

# Inverse Bridge Matching Distillation

Nikita Gushchin<sup>1 4</sup> David Li<sup>1 5 \*</sup> Daniil Selikhanovych<sup>1 2 3 \*</sup> Evgeny Burnaev<sup>1 4</sup>  
Dmitry Baranchuk<sup>2</sup> Alexander Korotin<sup>1 4</sup>

<sup>1</sup> Skolkovo Institute of Science and Technology

<sup>2</sup> Yandex Research

<sup>3</sup> HSE University

<sup>4</sup> Artificial Intelligence Research Institute

<sup>5</sup> Moscow Institute of Physics and Technology

\* Equal contribution

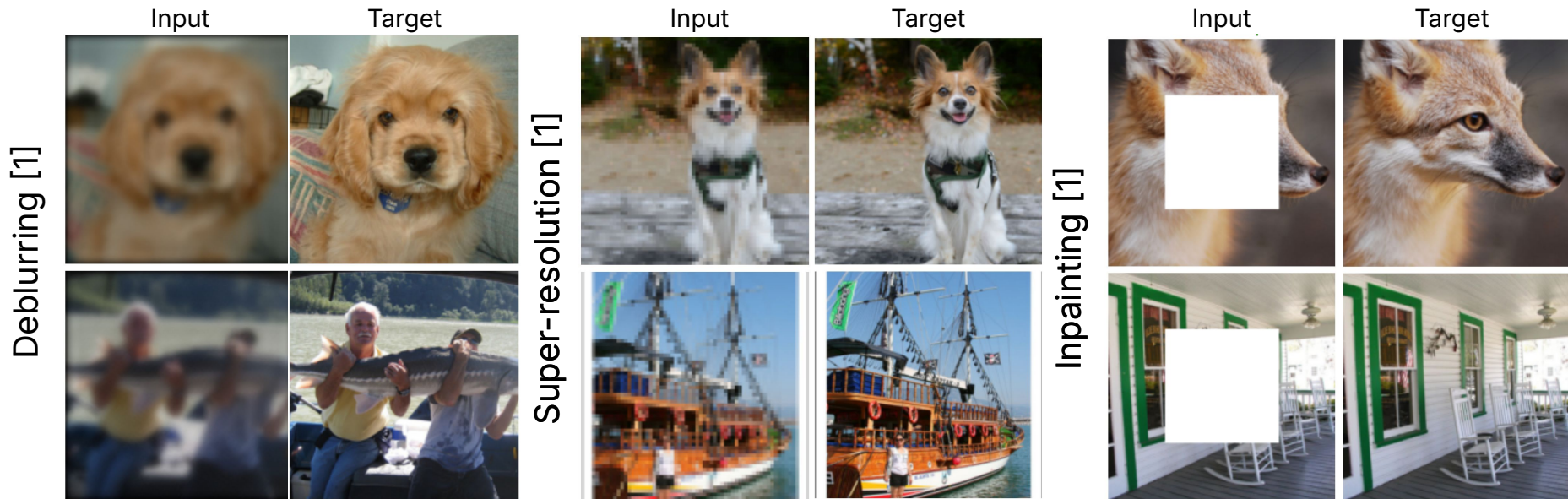


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# Paired image-to-image translation problem

