

Probabilistic Foundation and Opportunities of Diffusion Models

Mengdi Wang & Minshuo Chen

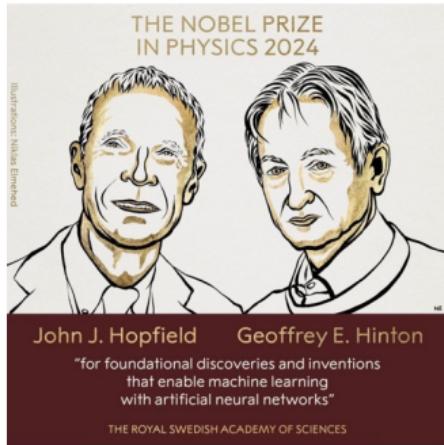


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AI Comes to The Nobels



"For me, this story highlights how far **AI** has come and
how much more potential there is to explore."

Millennium Growth of AI



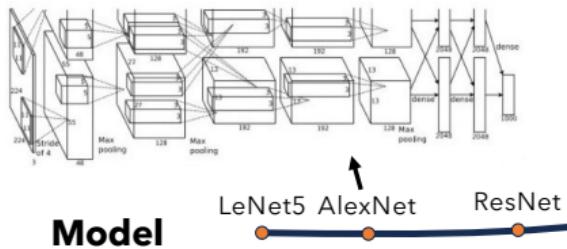
Discriminative AI



-- Thanks to blogs by Rockwell Anyoha , Toloka Team and Rick Merritt

Millennium Growth of AI

(Krizhevsky et al., 2012)

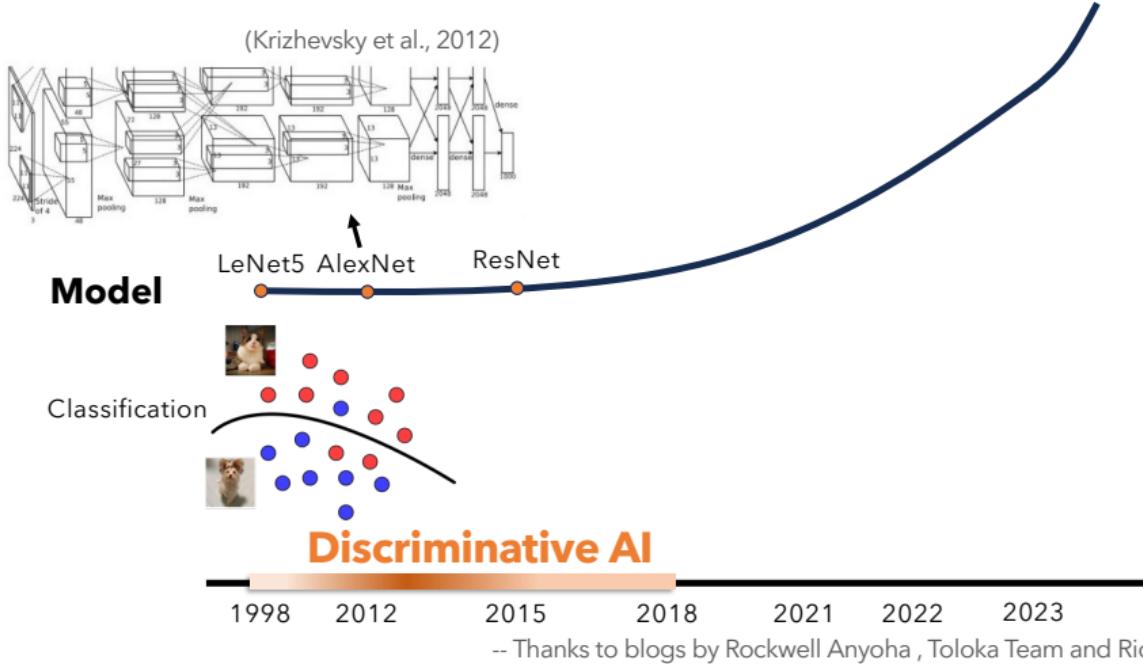


Discriminative AI

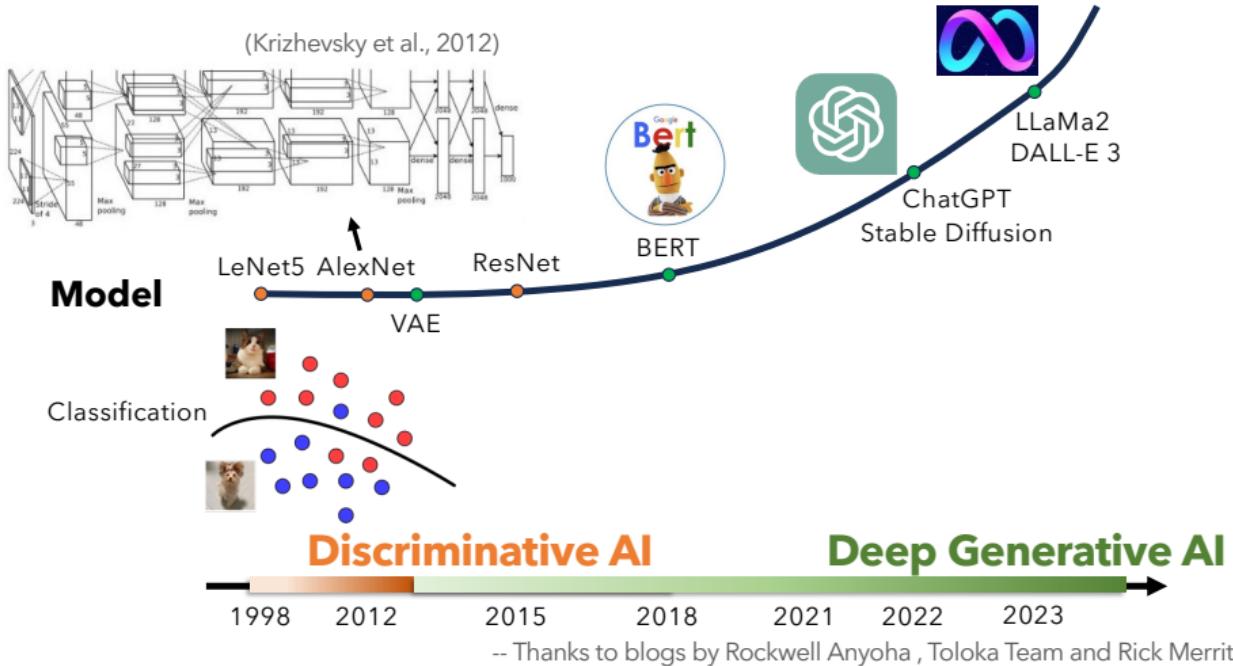
1998 2012 2015 2018 2021 2022 2023

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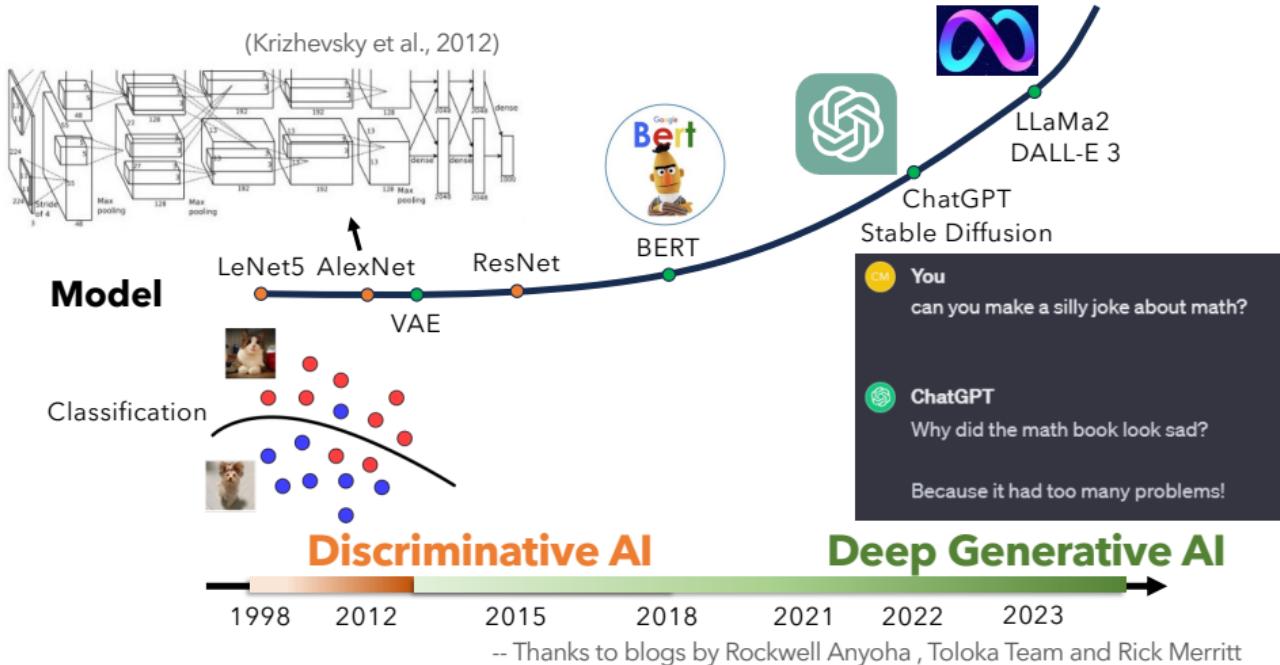
Millennium Growth of AI



Millennium Growth of AI



Millennium Growth of AI



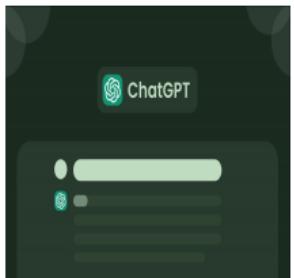
Transformative Power of Deep Generative AI



Transformative Power of Deep Generative AI



ChatGPT



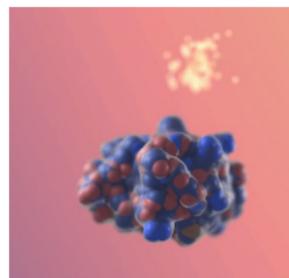
Language

Sora



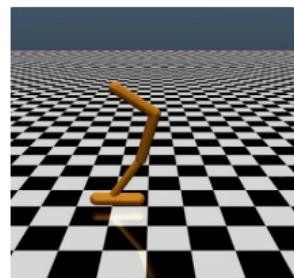
Video

RFDiffusion



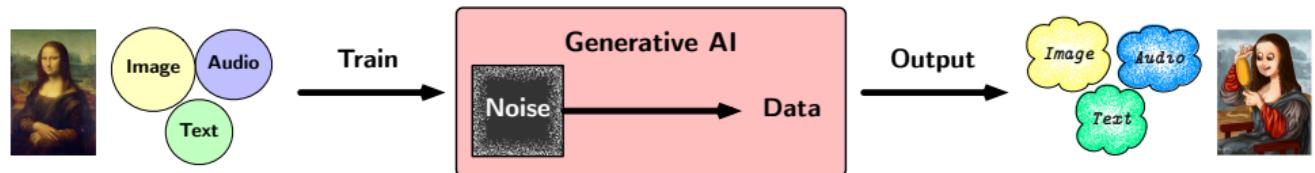
Biology

Decision Diffuser

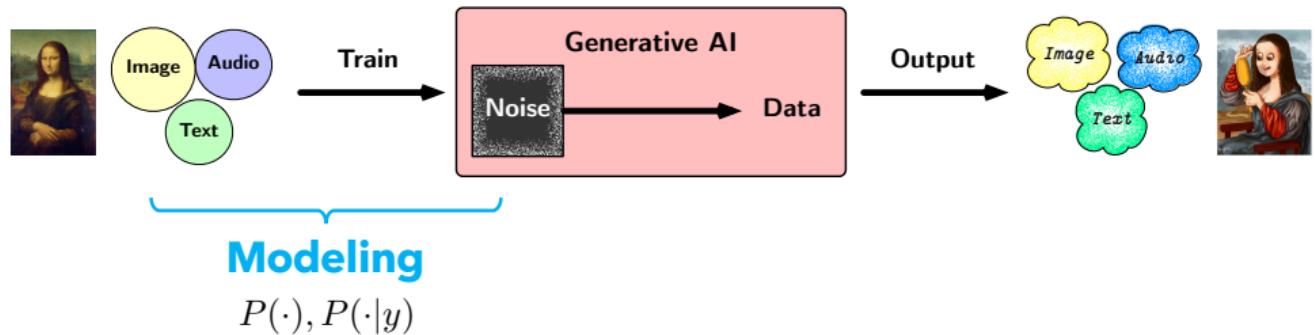


RL/control

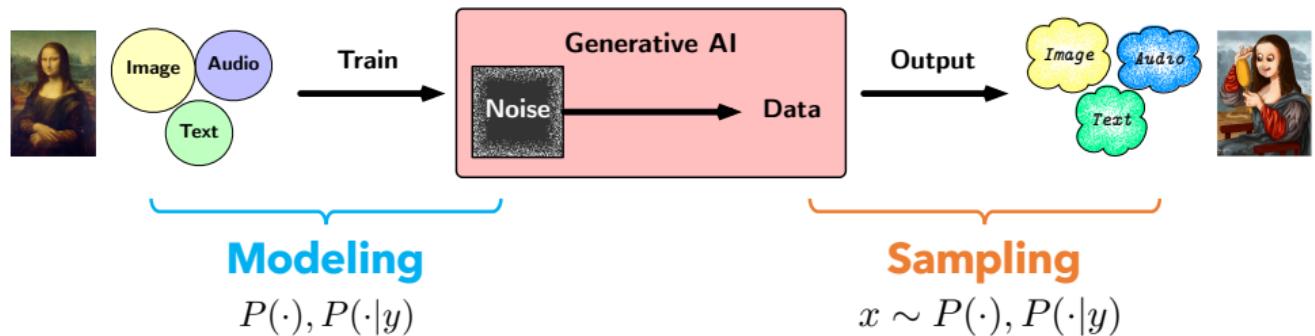
Constitution of Deep Generative AI



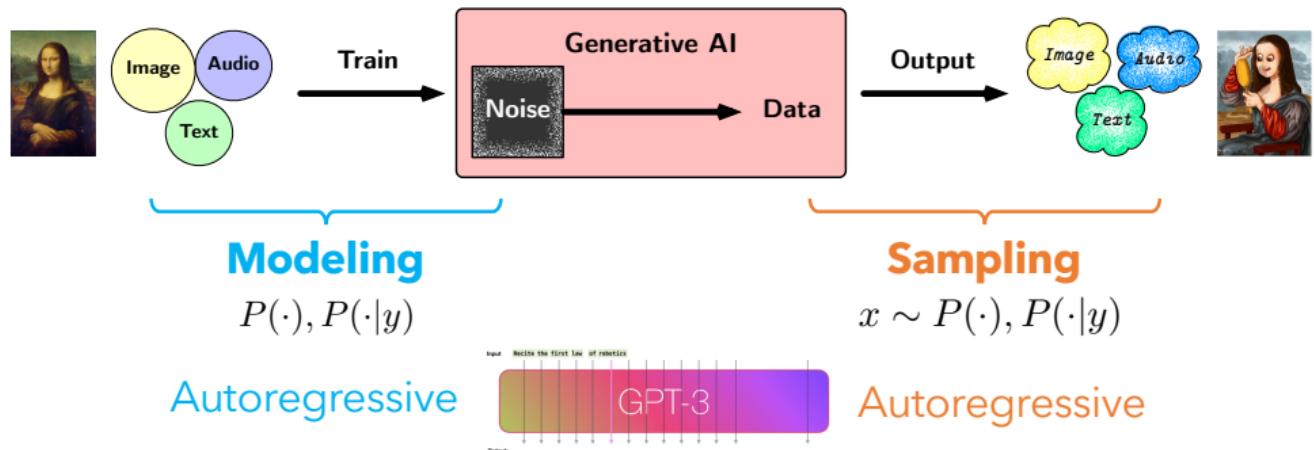
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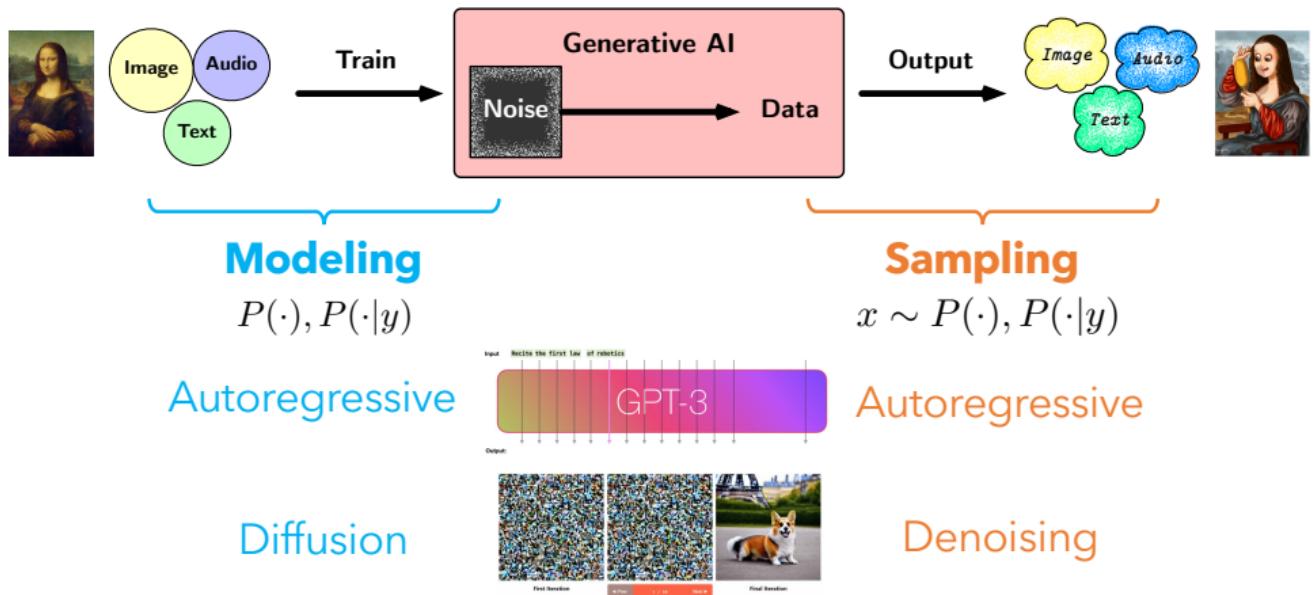
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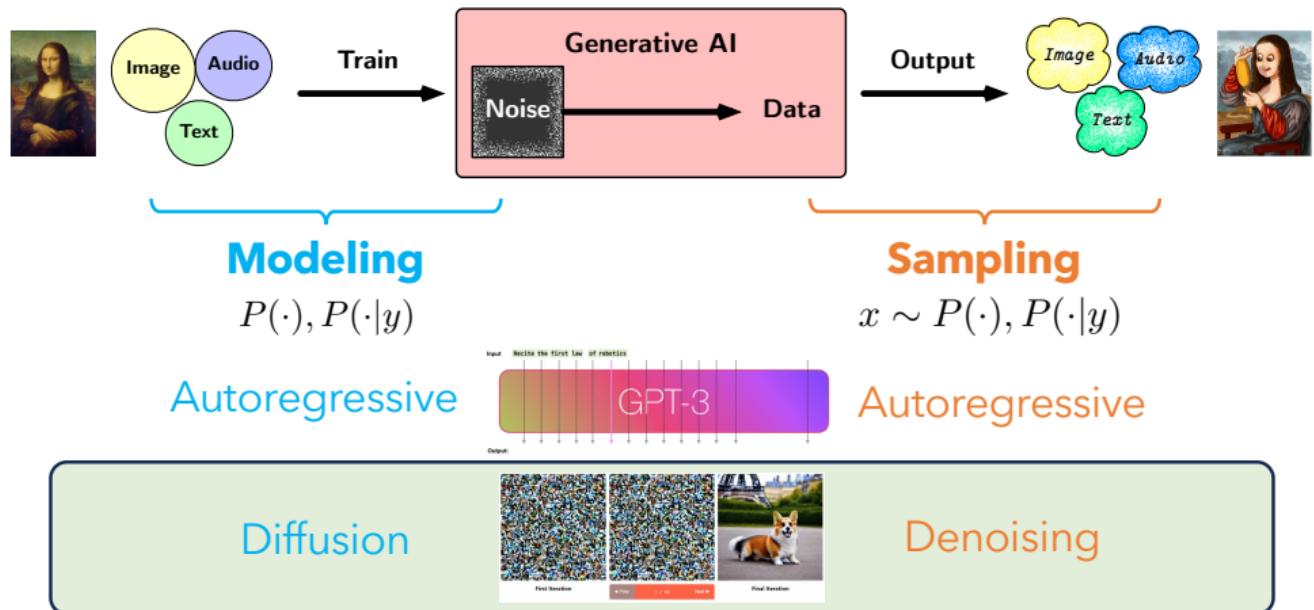
Constitution of Deep Generative AI



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Constitution of Deep Generative AI



New Promises of Diffusion Models

New Promises of Diffusion Models

Diffusion Models Beat GANs on Image Synthesis

Prafulla Dhariwal*
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SCORE-BASED GENERATIVE MODELING THROUGH STOCHASTIC DIFFERENTIAL EQUATIONS

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colorization. Combined with multiple architectural improvements, we achieve record-breaking performance for unconditional image generation on CIFAR-10 with an Inception score of 9.89 and FID of 2.20, a competitive likelihood of 2.99

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10M

Global users just two months after its release

270,000

Stable Diffusion's Discord channel Members

+170M

Images generated with Clipdrift SDK

400M

Images generated using Stability AI's API

New Promises of Diffusion Models

Generative AI imagines new protein structures

"FrameDiff" is a computational tool that uses generative AI to craft new protein structures, with the aim of accelerating drug development and improving gene therapy.

