

Enhancing Synchrotron Low-Dose Computed Tomography Image Quality Using Diffusion-Based Generative Models

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Supervisor: Prof. Dr. Zanoni Dias

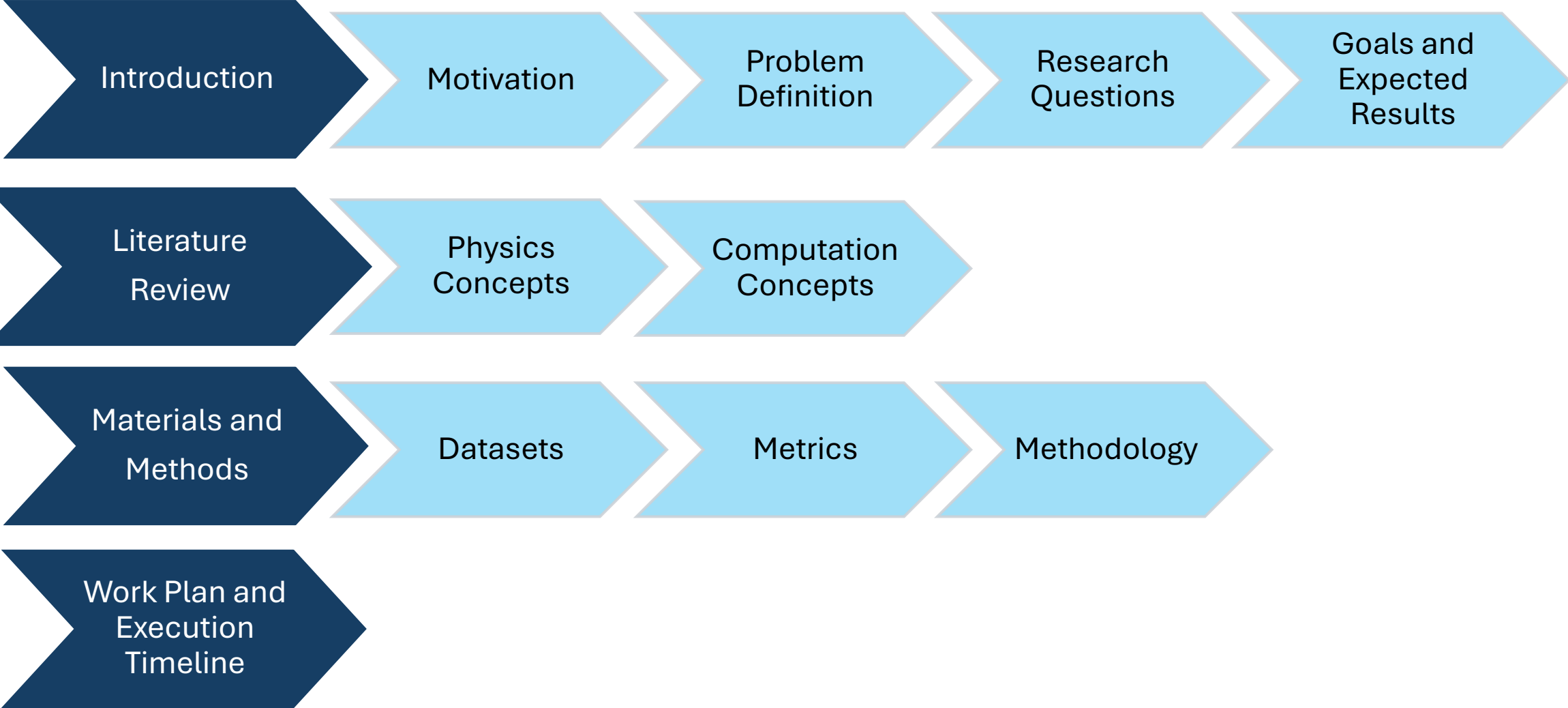
Co-supervisor: Prof. Dr. Hélio Pedrini

Master's Degree Proposal

Institute of Computing - University of Campinas

October 17th of 2024

Presentation Outline

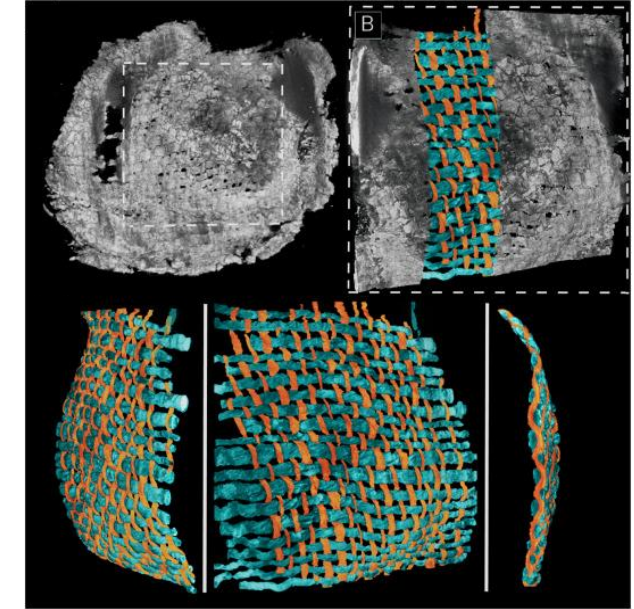


- X-Ray Computed Tomography (CT) offers a **non-invasive** technique for assessing internal structures of objects
- Applied over multiple domains:
 - Medical (golden standard for trauma assessment [43])
 - Archaeology [28]
 - Paleontology [23,44]
 - Material Science [50]
- Synchrotron facilities allow achieving **higher spatial and time resolutions** if compared to conventional X-Ray sources [16]

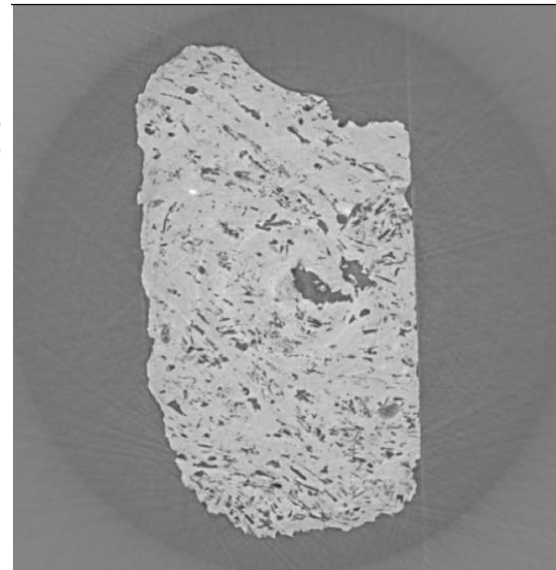


Source: Hamm et al. [23]

Tyrannosaurus rex left dentary

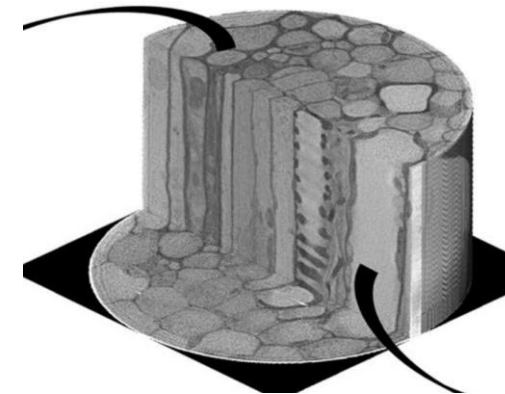


Source: Karjalainen et al. [28]

14th Century fabric

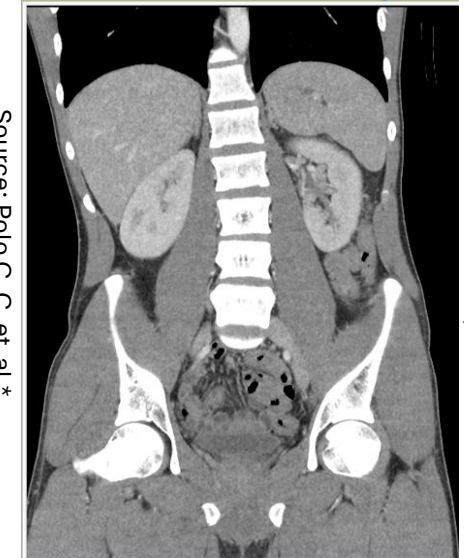
Source: MOGNO Beamline from Sirius

Carbonatic Rock



Plant Tissue

Source: Polo C. C. et al.*



Source: Wikipedia

Human body (Axial plane)

- High doses of radiation may be harmful for health [9,40] while for radiation-sensitive samples it may cause damages to it and directly impact the experimental results [36]
- Development of Low-dose Computed Tomography (LDCT) techniques is crucial
 - As Low As Reasonably Achievable (ALARA) principle
- Lower dose = lower CT image quality [17,49,51]
 - Higher noise
 - Lower contrast
- Methods to enhance LDCT image quality is crucial



