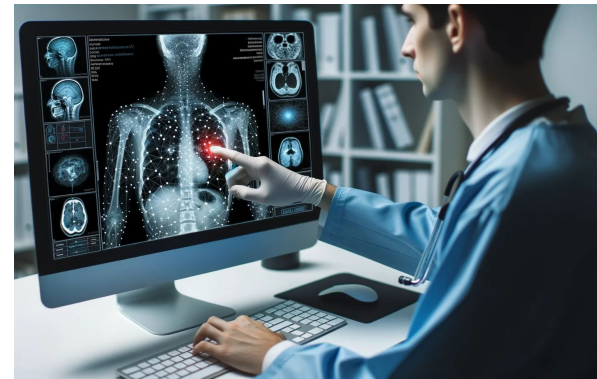
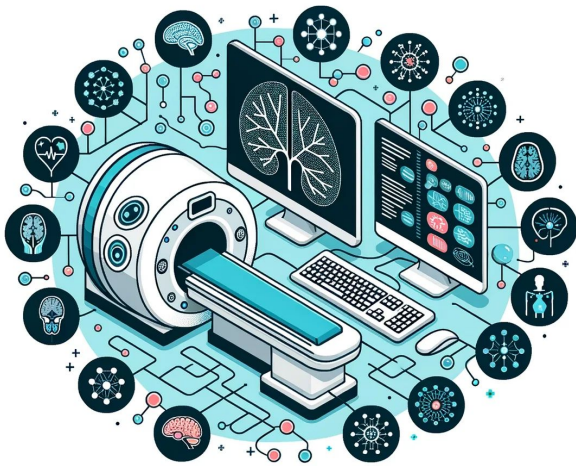


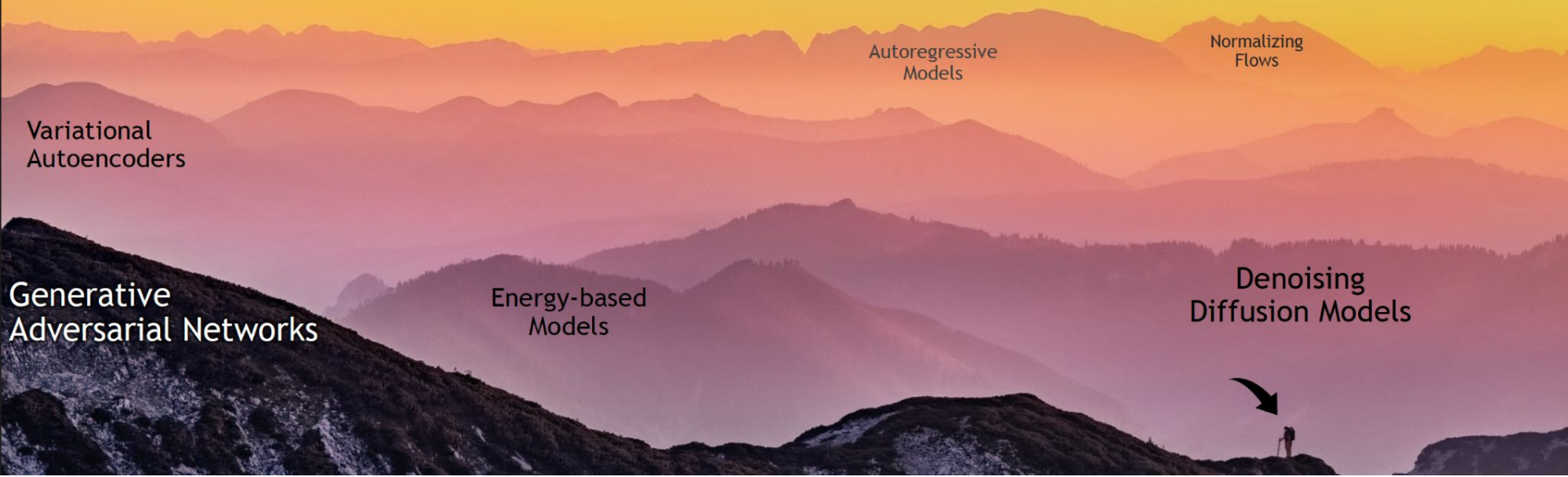
# Applied Deep Learning and Generative Models in Healthcare



**Session 5:** Medical Image Synthesis with Diffusion  
**Date:** Mar 08 2025

**Instructor:** Mahmoud E. Khani, Ph.D.

# The Landscape of Deep Generative Learning



Variational  
Autoencoders

Autoregressive  
Models

Normalizing  
Flows

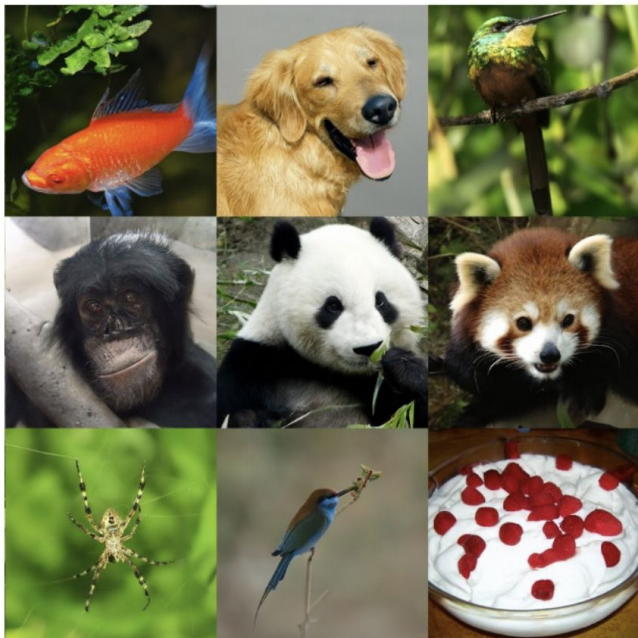
Generative  
Adversarial Networks

Energy-based  
Models

Denoising  
Diffusion Models



# Denoising Diffusion Models



[“Diffusion Models Beat GANs on Image Synthesis”](#)  
Dhariwal & Nichol, OpenAI, 2021



[“Cascaded Diffusion Models for High Fidelity Image Generation”](#)  
Ho et al., Google, 2021



# Text-to-Image Generation

## DALL·E 2

“a teddy bear on a skateboard in times square”



[“Hierarchical Text-Conditional Image Generation with CLIP Latents”](#)  
Ramesh et al., 2022