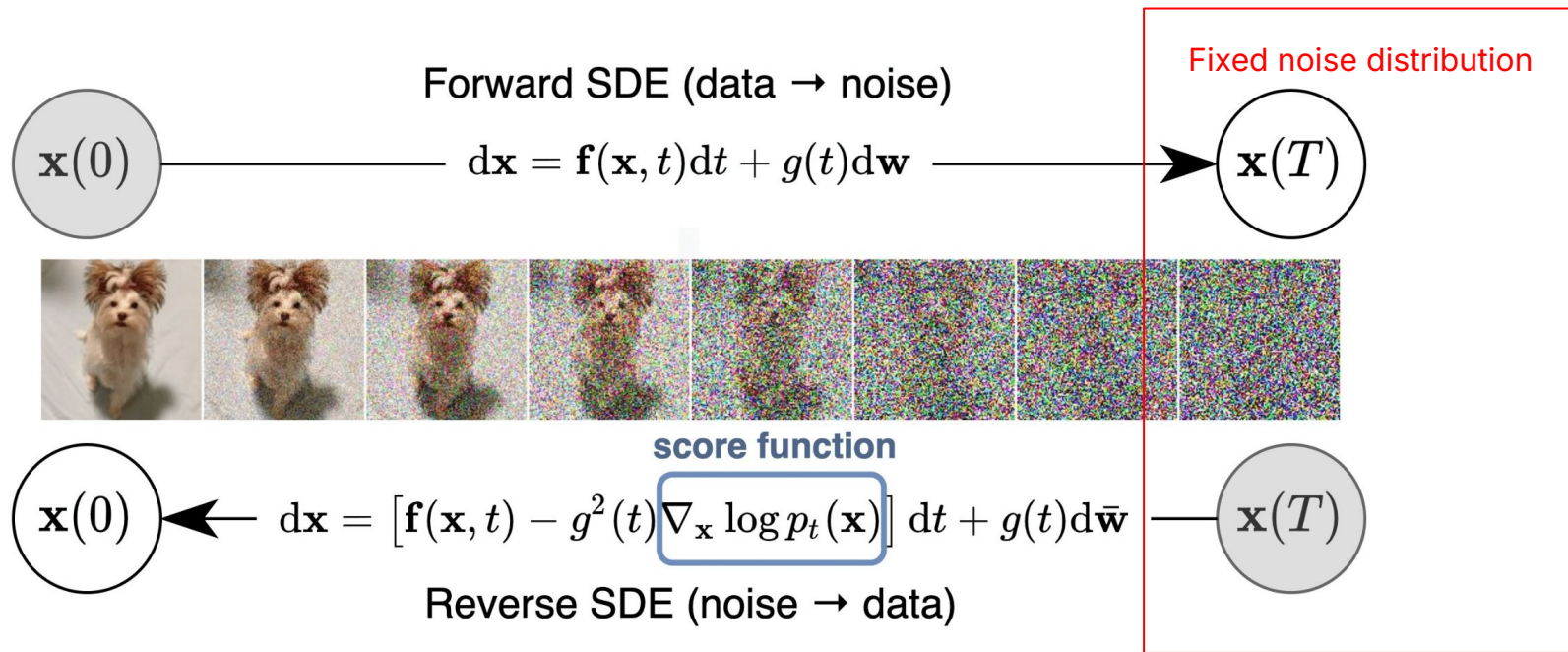
**Paired image-to-image translation problem:** given a paired dataset of corrupted/clean images — or of two distinct domains — the task is to construct a model that learns to transform inputs into aligned target images.



**The question:** Diffusion models are known for their superior image generation quality. Can we adapt them to solve image-to-image translation tasks?

# Recap on diffusion models

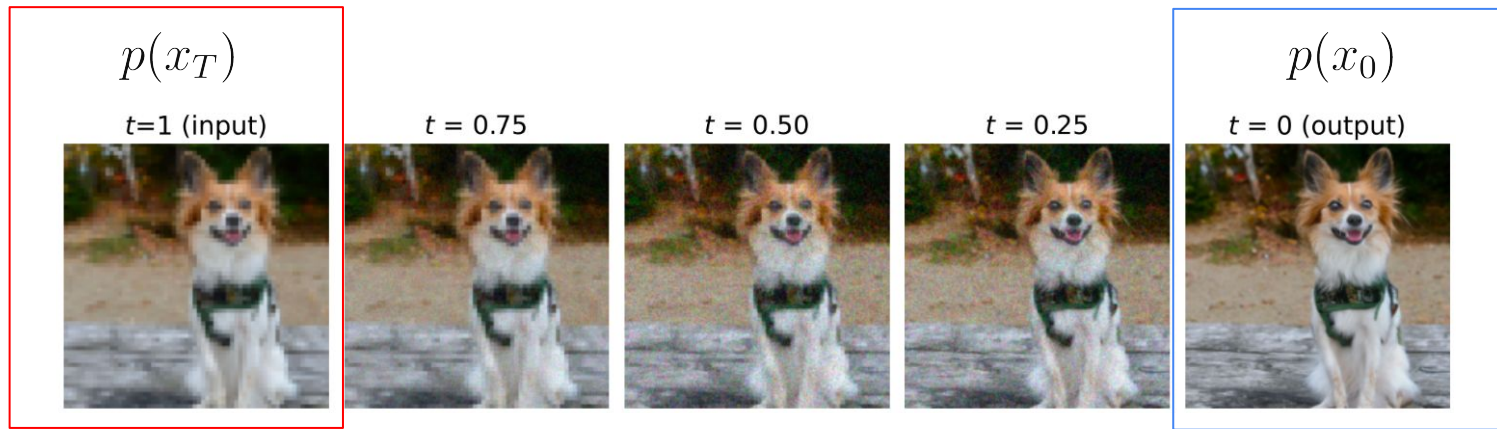
**General idea of diffusion models:** Define a forward SDE that gradually transforms data into noise. Then, learn the reverse process by estimating the score function using score matching.



**The original question remains:** Diffusion models are known for their superior image generation quality. Can we adapt them to image-to-image translation tasks?

# What exactly do we want to achieve?

**Motivation for diffusion bridge models:** Build a diffusion process that directly transforms the input distribution  $p(x_T)$  into the target distribution  $p(x_0)$ .