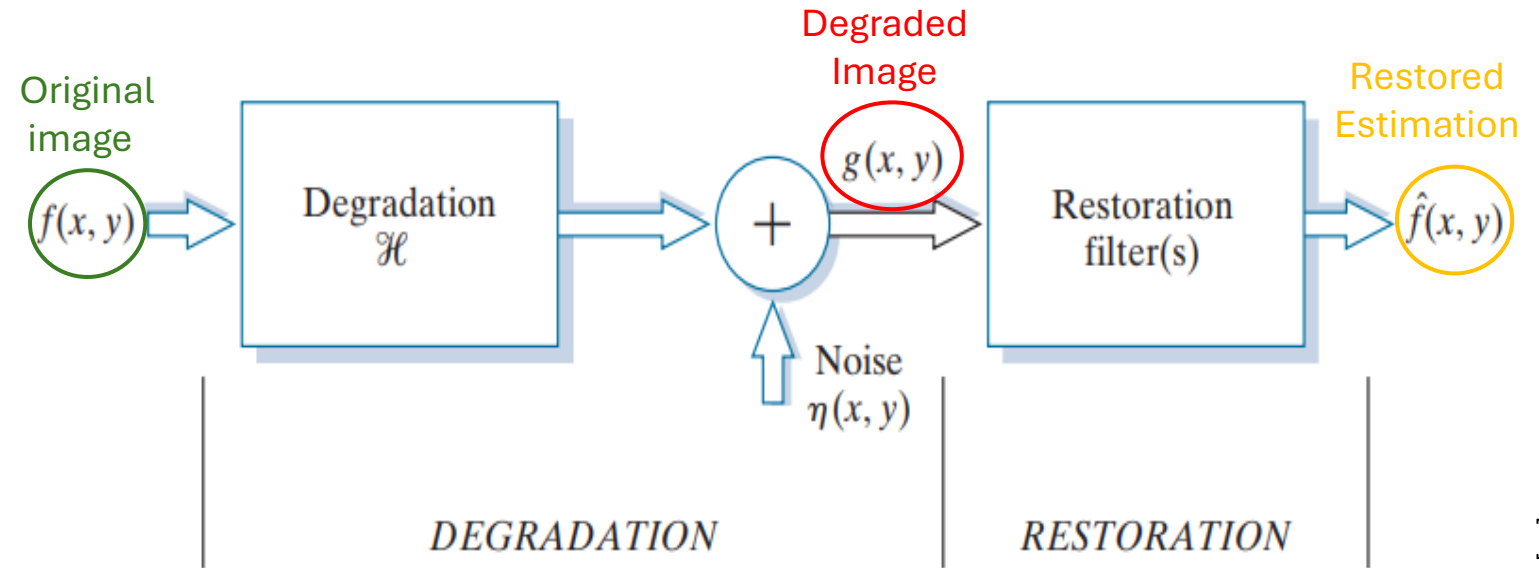
Low-dose



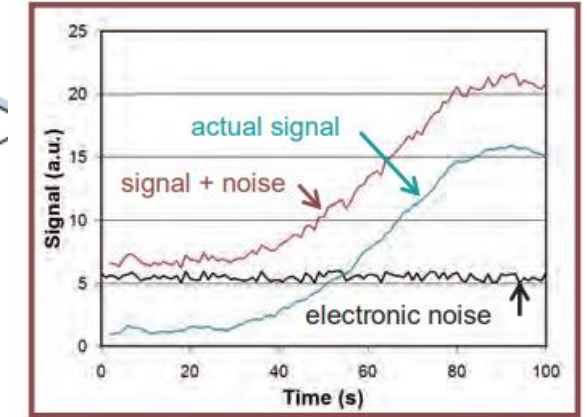
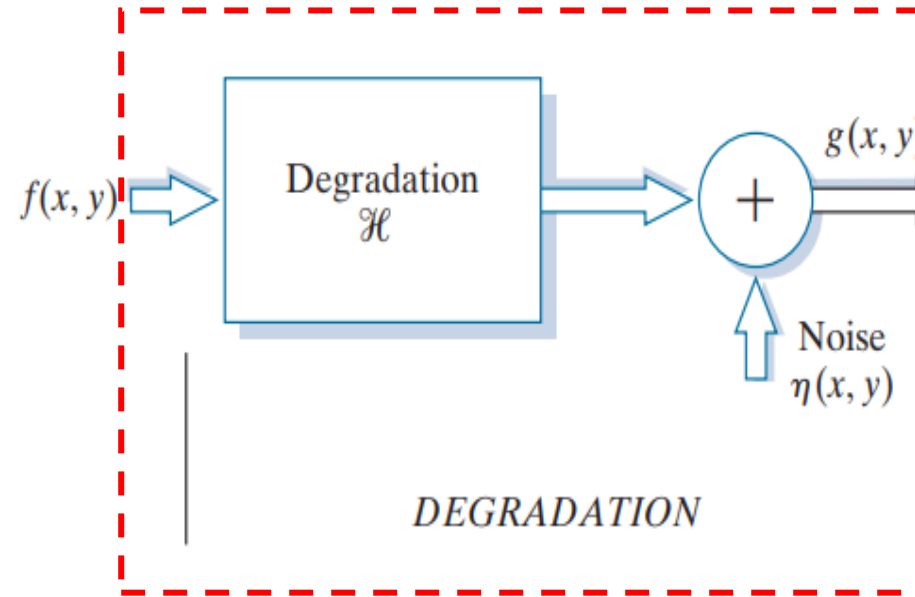
Normal-dose

Source: Stewart Bushong**

- The problem can be defined as a Degradation/Restoration problem
- Objective:** Developed a way to restore the image as similar as possible to the original



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- CT images are susceptible to various sources of noise [9]:
 - Grain noise
 - Quantum noise
 - Anatomical noise
 - Structure noise
 - Electronic noise
- Low-dose exposure worsens image quality



Source: Bushberg et al. [9]

Electronic Noise



Source: Bushberg et al. [9]

Structured Noise



Source: Bushberg et al. [9]

Anatomical Noise

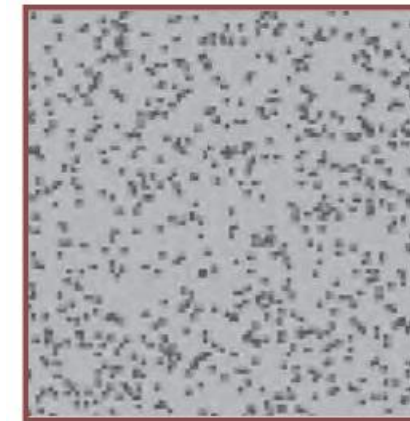


Quantum mottle



Optimal image

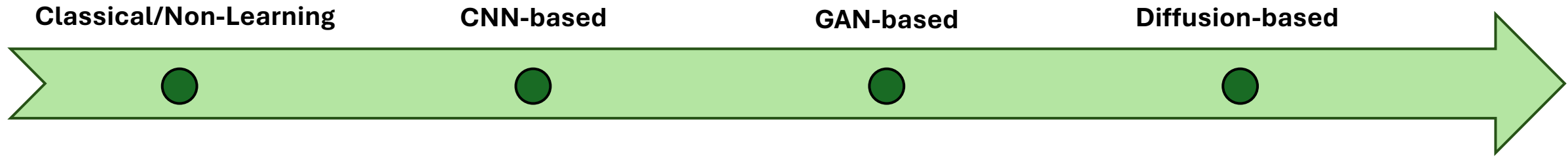
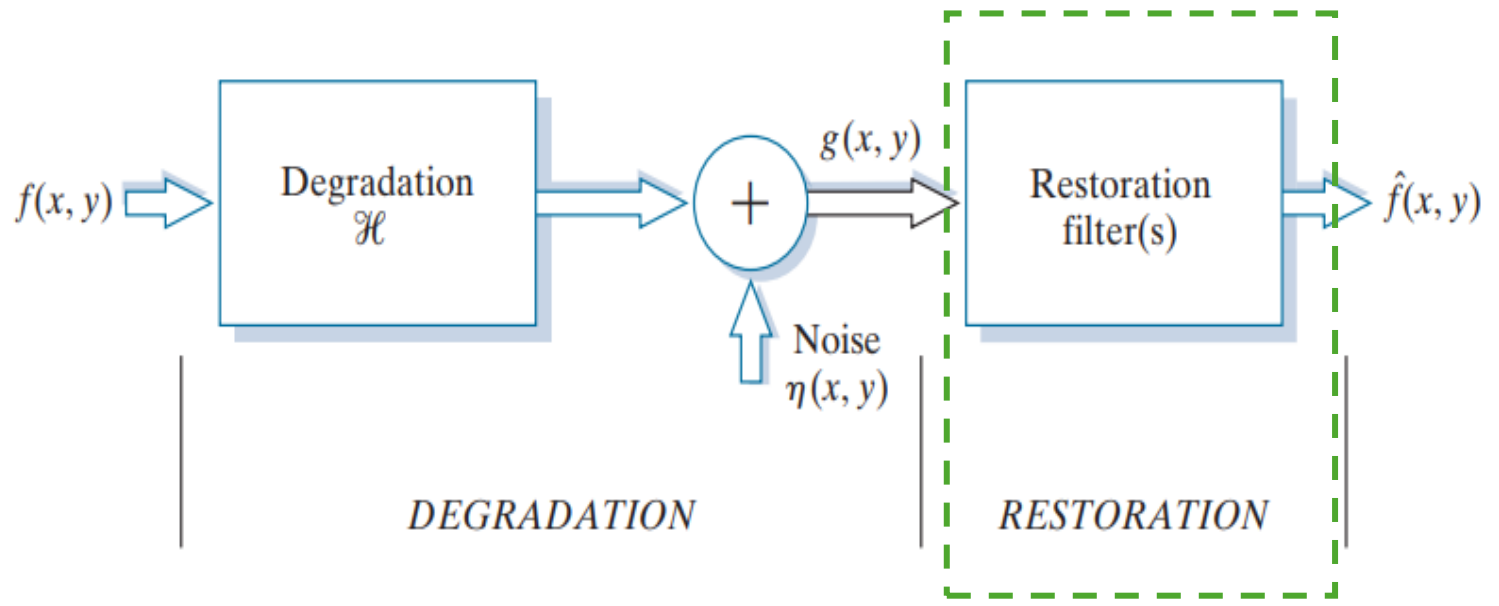
Source: Stewart Bushong*



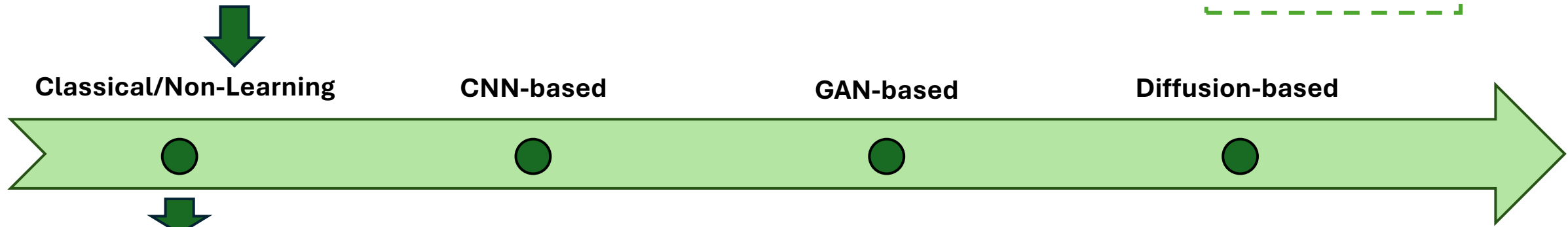
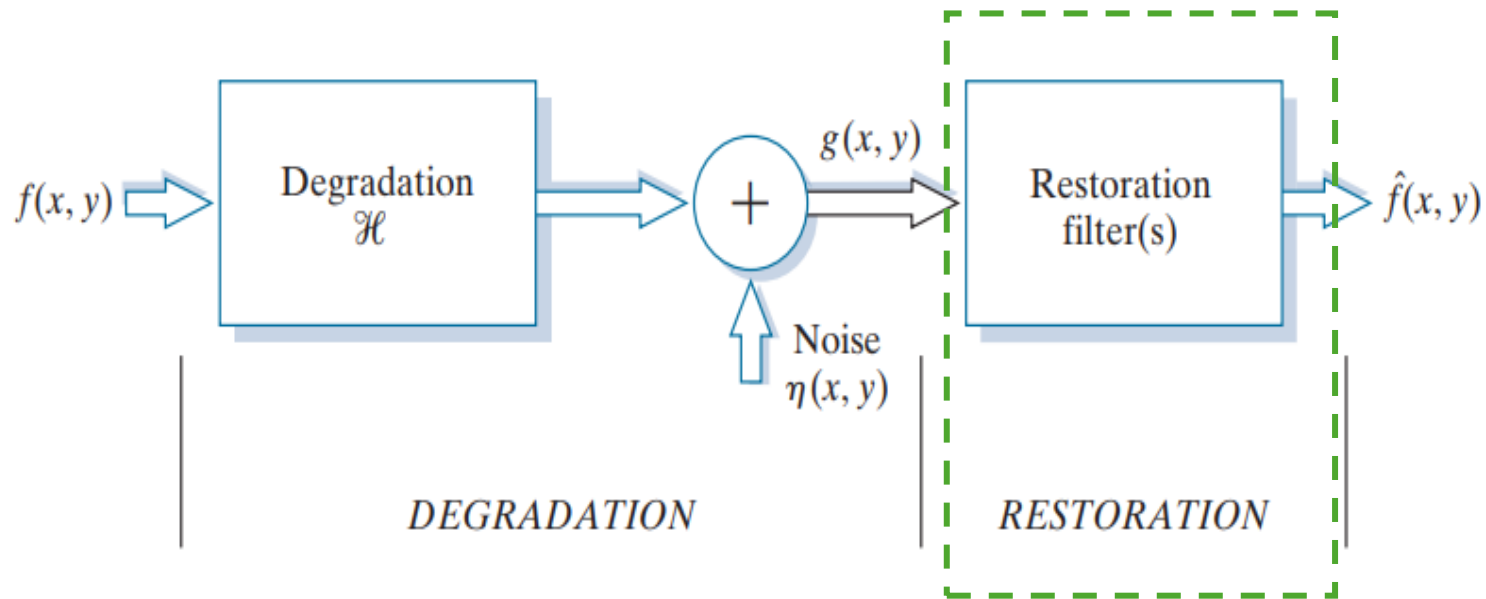
Source: Bushberg et al. [9]