## Imagen

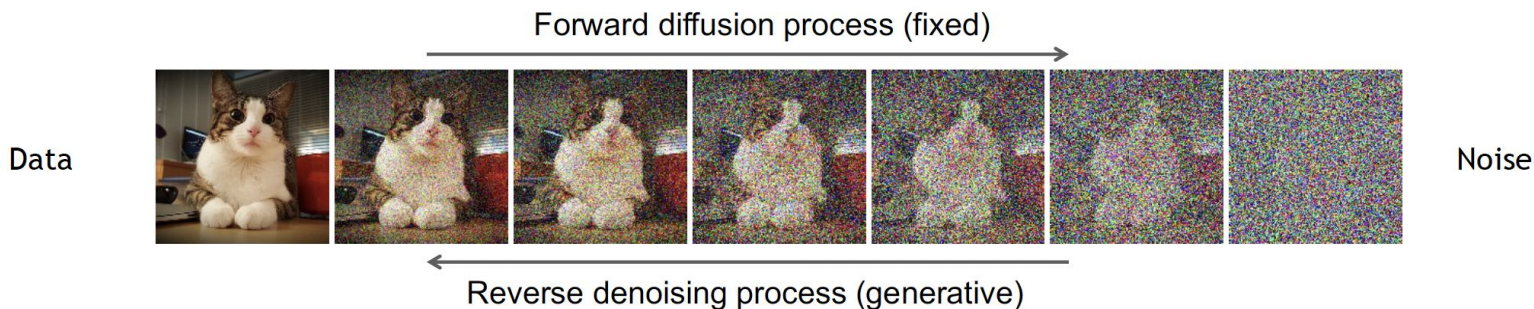
A group of teddy bears in suit in a corporate office celebrating the birthday of their friend. There is a pizza cake on the desk.



[“Photorealistic Text-to-Image Diffusion Models with Deep Language Understanding”](#), Saharia et al., 2022

# Learning to Generate by Denoising

- Denoising diffusion consists of two processes:
  - Forward diffusion to gradually add noise to input
  - Reverse denoising that learns to generate data by denoising



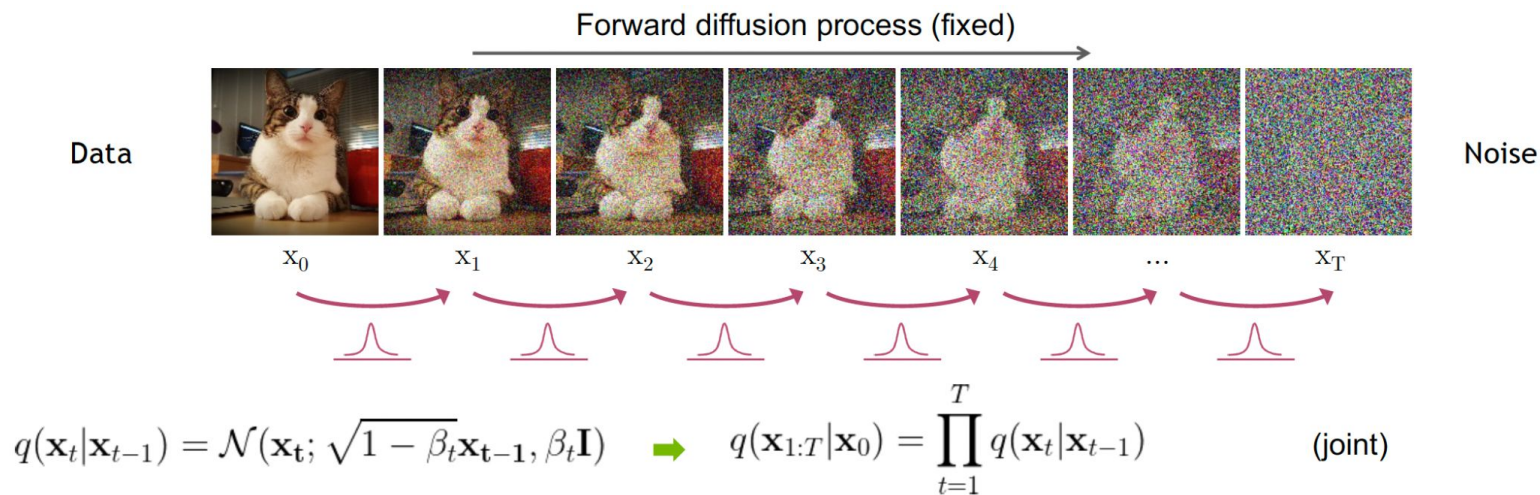
[Sohl-Dickstein et al., Deep Unsupervised Learning using Nonequilibrium Thermodynamics, ICML 2015](#)

[Ho et al., Denoising Diffusion Probabilistic Models, NeurIPS 2020](#)

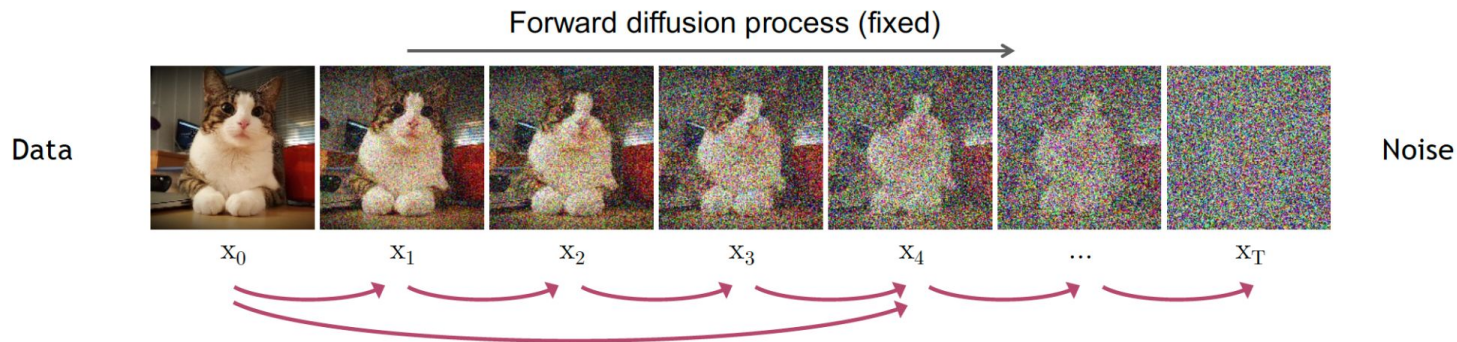
[Song et al., Score-Based Generative Modeling through Stochastic Differential Equations, ICLR 2021](#)