Generation process using the constructed diffusion:

$$dx_t = f_{\text{Bridge}}(x_t, t)dt + g(t)d\bar{w}_t, x_T \sim p(x_T).$$

**Unlike standard diffusion which starts from noise, here we start from structured input.**

# Diffusion bridge

Let  $Q$  be a forward diffusion process (the "Prior") over  $[0, T]$  in  $\mathbb{R}^D$ :

$$\text{Prior } Q : \quad dx_t = f(x_t, t)dt + g(t)dw_t,$$

Where  $f(x_t, t) : \mathbb{R}^D \times [0, T] \rightarrow \mathbb{R}^D$  is a drift function and  $g(t) : [0, T] \rightarrow \mathbb{R}^D$  is a noise schedule.

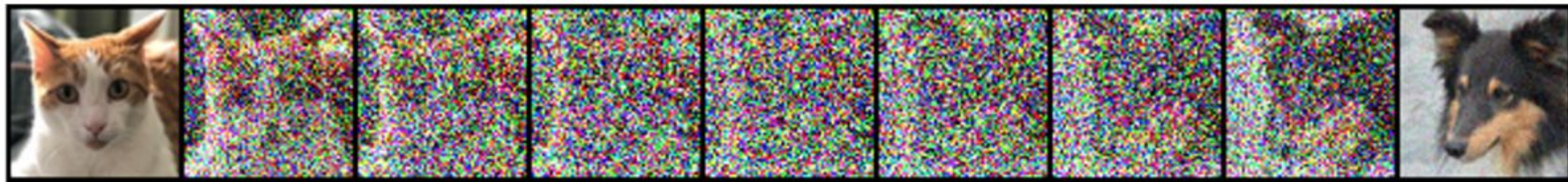
**Diffusion bridge** is the conditional stochastic process  $Q_{|x_0, x_T}$ , where the trajectory is fixed at given  $x_0$  and  $x_T$ . It can be derived using the Doob h-transform [2, 3].

**Example.** For  $f(x_t, t) = 0$  and  $g(t) = \sigma$  diffusion bridge is the interpolation combined with noising and denoising:

$x_0$

Diffusion Bridge  $Q_{|x_0, x_T}$

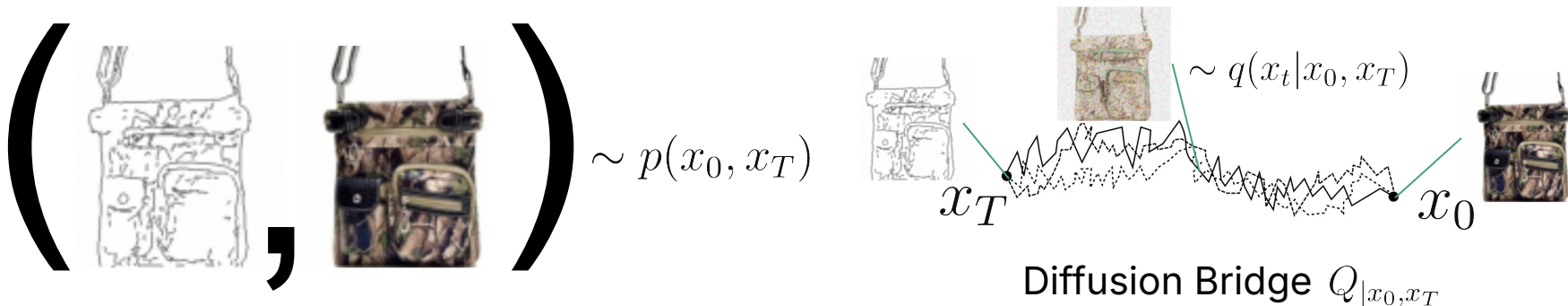
$x_T$





# Bridge Matching: General Approach to Build Diffusion Bridge Models

1. Build a mixture of diffusion bridges using paired data coupling  $p(x_0, x_T)$  and a diffusion bridge  $Q_{|x_0, x_T}$ :



2. Learn a diffusion that approximates this mixture by solving the Bridge Matching [1, 4, 5, 6] problem — a generalization of Flow Matching.

$$dx_t = \{f(x_t, t) - g^2(t)v^*(x_t, t)\}dt + g(t)d\bar{w}_t, x_T \sim p(x_T)$$

where the drift  $v^*(x_t, t)$  is learned via solving Bridge Matching problem:

$$\min_{\phi} \mathbb{E}_{x_0, t, x_t} [\|v_{\phi}(x_t, t) - \nabla_{x_t} \log q(x_t | x_0)\|^2], \quad \text{Known analytically for simple } Q, \text{ such as linear SDEs.}$$

$$(x_0, x_T) \sim p(x_0, x_T), \quad t \sim U([0, T]), \quad x_t \sim q(x_t | x_0, x_T).$$