Grain Noise

- The problem can be defined as a Degradation/Restoration problem
- **Objective:** Developed a way to restore the image as similar as possible to the original
- Solutions are typically classified as [36, 42]:
 - Raw data filtering
 - Iterative reconstruction
 - Post-processing algorithms

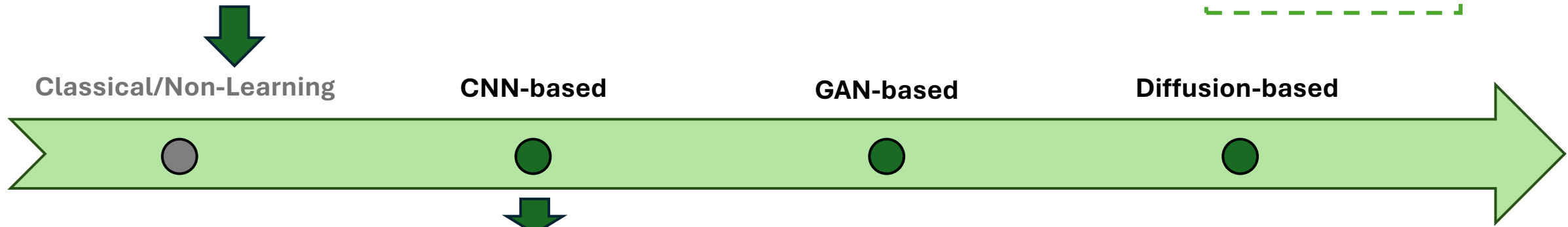
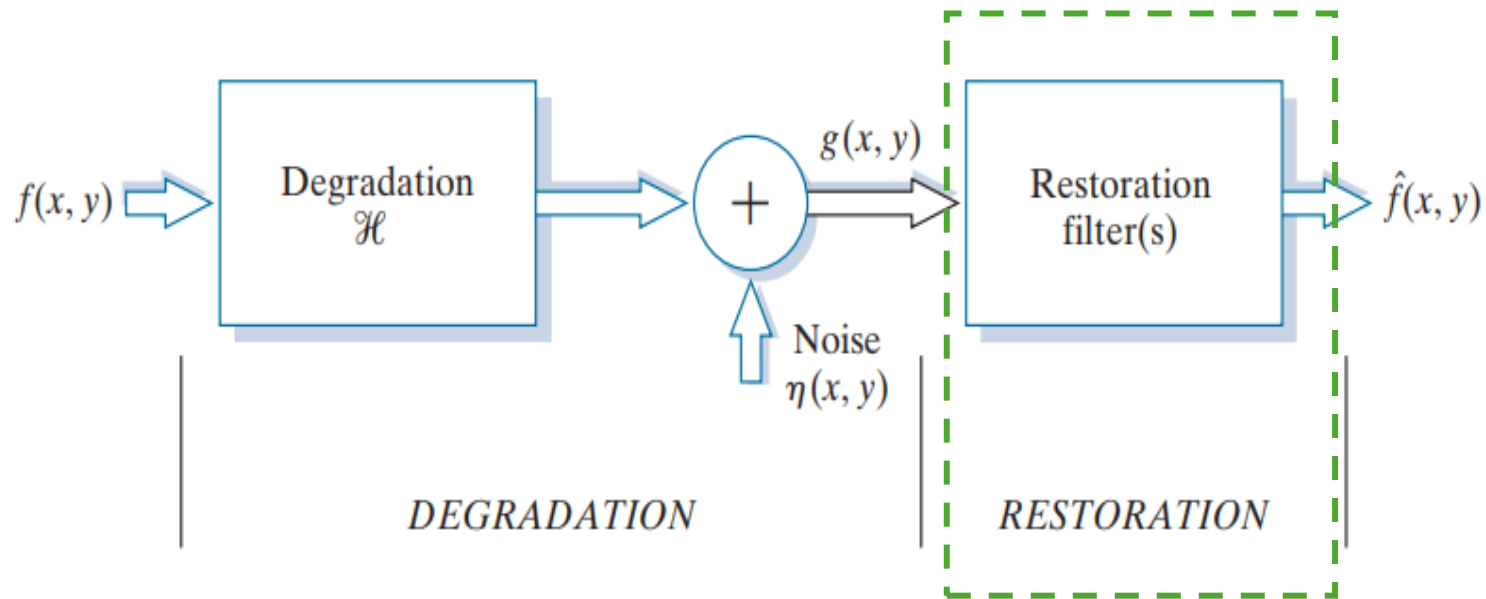


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1. Performance requires improvement [17, 56]

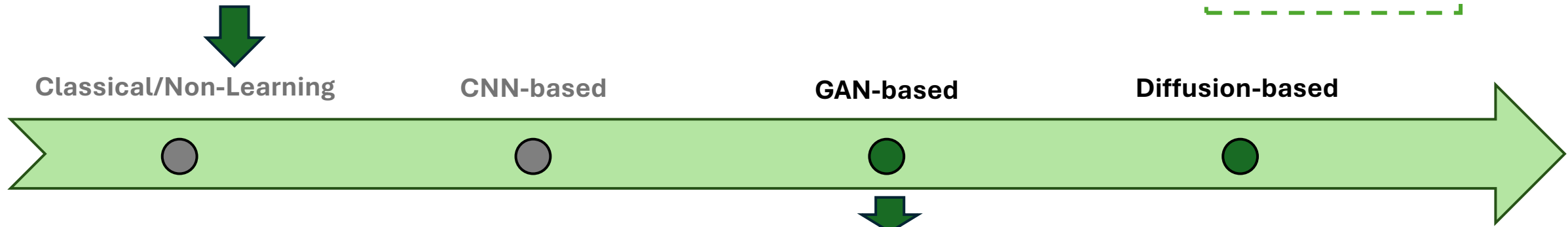
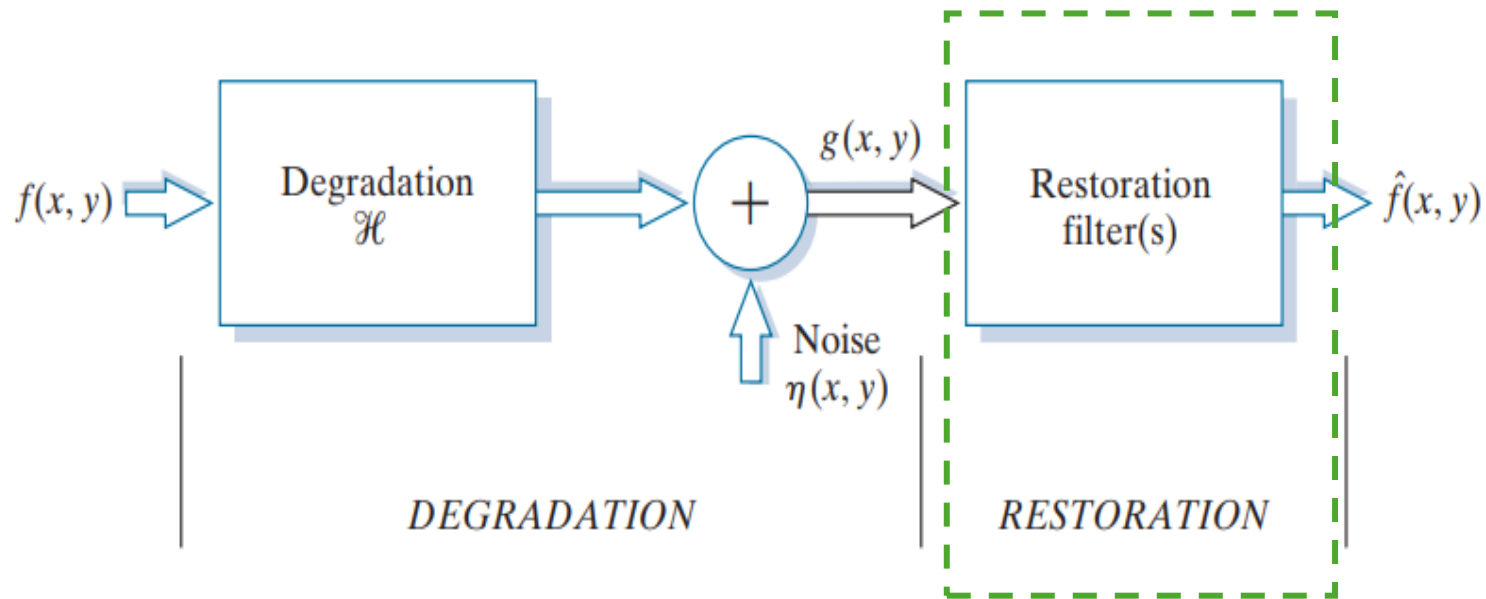
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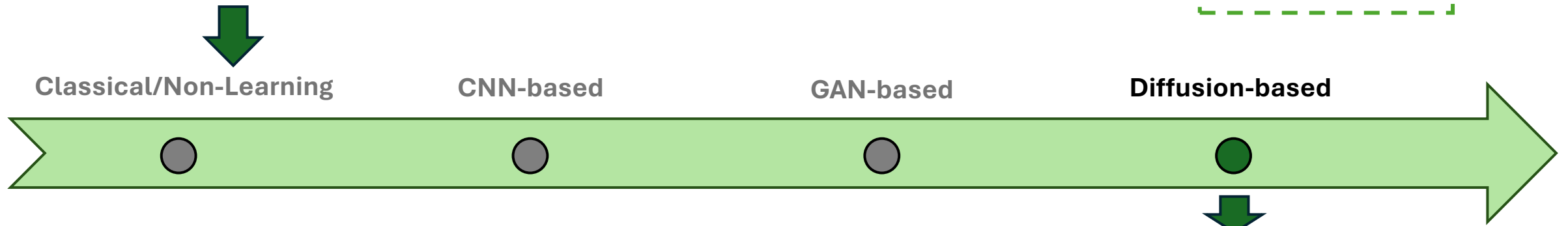
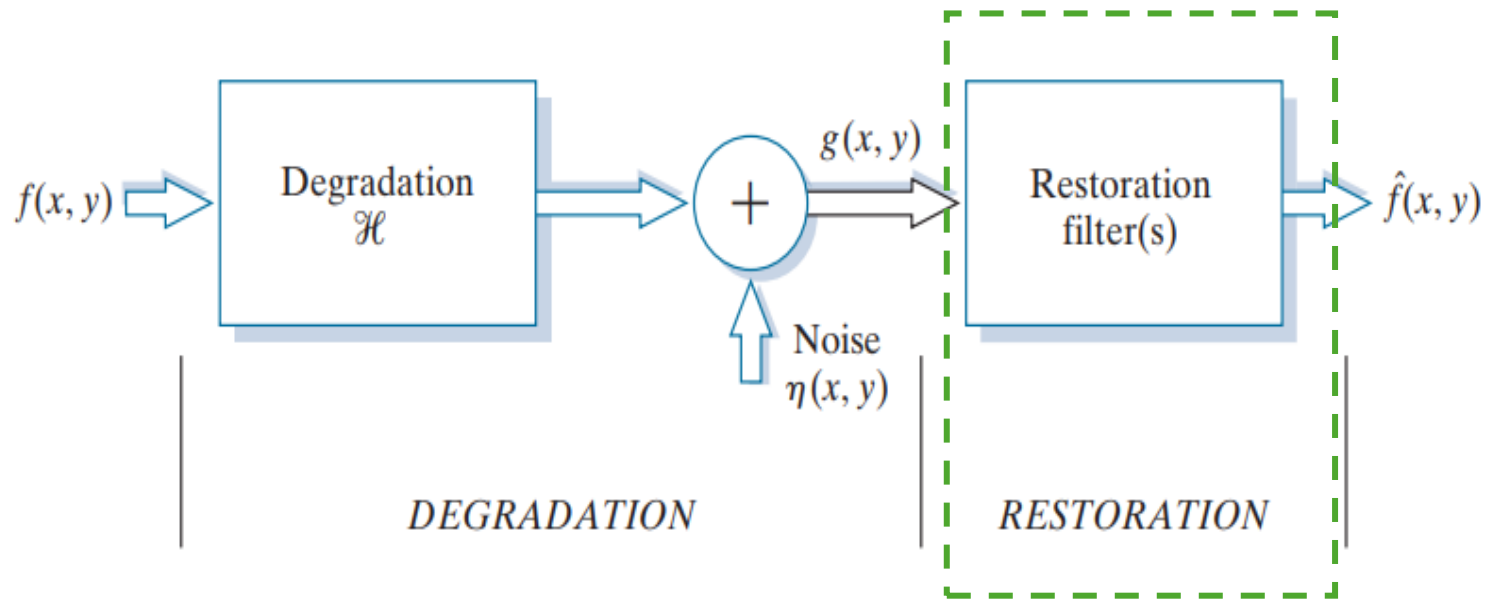