Rachel Gordon | MIT CSAIL

July 12, 2023



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Biology is a wondrous yet delicate tapestry. At the heart is DNA, the master weaver that encodes proteins, responsible for orchestrating the many biological functions that sustain life within the human body. However, our body is akin to a finely tuned instrument, susceptible to losing its harmony. After all, we're faced with an ever-changing and relentless natural world: pathogens, viruses, diseases, and cancer.

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New Promises of Diffusion Models

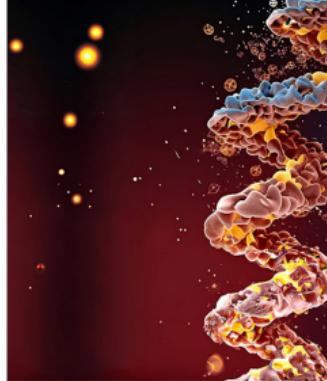
Generative AI imagines new Diffusion models are now turbocharging reinforcement learning systems

"FrameDiff" is a computational tool that uses protein structures, with the aim of accelerating improving gene therapy.

Rachel Gordon | MIT CSAIL

July 12, 2023

By Ben Dickson - March 4, 2024



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Biology is a wondrous yet delicate tapestry. At the heart of it all is the genome, which encodes proteins, responsible for orchestrating the millions of processes within the human body. However, our body is akin to a finely tuned machine that can lose its harmony. After all, we're faced with an ever-growing list of pathogens, viruses, diseases, and cancer. This article is part of our coverage of the latest in AI research.

Image generated with Bing Image Creator

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Discord Images generated with Clipdrift SDK. Images generated using Stability AI's API

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Rachel Gordon | MIT CSAIL
July 12, 2023



AniPortrait: Audio-Driven Synthesis of Photorealistic Portrait Animation

Published 3 days ago on May 3, 2024
By Kumal Kejriwal



Over the years, the creation of realistic and expressive portraits from static images and audio has found a range of applications including gaming, digital media, virtual reality, and a lot more. Despite its potential application, it is still difficult for developers to create frameworks capable of generating high-quality animations that maintain temporal consistency and are visually captivating. A major cause for the complexity is the need for intricate coordination of lip movements, head positions, and facial expressions to craft a visually compelling effect.

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Biology is a wondrous yet delicate tapestry. At the heart of this complexity lies the genome, which encodes proteins, responsible for orchestrating the myriad of processes within the human body. However, our body is akin to a finely tuned machine that requires constant maintenance and repair. This delicate balance can be disrupted by various factors, such as environmental pollutants, viruses, and mutations. When this balance is lost, it can lead to a range of diseases, from common colds to life-threatening conditions like cancer. The study of these complex systems is crucial for developing effective treatments and prevention strategies.

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New Promises of Diffusion Models

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turbocharging reinforcement learning systems

ARTIFICIAL INTELLIGENCE

AniPortrait: Audio-Driven Synthesis of Photorealistic Portrait Animation

Rachel Gordon | MIT CSAIL
July 12, 2023

By Ben Dickson - March 4, 2024

Published 3 days ago on May 3, 2024
By Kunal Kejriwal



Diffusion Models Are Real-Time Game Engines

Dani Valevski*

Google Research

Yaniv Leviathan*

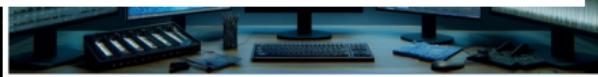
Google Research

Moab Arar*†

Tel Aviv University

Shlomi Fruchter*

Google DeepMind



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Biology is a wondrous yet delicate tapestry. At the heart of it all is the genome, which encodes proteins, responsible for orchestrating the myriad processes within the human body. However, our body is akin to a finely tuned machine; it needs to maintain a delicate balance between its various systems. Any disruption can lead to a loss of harmony. After all, we're faced with an ever-evolving threat from various pathogens, viruses, diseases, and cancer.

Over the years, the creation of realistic and expressive portraits animations from static images and audio has found a range of applications including gaming, digital media, virtual reality, and a lot more. Despite its potential application, it is still difficult for developers to create frameworks capable of generating high-quality animations that maintain temporal consistency and are visually captivating. A major cause for the complexity is the need for intricate coordination of lip movements, head positions, and facial expressions to craft a visually compelling effect.

This article is part of our coverage of the latest developments in AI and machine learning.

Outline

- A probabilistic foundation for diffusion models
- How diffusion models capture diverse data
- How to leverage diffusion models
- Inspirations and future directions

Outline



Chen et al., "Challenges and Opportunities of Diffusion Models for Generative AI", NSR 2024

- A probabilistic foundation for diffusion models
- How diffusion models capture diverse data
- How to leverage diffusion models
- Inspirations and future directions

Outline



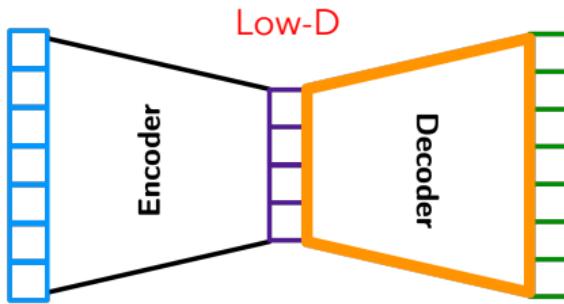
Chen et al., "Challenges and Opportunities of Diffusion Models for Generative AI", NSR 2024

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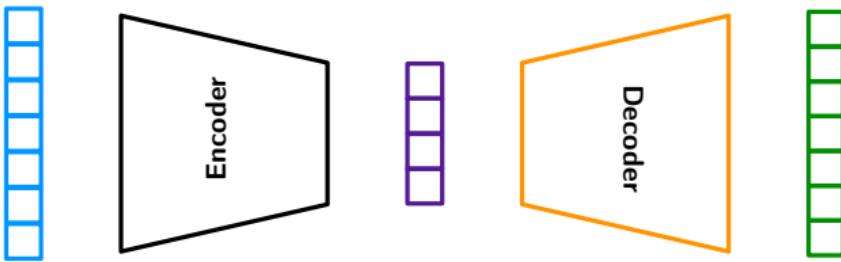
Foundation of Diffusion Models

Early Model of Deep Generative AI

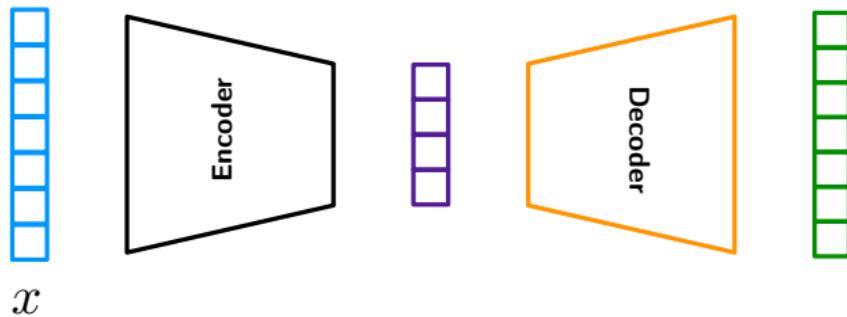


VAE (Kingma & Welling, 2013)

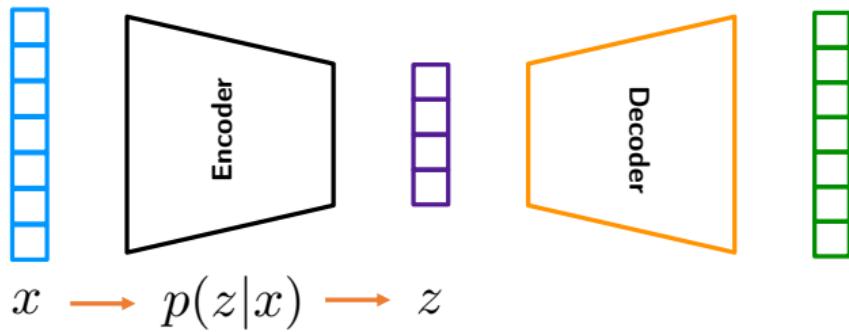
Dissemable VAE



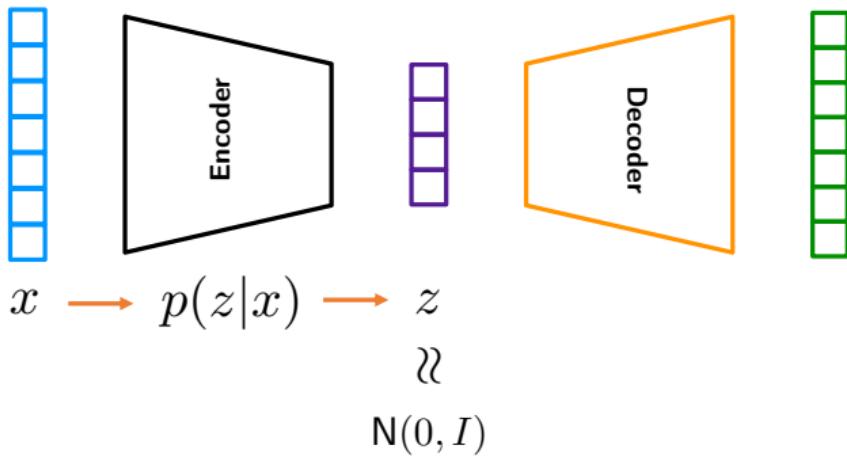
Dissemable VAE



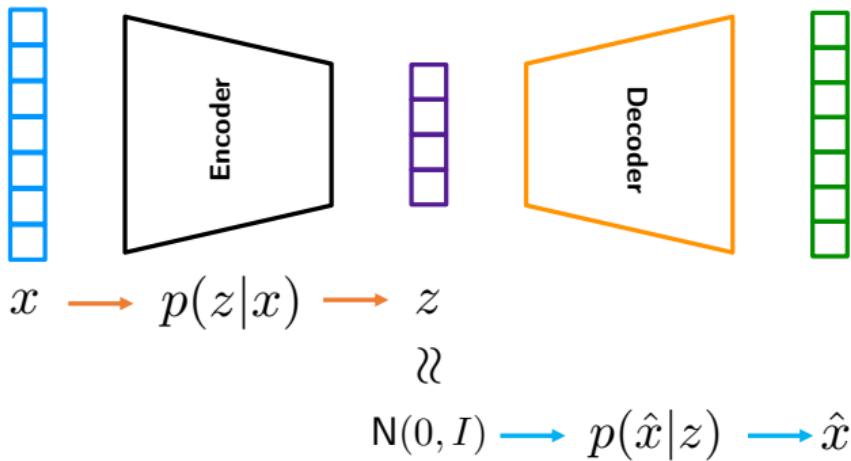
Dissemable VAE



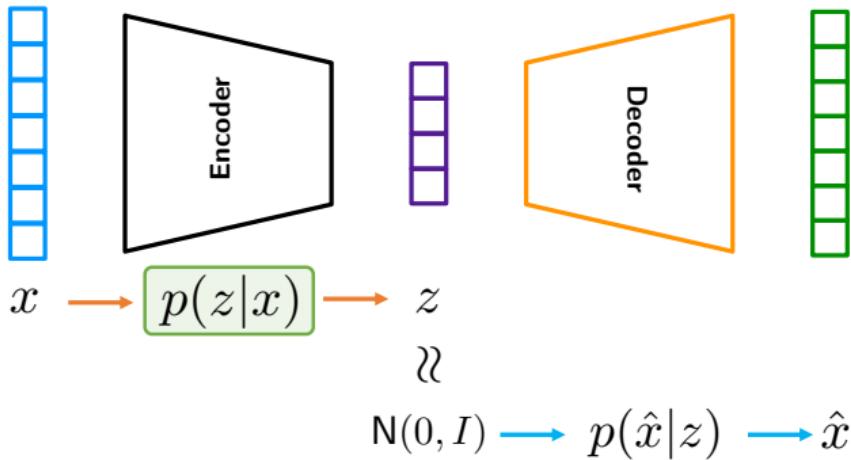
Dissemable VAE



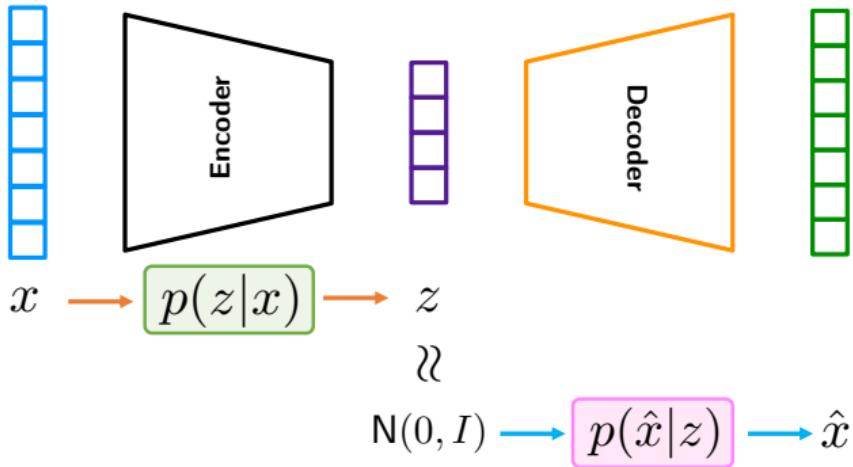
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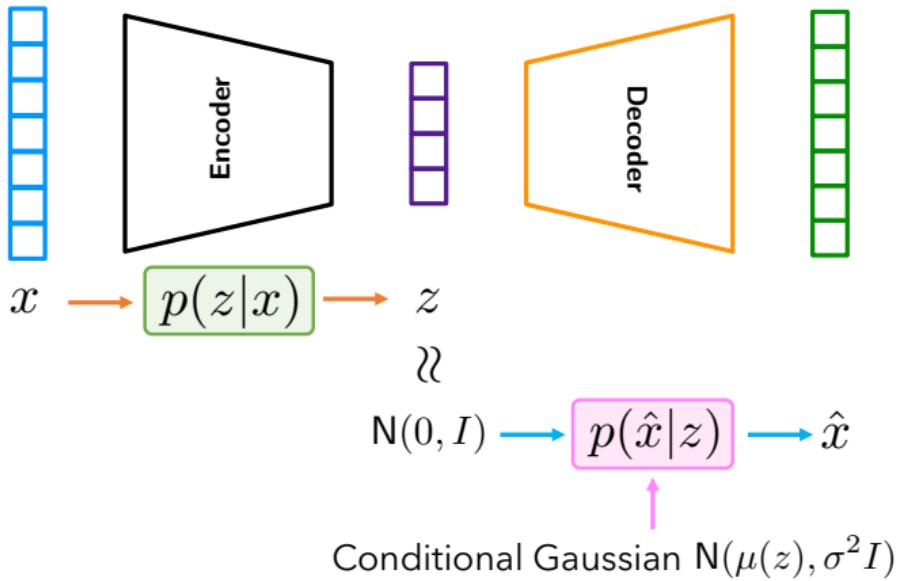
Dissemable VAE



