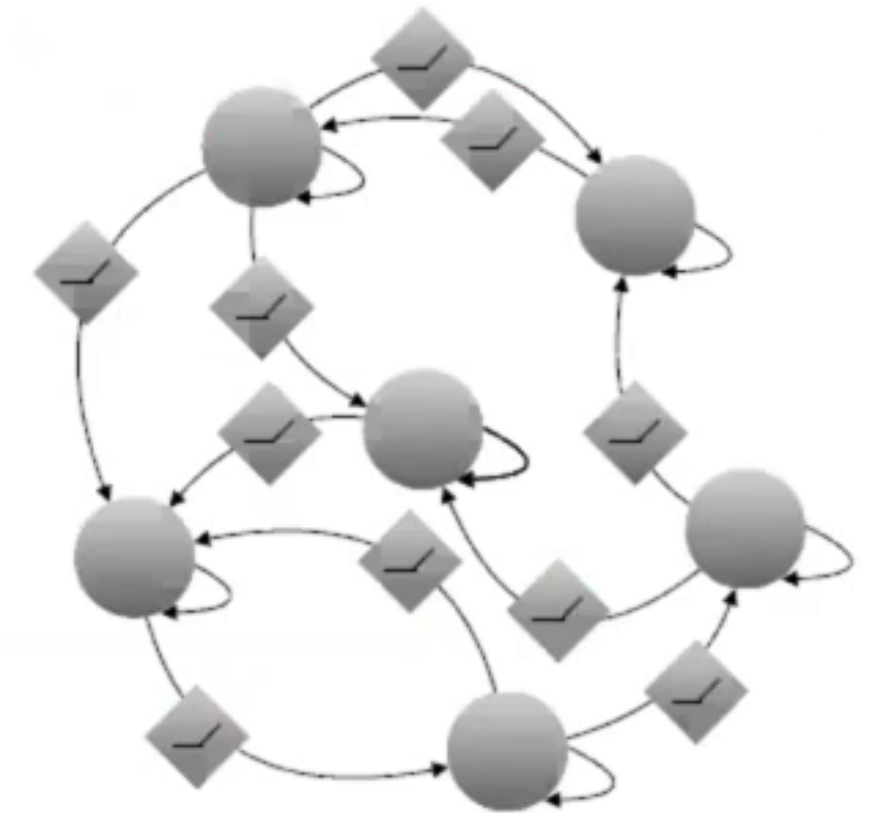
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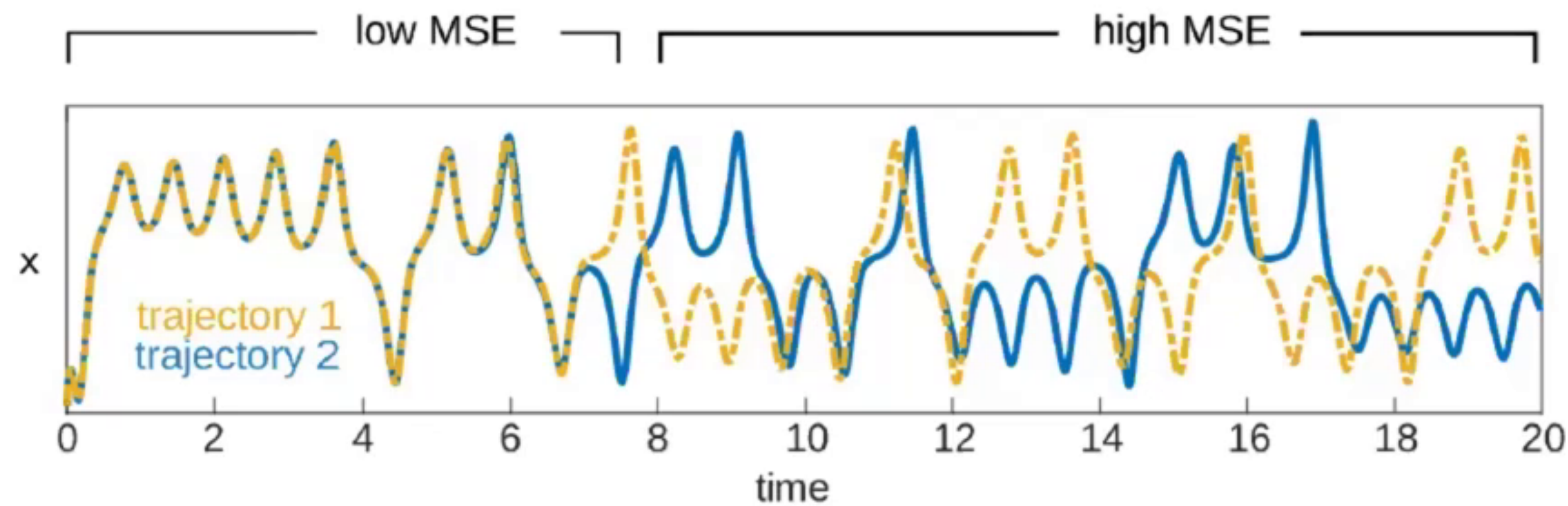
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Identifying nonlinear dynamical systems via generative recurrent neural networks with applications to fMRI. PLoS Computational Biology, 15.