# Forward Diffusion Process

- The formal definition of the forward process in  $T$  steps:



# Diffusion Kernel



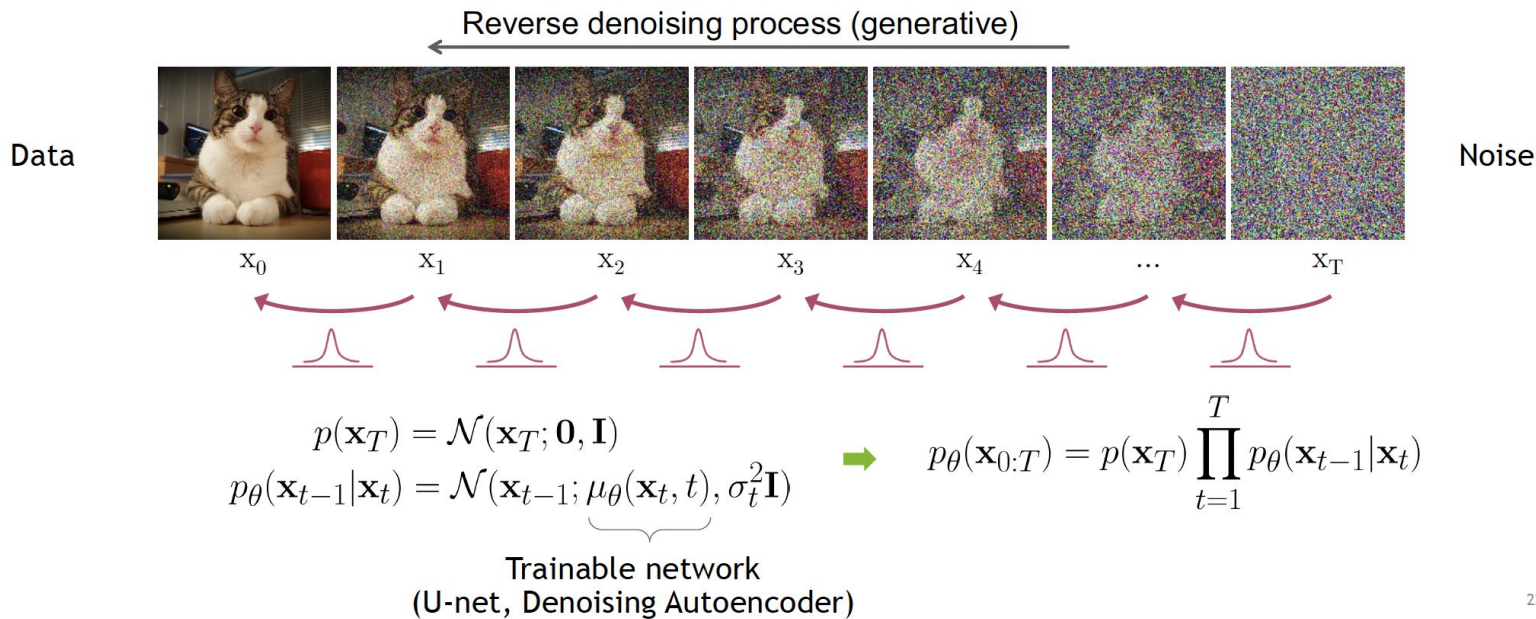
Define  $\bar{\alpha}_t = \prod_{s=1}^t (1 - \beta_s)$   $\rightarrow q(\mathbf{x}_t | \mathbf{x}_0) = \mathcal{N}(\mathbf{x}_t; \sqrt{\bar{\alpha}_t} \mathbf{x}_0, (1 - \bar{\alpha}_t) \mathbf{I})$  (Diffusion Kernel)

For sampling:  $\mathbf{x}_t = \sqrt{\bar{\alpha}_t} \mathbf{x}_0 + \sqrt{(1 - \bar{\alpha}_t)} \epsilon$  where  $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$

$\beta_t$  values schedule (i.e., the noise schedule) is designed such that  $\bar{\alpha}_T \rightarrow 0$  and  $q(\mathbf{x}_T | \mathbf{x}_0) \approx \mathcal{N}(\mathbf{x}_T; \mathbf{0}, \mathbf{I})$

# Reverse Denoising Process

- Formal definition of reverse process in  $T$  steps:





# Training and Sampling Algorithms

---

## Algorithm 1 Training

---

```
1: repeat  
2:    $\mathbf{x}_0 \sim q(\mathbf{x}_0)$   
3:    $t \sim \text{Uniform}(\{1, \dots, T\})$   
4:    $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$   
5:   Take gradient descent step on  
        $\nabla_{\theta} \|\epsilon - \epsilon_{\theta}(\sqrt{\bar{\alpha}_t} \mathbf{x}_0 + \sqrt{1 - \bar{\alpha}_t} \epsilon, t)\|^2$   
6: until converged
```