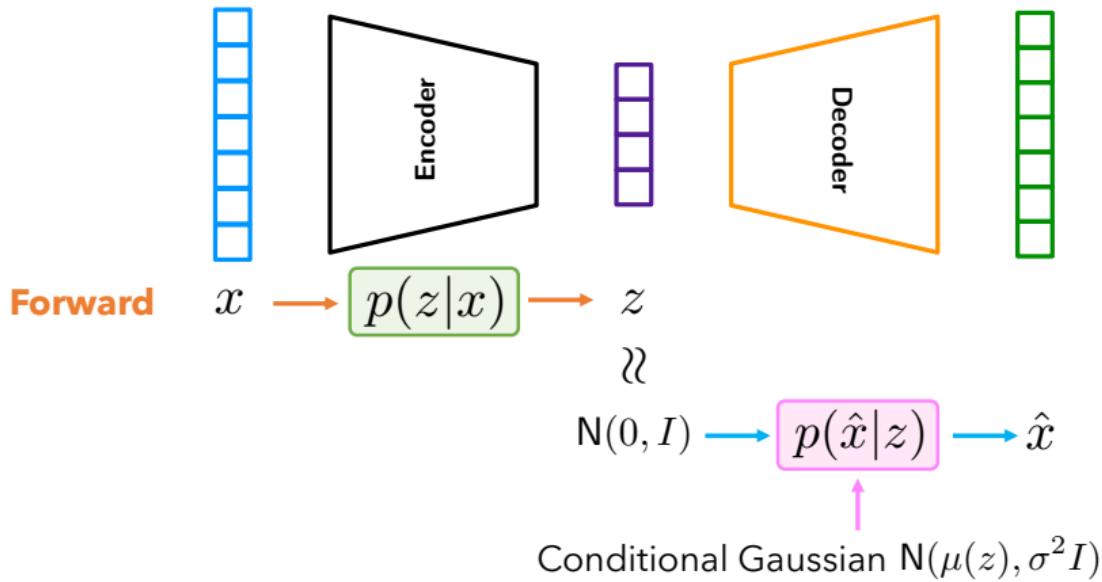
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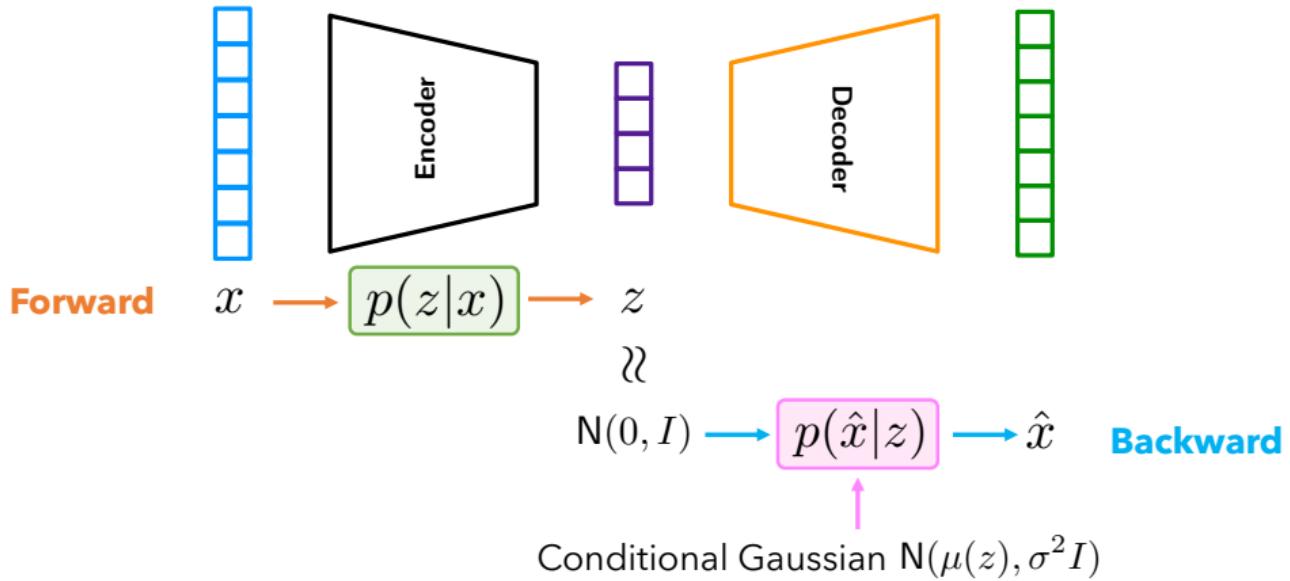
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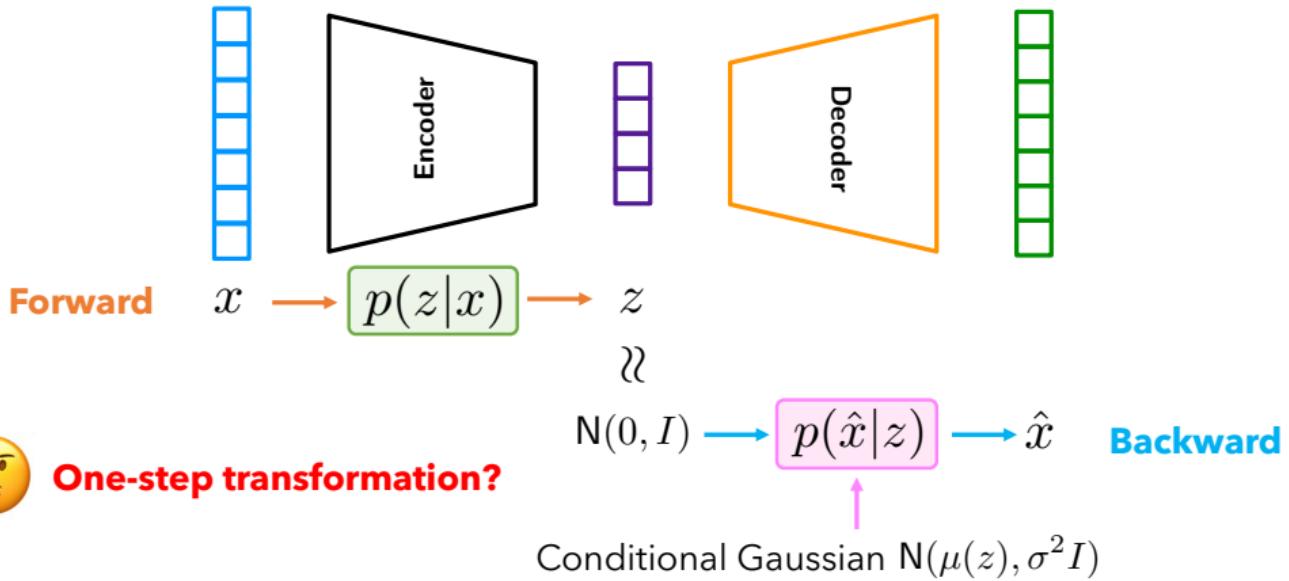
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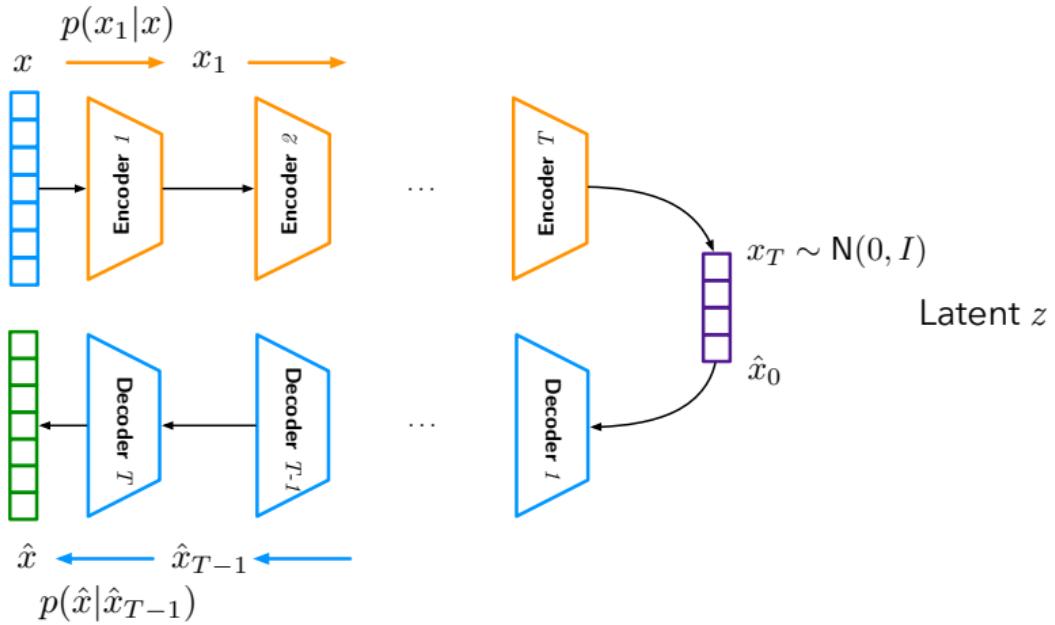
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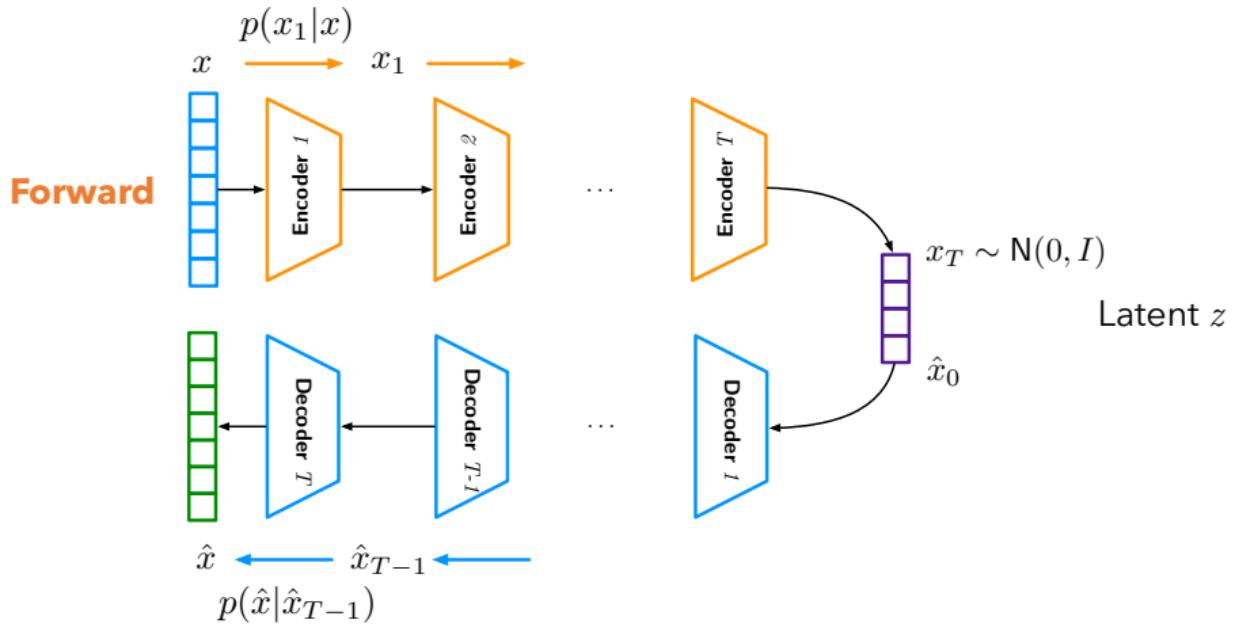
Dissemable VAE



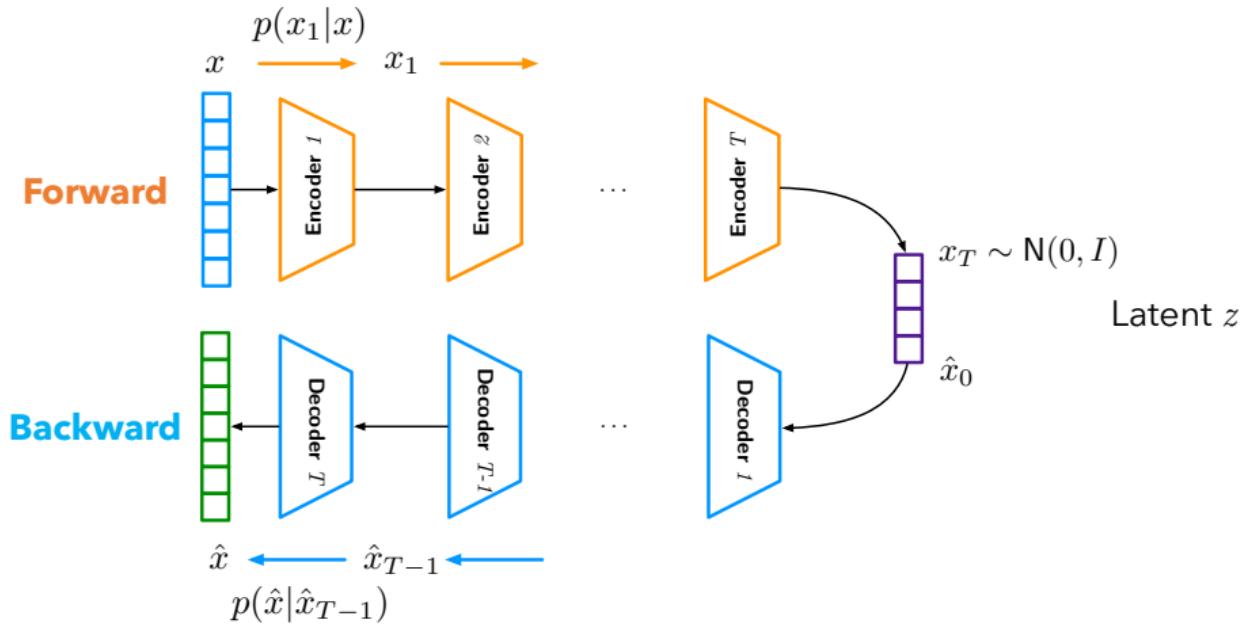
Let's Insert Some Intermediate Layers



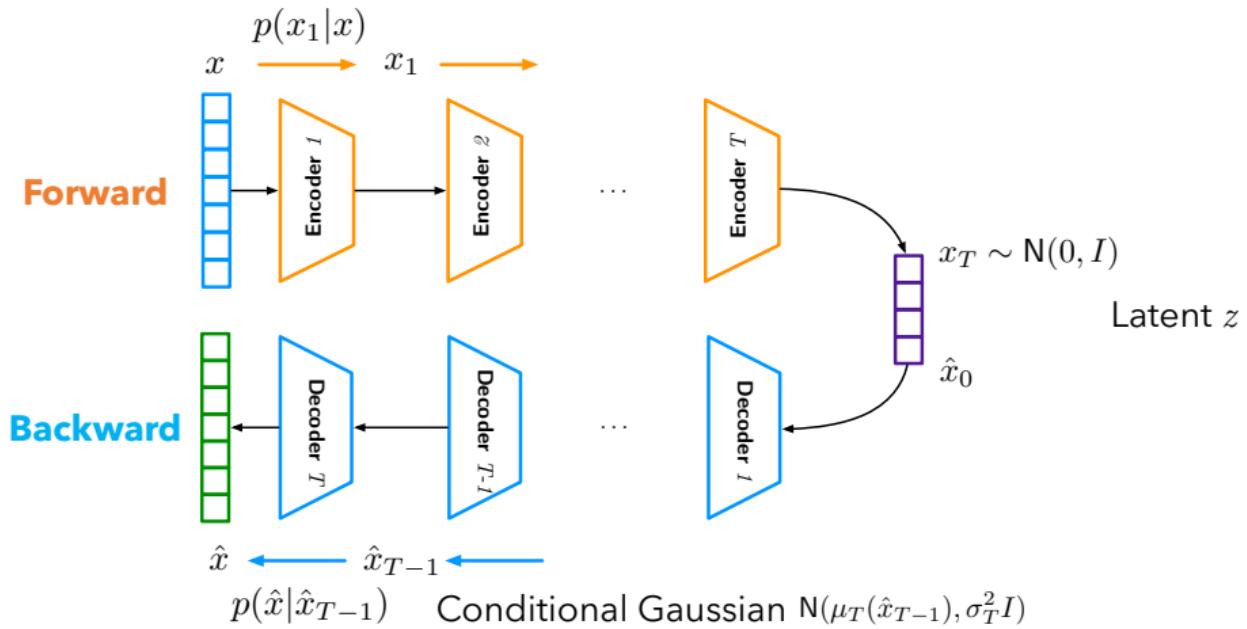
Let's Insert Some Intermediate Layers



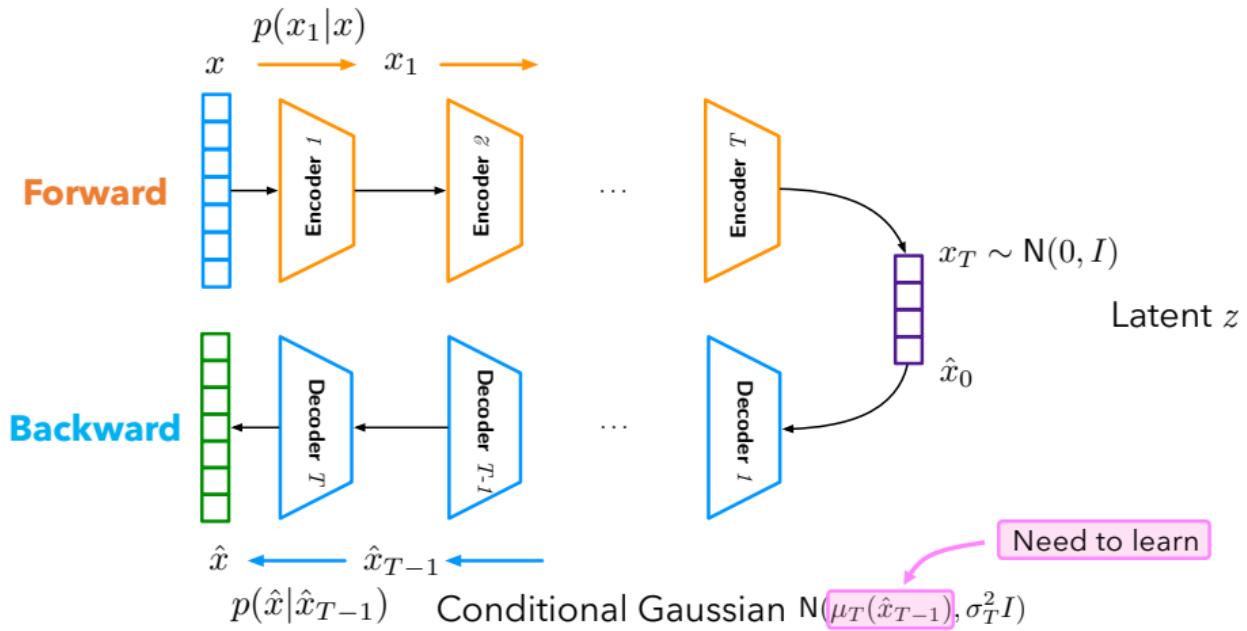
Let's Insert Some Intermediate Layers



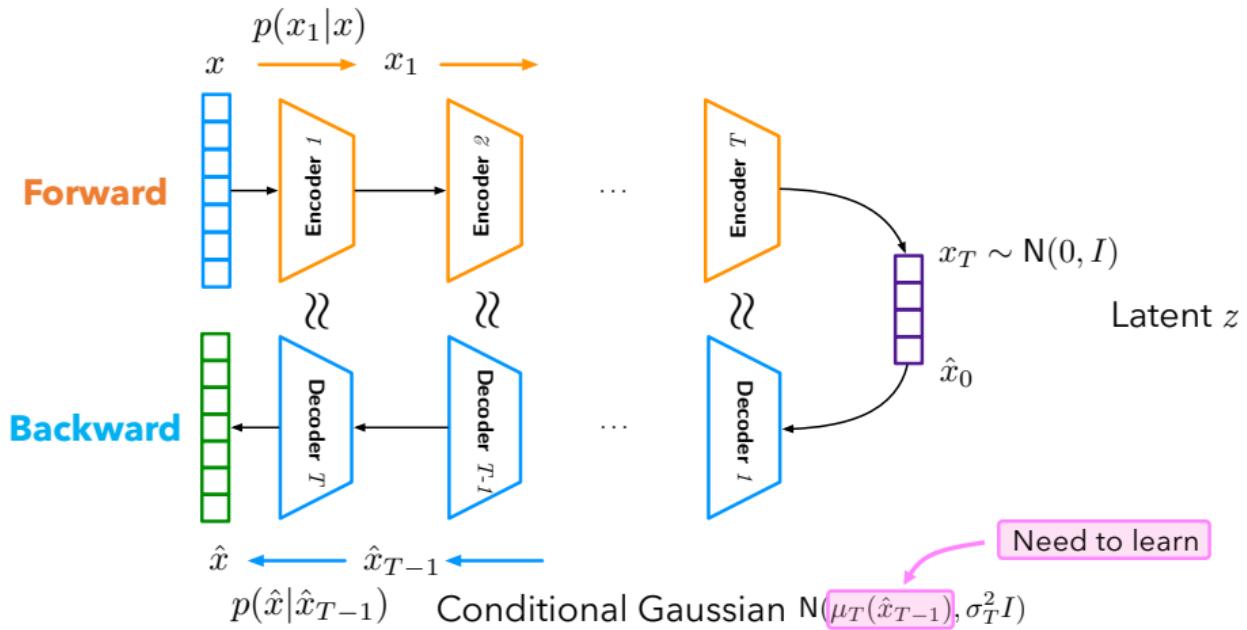
Let's Insert Some Intermediate Layers



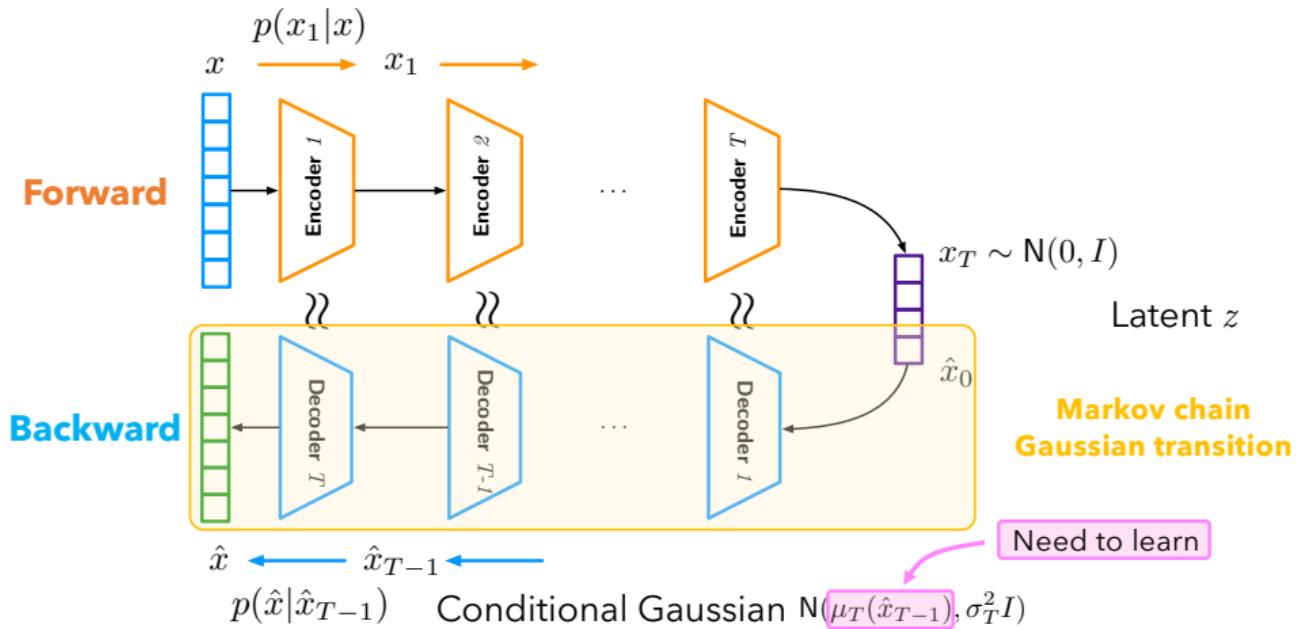
Let's Insert Some Intermediate Layers



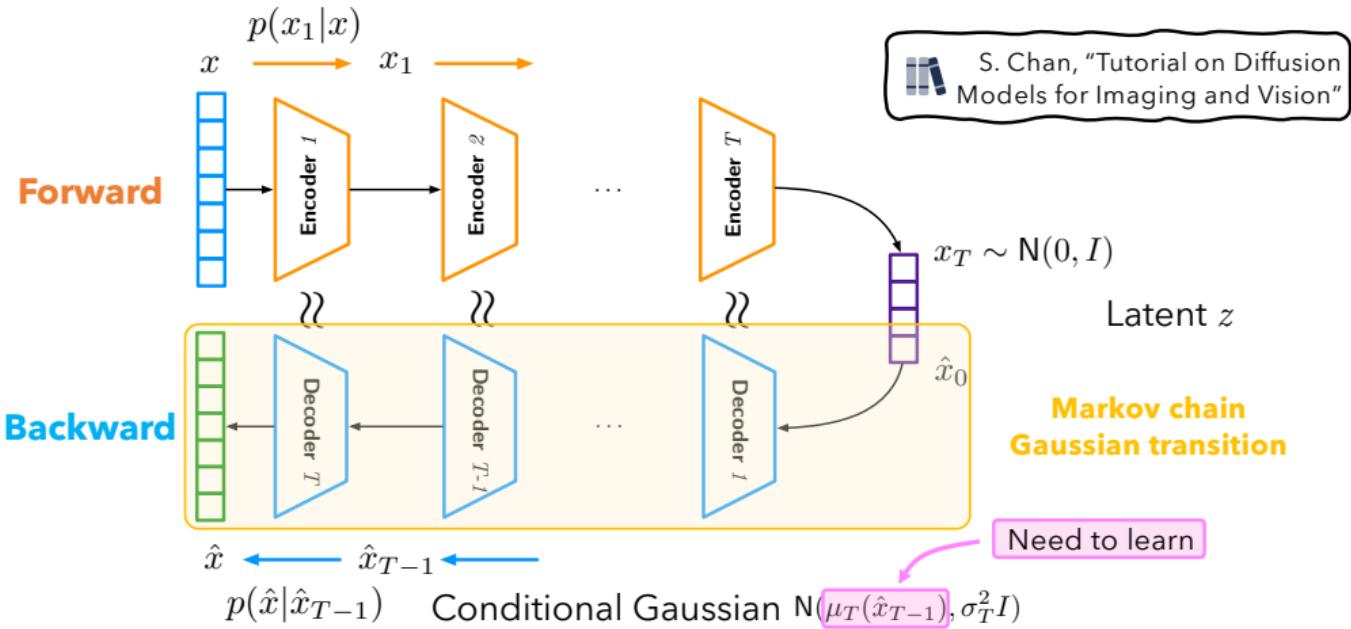
Let's Insert Some Intermediate Layers



Let's Insert Some Intermediate Layers



Let's Insert Some Intermediate Layers



A Revolution - Diffusion Model

- Sequential transformation in high-D

Noise

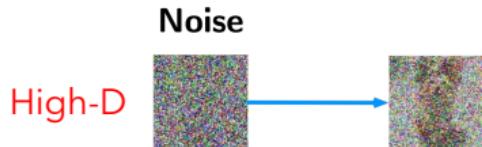
High-D



(Sohl-Dickstein et al., 2015)
(Song and Ermon, 2019)
(Ho et al., 2020)