# Diffusion bridge models

The **diffusion bridge model** is provided by the reverse-time SDE:

$$dx_t = \{f(x_t, t) - g^2(t)v^*(x_t, t)\}dt + g(t)d\bar{w}_t, x_T \sim p(x_T)$$

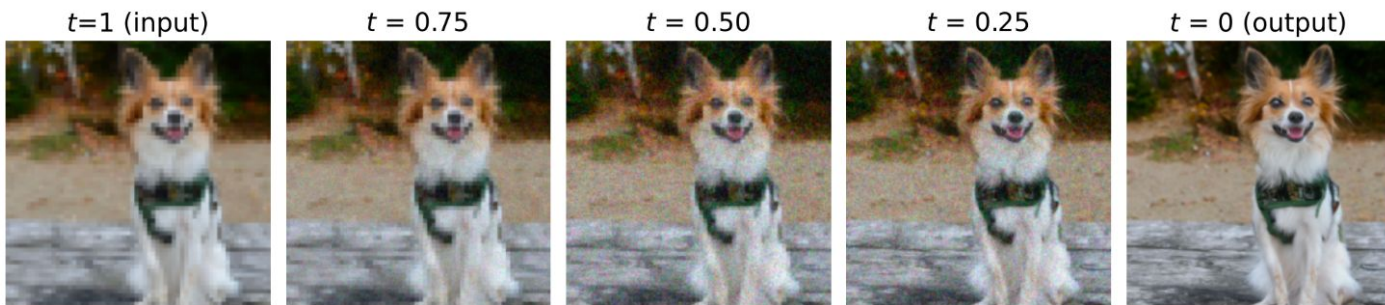
The drift  $v^*(x_t, t)$  is learned via solving Bridge Matching problem:

$$\min_{\phi} \mathbb{E}_{x_0, t, x_t} [\|v_{\phi}(x_t, t) - \nabla_{x_t} \log q(x_t|x_0)\|^2],$$

Known analytically for simple  $Q$ , such as linear SDEs.

$$(x_0, x_T) \sim p(x_0, x_T), t \sim U([0, T]), x_t \sim q(x_t|x_0, x_T).$$

After learning one can simulate the reverse time SDE using numerical solvers (10-1000 steps) to transform input (corrupted) image to the clean image:



Generation process

# Conditional and unconditional diffusion bridge models [7].

There are **two types** of diffusion bridge models (DBMs) which differ only by introducing additional conditioning on the final point  $x_T$ .

## Unconditional DBMs:

The goal is to learn SDE drift  $v(x_t, t)$  by optimizing:

$$\min_{\phi} \mathbb{E}_{x_0, t, x_t} [\|v_{\phi}(x_t, t) - \nabla_{x_t} \log q(x_t|x_0)\|^2],$$
$$(x_0, x_T) \sim p(x_0, x_T), \quad t \sim U([0, T]), \quad x_t \sim q(x_t|x_0, x_T).$$

Examples: [I2SB \[1\]](#), [Stochastic Interpolants \[8\]](#).

## Conditional DBMs:

The goal is to learn SDE drift  $v(x_t, t, x_T)$  by optimizing:

$$\min_{\phi} \mathbb{E}_{x_0, t, x_t, \mathbf{x}_T} [\|v_{\phi}(x_t, t, \mathbf{x}_T) - \nabla_{x_t} \log q(x_t|x_0)\|^2]$$
$$(x_0, x_T) \sim p(x_0, x_T), \text{ and } x_t \sim q(x_t|x_0, x_T).$$

Examples: [DDBM \[2\]](#), [GOUB \[9\]](#), [ResShift \[10\]](#).

**Theoretical aspects:** Conditional DBMs can be represented as classical diffusion models with a modified forward process. This allows adaptation of acceleration techniques developed for classical DMs. In contrast, unconditional DBMs cannot be expressed this way.



# Bridge Matching in practice and $x_0$ reparametrization

In practice, the Prior process  $Q$  is chosen such, that both transitional density  $q(x_t|x_0) = \mathcal{N}(x_t|\alpha_t x_0, \sigma^2 I)$  and bridge density  $q(x_t|x_0, x_T)$  are Gaussian. In this case  $\nabla_{x_t} \log q(x_t|x_0) = -\frac{x_t - \alpha_t x_0}{\sigma_t^2}$ .