---

---

## Algorithm 2 Sampling

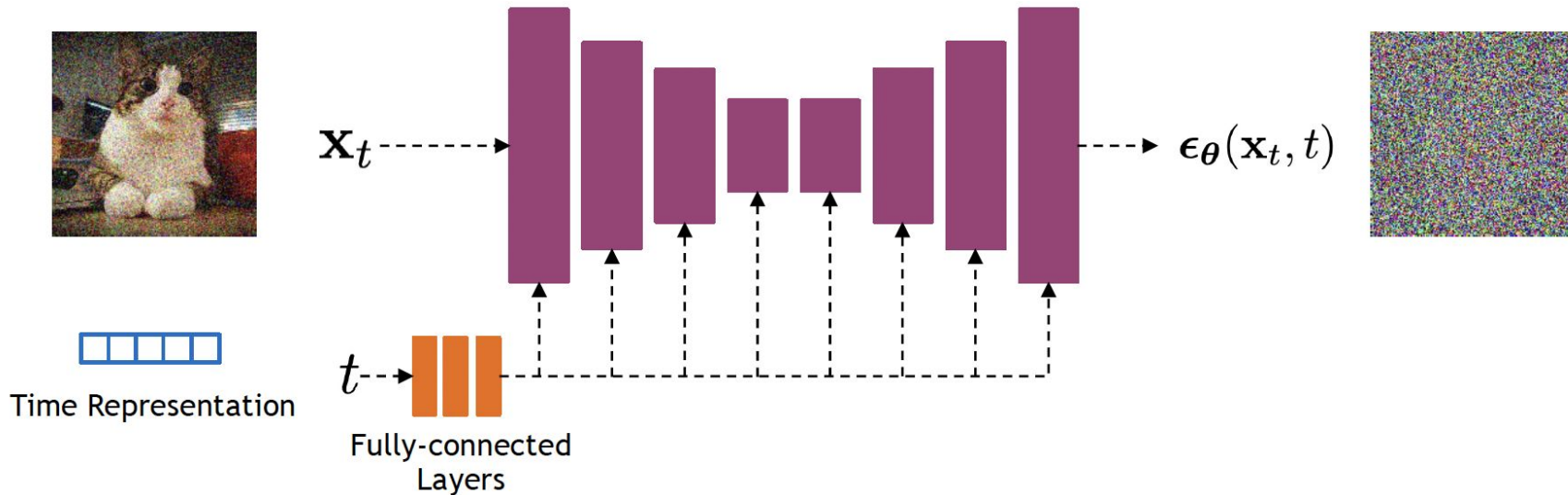
---

```
1:  $\mathbf{x}_T \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$   
2: for  $t = T, \dots, 1$  do  
3:    $\mathbf{z} \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$   
4:    $\mathbf{x}_{t-1} = \frac{1}{\sqrt{\alpha_t}} \left( \mathbf{x}_t - \frac{1 - \alpha_t}{\sqrt{1 - \bar{\alpha}_t}} \epsilon_{\theta}(\mathbf{x}_t, t) \right) + \sigma_t \mathbf{z}$   
5: end for  
6: return  $\mathbf{x}_0$ 
```

---

# Network Architectures

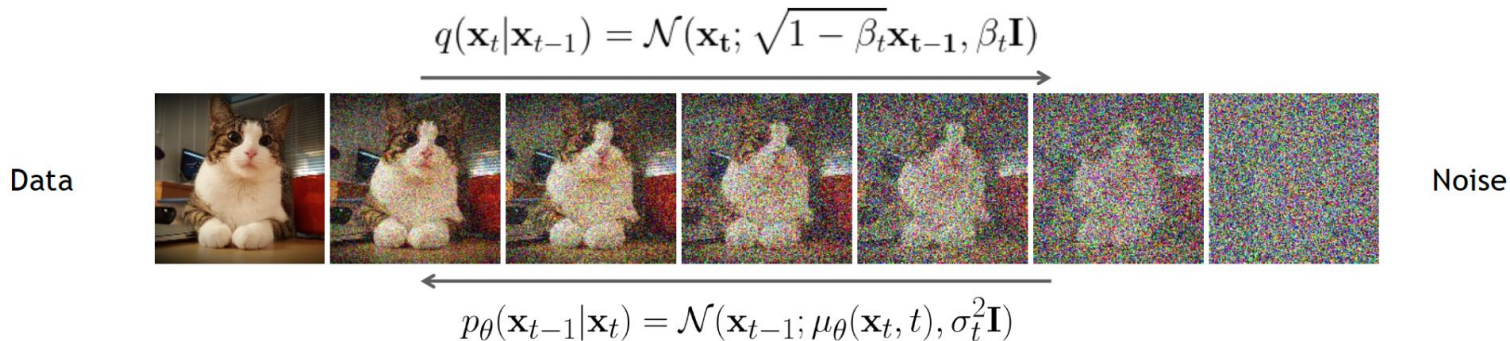
- Diffusion models often use **U-Net** architectures with **ResNet** blocks and **self-attention** layers



Time representation: sinusoidal positional embeddings or random Fourier features.

# Diffusion Parameters

- The noise schedule



Above,  $\beta_t$  and  $\sigma_t^2$  control the variance of the forward diffusion and reverse denoising processes respectively.

Often a linear schedule is used for  $\beta_t$ , and  $\sigma_t^2$  is set equal to  $\beta_t$ .

# Fourier Analysis of the Forward Process

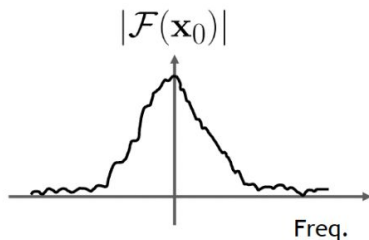
Recall that sampling from  $q(\mathbf{x}_t|\mathbf{x}_0)$  is done using  $\mathbf{x}_t = \sqrt{\bar{\alpha}_t} \mathbf{x}_0 + \sqrt{(1 - \bar{\alpha}_t)} \epsilon$  where  $\epsilon \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$

$$\mathbf{x}_t = \sqrt{\bar{\alpha}_t} \mathbf{x}_0 + \sqrt{(1 - \bar{\alpha}_t)} \epsilon$$

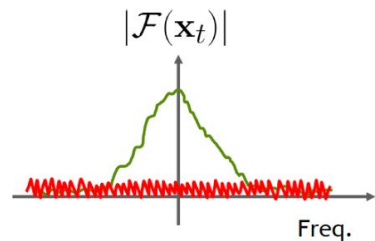


Fourier Transform

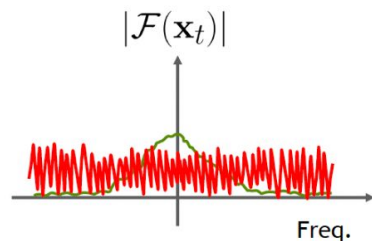
$$\mathcal{F}(\mathbf{x}_t) = \sqrt{\bar{\alpha}_t} \mathcal{F}(\mathbf{x}_0) + \sqrt{(1 - \bar{\alpha}_t)} \mathcal{F}(\epsilon)$$