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 3. Unstable train
 4. Value shift [17, 18]

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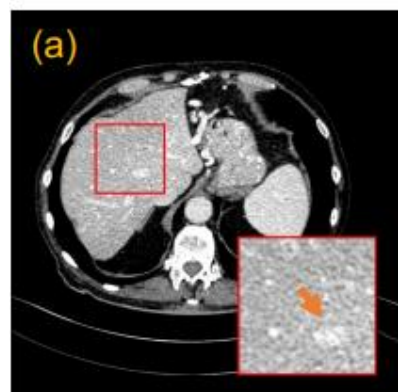


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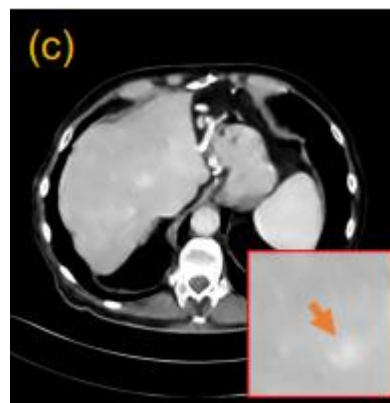
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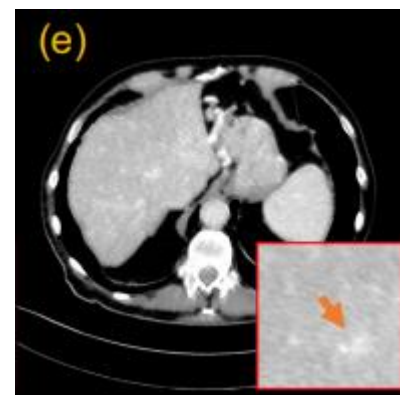
1. Promising results for denoising in other domains when compared to CNN and GANS
 2. Already explored for LDCT [17, 18, 35, 56]
 3. Not extensively explored yet
 4. Never applied to synchrotron LDCT denoising



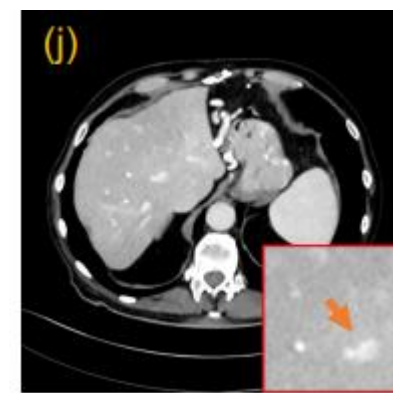
NDCT



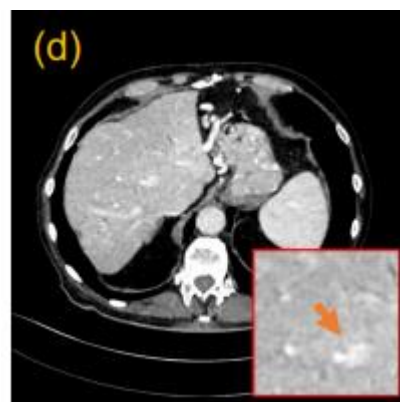
RED-CNN



WGAN-VGG



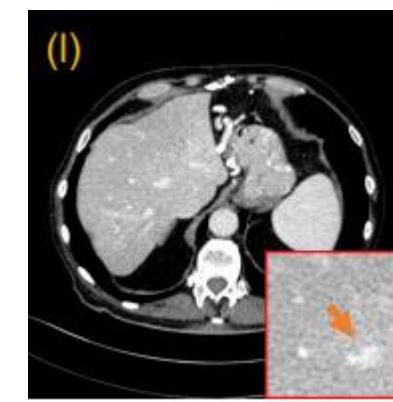
CoreDiff-10



PDF-RED-CNN



DU-GAN



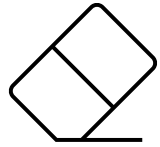
CoreDiff+OSLu-10

↑
CNN-based

↑
GAN-based

↑
Diffusion-based

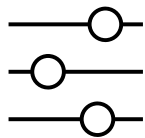




Can diffusion models perform denoising tasks on LDCT reconstructed data toward increasing the quality of synchrotron LDCT?



Can diffusion models trained over CT medical images be directly repurposed to perform denoising tasks on LDCT synchrotron images?



Does finetuning a model trained over CT medical images with synchrotron CT images enhances the acquired results?

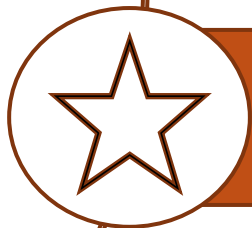




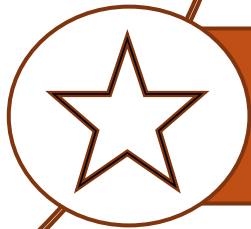
Evaluate diffusion-based generative models for denoising synchrotron LDCT



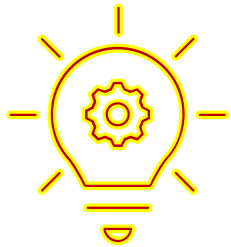
Assess the generalization of diffusion models trained on medical LDCT images for denoising synchrotron LDCT images



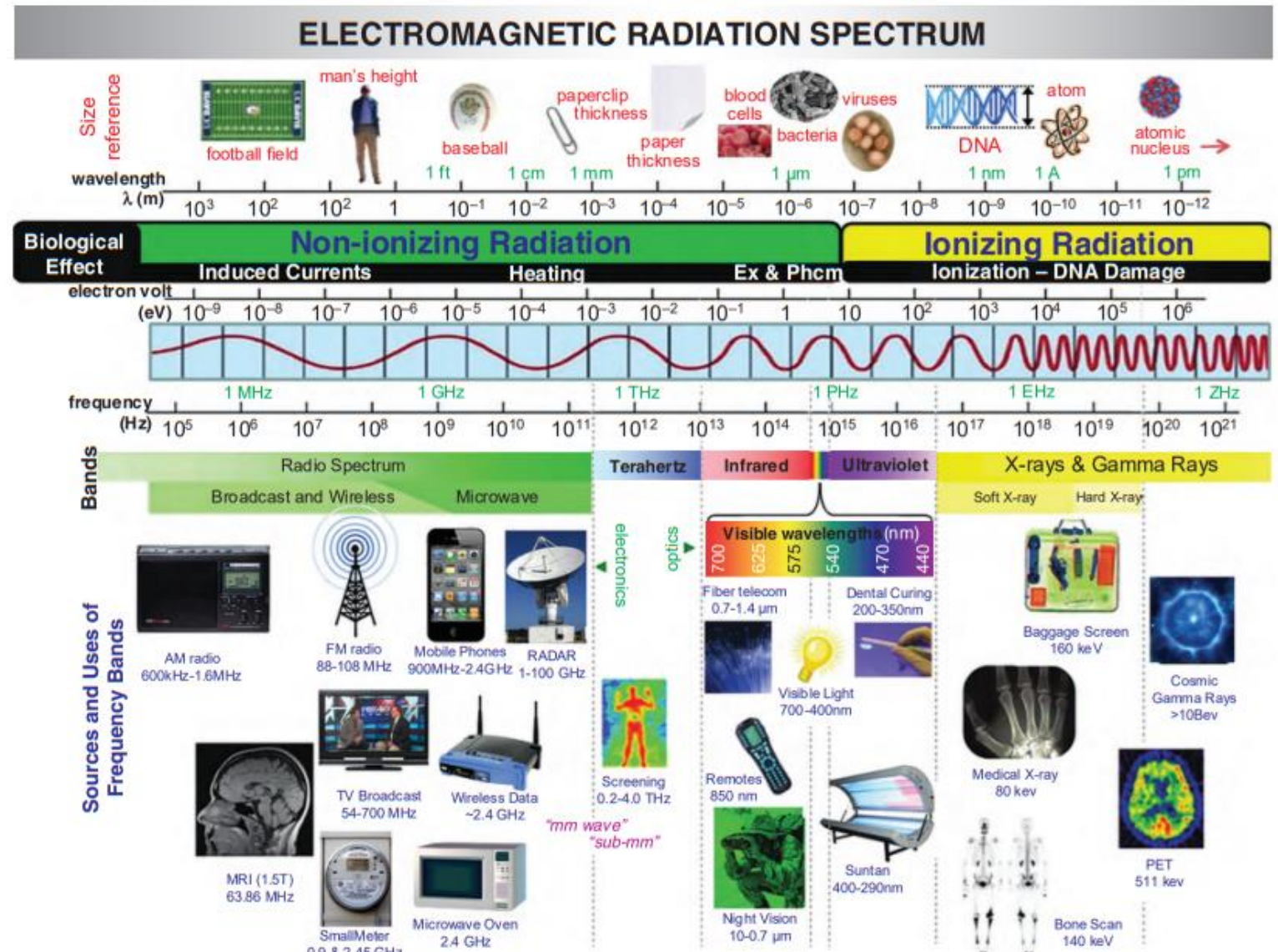
Contribute with a methodology based on diffusion generative models for enhancing synchrotron LDCT image quality



Explore taking advantage of medical LDCT datasets to train models to be used over synchrotron LDCT images