A Revolution - Diffusion Model

- Sequential transformation in high-D



(Sohl-Dickstein et al., 2015)
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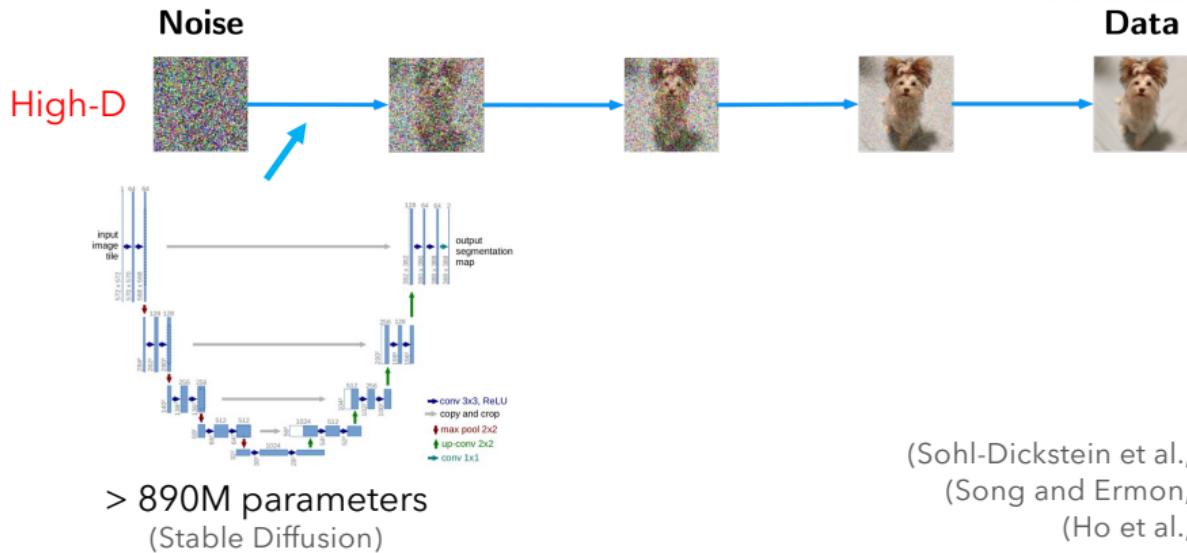
- Sequential transformation in high-D



(Sohl-Dickstein et al., 2015)
(Song and Ermon, 2019)
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A Revolution - Diffusion Model

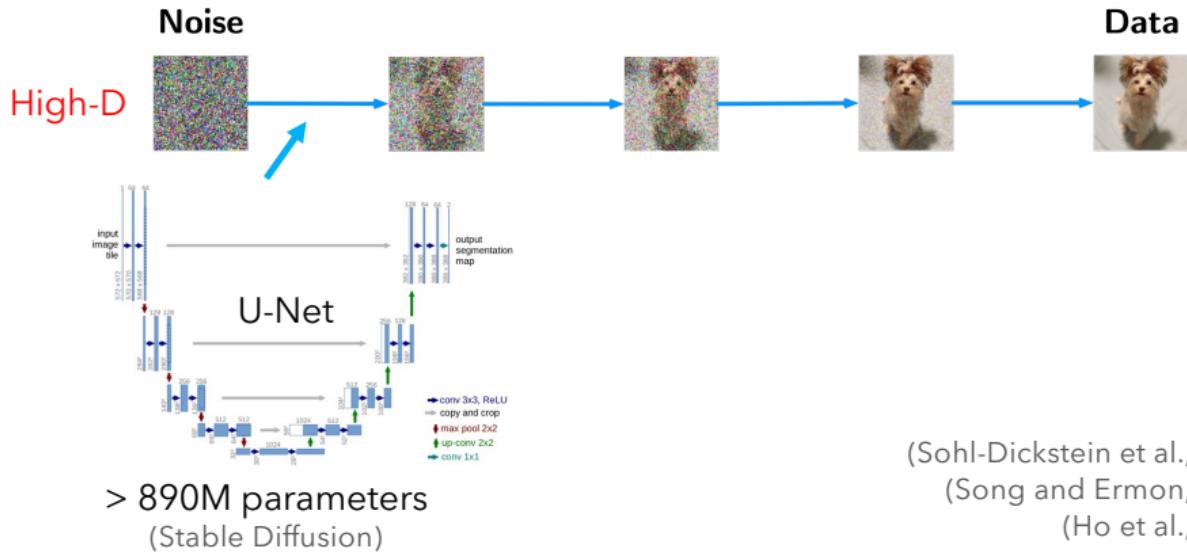
- Sequential transformation in high-D



(Sohl-Dickstein et al., 2015)
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A Revolution - Diffusion Model

- Sequential transformation in high-D



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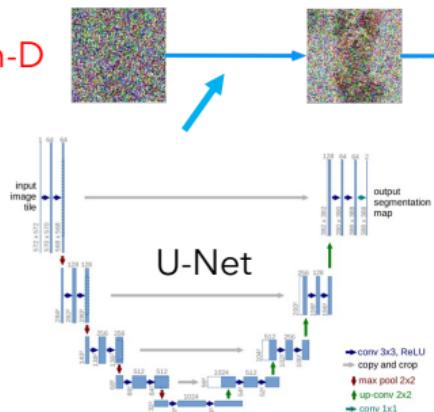
A Revolution - Diffusion Model

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Noise

High-D

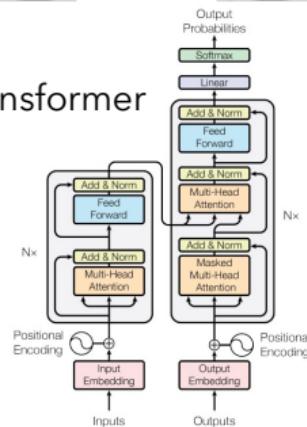


> 890M parameters
(Stable Diffusion)

Data

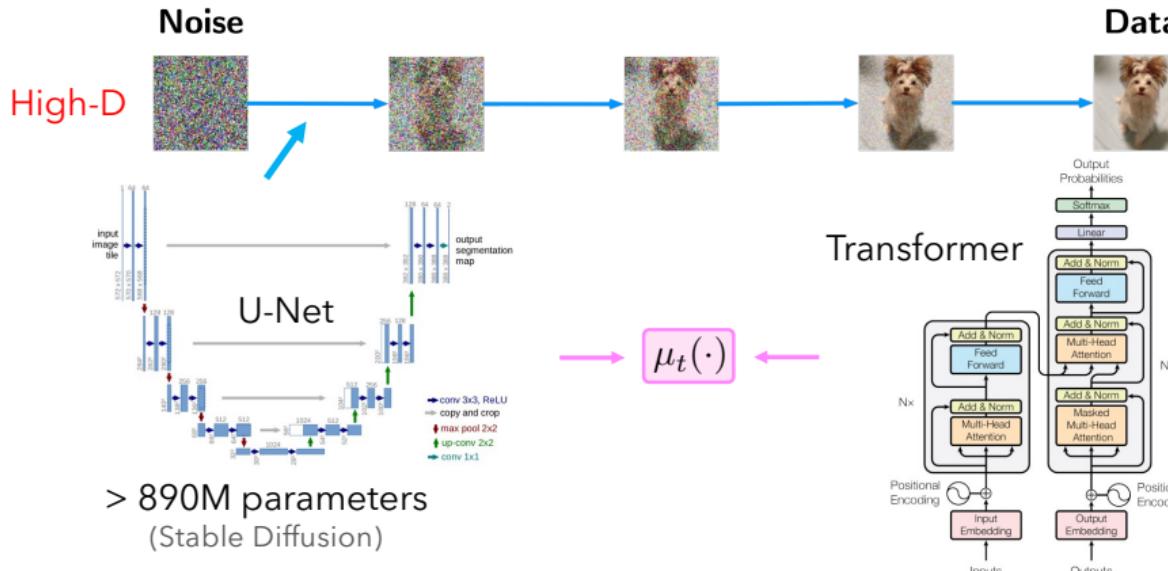
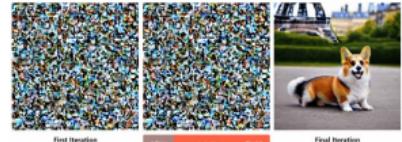


Transformer



A Revolution - Diffusion Model

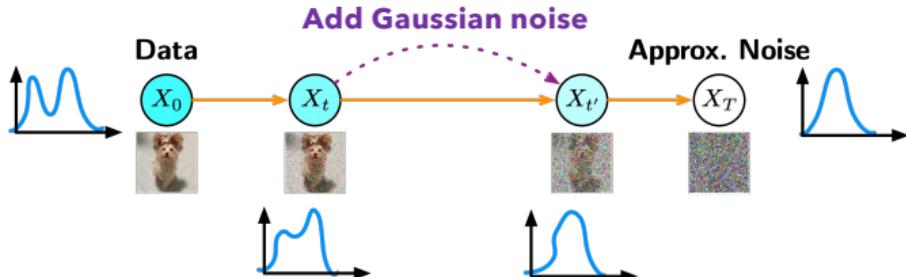
- Sequential transformation in high-D



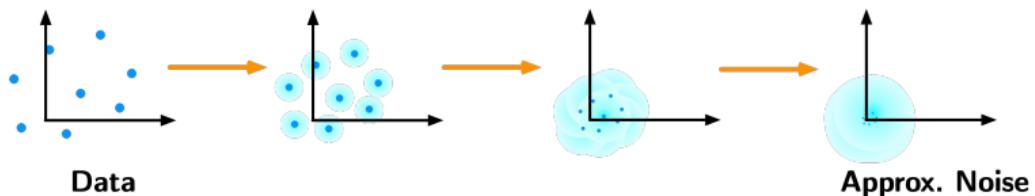
Forward Process - Noise Corruption

- Noise corruption process $dX_t = -\frac{1}{2}X_t dt + dW_t$

Insert infinitely many intermediate layers!

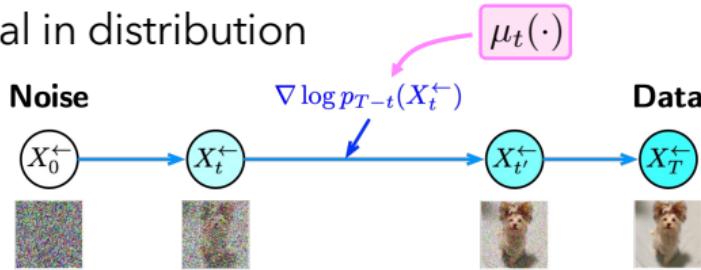


- The noise corruption



Backward Process - Sample Generation

- Time reversal in distribution



- The math (Anderson, 1982; Haussmann and Pardoux, 1986)

Forward

$$dX_t = -\frac{1}{2}X_t dt + dW_t$$

Backward

$$dX_t^{\leftarrow} = \left[\frac{1}{2}X_t^{\leftarrow} + \underbrace{\nabla \log p_{T-t}(X_t^{\leftarrow})}_{\text{Score Function}} \right] dt + d\bar{W}_t$$

Brownian

Theorem. Let x, u be the process described by (3.3), and suppose $f(\cdot, \cdot)$ and $g(\cdot, \cdot)$ are such as to guarantee the existence of the probability density $p(x, t)$ for $0 < t < T$ as a smooth and unique solution of its associated Kolmogorov equation. Suppose further that an n -vector process u_t is defined by $u_n = 0$ and

$$du_t^n = du_t^{n-1} + \frac{1}{p(x, t)^2} \frac{\partial}{\partial x_i} [p(x, t)u_i^n(x, t)] dt, \quad (3.10)$$

and that the forward Kolmogorov equation associated with the joint process (x, u) yields a smooth and unique solution in $L^2(\Omega)$ for $p(x, u, t)$ and in $L^1(\Omega)$ for $p(x, u, t|u_n, t)$. Then

- x and $u - u_n$ are independent for all $t > n > 0$.
- (x, u) is \mathcal{F}_t -adapted with respect to which x_i for $i > t$ and u_i for $i > t$ are random variables (3.4) and (3.5) hold.
- A reverse time model for x is defined by

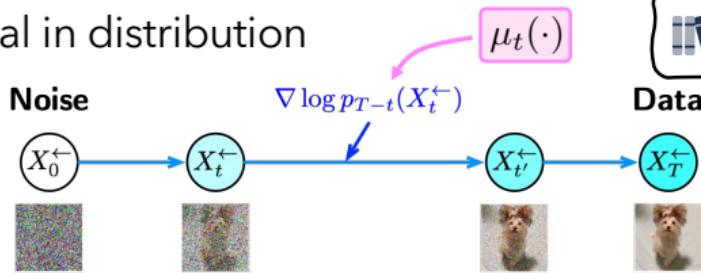
$$dx_i = f(x_i, t) dt + g(x_i, t) d\bar{W}_i, \quad (3.11)$$

where

$$\bar{f}'(x_n, t) = f'(x_n, t) - \frac{1}{p(x_n, t)} \frac{\partial}{\partial x_i} [p(x_n, t)g^B(x_n, t)g^B(x_n, t)], \quad (3.12)$$

Backward Process - Sample Generation

- Time reversal in distribution



Tang & Zhao, "Score-based Diffusion Models via SDE"

- The math (Anderson, 1982; Haussmann and Pardoux, 1986)

Forward

$$dX_t = -\frac{1}{2}X_t dt + dW_t$$

Backward

$$dX_t^{\leftarrow} = \left[\frac{1}{2}X_t^{\leftarrow} + \underbrace{\nabla \log p_{T-t}(X_t^{\leftarrow})}_{\text{Score Function}} \right] dt + d\bar{W}_t$$

Brownian

Theorem. Let x, u be the process described by (3.3), and suppose $f(\cdot, \cdot)$ and $g(\cdot, \cdot)$ are such as to guarantee the existence of the probability density $p(x, t)$ for $t < t$ as a smooth and unique solution of its associated Kolmogorov equation. Suppose further that an n -vector process u_t is defined by $u_n = 0$ and

$$du_t^n = du_t^{n-1} + \frac{1}{p(x, t)} \frac{\partial}{\partial x_i} [p(x, t)u_i^n(x, t)] dt, \quad (3.10)$$

and that the forward Kolmogorov equation associated with the joint process (x, u) yields a smooth and unique solution in $L^2(\Omega)$ for $p(x, u, t)$ and in $L^1(\Omega)$ for $p(x, u, t|u_n, t)$. Then

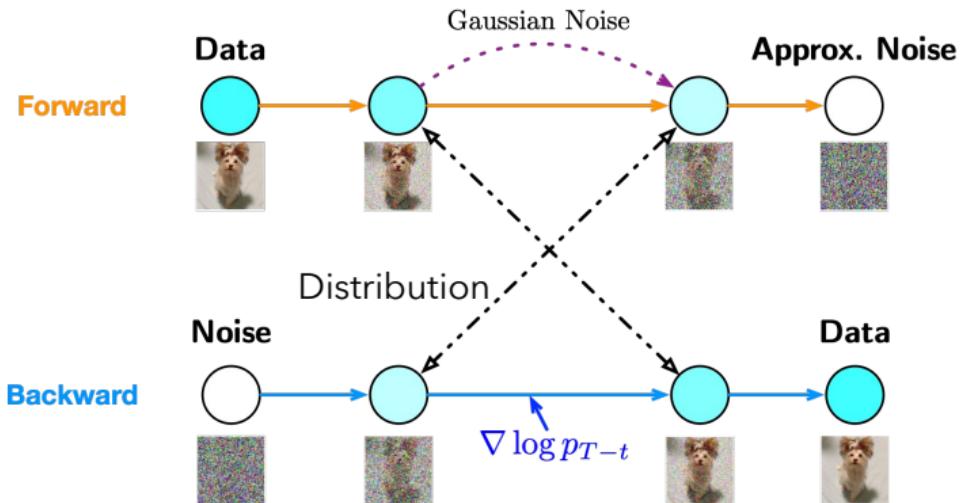
- x and $u_n - u_i$ are independent for all $i > n > n_0$.
- x and $u_n - u_i$ are independent with respect to which x_i , for $i > n$ and u_i , for $i > n$ are random variables.
- A reverse time model for x is defined by

$$dx_i = f(x_i, t) dt + g(x_i, t) d\tilde{W}_i, \quad (3.11)$$

where

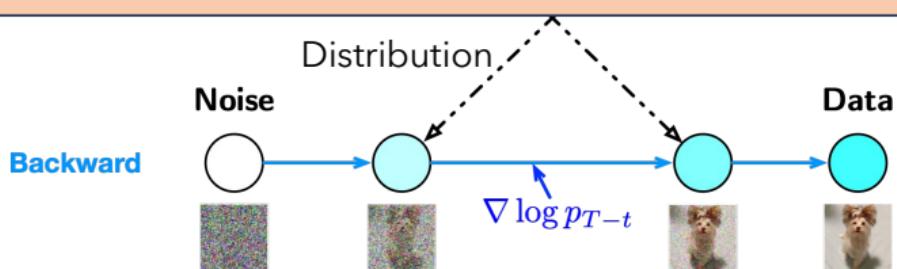
$$\tilde{f}(x_n, t) = f'(x_n, t) - \frac{1}{p(x_n, t)} \frac{\partial}{\partial x_i} [p(x_n, t)g^n(x_n, t)g^B(x_n, t)g^B(x_n, t)], \quad (3.12)$$

Forward and Backward Coupling



Forward and Backward Coupling

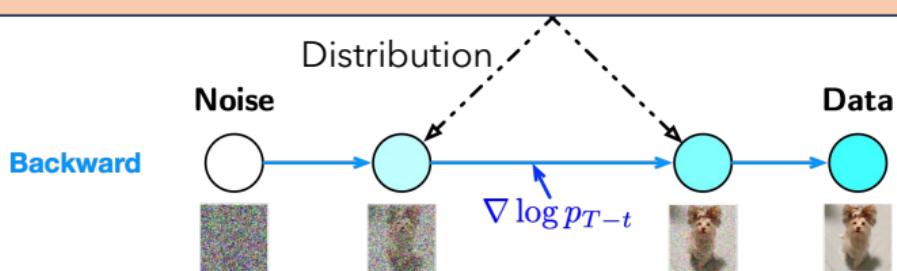
Training $\int_0^T \mathbb{E}_{x_t} [\|\nabla \log p_t(x_t) - s(x_t, t)\|_2^2] dt$