Small  $t$   
 $\bar{\alpha}_t \sim 1$



Large  $t$   
 $\bar{\alpha}_t \sim 0$

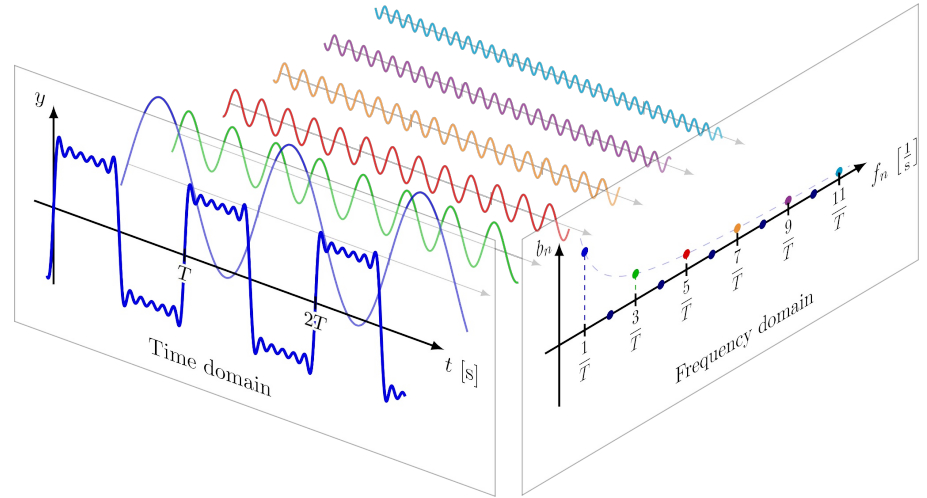
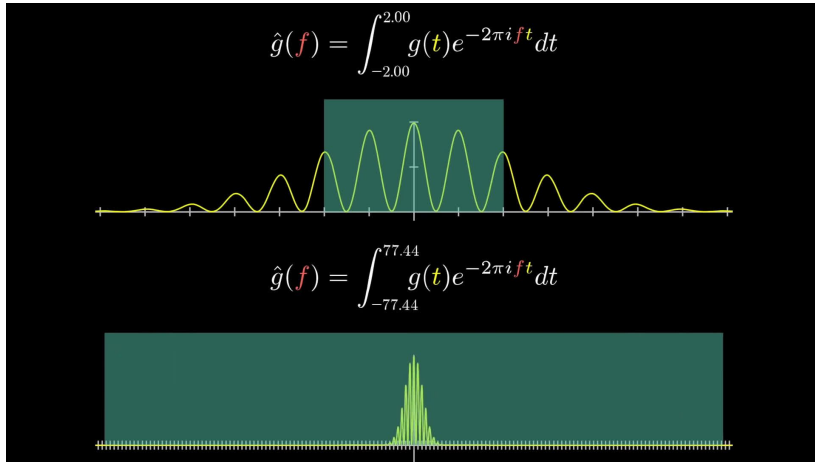


In the forward diffusion, the high frequency content is perturbed faster.