Hence, one can use change of variables:  $v(x_t, t, x_T) = -\frac{x_t - \alpha_t \hat{x}_0(x_t, t, x_T)}{\sigma_t^2}$

Original problem:

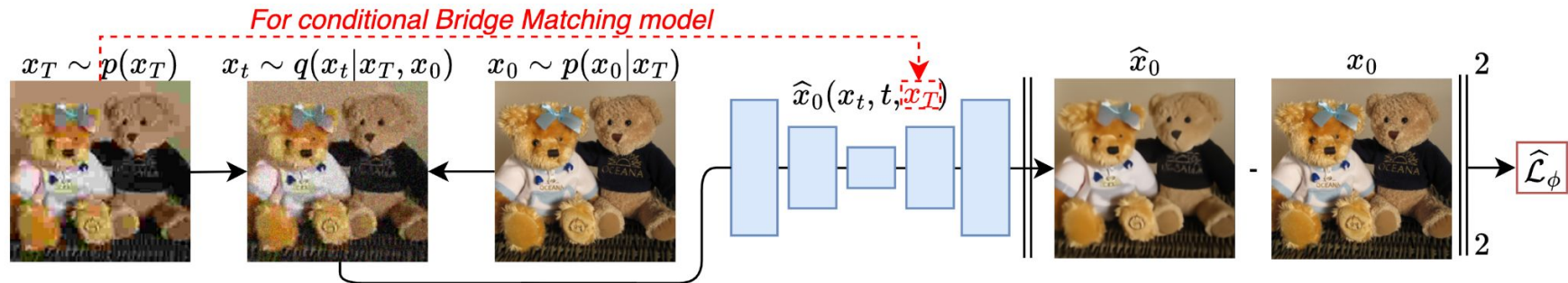
$$\min_{\phi} \mathbb{E}_{x_0, t, x_t, \mathbf{x}_T} [\|v_{\phi}(x_t, t, \mathbf{x}_T) - \nabla_{x_t} \log q(x_t|x_0)\|^2] \\ (x_0, x_T) \sim p(x_0, x_T), \text{ and } x_t \sim q(x_t|x_0, x_T).$$



Reparameterized problem:

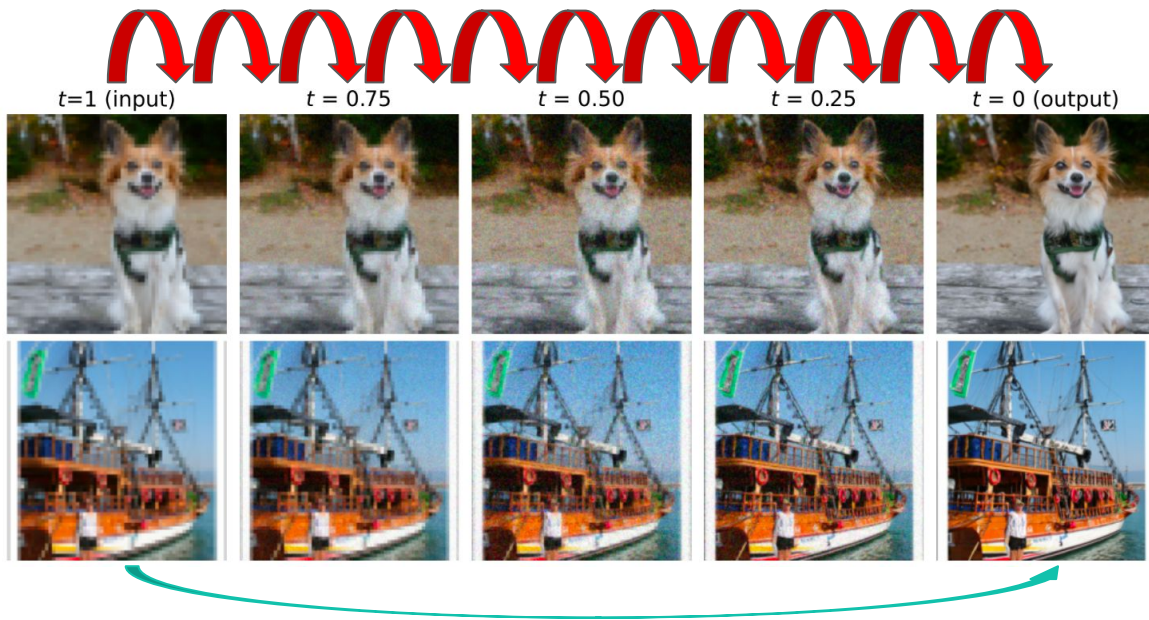
$$\min_{\phi} \mathbb{E}_{x_0, t, x_t, \mathbf{x}_T} [\lambda(t) \|\hat{x}_0^{\phi}(x_t, t, \mathbf{x}_T) - x_0\|^2], \quad (7) \\ (x_0, x_T) \sim p(x_0, x_T), t \sim U([0, T]), x_t \sim q(x_t|x_0, x_T),$$

## Learning pipeline for the reparameterized Diffusion Bridge model (Bridge Matching model)



# Distillation of diffusion bridge models

**Long inference problem:** Diffusion bridge models produce high-quality translations, but — like classical diffusion models — they require 10 to 1000 of steps to simulate the reverse SDE.