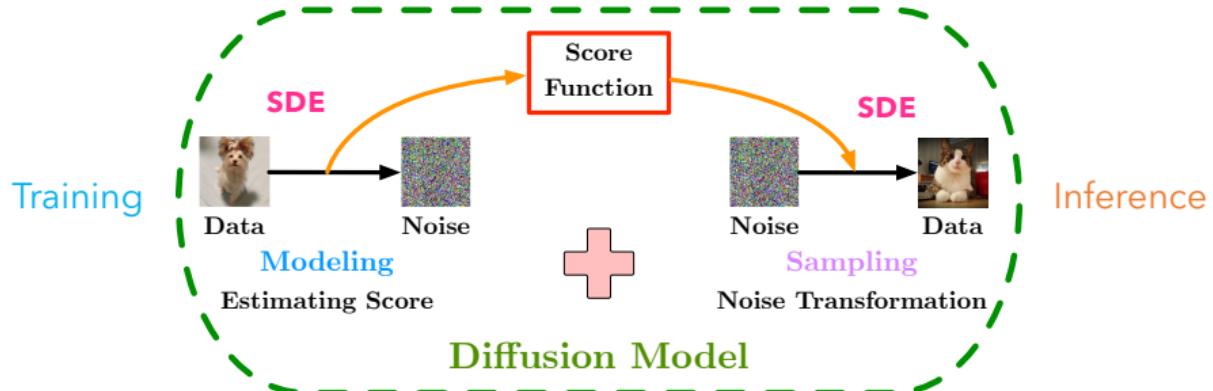
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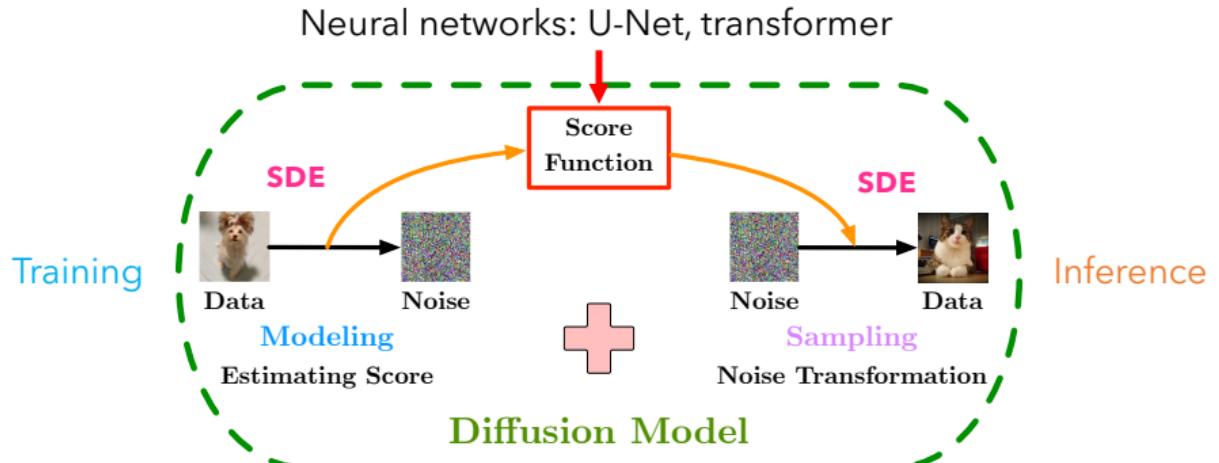
Unsupervised learning \longrightarrow **Regression!**



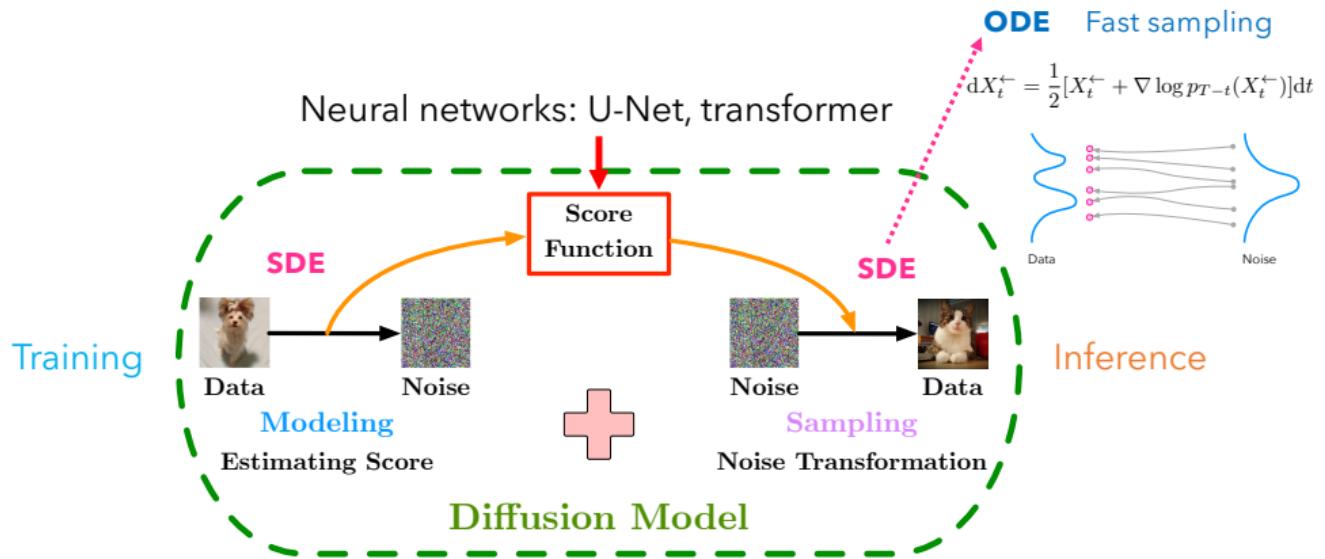
Decomposition of Diffusion Models



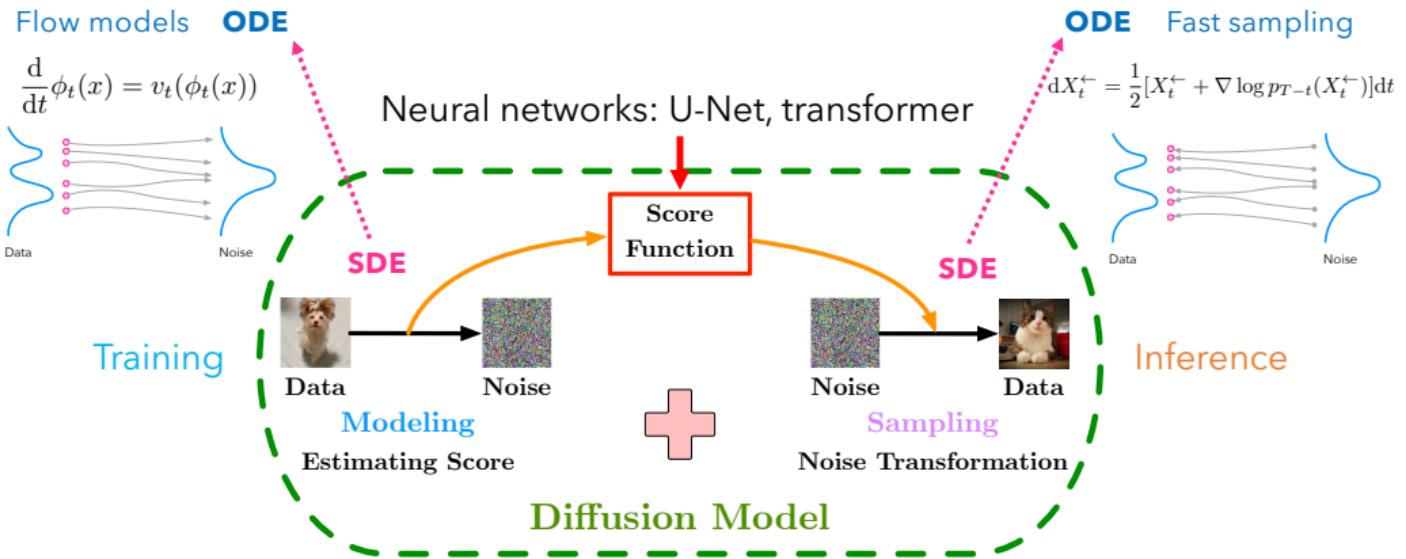
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Decomposition of Diffusion Models

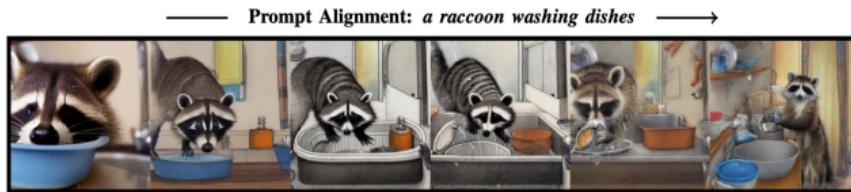


Decomposition of Diffusion Models

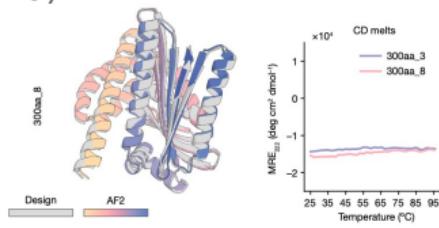


From $P(x)$ to $P(x|y)$

- Text-to-image generation (Black et al., 2023)

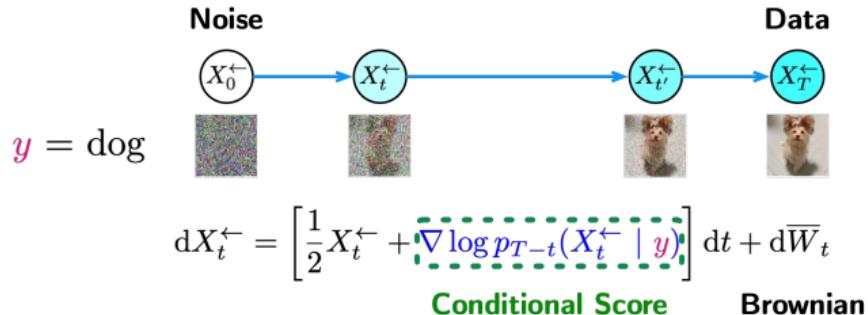


- Protein generation with biochemical properties (Watson et al., 2023; Gruver et al., 2023)



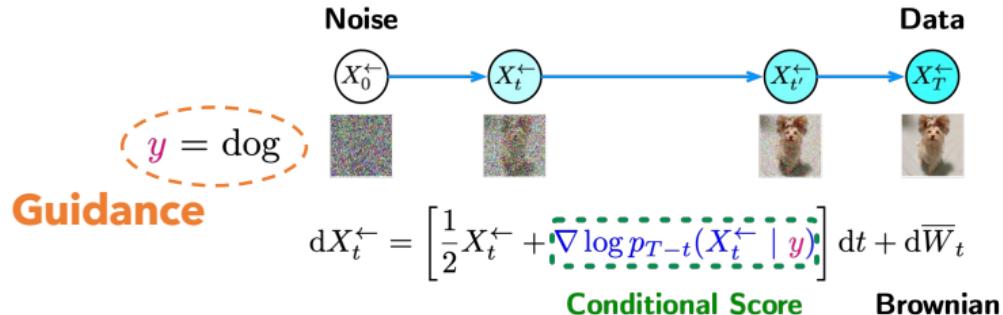
Adding Guidance to Diffusion Models

- Conditioned sample generation



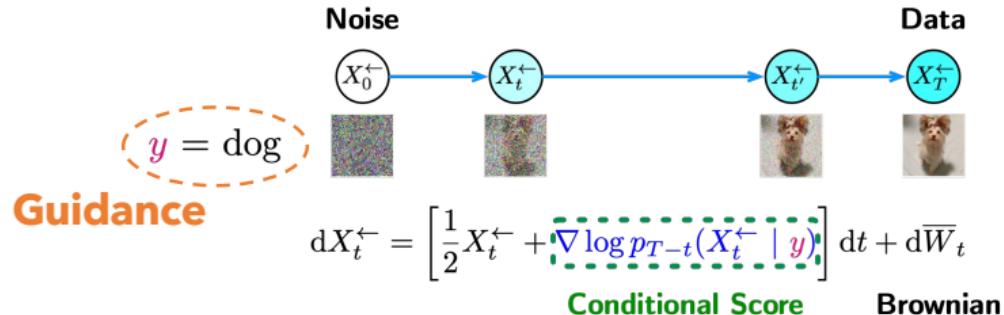
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Adding Guidance to Diffusion Models

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Classifier guidance (Dhariwal & Nichol, 2021)
Classifier-free guidance (Ho & Salimans, 2022)

The Bayes Rule - Classifier Guidance

- Discrete label

$$\nabla \log p_t(x_t | y) = \nabla \log p_t(x_t) + \nabla \log c_t(y | x_t)$$

Unconditioned Logit

**External
Classifier**



The Bayes Rule - Classifier Guidance

- Discrete label

$$\nabla \log p_t(x_t | y) = \nabla \log p_t(x_t) + \textcolor{blue}{\nabla \log c_t(y | x_t)}$$

Unconditioned Logit

**External
Classifier**



- The **magic** in practical implementation

$$s_{\text{practice}}(x_t, y, t) = \nabla \log p_t(x_t) + \textcolor{red}{\eta} \nabla \log c_t(y | x_t)$$

The Bayes Rule - Classifier Guidance

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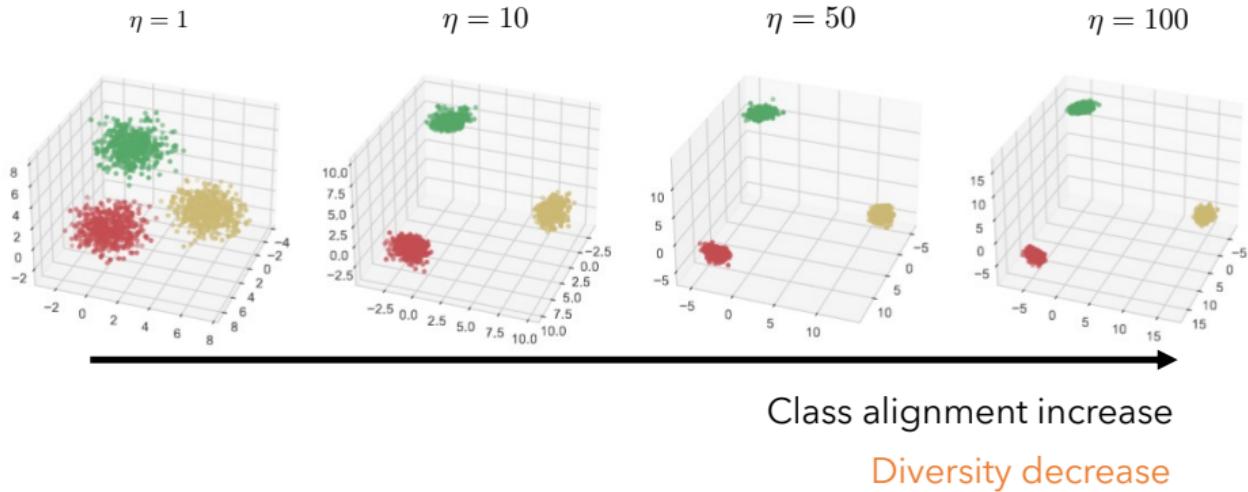


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A Glimpse of Influence in 3D Gaussian Mixture



-- Y. Wu, M. Chen, Z. Li, M. Wang, Y. Wei. "Theoretical Insights for Diffusion Guidance: A Case Study for Gaussian Mixture Models", ICML 2024.

Classifier-Free Guidance

- Limitations of classifier guidance

Discrete label and external training

Classifier-Free Guidance

- Limitations of classifier guidance
 - Discrete label and external training
- Classifier-free guidance introduces a mask signal

$$\tau \in \{\emptyset, \text{id}\}$$

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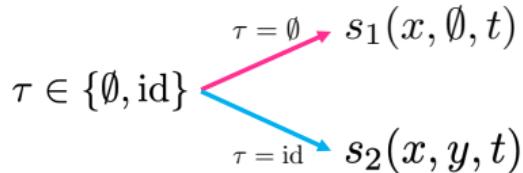
$$\tau \in \{\emptyset, \text{id}\} \xrightarrow{\tau = \emptyset} s_1(x, \emptyset, t)$$

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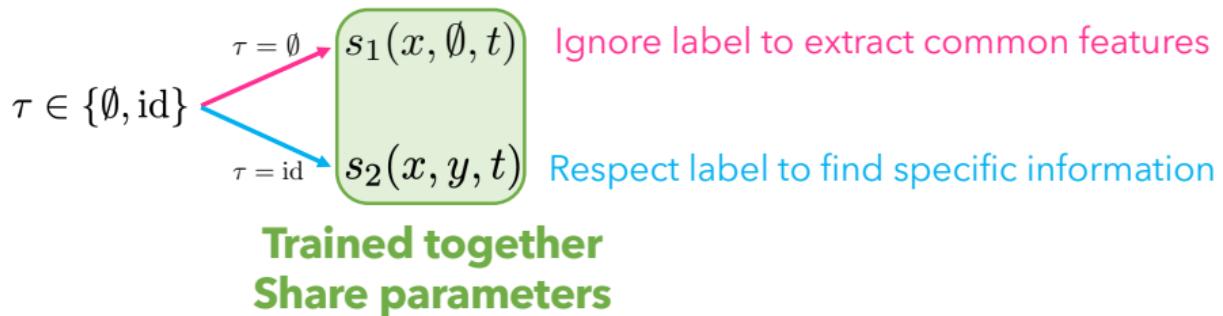


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Modeling Diverse Data

Practical Data Is High-D And Complex

- ImageNet resolution: $D = 224 \times 224 \times 3$



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Curse of Dimensionality 🤔



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Curse of Dimensionality 🤔



- Sequential data enlarges the dimension heavily

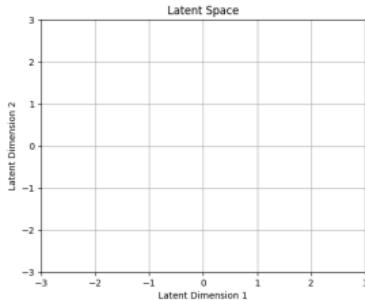
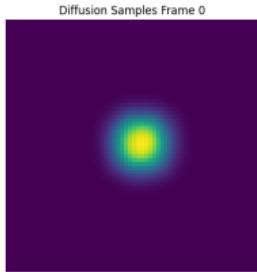
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2D bouncing ball

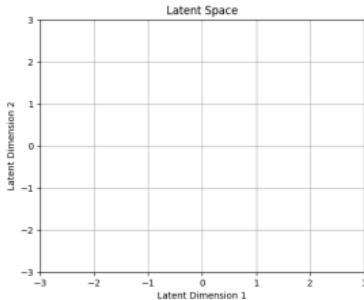
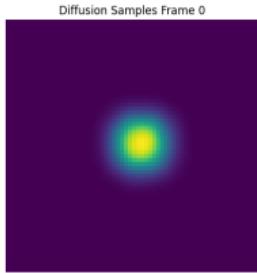
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- also introduces **spatial-temporal dependencies**



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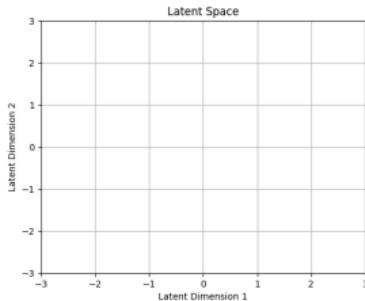
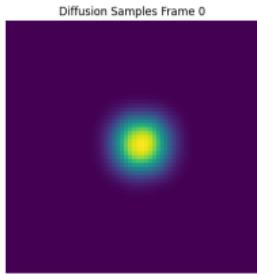
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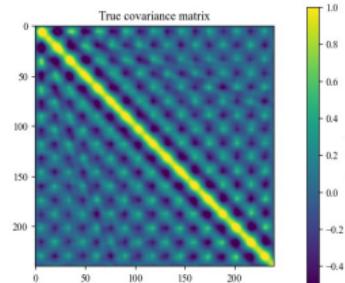
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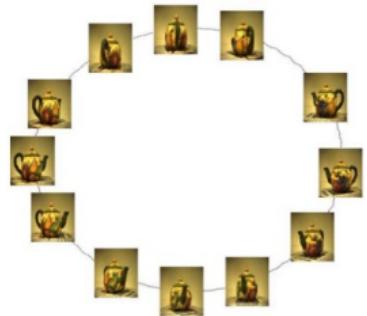


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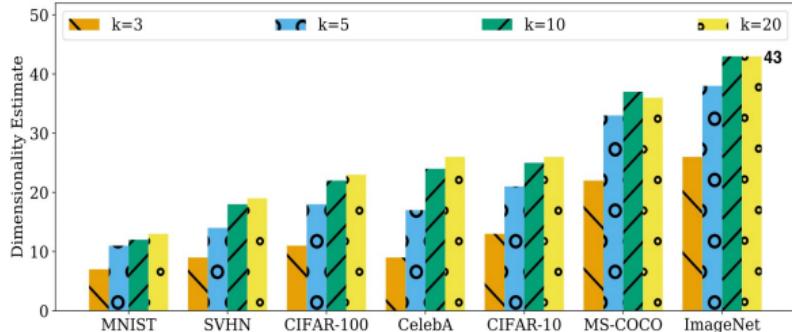
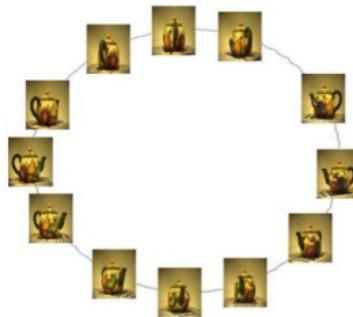


Correlation between frames

Practical High-D Data Is Low-Dimensional



Practical High-D Data Is Low-Dimensional



$224 \times 224 \times 3$ v.s. ≤ 43

-- Figure credit: (Weinberger & Saul, 2006; P. Pope et al., 2021)

How Diffusion Model Finds Structures?