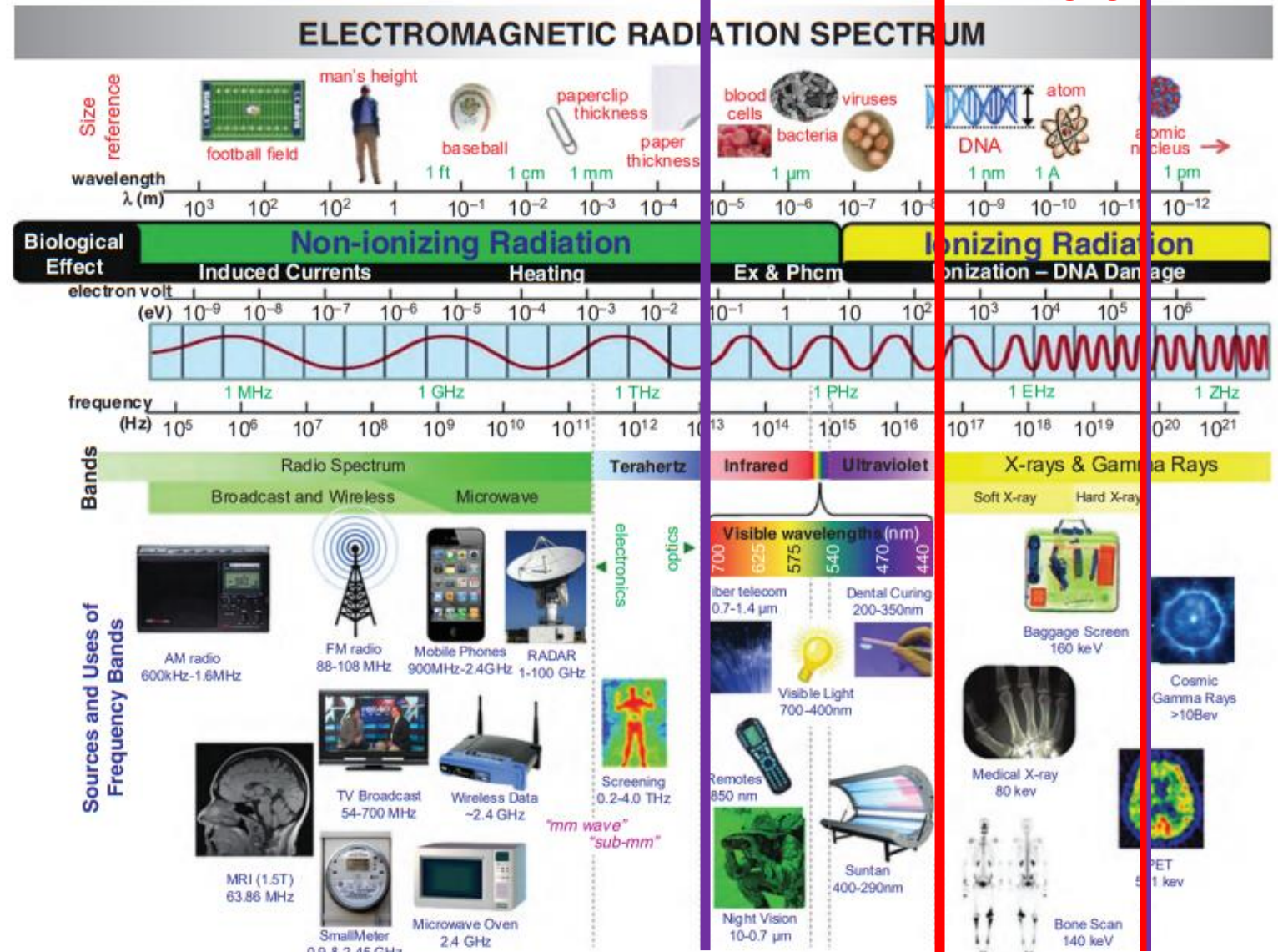
- Radiation is “energy that travels through space or matter” [9]
 - Due to the “wave-particle duality” from quantum mechanics EM can be described as both waves and particles called *photon*
 - Characterized by:
 - Wavelength (λ)
 - Frequency (ν)
 - Energy (E)
- $$E = h\nu = \frac{hc}{\lambda}$$
- Divided in groups according to those characteristics
 - Ionization may occur when *photon* interacts with molecules/atom depending on:
 - Photon energy
 - Target molecule/atom



Source: Bushberg et al. [9]

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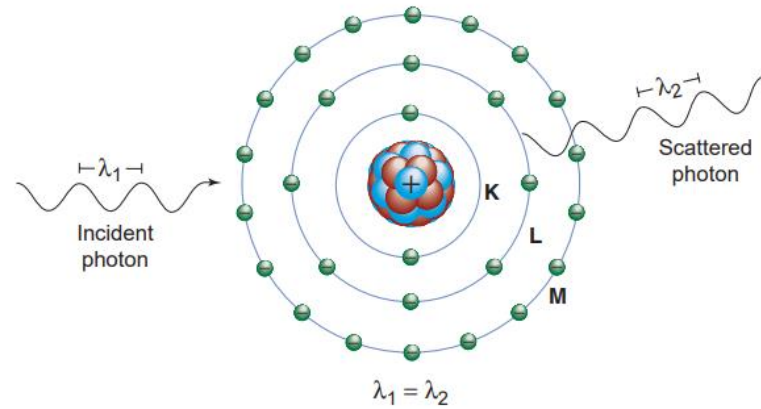
Synchrotron Radiation Spectrum CT Imaging



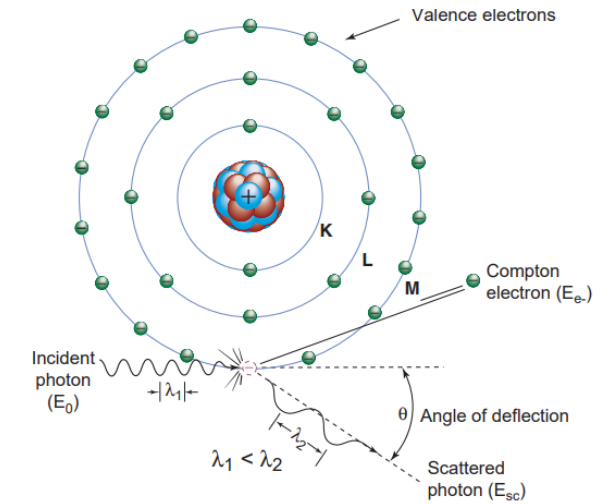
Source: Bushberg et al. [9]

- Types of interaction between EM radiation and matter [9]:

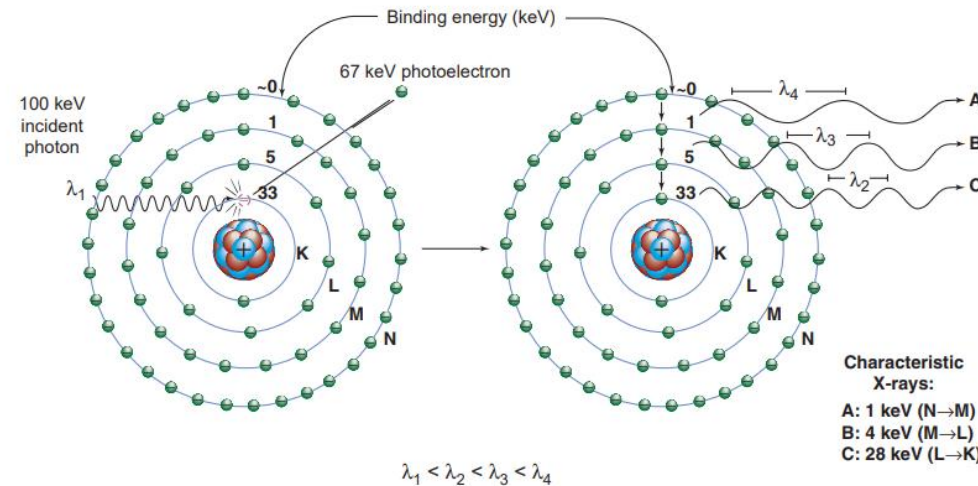
- Rayleigh Scattering
- Compton Scattering
- Photoelectric Absorption
- Pair Production (Only on high-energies)



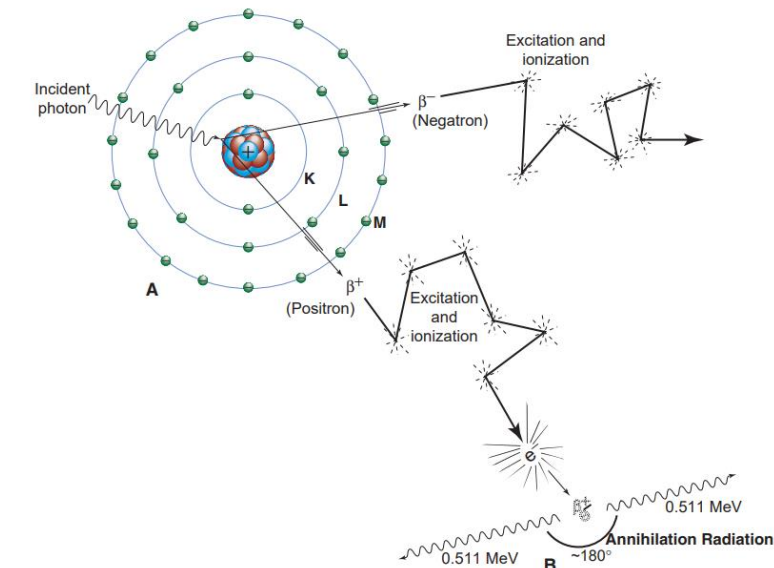
Rayleigh Scattering



Compton Scattering



Photoelectric Absorption



Pair Production

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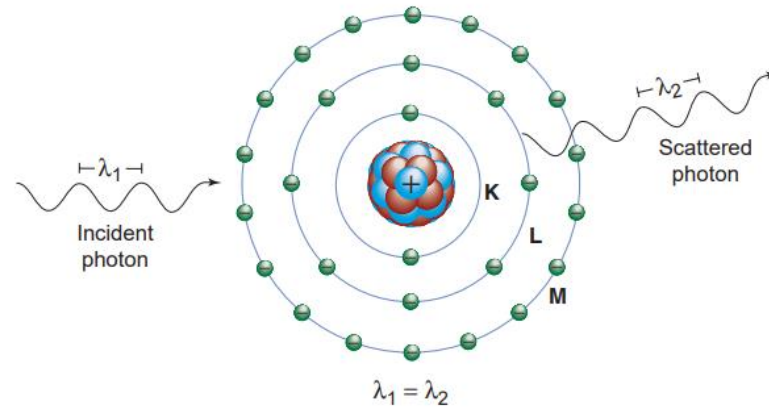
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Combined occurrence

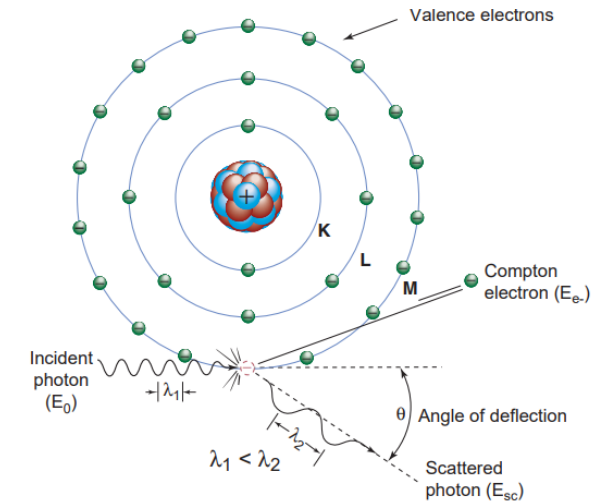
EM Attenuation

Varies with material
composition and thickness

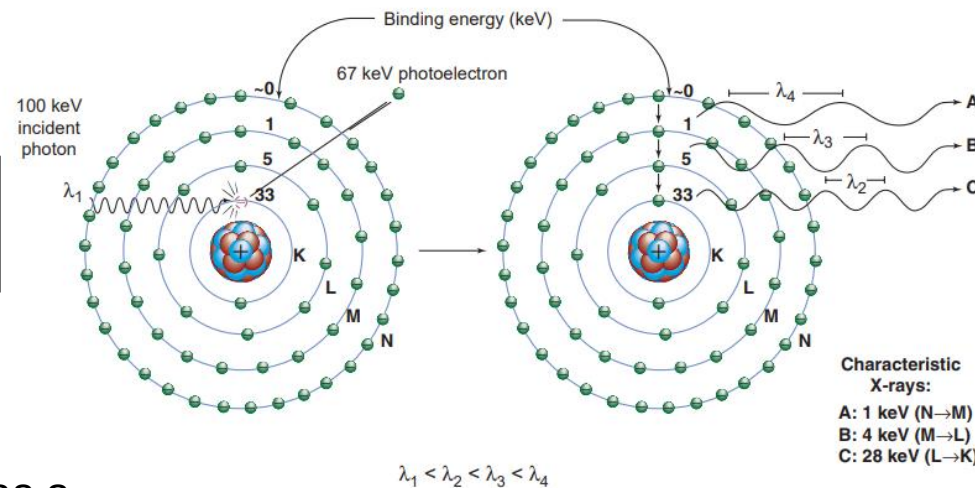
EM Attenuation can be defined as a
Proportion of atoms per volume [9]