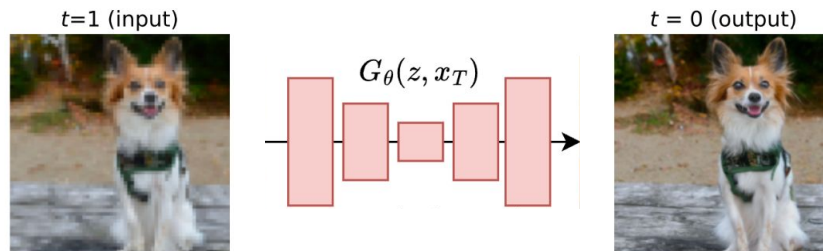
The goal of distillation is to train a new 1-step or few-step generator that mimics the full diffusion bridge model.

# Inverse Bridge Matching Distillation [11]

The core idea is to learn a one-step generator  $G_\theta$  of clean images from corrupted such that Diffusion Bridge Model for generated data coupling  $p_\theta(x_0, x_T)$  matches the teacher Diffusion Bridge Model learned on the ground-truth data:

1. We parameterize stochastic map  $p_\theta(x_0|x_T)$  by the one step stochastic generator  $G_\theta(x_T, z)$ ,

$$z \sim N(0, I), x_T \sim p(x_T).$$



2. We learn the generator  $G_\theta(x_T, z)$  in such way, that diffusion model for the generated data  $x_0^\theta$  matches the pre-trained teacher data  $x_0^*$ .

$$\begin{aligned} & \min_{\theta} \mathbb{E}_{x_t, t, x_0, \mathbf{x}_T} [\lambda(t) \|\hat{x}_0^*(x_t, t, \mathbf{x}_T) - \hat{x}_0^\theta(x_t, t, \mathbf{x}_T)\|^2], \\ & \text{s.t. } \hat{x}_0^\theta = \arg \min_{\hat{x}_0} \mathbb{E}_{x_t, t, x_0, \mathbf{x}_T} [\lambda(t) \|\hat{x}_0(x_t, t, \mathbf{x}_T) - x_0\|^2], \\ & (x_0, x_T) \sim p_\theta(x_0, x_T), \quad t \sim U([0, T]), \quad x_t \sim q(x_t|x_0, x_T), \\ & \text{Data from } G_\theta \end{aligned}$$

**Intractable gradients problem** since the objective includes term with argminimum.

# Tractable objective for the inverse bridge matching problem.

**Solution:** we show that the original constrained problem can be reformulated in the unconstrained one and used in practise:

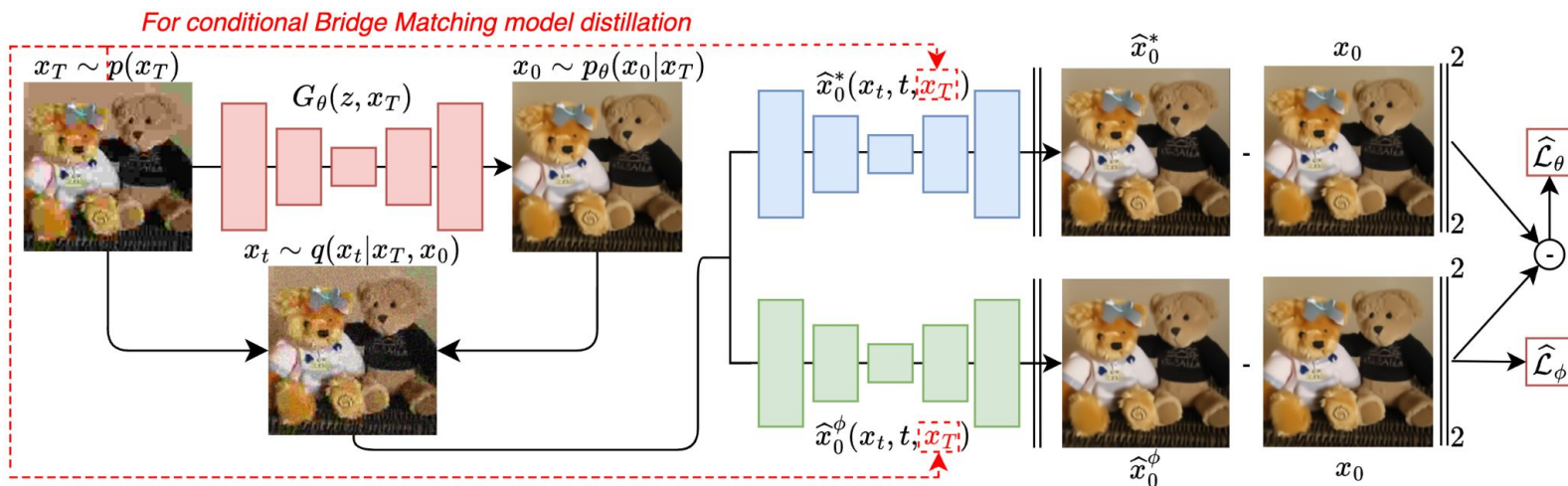
$$\min_{\theta} \left[ \mathbb{E}_{x_t, t, x_0, \mathbf{x}_T} [\lambda(t) \|\hat{x}_0^*(x_t, t, \mathbf{x}_T) - x_0\|^2] \right] - \text{Teacher model}$$

$$\min_{\phi} \mathbb{E}_{x_t, t, x_0, \mathbf{x}_T} [\lambda(t) \|\hat{x}_0^{\phi}(x_t, t, \mathbf{x}_T) - x_0\|^2] , \quad \text{Model for } G_{\theta}$$