- Dynamic evolution

$$\text{Backward} \quad dX_t^\leftarrow = \left[\frac{1}{2} X_t^\leftarrow + \boxed{\nabla \log p_{T-t}(X_t^\leftarrow)} \right] dt + d\bar{W}_t$$

Score Function Brownian

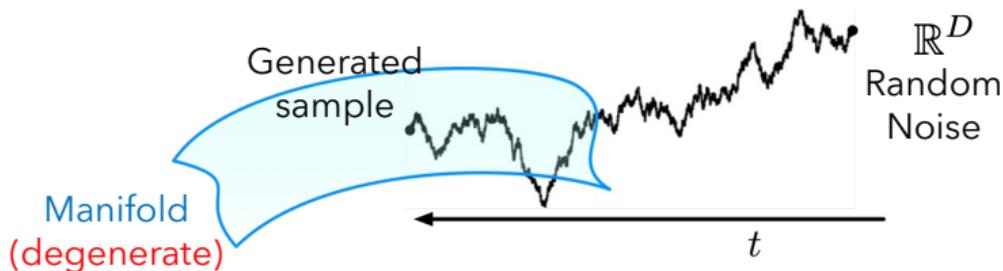
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Score Function Brownian

- Start from high-D but land in a manifold?



Score Function Adapts to Data Structures

- Linear subspace data

$$x = Az \quad \text{with} \quad z \sim P_z \quad z \in \mathbb{R}^d$$

- Score decomposition

$$\nabla \log p_t(x) = \boxed{A \nabla \log p_t^z(A^\top x)} - \boxed{\frac{1}{1 - e^{-t}} (I_D - AA^\top) x}$$

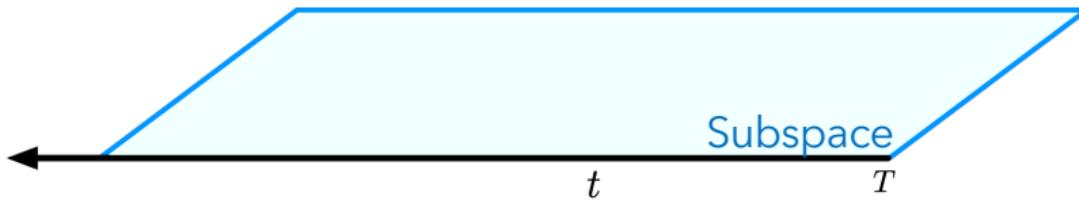
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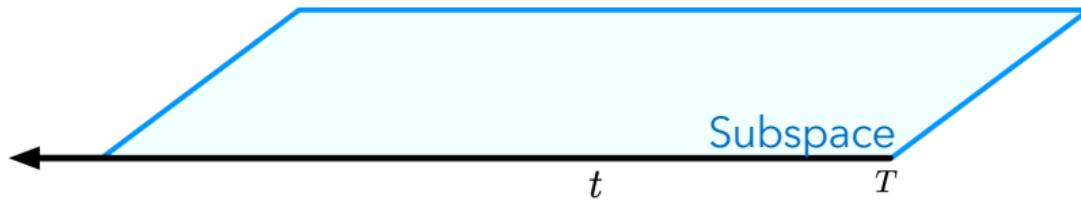
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• x_{t_1}



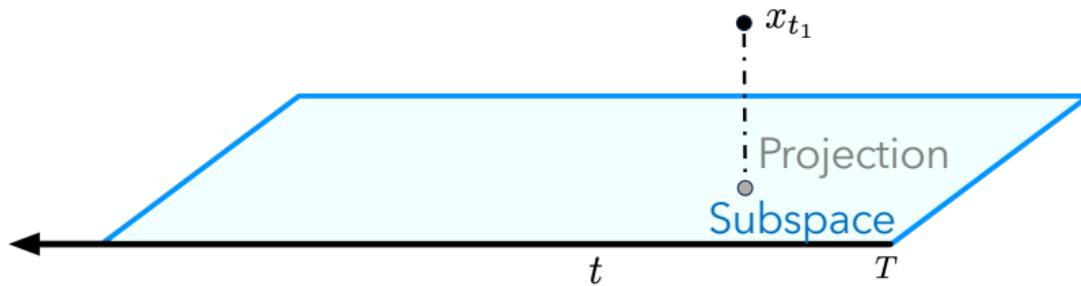
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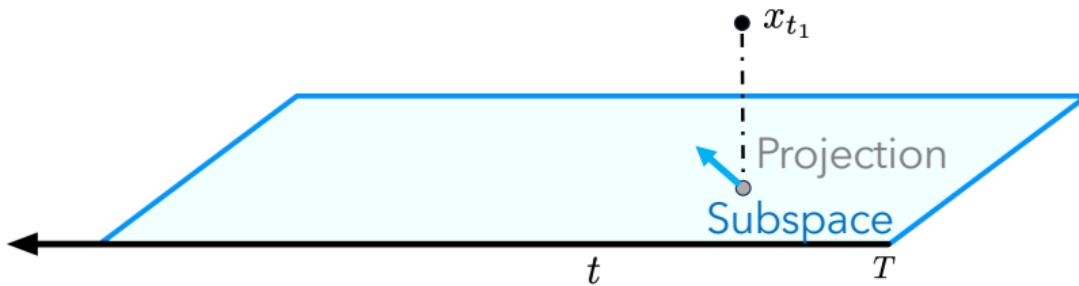
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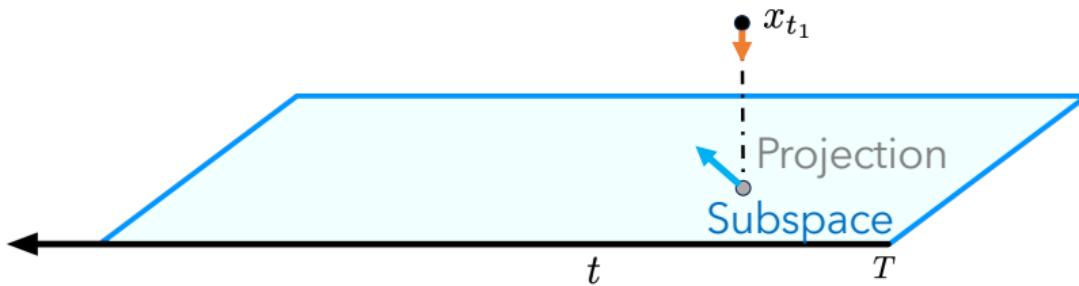
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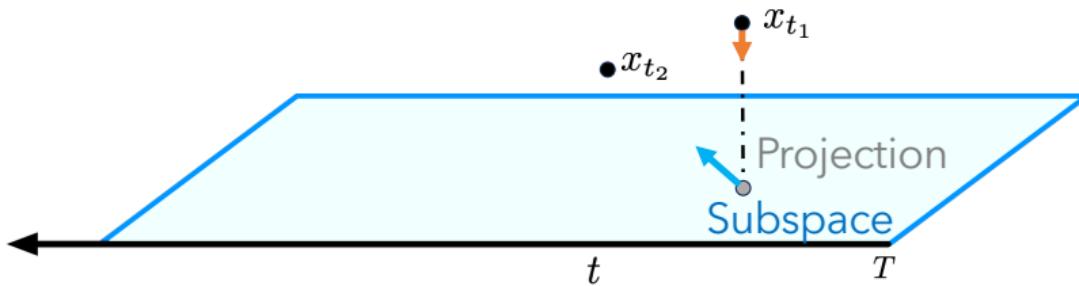
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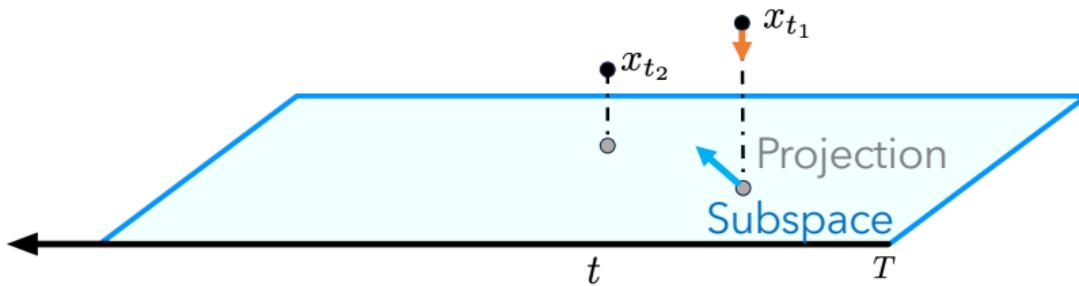
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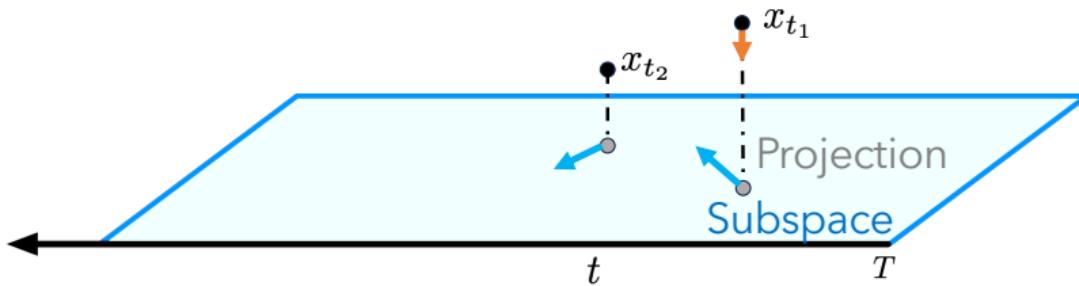
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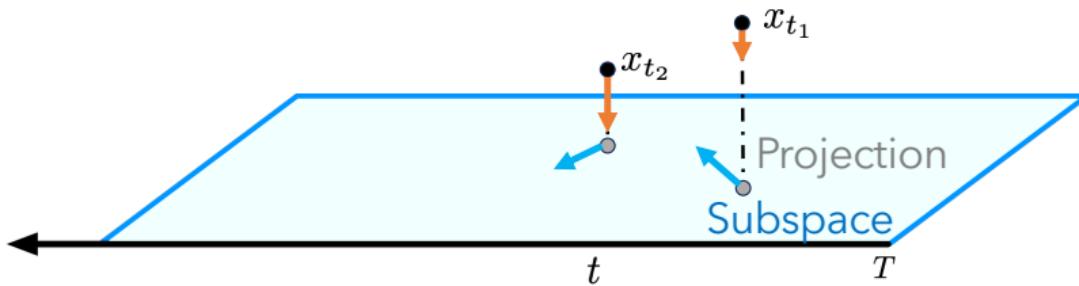
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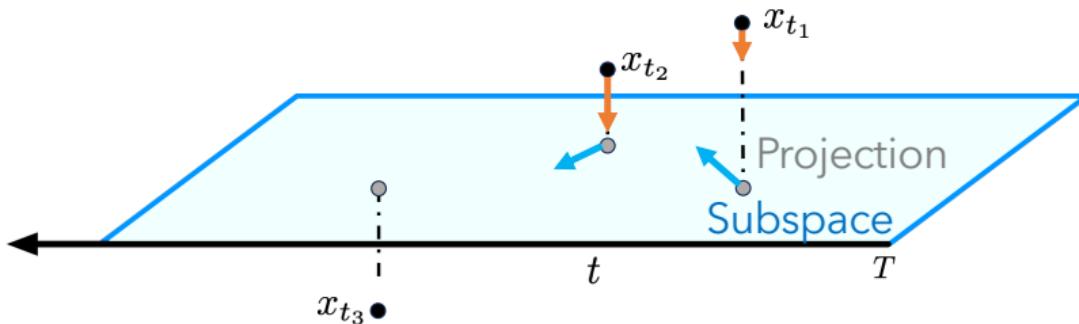
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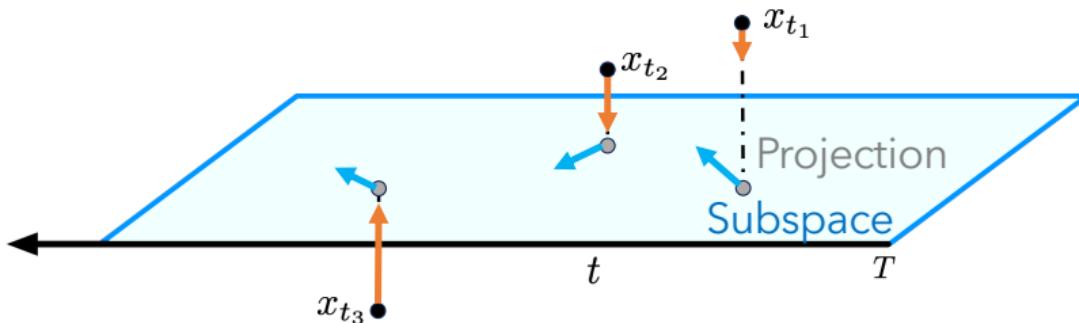
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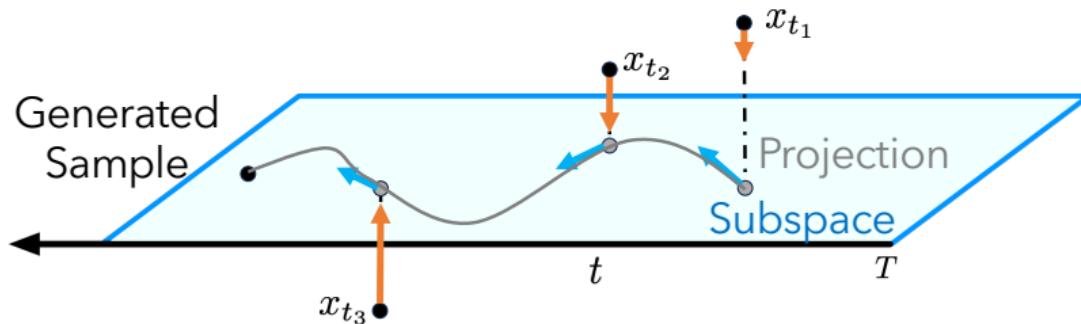
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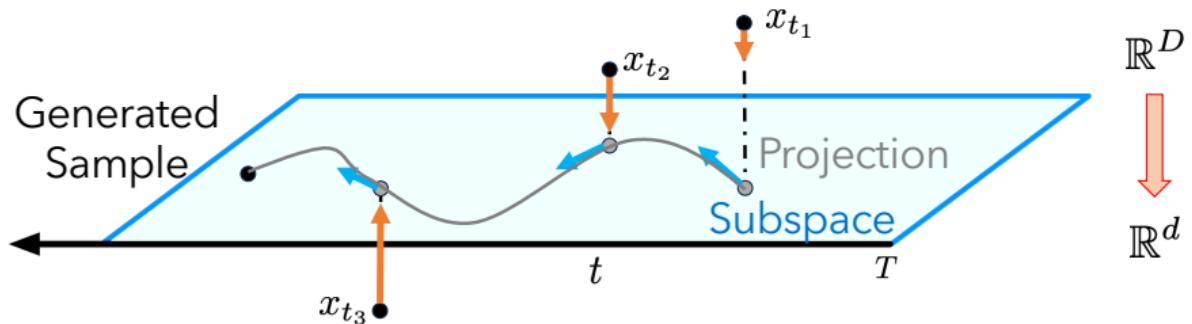
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Diffusion Model Efficiently Learns Low-D Data

Theorem

- ✓ Score function can be learned **efficiently** at the rate

$$\tilde{\mathcal{O}}\left(n^{-\frac{1}{2(\textcolor{red}{d}+5)}}\right)$$

- ✓ Underlying distribution is learned at the same rate.

-- M. Chen, K. Huang, T. Zhao, M. Wang. "Score Approximation, Estimation and Distribution Recovery of Diffusion Models on Low-Dimensional Data", ICML 2023

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Take-home message:

- ✓ **Accurate** in learning data distributions
- ✓ **Efficient**: no curse of dimensionality
- ✓ **Generalizable** to manifold data (Tang and Yang, 2024)

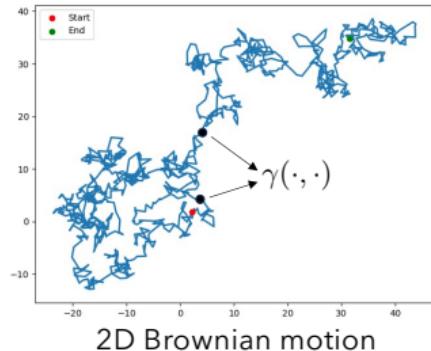
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Sequence Data with Dependencies

- We consider

$$X_1, \dots, X_{h_N} \sim \mathcal{GP}(\mu(\cdot), \gamma(\cdot, \cdot), \Lambda) \in \mathbb{R}^D$$

- $0 = h_1 < \dots < h_N = H$ sampling times
- $\mu(\cdot)$ time varying mean function
- $\gamma(\cdot, \cdot)$ covariance function (kernel)
- $\Lambda = \text{Cov}[X_h]$ marginal covariance matrix

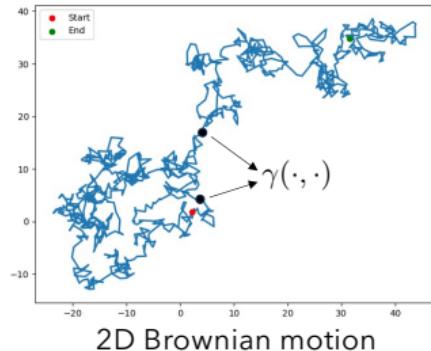


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Simplification

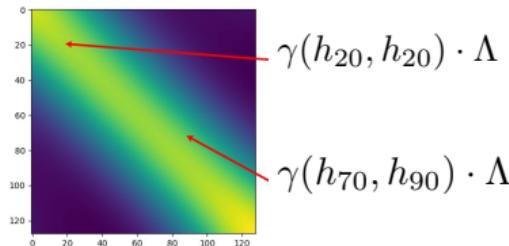
- ✓ $\gamma(\cdot, \cdot)$ only depends on time gaps, i.e.,

$$\gamma(t_1, t_2) = g(|t_1 - t_2|)$$

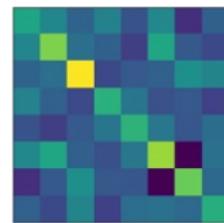
Description of Spatial-Temporal Dependencies

- We stack data together as a vector in \mathbb{R}^{DN} , whose distribution is Gaussian

$$\mathcal{N}\left(\begin{bmatrix} \mu(h_1) \\ \vdots \\ \mu(h_N) \end{bmatrix}, \Gamma \otimes \Lambda = \begin{bmatrix} \gamma(h_1, h_1)\Lambda, \dots, \gamma(h_1, h_N)\Lambda \\ \vdots \\ \gamma(h_N, h_1)\Lambda, \dots, \gamma(h_N, h_N)\Lambda \end{bmatrix}\right)$$



Temporal dependencies



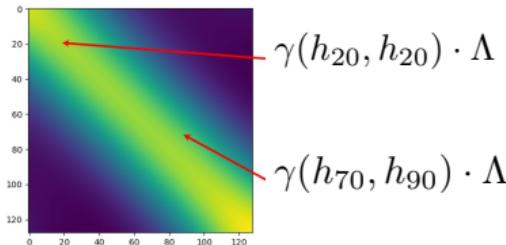
Spatial dependencies

Description of Spatial-Temporal Dependencies

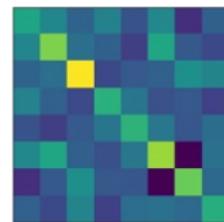
- We stack data together as a vector in \mathbb{R}^{DN} , whose distribution is Gaussian

Huge dimension?

$$\mathcal{N}\left(\begin{bmatrix} \mu(h_1) \\ \vdots \\ \mu(h_N) \end{bmatrix}, \Gamma \otimes \Lambda = \begin{bmatrix} \gamma(h_1, h_1)\Lambda, \dots, \gamma(h_1, h_N)\Lambda \\ \vdots \\ \gamma(h_N, h_1)\Lambda, \dots, \gamma(h_N, h_N)\Lambda \end{bmatrix}\right)$$