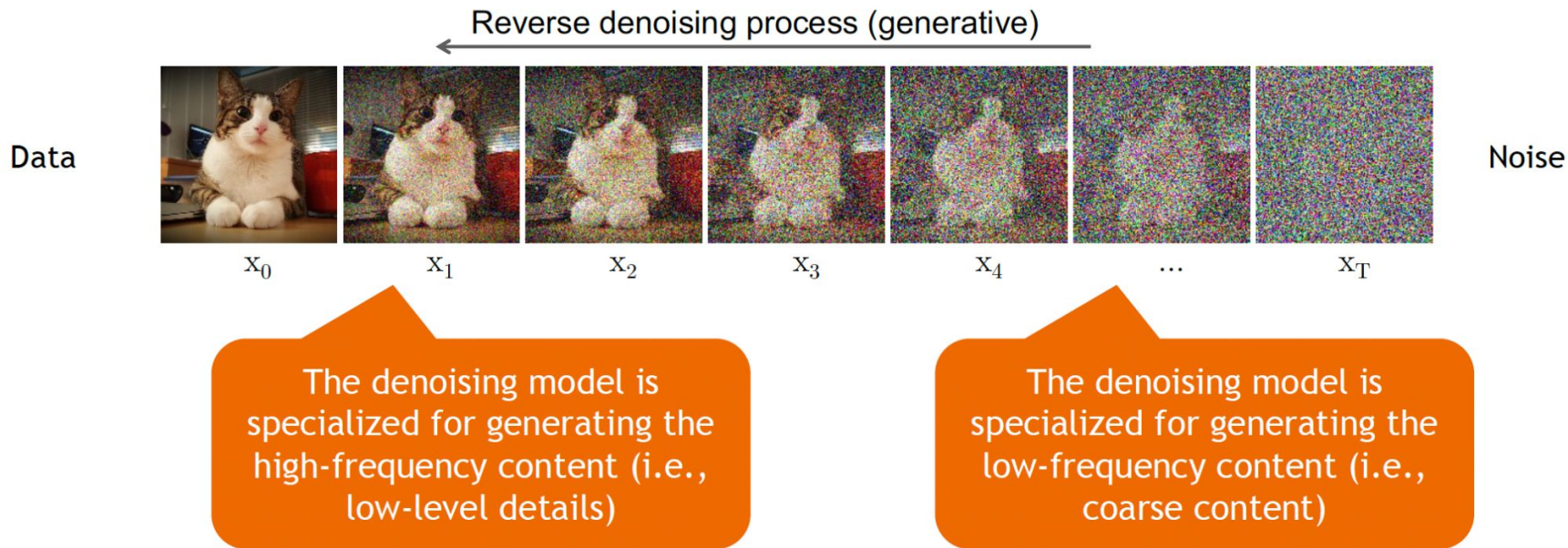
# Fourier Analysis

- Decomposition of signal into **frequency** components



<https://dibsmethodsmeetings.github.io/fourier-transforms/>

# Content-Detail Tradeoff

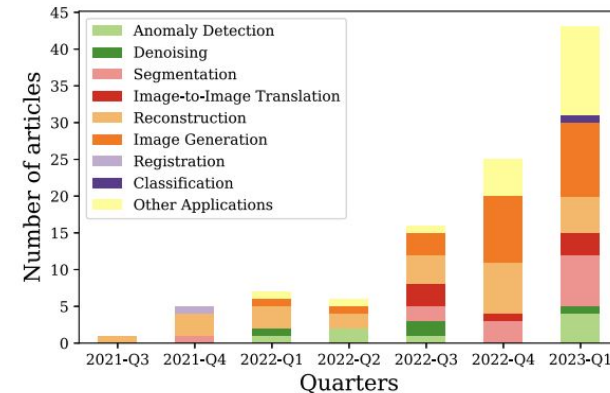
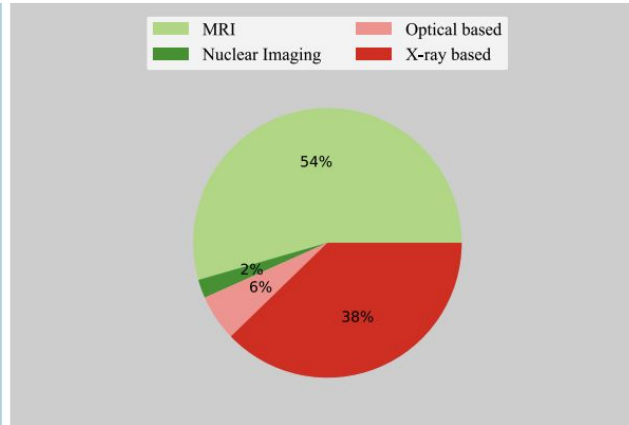
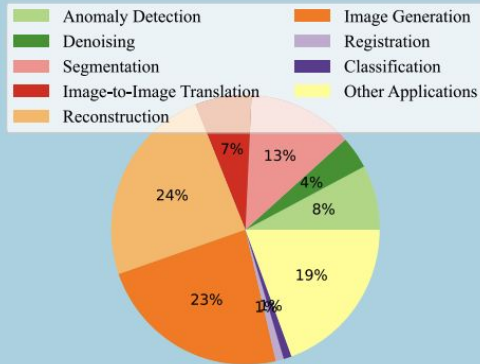


The weighting of the training objective for different timesteps is important!



# Diffusion Models in Medical Imaging

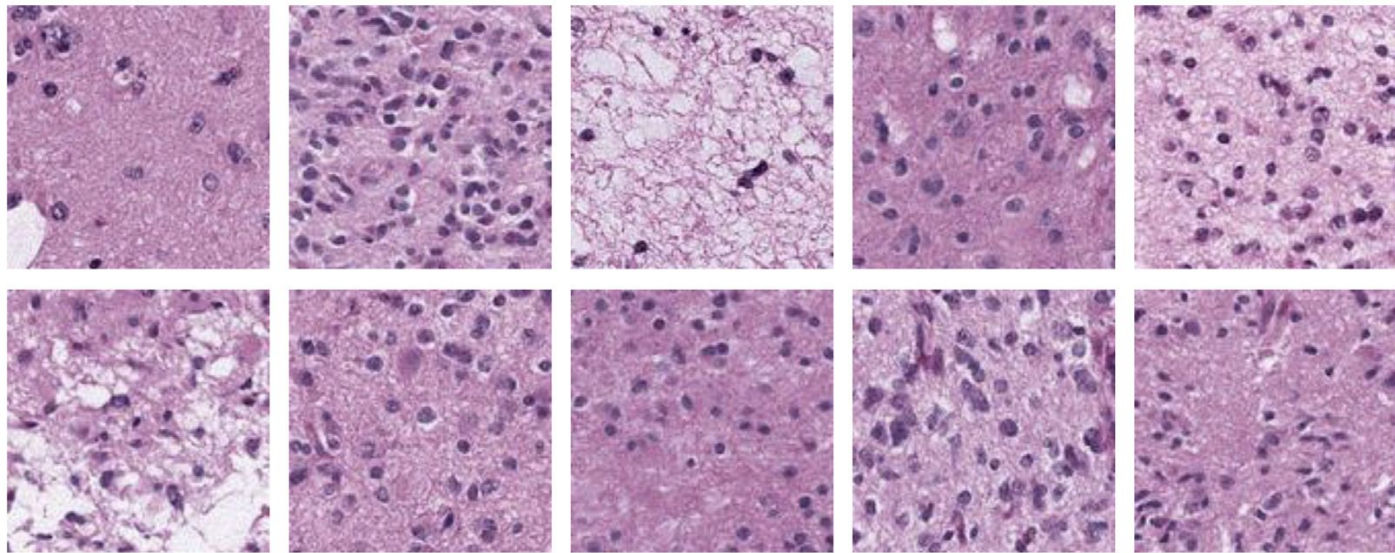
- Diffusion models have emerged as **powerful generative models** in medical imaging
  - improved sample quality, mode coverage, and versatility across various applications.





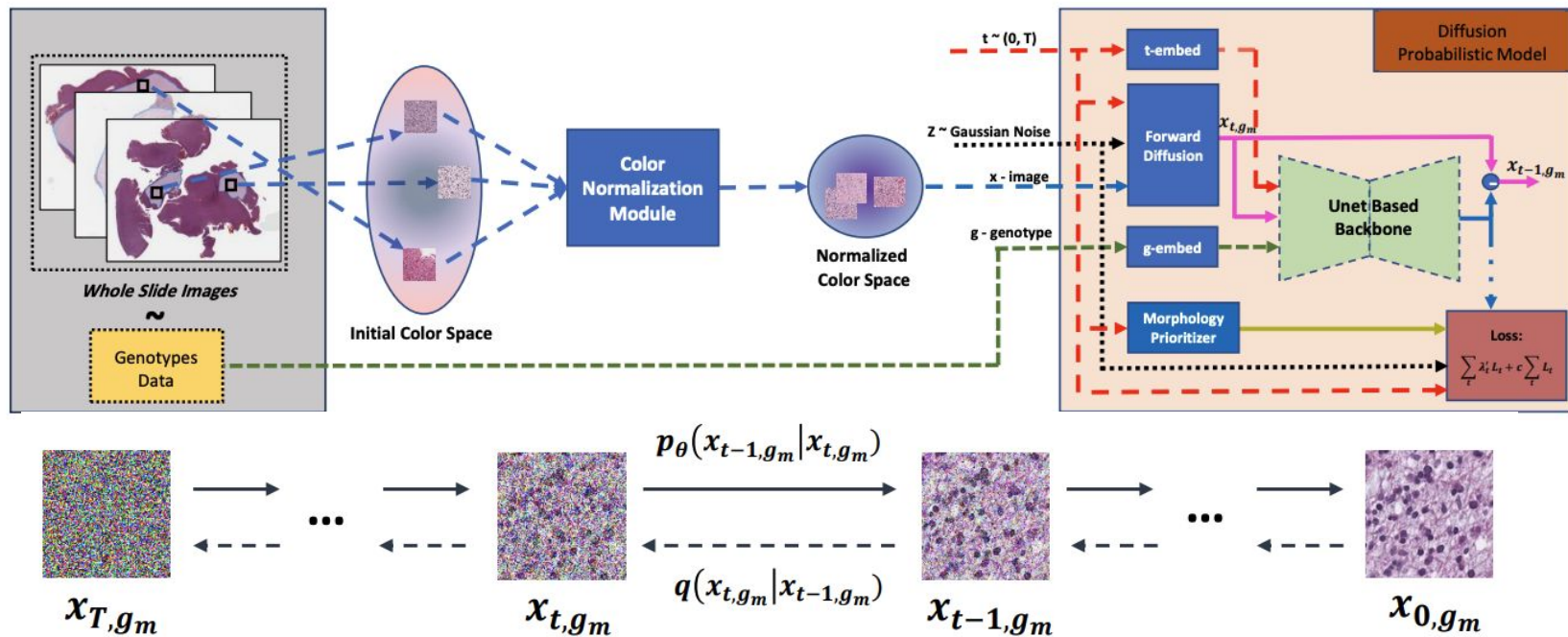
# Applications in medical image generation