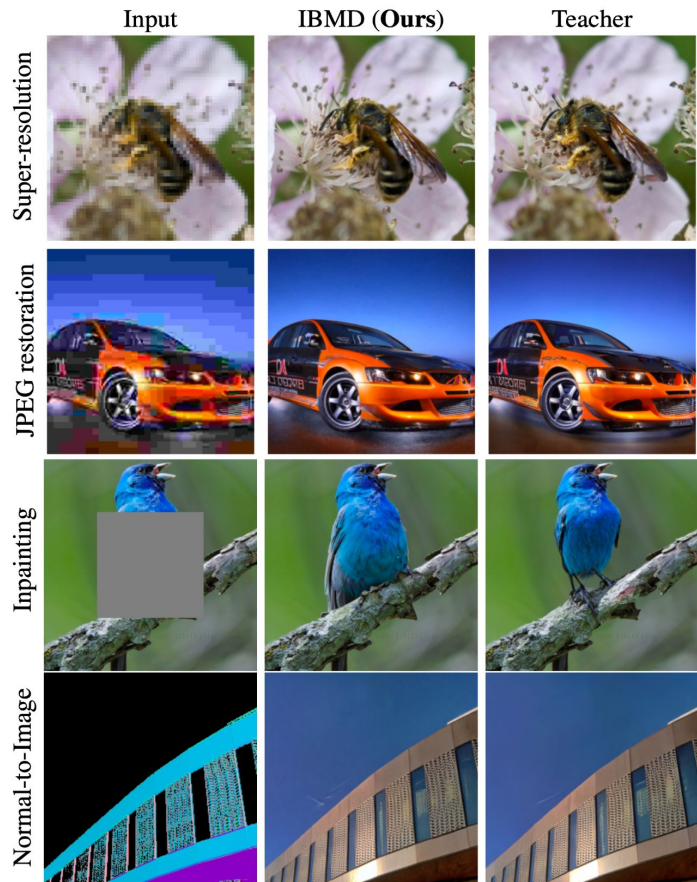
$$(x_0, x_T) \sim p_{\theta}(x_0, x_T), \quad t \sim U([0, T]), \quad x_t \sim q(x_t | x_0, x_T).$$

Data from  $G_{\theta}$

## Learning pipeline for the inverse bridge matching distillation



# The results of Inverse Bridge Matching Distillation (IBMD, ours).



Unconditional model,  
1-step IBMD (ours)  
beats **1000-step**  
teacher model.

Unconditional model,  
1-step IBMD (ours)  
accelerate inference  
up to **100 times**.

Conditional model,  
IBMD (ours)  
accelerate inference  
up to **5 times**.

Conditional model,  
IBMD (ours)  
accelerate inference  
in **50 times**.

We show the applicability of our IBMD distillation method in both *unconditional* and *conditional* settings.

We evaluate IBMD on *diverse tasks*, including super-resolution, JPEG restoration, inpainting, and sketch-to-image translation.

In all settings, IBMD achieves *significant inference speedups* (up to 100×) and sometimes *improves generation quality over the teacher*.



# IBMD (ours) outperforms previous approaches for DBMs acceleration

| 4× super-resolution (bicubic)         | ImageNet (256 × 256) |            |             |
|---------------------------------------|----------------------|------------|-------------|
|                                       | NFE                  | FID ↓      | CA ↑        |
| DDRM (Kawar et al., 2022)             | 20                   | 21.3       | 63.2        |
| DDNM (Wang et al., 2023)              | 100                  | 13.6       | 65.5        |
| PIGDM (Song et al., 2023)             | 100                  | 3.6        | 72.1        |
| ADM (Dhariwal & Nichol, 2021)         | 1000                 | 14.8       | 66.7        |
| CDSB (Shi et al., 2022)               | 50                   | 13.6       | 61.0        |
| I <sup>2</sup> SB (Liu et al., 2023a) | 1000                 | 2.8        | 70.7        |
| <b>IBMD-I<sup>2</sup>SB (Ours)</b>    | <b>1</b>             | <b>2.5</b> | <b>72.4</b> |

Table 2. Results on the image JPEG restoration task with QF=5. Baseline results are taken from I<sup>2</sup>SB (Liu et al., 2023a).