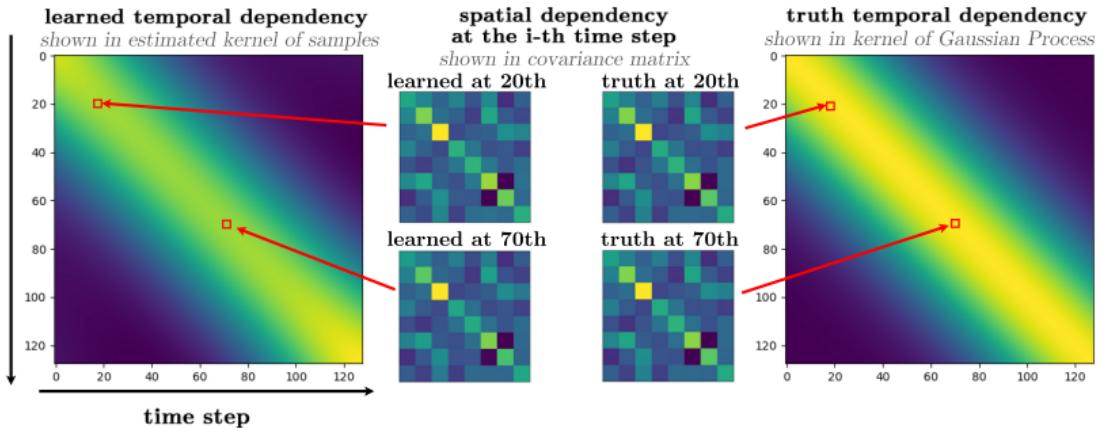
Temporal dependencies



Spatial dependencies

Proper Score Network Learns Dependencies

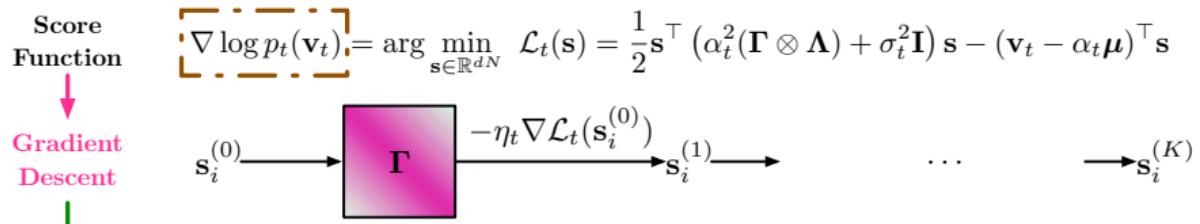
- We simulate a Gaussian process with 128 length
- Diffusion model with **transformer learns very well!**



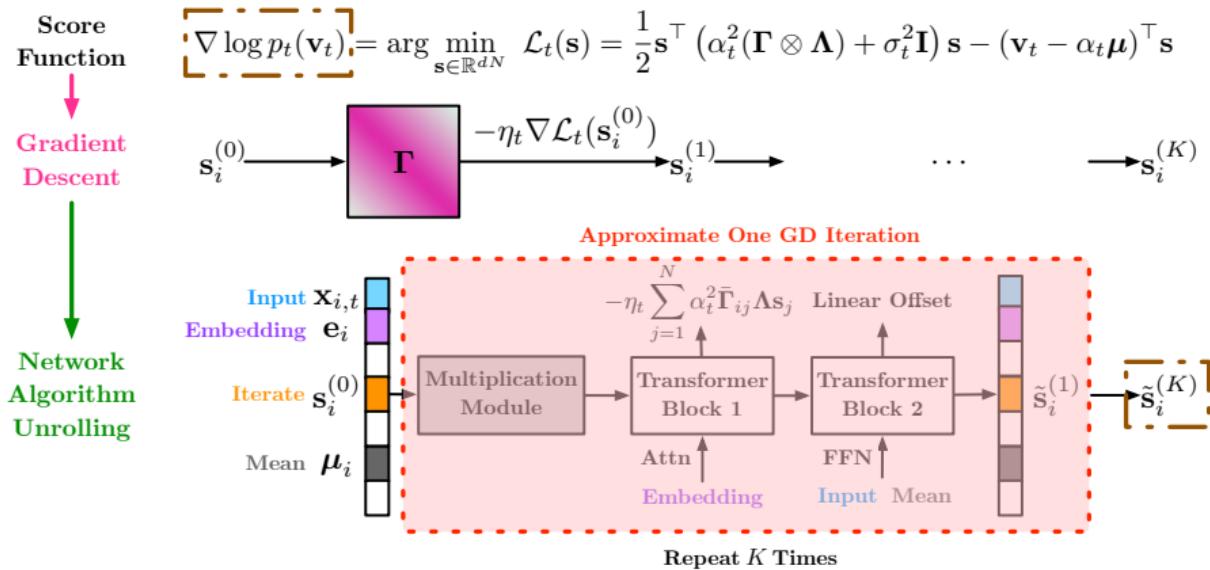
Represent Score via Algorithm Unrolling

Score Function $\boxed{\nabla \log p_t(\mathbf{v}_t)}$ = $\arg \min_{\mathbf{s} \in \mathbb{R}^{dN}} \mathcal{L}_t(\mathbf{s}) = \frac{1}{2} \mathbf{s}^\top (\alpha_t^2 (\boldsymbol{\Gamma} \otimes \boldsymbol{\Lambda}) + \sigma_t^2 \mathbf{I}) \mathbf{s} - (\mathbf{v}_t - \alpha_t \boldsymbol{\mu})^\top \mathbf{s}$

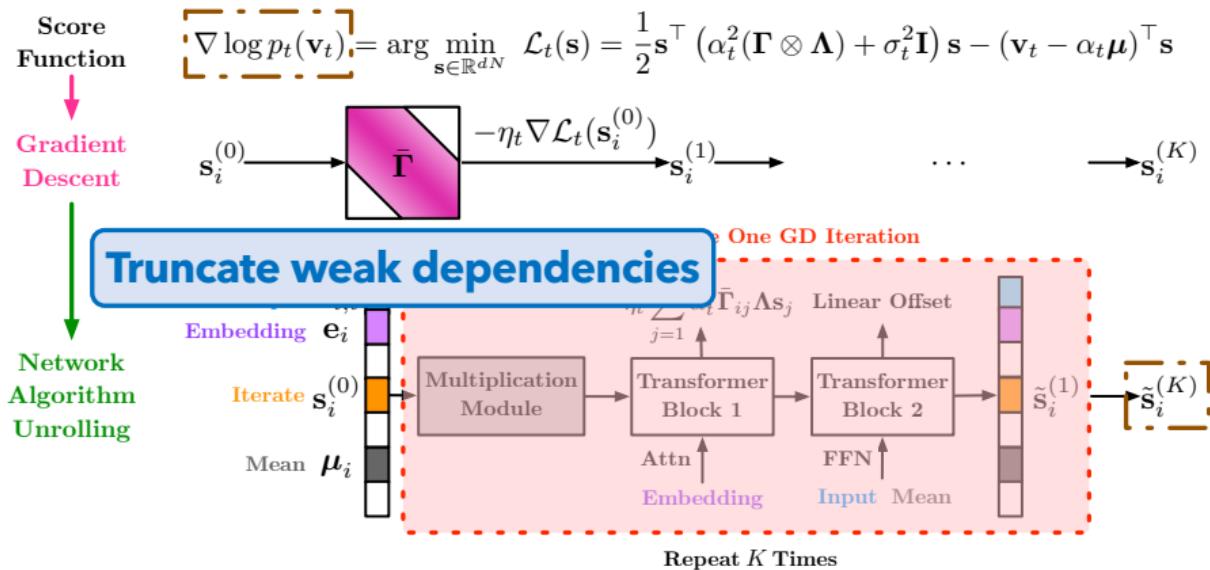
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Represent Score via Algorithm Unrolling



Diffusion Model Learns Gaussian Process

Theorem

- ✓ Score function can be learned **efficiently** at the rate

$$\tilde{\mathcal{O}}\left(\sqrt{\frac{\text{dependency-decay} \cdot \text{sequence-length} \cdot D^3}{n}}\right)$$

- ✓ Gaussian process distribution is learned at the same rate.

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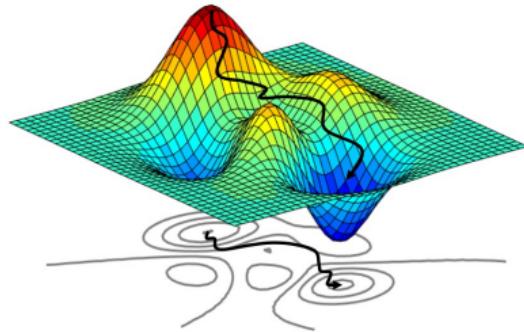
Take-home message:

- ✓ **Weak** dependence on the length of sequence
- ✓ **Adaptive** to spatial-temporal dependencies

Leverage Diffusion Models

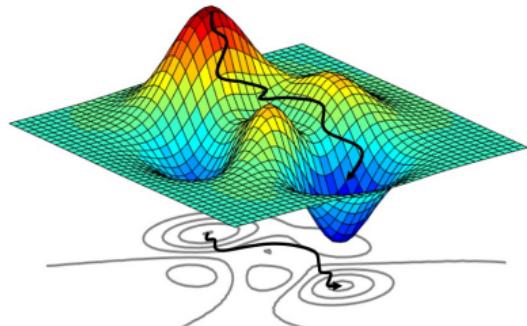
Rethinking Optimization

$$x \in \arg \max f^*(\cdot)$$



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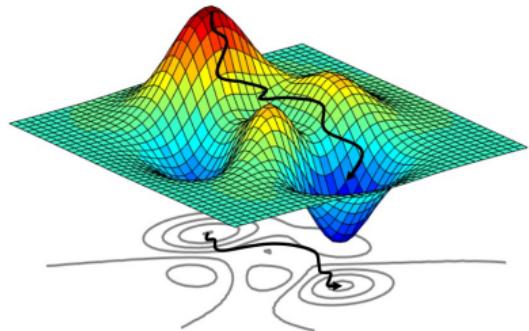


High-D

Nonconvex

Rethinking Optimization

$$x \in \arg \max f^*(\cdot)$$



High-D

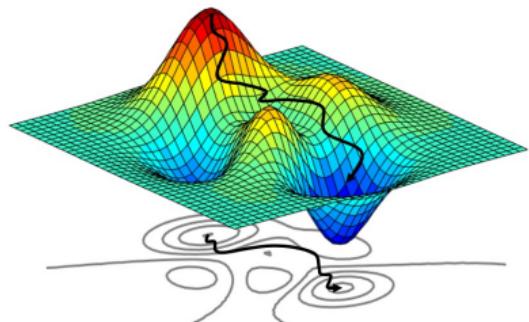
Nonconvex

Generate solution

$$x \sim \mathbb{P}(\cdot | f^*(\cdot) \geq a)$$

Rethinking Optimization

$$x \in \arg \max f^*(\cdot)$$



High-D

Nonconvex

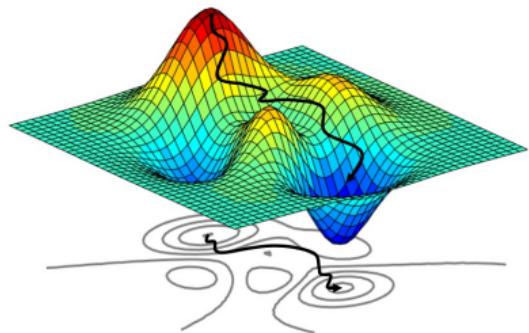
Generative Optimization

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Rethinking Optimization

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High-D

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Generate solution

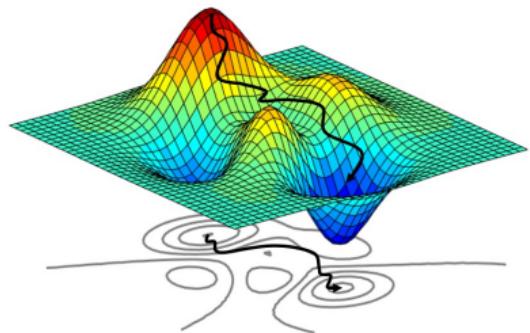
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High-D

Conditional distribution

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Guidance

High-D

Conditional distribution

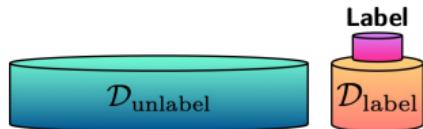
Problem Setup: Offline Reward Maximization

- Given a training data set, generate new x
- Training data set

$$\mathcal{D}_{\text{unlabel}} = \{x_j\}_{j=1}^{n_{\text{unlabel}}}$$

$$\mathcal{D}_{\text{label}} = \{x_i, y_i = f^*(x_i) + \epsilon_i\}_{i=1}^{n_{\text{label}}}$$

- ϵ_i is observation noise
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- ❑ **Example:** a large collection of unlabeled protein structures; only a few has measured properties.

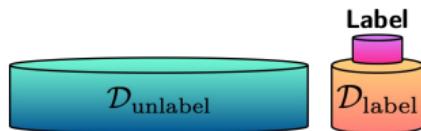
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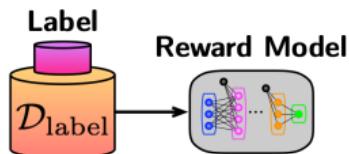


Off-policy bandit problem

(Jin et al., 2021; Nguyen-Tang et al., 2021)

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Meta Algorithm

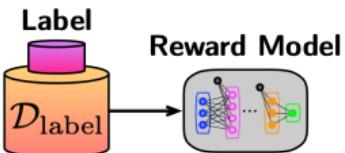


Step 1: Reward Learning

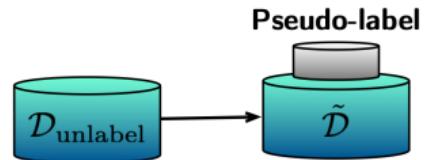
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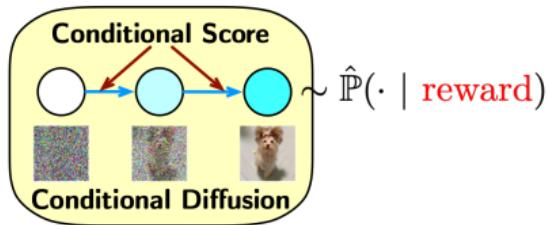
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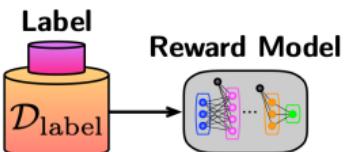


Step 2: Pseudo Labeling

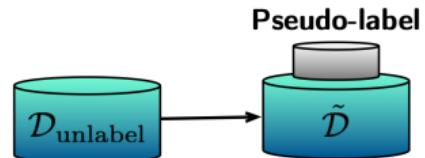


Step 3: Conditional Diffusion Training

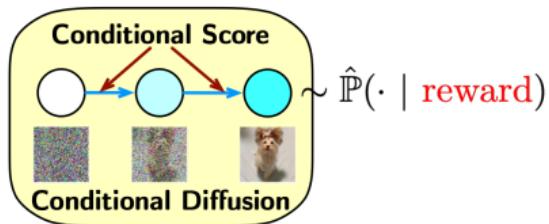
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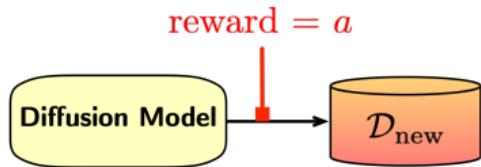
Step 1: Reward Learning



Step 2: Pseudo Labeling



Step 3: Conditional Diffusion Training



Step 4: Guided Generation

How Far Are We from The Targeted Reward

- Let a be the target reward of generation

$$\text{SubOpt}(a) = a - \text{Generated Average Reward}$$

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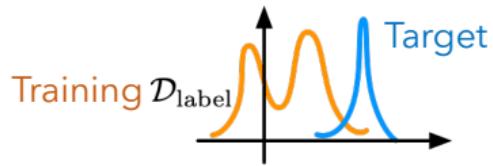
(Reward estimation error)

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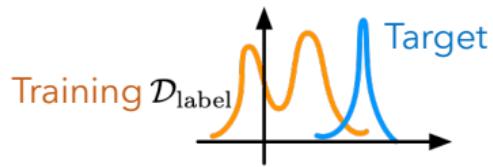
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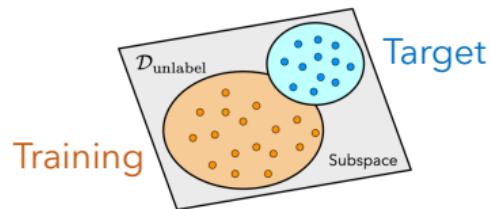
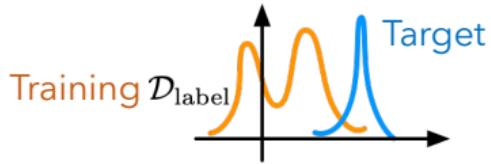
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(Conditional diffusion error) \circ (Diffusion distribution shift)



Case Study: Subspace Data + Linear Reward

Theorem

- ✓ The sub-optimality satisfies

$$\text{SubOpt}(a) = \tilde{\mathcal{O}} \left(\sqrt{\text{Trace}(\hat{\Sigma}_\lambda^{-1} \Sigma_a)} \cdot \sqrt{\frac{d \log(n_{\text{label}})}{n_{\text{label}}}} + \min\{a, d\} \cdot \frac{a \cdot \text{poly}(D, d)}{n_{\text{unlabel}}^{1/6}} \right)$$

where $\hat{\Sigma}_\lambda = (X^\top X + \lambda I)/n_{\text{label}}$ for X the data matrix, $\lambda > 0$, and Σ_a is the covariance matrix of $P_a(\cdot \mid \text{reward} = a)$.

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- ❖ Match optimal off-policy bandit learning with representation learning (Jin et al., 2021; Nguyen-Tang et al., 2021)

-- Z. Li, H. Yuan, K. Huang, C. Ni, Y. Ye, M. Chen, M. Wang. "Diffusion Model for Data-Driven Black-Box Optimization", NeurIPS 2023

Advantages of Generative Optimization

- ✓ Meta algorithm provably generates samples of high reward and fidelity, in **nonparametric settings**.

$$\text{SubOpt}(a) = \tilde{O} \left(\kappa_1(a) \cdot n_{\text{label}}^{-\frac{\alpha}{d+2\alpha}} + \kappa_2(a) \cdot n_{\text{unlabel}}^{-\frac{2}{3(d+6)}} \right)$$

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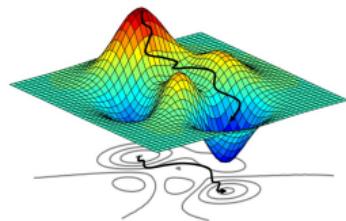
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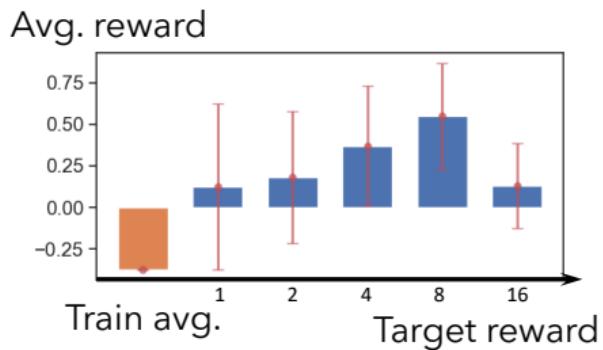
Generative optimization in offline:

- ✓ **Off-policy bandit optimality**
- ✓ **High-fidelity** to intrinsic structures
- ✓ **Efficiency**: no curse of dimensionality
- ✓ **Generalizable** to human preferences



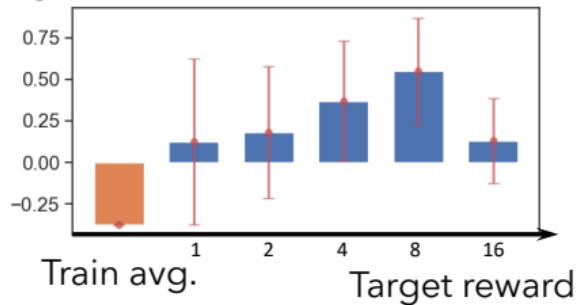
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Application 1: CIFAR Reward Optimization



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Avg. reward



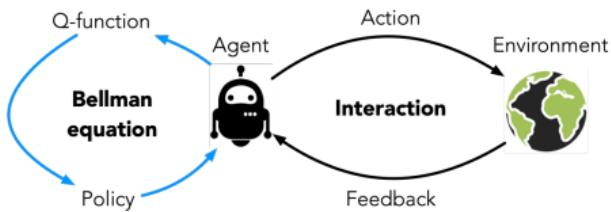
Quality degrades



Reward improves

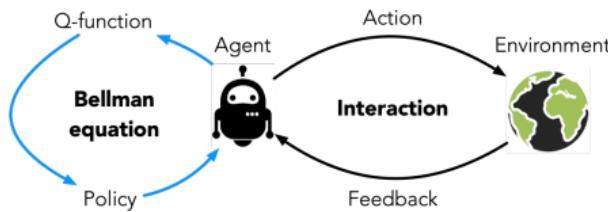
Application 2: Generative Optimization in RL