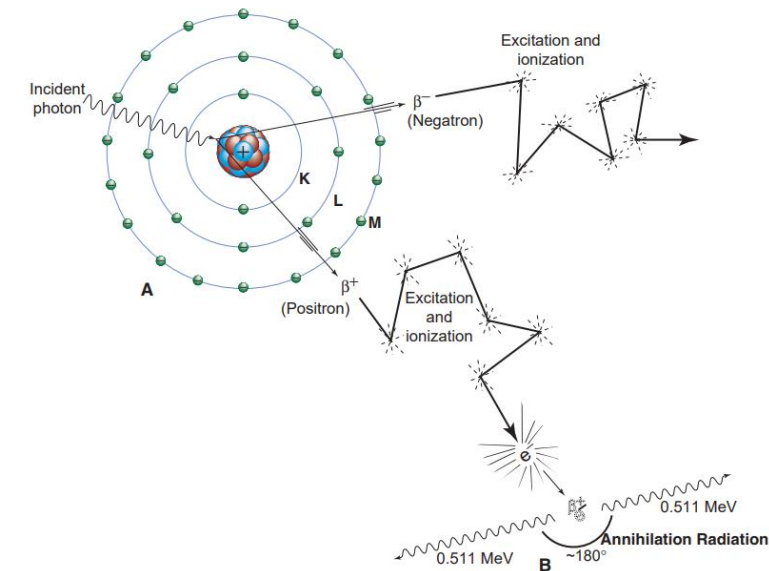
Rayleigh Scattering



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Proportion of atoms per volume [9]

$$\mu = \mu_{\text{Rayleigh scatter}} + \mu_{\text{photoelectric effect}} + \mu_{\text{Compton scatter}} + \mu_{\text{pair production}}$$

Mass Attenuation Coefficient = $\frac{\mu}{\rho}$ → Material density

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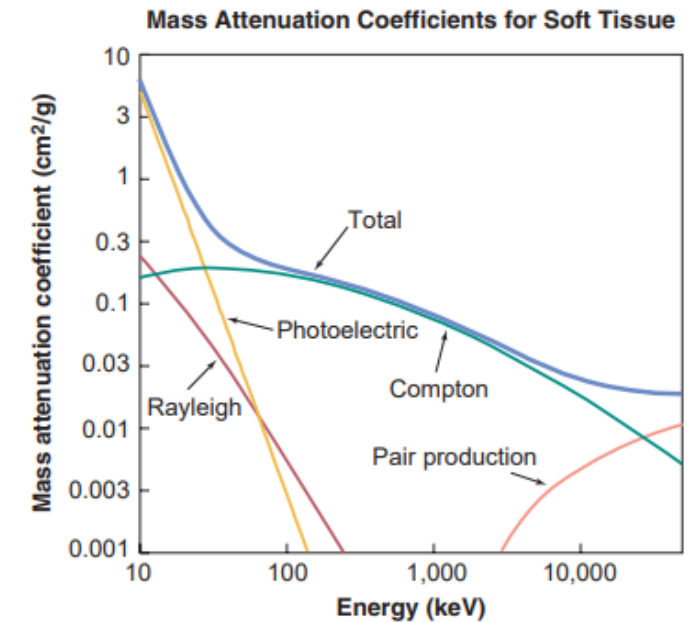
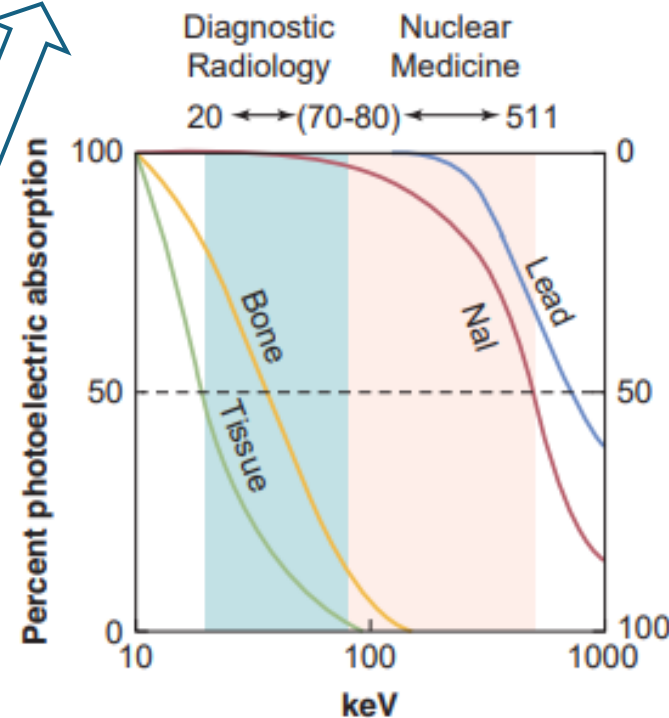
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$$\text{Mass Attenuation Coefficient} = \frac{\mu}{\rho} \quad \text{Material density}$$



Source: Bushberg et al. [9]

EM Attenuation can be defined as a
Proportion of atoms per volume [9]

- Radiation dose is the energy deposited in the irradiated material [9]

- Fluence (ϕ) and flux (ψ)

- Synchrotron facilities have a higher flux than medical X-Ray sources

- Absorbed Dose (D)

- Radiation protection and Medical Imaging application commonly uses:

- Equivalent Dose (H)

- Adds a weighting factor based on the radiation type

- Effective Dose (E_{dose})

- Adds a weighting factor based on each material being irradiated

$$\phi = \frac{\text{Photons}}{\text{Area}}$$

$$\psi = \phi \times E$$

$$D = \frac{E}{m}$$

Irradiated material mass

$$H = D \times w_R$$

Radiation effectiveness on causing biological damage

$$E_{dose} = \sum_T (w_T \times H_T)$$

Material sensitivity

Very simplified way to calculate, as it don't consider:

- Poly-energetic behavior of the radiation source
- Material heterogeneity and morphology
- Radiation-matter interaction variation
- Scattering causing re-interaction
- Surroundings

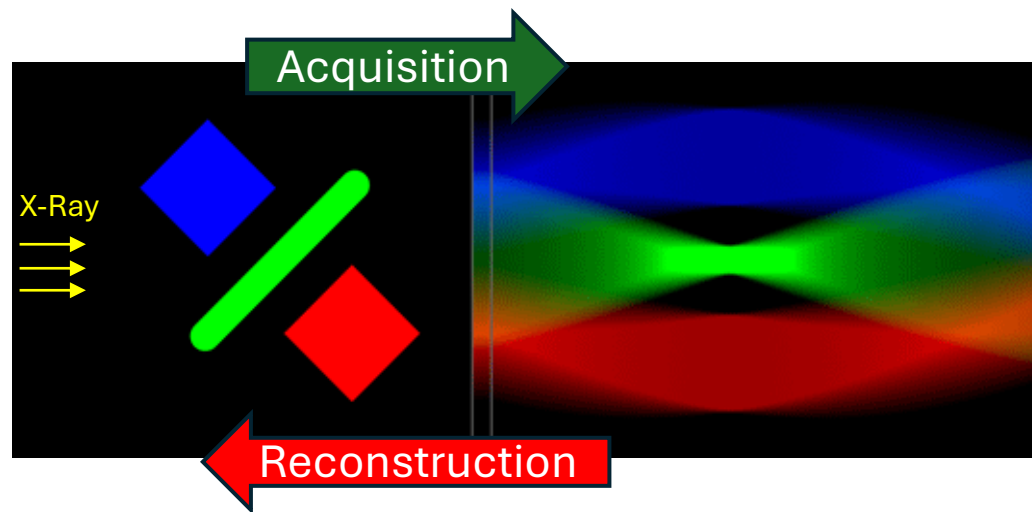
High complexity on precisely calculating dose

Due to that complexity, dose calculation varies depending on application

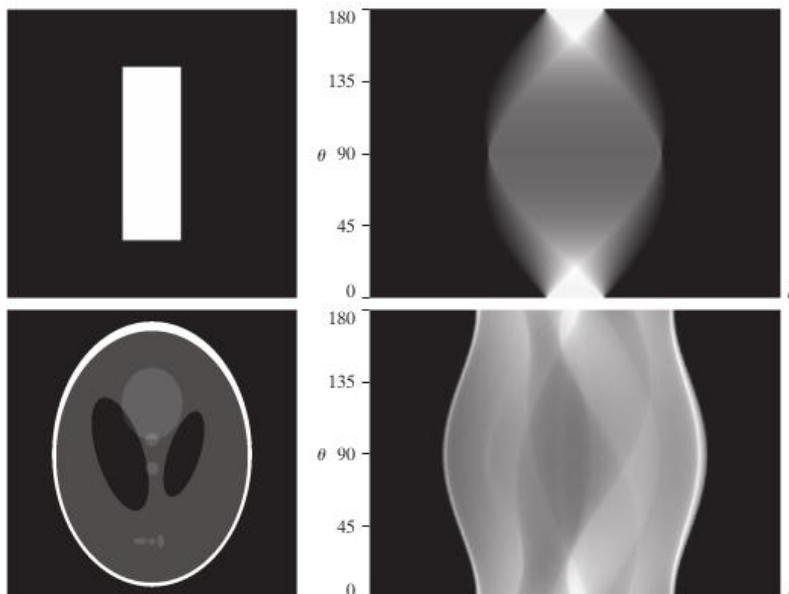
- X-Ray Computer Tomography can be summarized in two stages acquisition and reconstruction [31 , 36]

- Acquisition:** Capture 2D transmission projection images of an object from various angles around a common axis

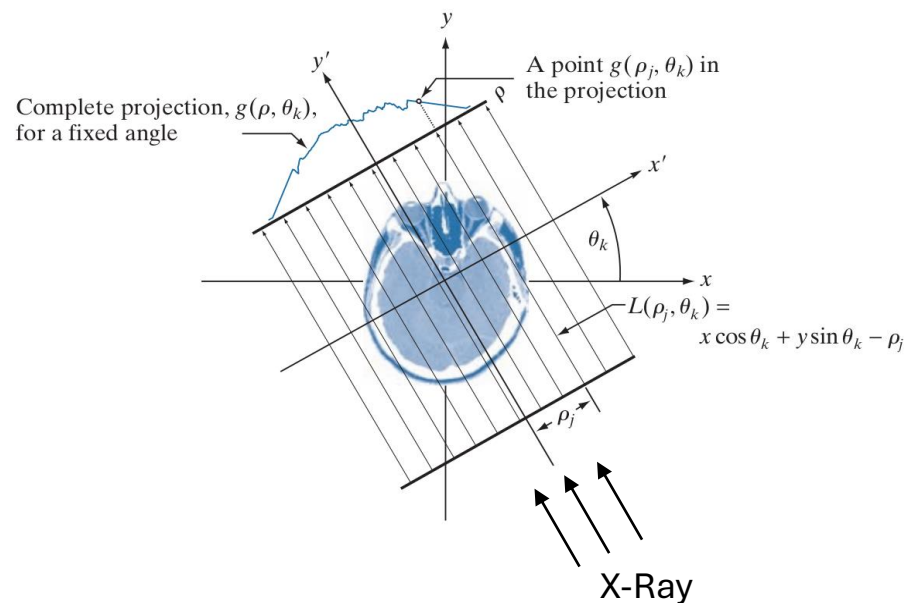
- Reconstruction:** Apply a computational reconstruction method to restore object's 3D morphology