Global vs. Local Metrics

Kullback-Leibler Divergence vs. Mean Squared Error



low $\tilde{K}L_x = .06$, high MSE = 2.48

high $\tilde{K}L_x = .71$, low MSE = 1.40

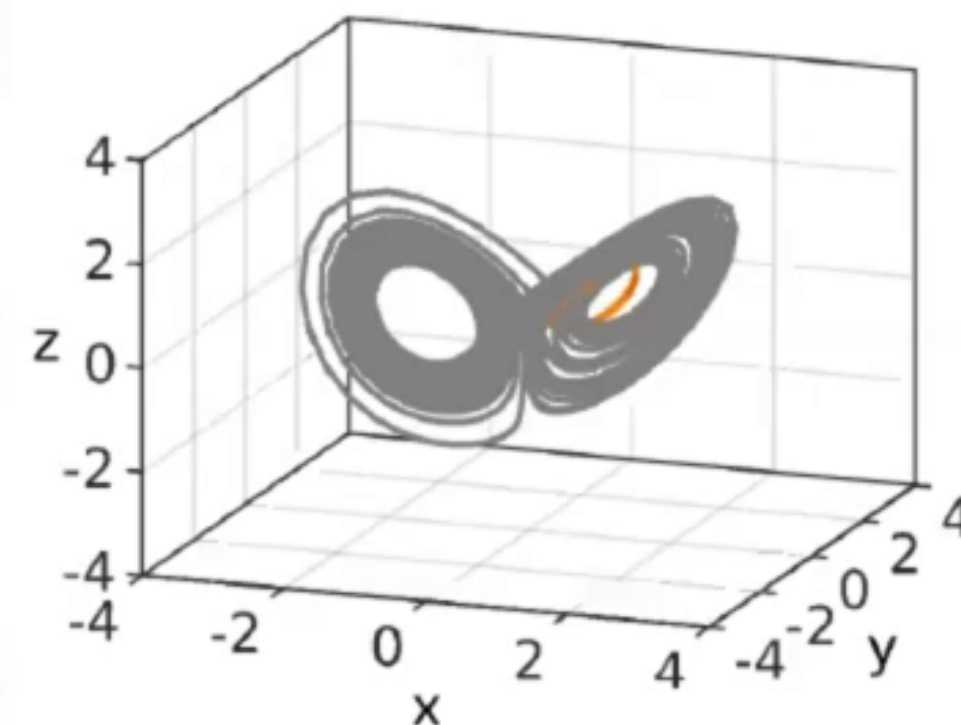