- Reinforcement learning

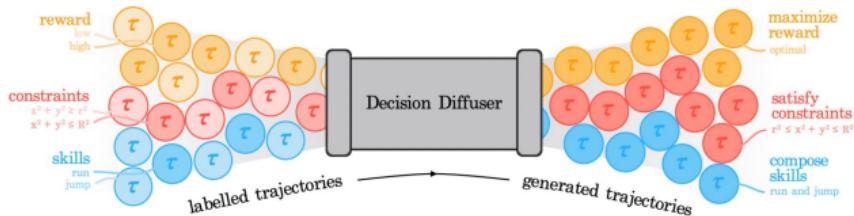


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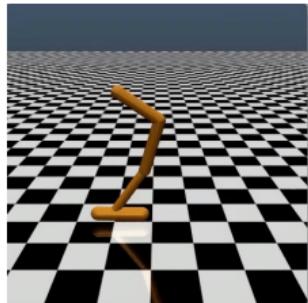
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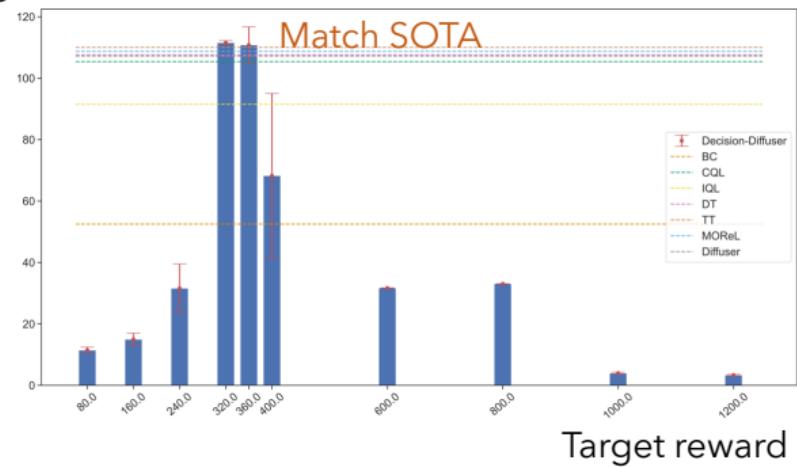
- Generative optimization (Decision diffuser, Ajay et al., 2023)



Hopper Control



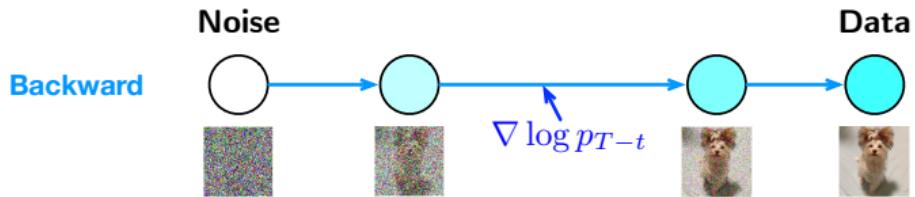
Avg. reward



Inspirations and Future Directions

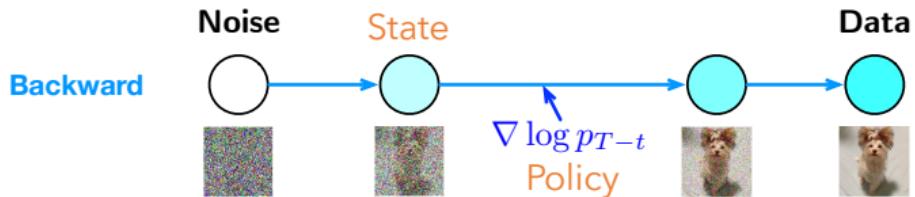
Control/RL Perspective on Diffusion Model

- We design backward process to be Markovian



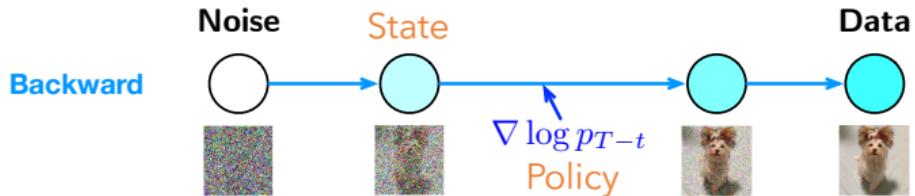
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Control/RL Perspective on Diffusion Model

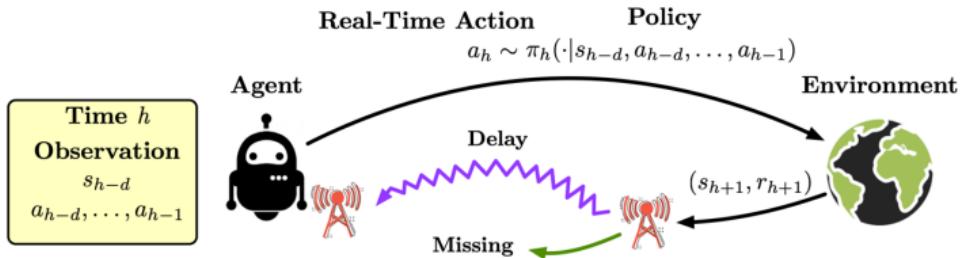
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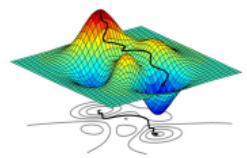
- Reward choice is task dependent
 - Fidelity metric for image generation
 - Satisfactory level for product design
 - Biochemical property for protein synthesis
 - Etc.

Diffusion Model for Control/RL

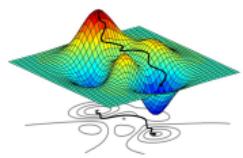
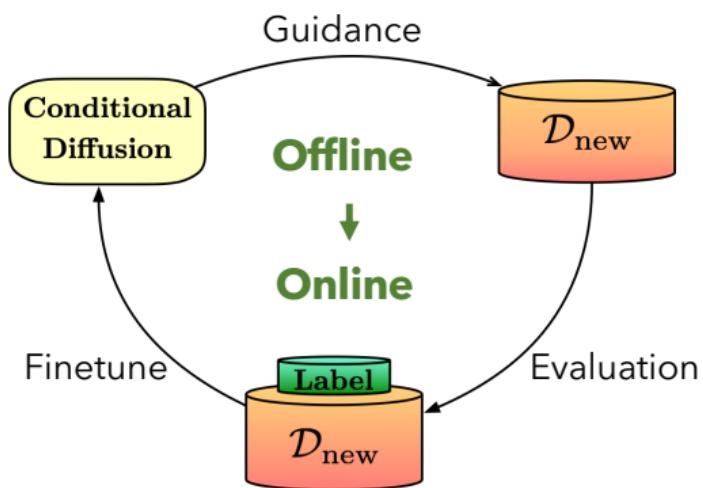
- Conditional diffusion models as a rich class for parameterizing **policies** and **transition kernels**
- Diffusion for practical RL with impaired observability (Chen et al., 2023)



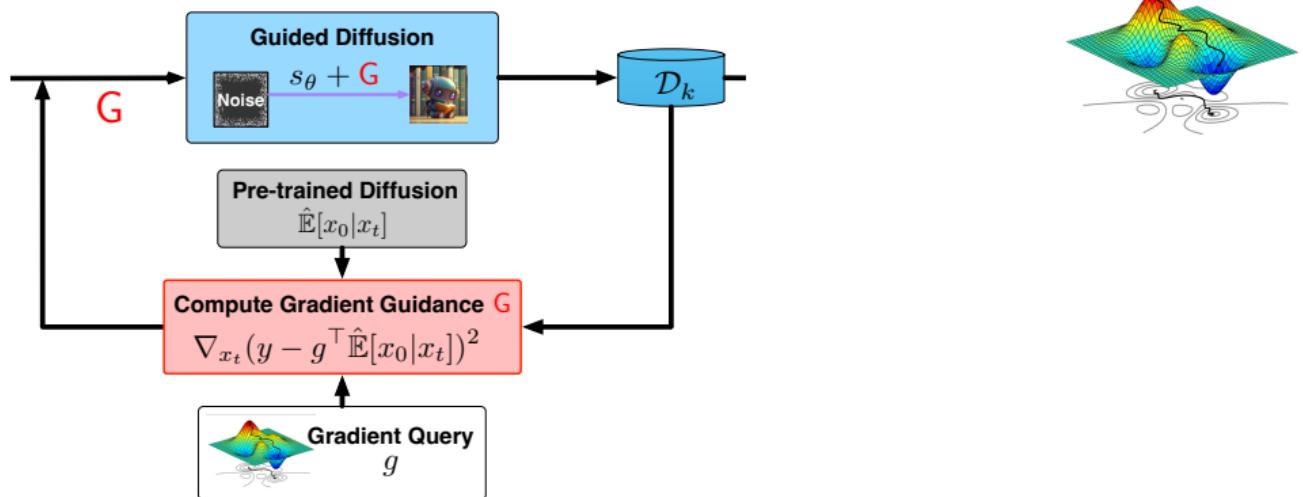
Diffusion Model for Generative Optimization



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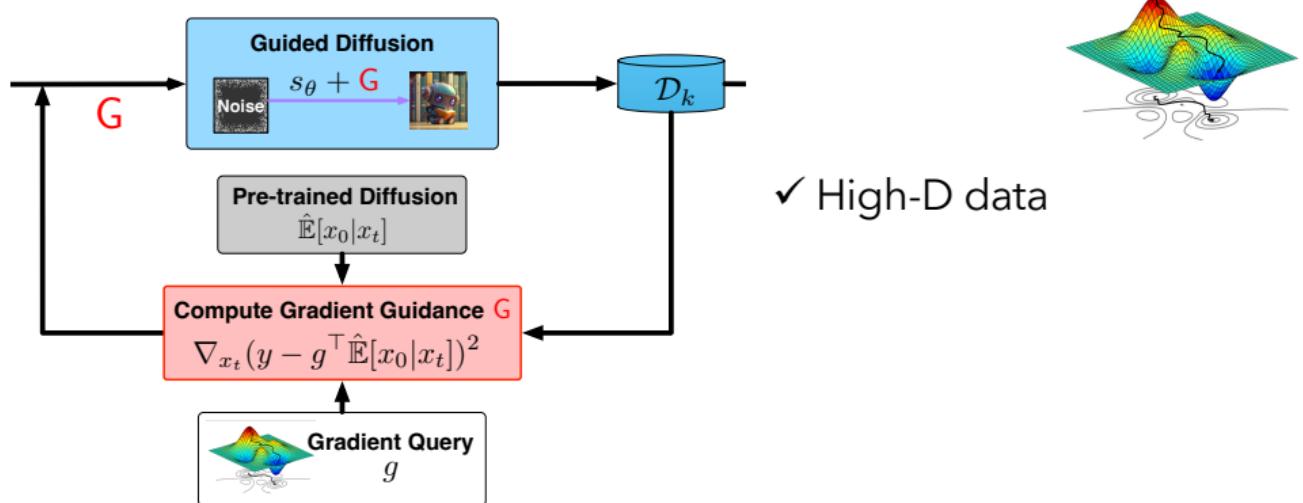


Diffusion Model for Generative Optimization



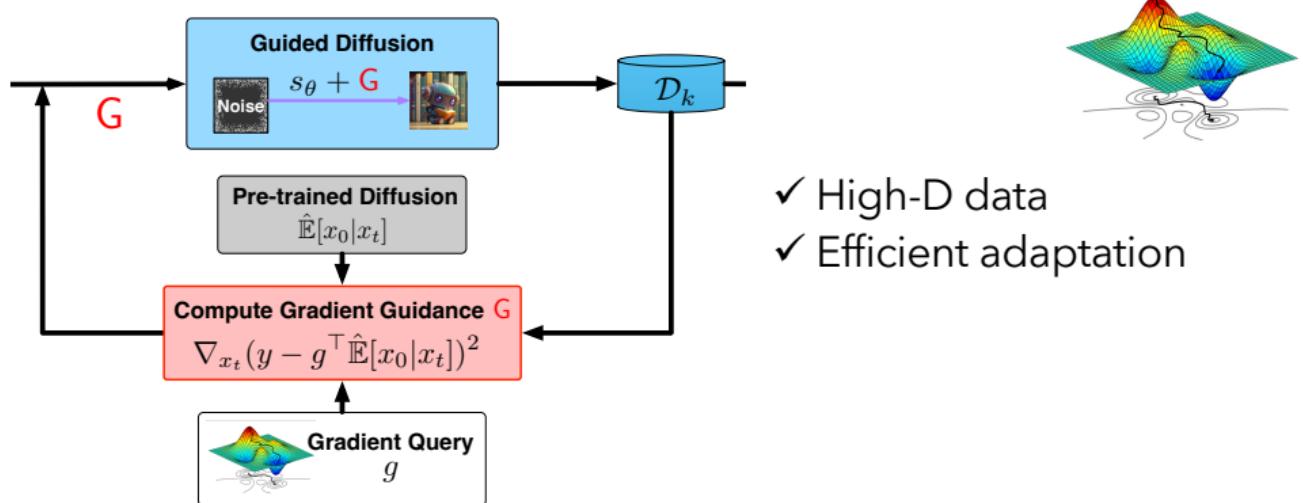
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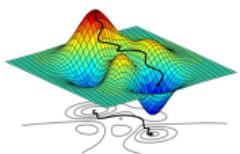
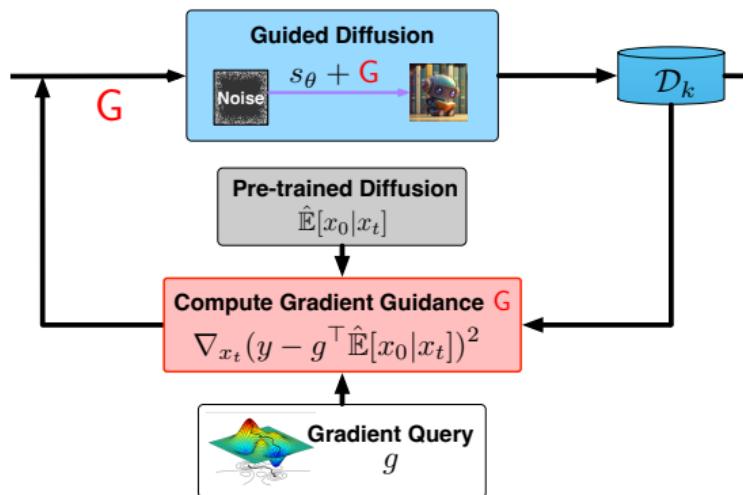
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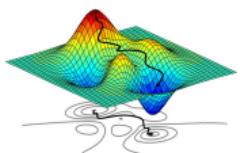
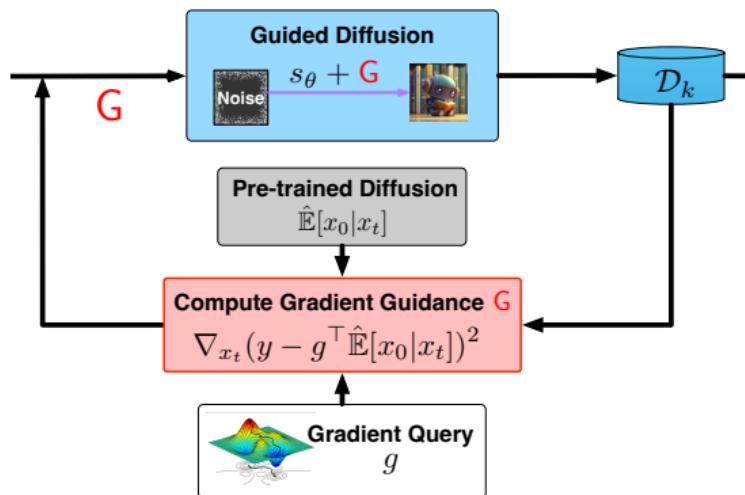
Diffusion Model for Generative Optimization



- ✓ High-D data
- ✓ Efficient adaptation
- ✓ Escape from bad local optima and saddle points

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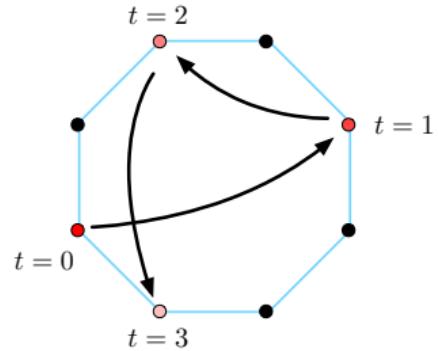
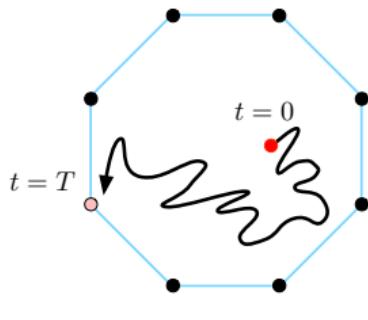


- ✓ High-D data
- ✓ Efficient adaptation
- ✓ Escape from bad local optima and saddle points
- ✓ Connecting to DRO

-- Y. Guo, H. Yuan, Y. Yang, M. Chen, M. Wang. "Gradient Guidance for Diffusion Models: An Optimization Perspective", NeurIPS 2024

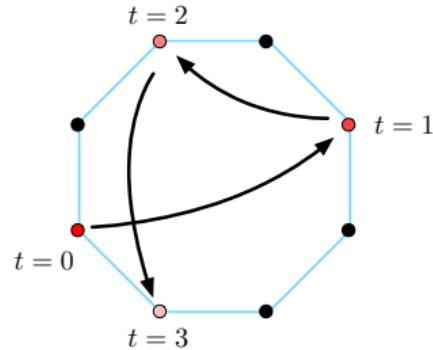
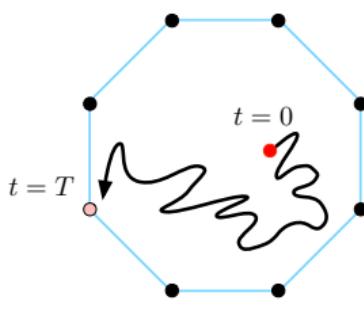
Diffusion Model for Discrete Data

- Gaussian noise may not be suitable for discrete data



Diffusion Model for Discrete Data

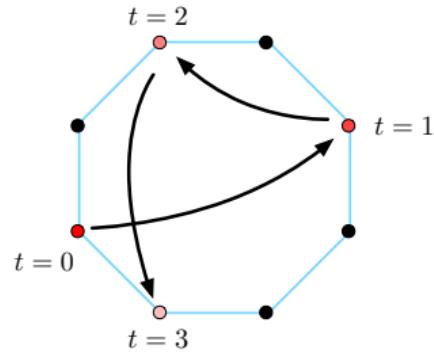
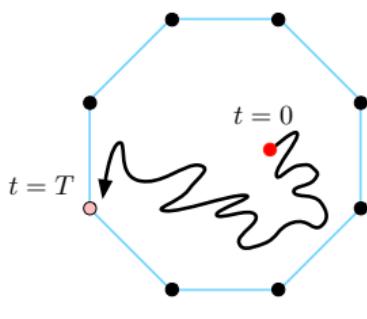
- Gaussian noise may not be suitable for discrete data



- Discrete diffusion jumps inside data support as “corruption”

Diffusion Model for Discrete Data

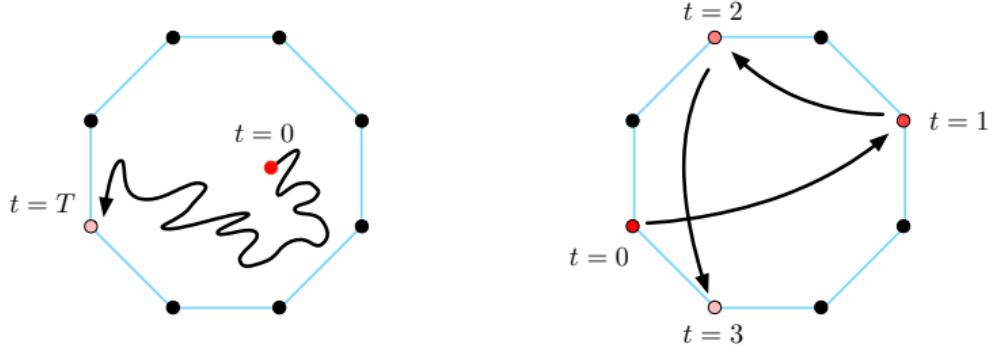
- Gaussian noise may not be suitable for discrete data



- Discrete diffusion jumps inside data support as “corruption”
 - Integer optimization

Diffusion Model for Discrete Data

- Gaussian noise may not be suitable for discrete data



- Discrete diffusion jumps inside data support as “corruption”
 - Integer optimization
 - Protein generation

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Thank You!