Source: Wikipedia



Source: Gonzales R. C. and Woods R. E. [20]

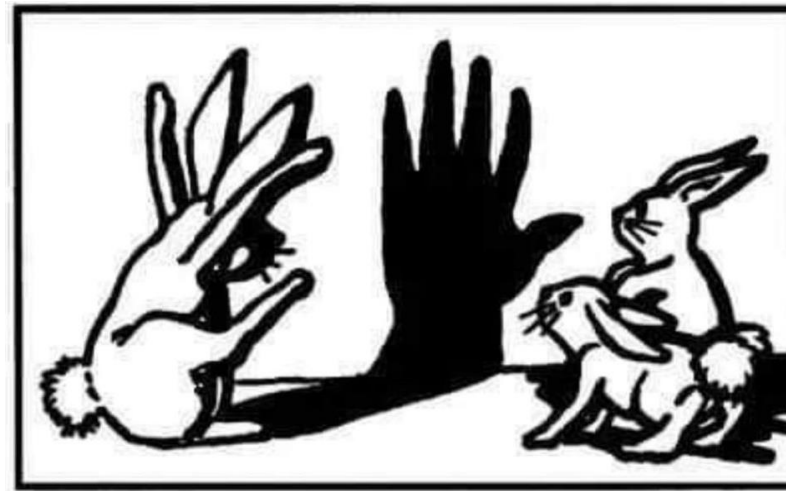


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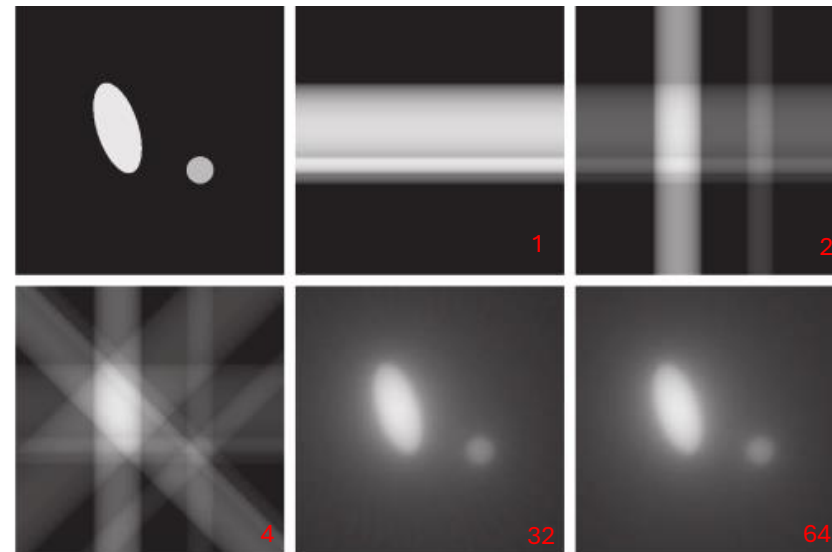
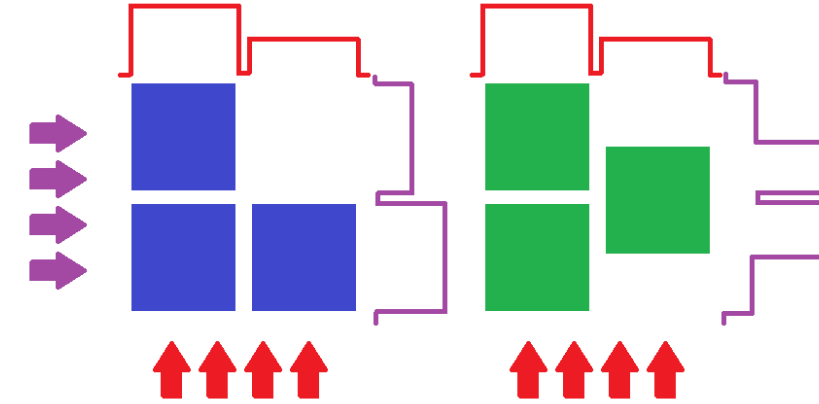


Credits: MOGNO beamline staff

- X-Ray Computer Tomography can be summarized in two stages acquisition and reconstruction [31 , 36]
- **Acquisition:** Capture 2D transmission projection images of an object from various angles around a common axis
- **Reconstruction:** Apply a computational reconstruction method to restore object's 3D morphology
 - Inverse problem
 - Different object may cause same projection
- The more acquired projections the better the reconstruction
 - In theory, with ∞ projections, it would be possible to invert the Radon transform exactly



Source: "La Découverte de l'ombre" (Roberto Casati)



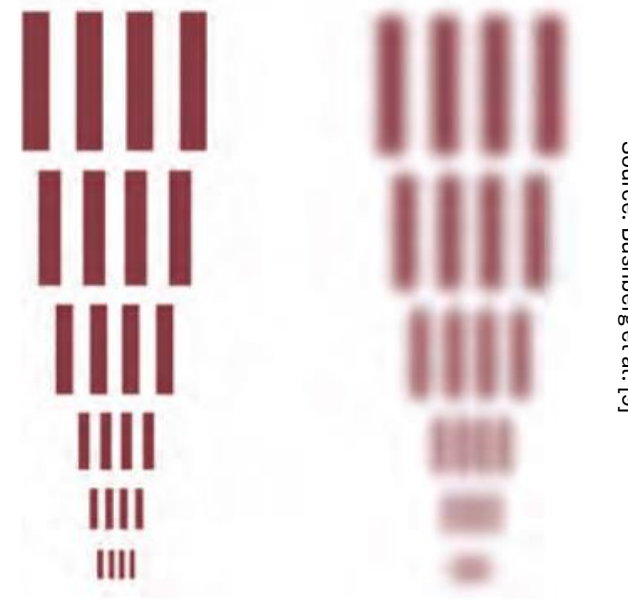
Source: Gonzales R. C. and Woods R. E. [20]

- According to Bushberg et al. [9], CT image quality is strongly bounded to:

- Spatial Resolution
- Contrast Resolution
- Temporal Resolution

- **Spatial Resolution:**

- Ability to distinguish two objects of different densities
- Determines edge sharpness and detail clarity
- Related to how much of real space is represented by a pixel/voxel



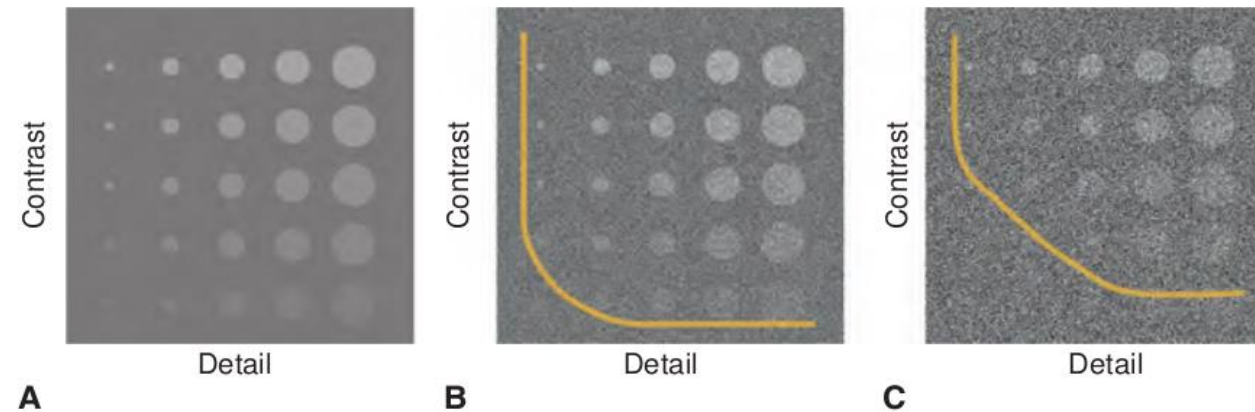
Source: Bushberg et al. [9]

- **Contrast Resolution:**

- Ability to differentiate objects with similar densities using grayscale values
- Emphasizes distinction between similarly shaded objects

- **Temporal Resolution:**

- How long CT image acquisition takes
- Crucial for imaging moving objects



Source: Bushberg et al. [9]

- **Discriminative Models:**

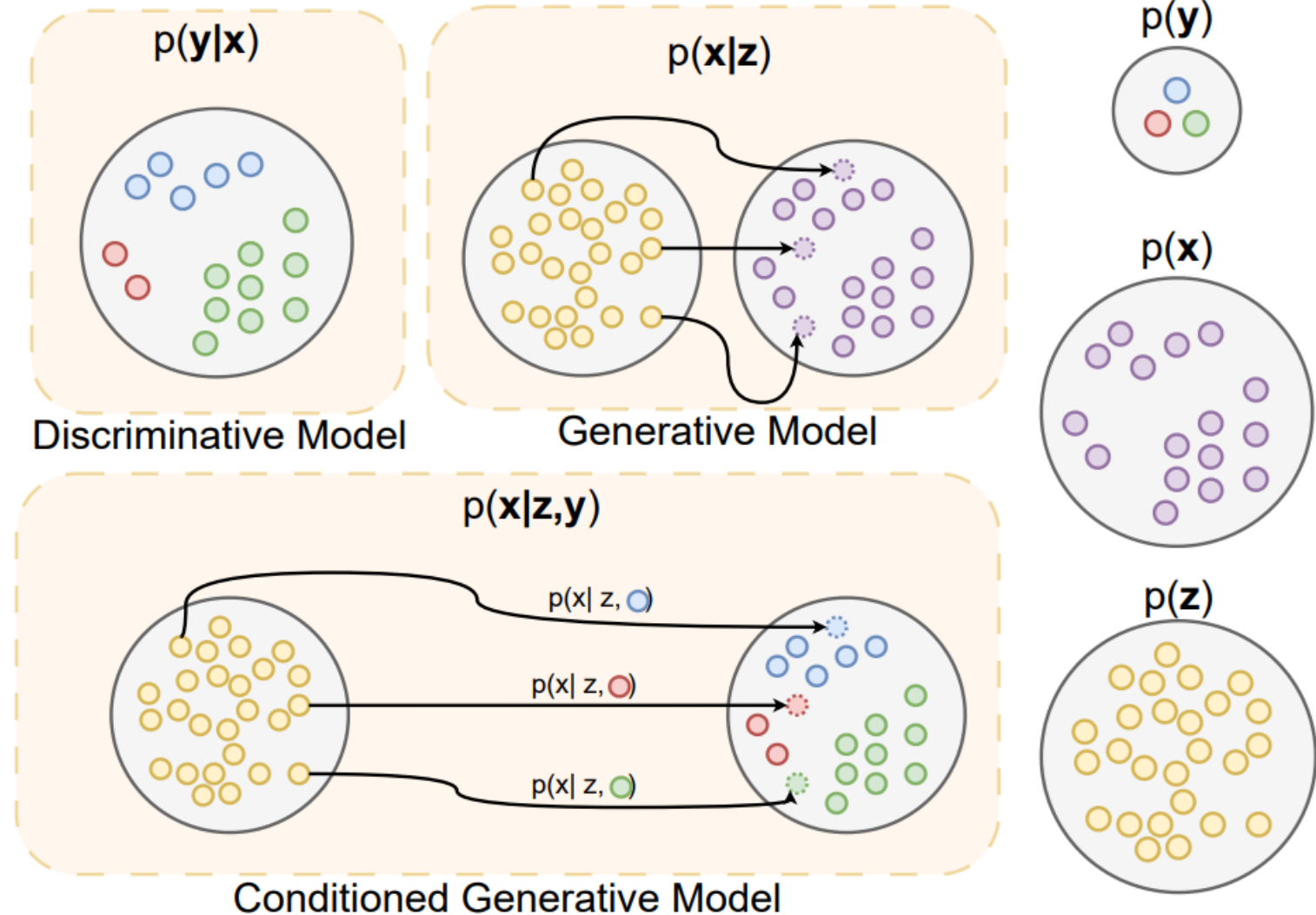
- Learn a function capable of defining boundaries that distinguish which class a sample fits