- Diffusion models have remarkable performance in generating **synthetic medical images**
  - aiding data augmentation and rare disease representation



# Applications in medical image generation

- Generating **histopathology** images with **genotype** guidance



# Genotype-conditioned Image Generation

## How Are IDH Mutation and 1p/19q Co-Deletion Extracted for Training Data?

The training data for the diffusion model includes **paired histopathology images and genotype information**. The genotype labels (IDH mutation status and 1p/19q co-deletion) are extracted through **molecular and genetic tests**, including:

### 1. IDH Mutation Detection

- **Immunohistochemistry (IHC)**: Uses **antibodies** to detect **mutant IDH1 protein** in tumor samples.
- **Next-Generation Sequencing (NGS)**: Directly sequences the **IDH1/IDH2 genes** to detect mutations.

### 2. 1p/19q Co-Deletion Detection

- **Fluorescence In Situ Hybridization (FISH)**: A cytogenetic test that detects the **loss of chromosome arms 1p and 19q** in tumor cells.
- **Comparative Genomic Hybridization (CGH)**: Identifies chromosomal deletions.
- **PCR-based methods**: Detects loss of heterozygosity (LOH) in 1p and 19q regions.

Once the genetic data is obtained, it is **paired with corresponding histopathology images** to create a **genotype-labeled dataset** for training.

# Genotype-conditioned Image Generation

- **What is an IDH Mutation?**

- IDH (Isocitrate Dehydrogenase) Mutation refers to genetic alterations in the IDH1 or IDH2 genes.
- These mutations are commonly found in gliomas (brain tumors) and are important for tumor classification, prognosis, and treatment decisions.
- IDH-mutant gliomas tend to have better survival rates compared to IDH-wildtype gliomas, which are more aggressive.

- **What is 1p/19q Co-Deletion?**

- The 1p/19q co-deletion is a chromosomal alteration where parts of chromosomes 1p (short arm of chromosome 1) and 19q (long arm of chromosome 19) are missing.
- This is a key molecular marker used to classify gliomas.
- Gliomas with 1p/19q co-deletion are almost always oligodendrogliomas, which respond well to chemotherapy and radiation therapy.
- If a glioma has IDH mutation but no 1p/19q co-deletion, it is classified as an astrocytoma instead of an oligodendroglioma.

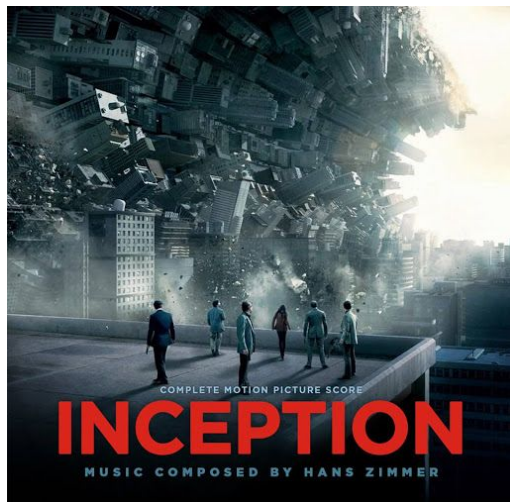


# Inception score

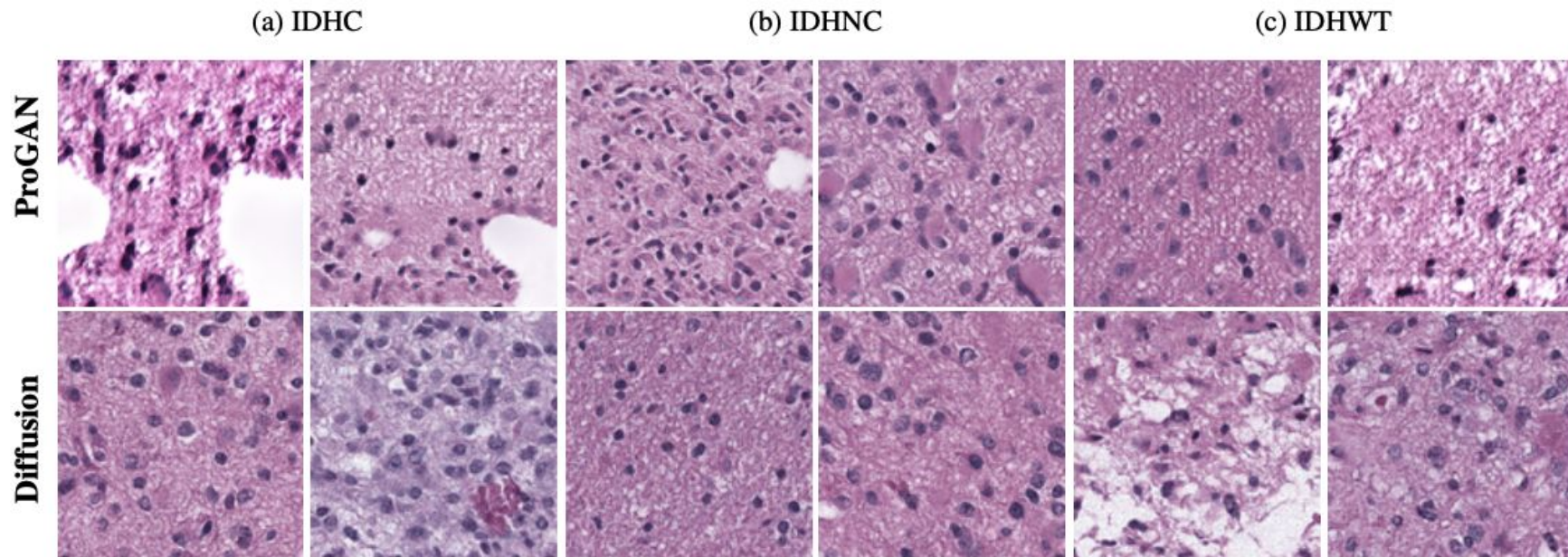
**What it measures:** The **quality** and **diversity** of generated images.