| JPEG restoration, QF= 5.              | ImageNet (256 × 256) |            |      |
|---------------------------------------|----------------------|------------|------|
|                                       | NFE                  | FID ↓      | CA ↑ |
| DDRM (Kawar et al., 2022)             | 20                   | 28.2       | 53.9 |
| PIGDM (Song et al., 2023)             | 100                  | 8.6        | 64.1 |
| Palette (Saharia et al., 2022)        | 1000                 | 8.3        | 64.2 |
| CDSB (Shi et al., 2022)               | 50                   | 38.7       | 45.7 |
| I <sup>2</sup> SB (Liu et al., 2023a) | 1000                 | <b>4.6</b> | 67.9 |
| I <sup>2</sup> SB (Liu et al., 2023a) | 100                  | 5.4        | 67.5 |
| <b>IBMD-I<sup>2</sup>SB (Ours)</b>    | <b>1</b>             | <b>5.3</b> | 67.2 |

| 4× super-resolution (pool)            | ImageNet (256 × 256) |            |             |
|---------------------------------------|----------------------|------------|-------------|
|                                       | NFE                  | FID ↓      | CA ↑        |
| DDRM (Kawar et al., 2022)             | 20                   | 14.8       | 64.6        |
| DDNM (Wang et al., 2023)              | 100                  | 9.9        | 67.1        |
| PIGDM (Song et al., 2023)             | 100                  | 3.8        | 72.3        |
| ADM (Dhariwal & Nichol, 2021)         | 1000                 | 3.1        | 73.4        |
| CDSB (Shi et al., 2022)               | 50                   | 13.0       | 61.3        |
| I <sup>2</sup> SB (Liu et al., 2023a) | 1000                 | 2.7        | 71.0        |
| <b>IBMD-I<sup>2</sup>SB (Ours)</b>    | <b>1</b>             | <b>2.6</b> | <b>72.7</b> |

Table 4. Results on the image JPEG restoration task with QF=10. Baseline results are taken from I<sup>2</sup>SB (Liu et al., 2023a).

| JPEG restoration, QF= 10.             | ImageNet (256 × 256) |            |      |
|---------------------------------------|----------------------|------------|------|
|                                       | NFE                  | FID ↓      | CA ↑ |
| DDRM (Kawar et al., 2022)             | 20                   | 16.7       | 64.7 |
| PIGDM (Song et al., 2023)             | 100                  | 6.0        | 71.0 |
| Palette (Saharia et al., 2022)        | 1000                 | 5.4        | 70.7 |
| CDSB (Shi et al., 2022)               | 50                   | 18.6       | 60.0 |
| I <sup>2</sup> SB (Liu et al., 2023a) | 1000                 | <b>3.6</b> | 72.1 |
| I <sup>2</sup> SB (Liu et al., 2023a) | 100                  | 4.4        | 71.6 |
| <b>IBMD-I<sup>2</sup>SB (Ours)</b>    | <b>1</b>             | <b>3.8</b> | 72.4 |

| Inpainting, Center (128 × 128)        | ImageNet (256 × 256) |             |             |
|---------------------------------------|----------------------|-------------|-------------|
|                                       | NFE                  | FID ↓       | CA ↑        |
| DDRM (Kawar et al., 2022)             | 20                   | 24.4        | 62.1        |
| PIGDM (Song et al., 2023)             | 100                  | 7.3         | <b>72.6</b> |
| DDNM (Wang et al., 2022)              | 100                  | 15.1        | 55.9        |
| Palette (Saharia et al., 2022)        | 1000                 | 6.1         | 63.0        |
| I <sup>2</sup> SB (Liu et al., 2023a) | 10                   | 5.4         | 65.97       |
| DBIM (Zheng et al., 2024)             | 50                   | 3.92        | 72.4        |
| DBIM (Zheng et al., 2024)             | 100                  | <b>3.88</b> | <b>72.6</b> |
| CBD (He et al., 2024)                 | 4                    | 5.34        | 69.6        |
| CBT (He et al., 2024)                 |                      | 4.77        | 70.3        |
| IBMD-I <sup>2</sup> SB (Ours)         |                      | 5.1         | 70.3        |
| IBMD-DDBM (Ours)                      |                      | <b>4.03</b> | <b>72.2</b> |
| CBD (He et al., 2024)                 | 2                    | 5.65        | 69.6        |
| CBT (He et al., 2024)                 |                      | 5.34        | 69.8        |
| IBMD-I <sup>2</sup> SB (Ours)         |                      | 5.3         | 65.7        |
| IBMD-DDBM (Ours)                      |                      | <b>4.23</b> | <b>72.3</b> |
| IBMD-I <sup>2</sup> SB (Ours)         | 1                    | 6.7         | 65.0        |
| IBMD-DDBM (Ours)                      |                      | <b>5.87</b> | <b>70.6</b> |

Our method (IBMD) outperforms previous acceleration methods based on consistency distillation (CBD/CBT) and more advanced sampling techniques (DBIM).



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