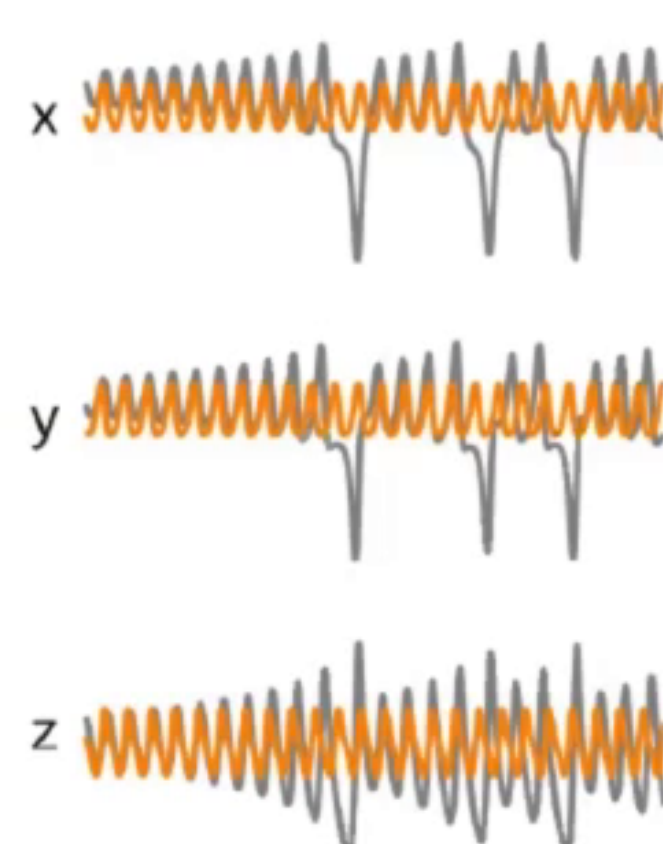
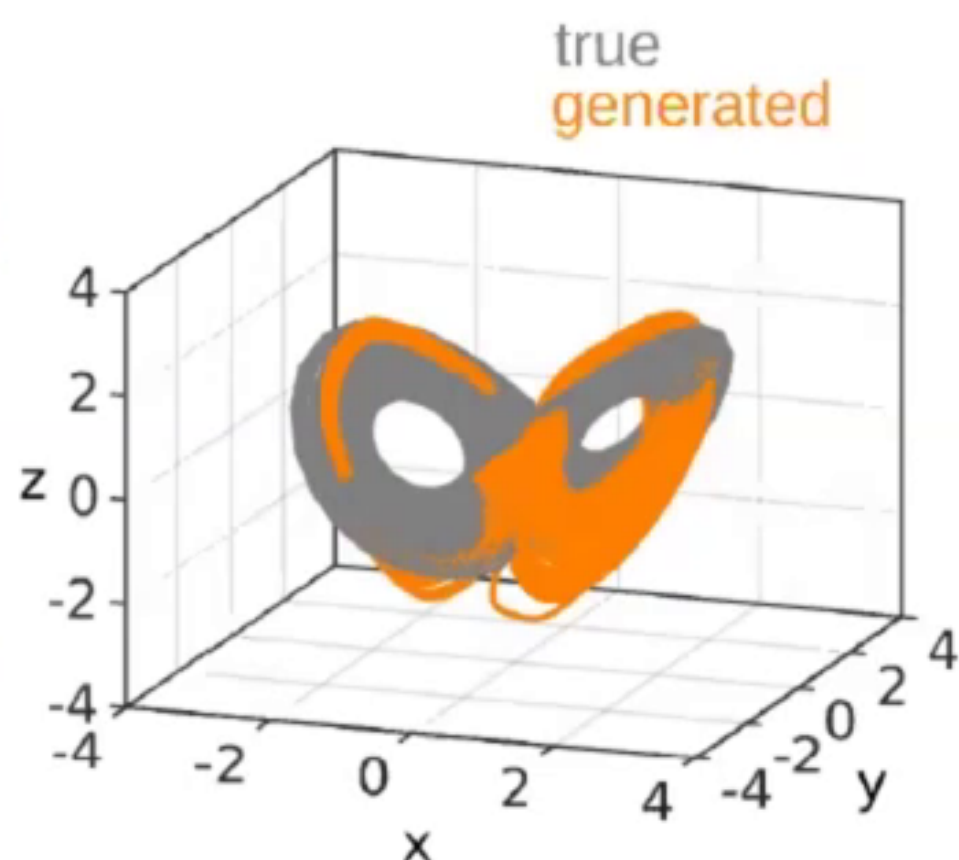
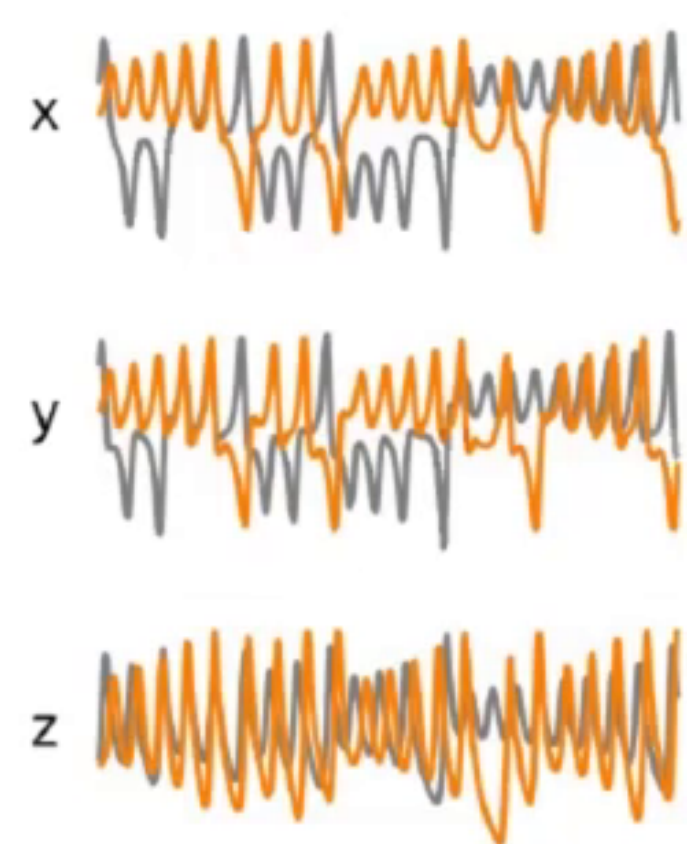


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