**How it works:**

- Uses a pre-trained **Inception network** (e.g., Inception v3) to classify generated images.
- Measures:
  - **Quality:** If a generated image is highly classifiable (i.e., strong class predictions).
  - **Diversity:** If the generated images cover multiple classes.
- Requires a **well-trained classifier on medical data**, unlike natural images (e.g., CIFAR, ImageNet).



# Applications in medical image generation



	ProGAN	Diffusion Model
Improved Recall	0.4816	<b>0.8528</b>
Improved Precision	0.0078	<b>0.2573</b>

	ProGAN	Diffusion Model
Inception Score	1.67	<b>2.08</b>
FID	53.85	<b>20.11</b>
sFID	24.37	<b>6.32</b>

**MedAI  
Group**



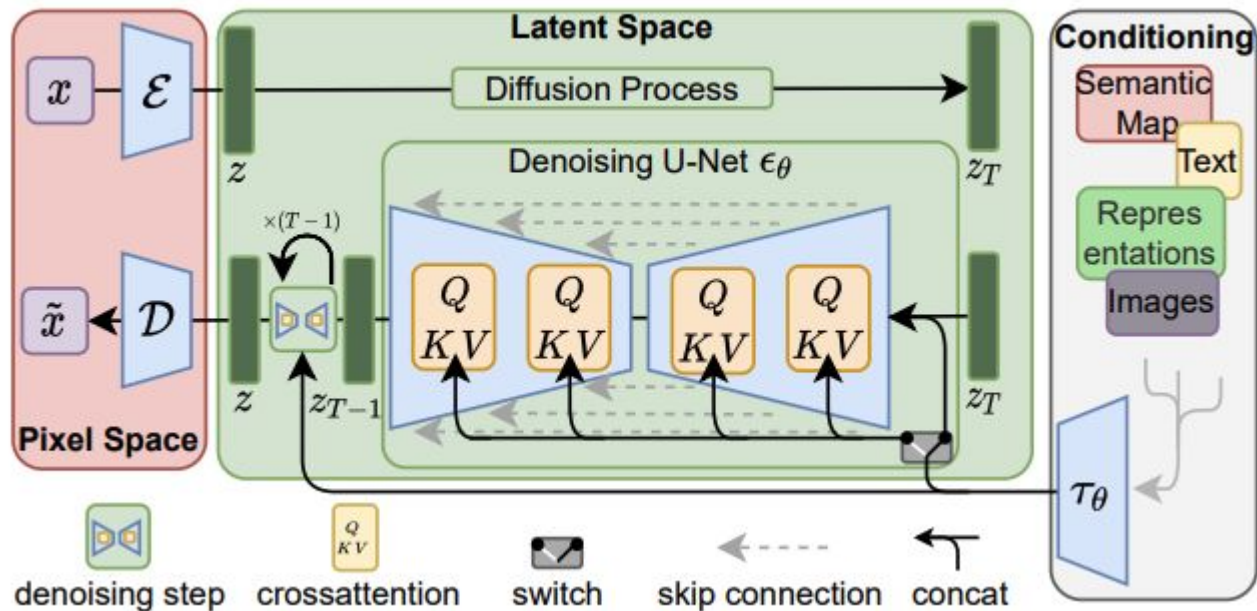
## Session #96: Denoising Diffusion Models for Medical Image Analysis

Julia Wolleb



<https://youtu.be/US9CzPrT2H8?si=Rrlt7CYwKE55rd5H>

# Conditional Diffusion



<https://arxiv.org/abs/2112.10752>

# Self-Attention Module

- **Purpose:** Captures long-range dependencies within feature maps.

## **Mechanism:**

- Flattens spatial dimensions into “tokens.”
- Applies multi-head self-attention (along with layer normalization and a feedforward network) to let every part of the feature map attend to every other part.
- **Role in Diffusion:** Helps preserve global structure during the denoising process.



# Self-attention vs Cross-attention

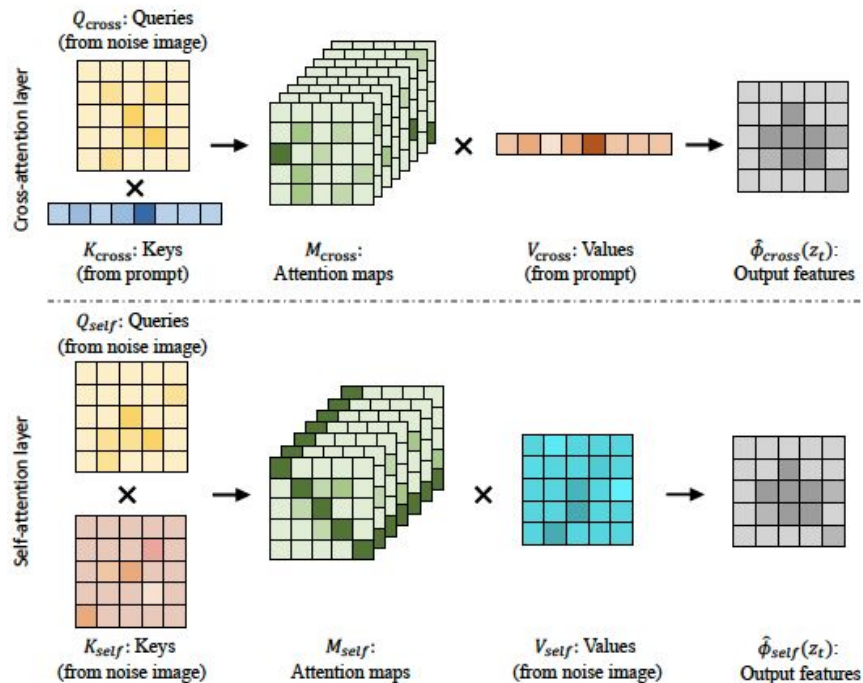


Figure 2. Cross and self-attention layers in Stable Diffusion.