- **Generative Models :**

- Learn how to transform a latent space variable (z) into a data space variable (x)

- **Conditioned Generative Models :**

- The same as generative models but condition the transformation to guide the transformation



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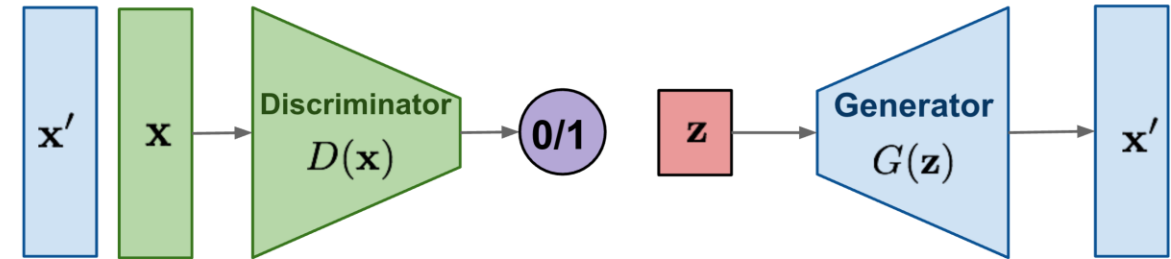
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- **Conditioned Generative Models :**

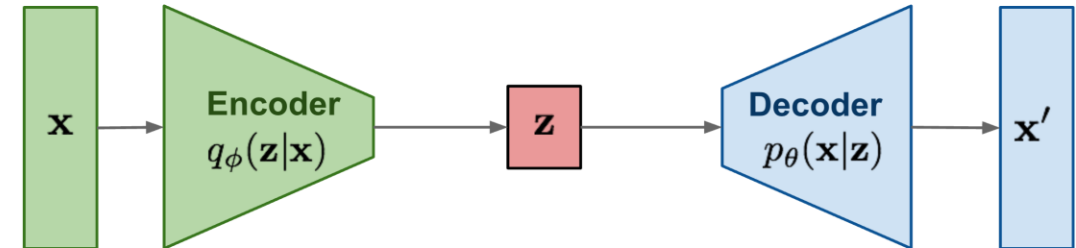
- The same as generative models but condition the transformation to guide the transformation

- Learning to map latent space variables (\mathbf{z}) to data space variables (\mathbf{x}) is the core idea of many generative models.

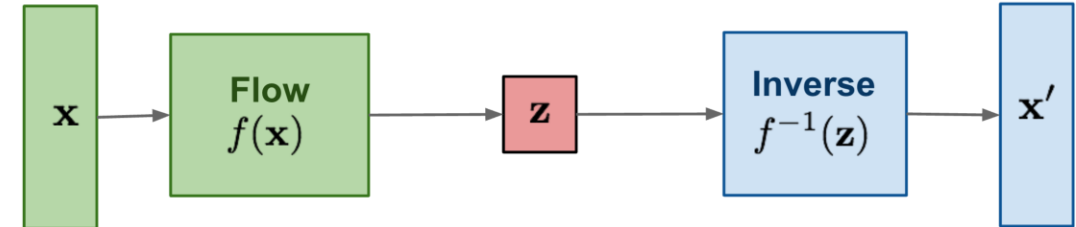
GAN: Adversarial training



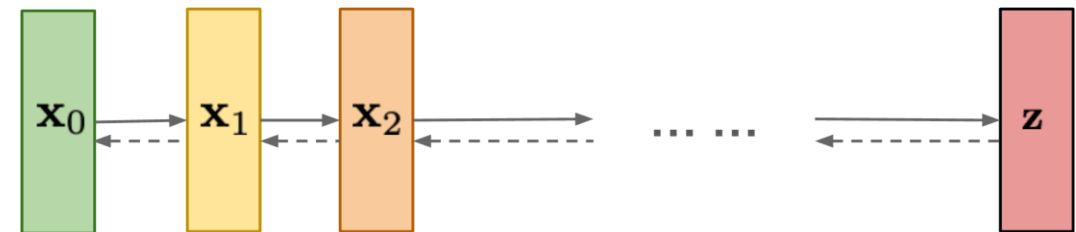
VAE: maximize variational lower bound



Flow-based models:
Invertible transform of distributions



Diffusion models:
Gradually add Gaussian noise and then reverse



- Generative Adversarial Networks (GANs) [21]

- Generator**

- Generate realistic samples by transforming data from latent space $p(\mathbf{z})$ to samples from data space $p(\mathbf{x})$

- Discriminator**

- Distinguish fake and real samples

- Adversarial training:

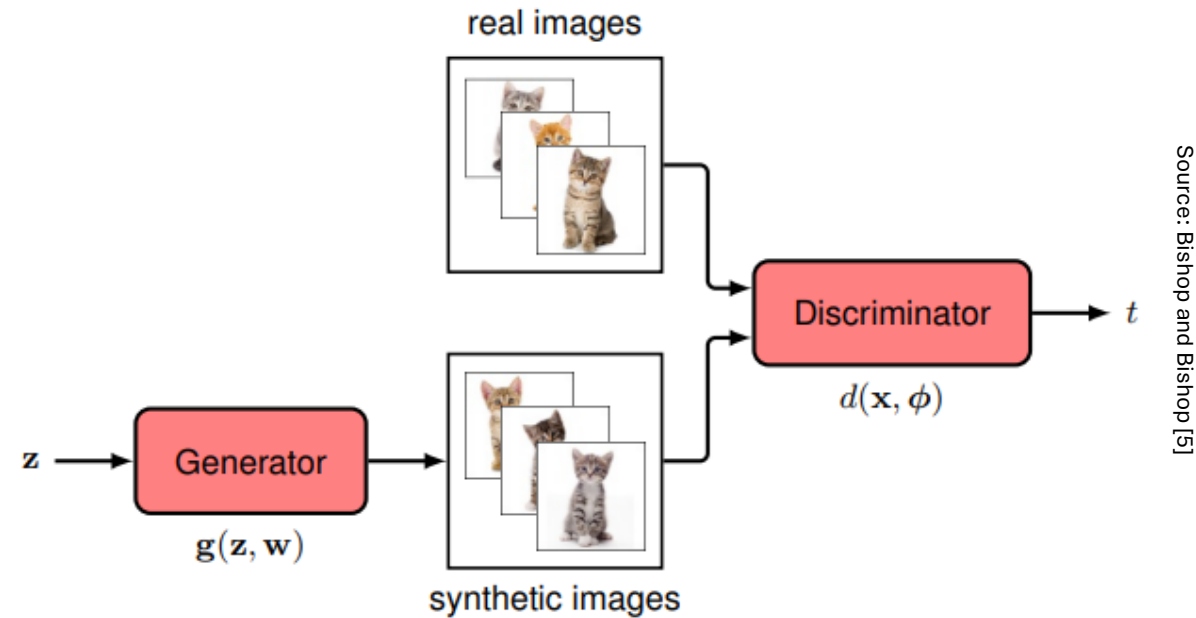
- Shared loss between models
 - Discriminator \rightarrow Minimizes the error
 - Generator \rightarrow Maximizes the error
 - Force both models to improve iteratively

- Training GANs can be challenging

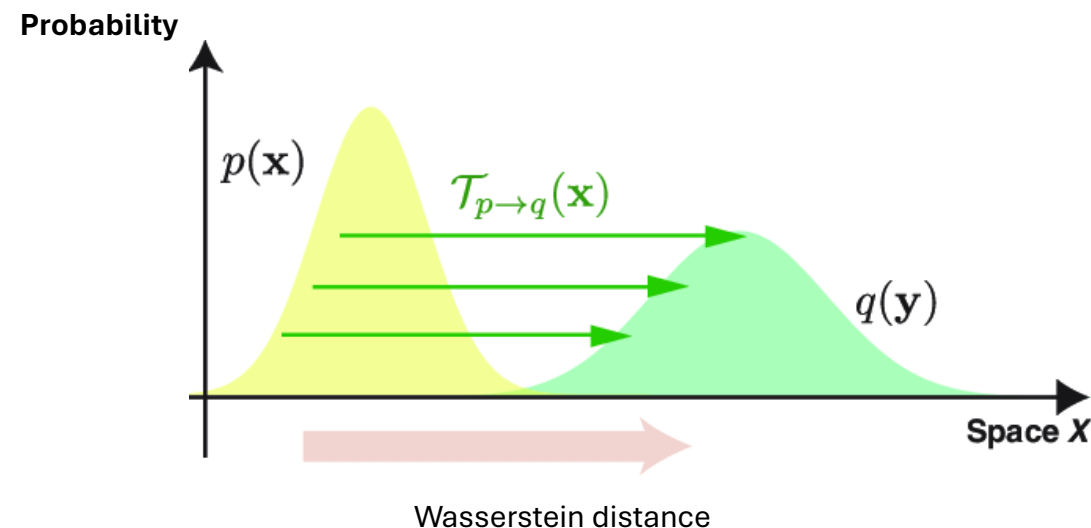
- Model collapse: Generator map $p(\mathbf{z})$ to only a subgroup of $p(\mathbf{x})$
 - No clear progression metric

- Wasserstein Generative Adversarial Network (WGAN) [37]

- More stability
 - Ensure the generator is Moving toward the desired distribution
 - Meaningful loss to show training progression



Source: Bishop and Bishop [5]

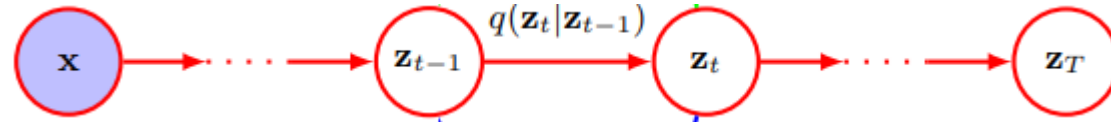
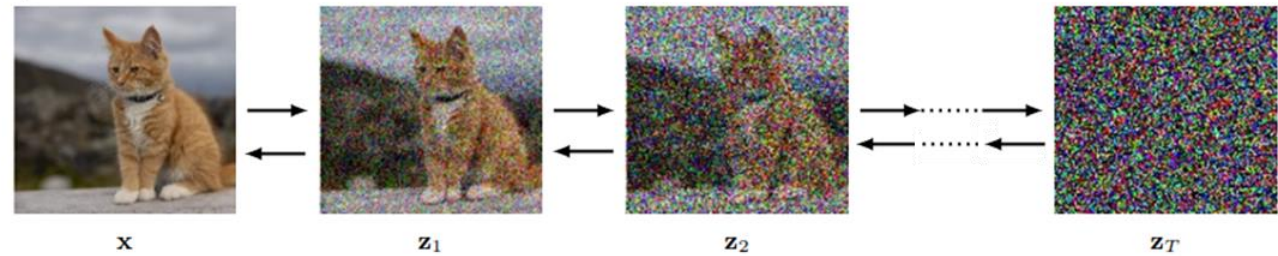


Source: Nakazato and Ito ***

- Diffusion models [46] typically involve two stages:

- Forward encoder**

- Gradually corrupts input data by adding noise to it
 - After several steps input data is transformed on a known noise-like distribution



$$\mathbf{z}_t = \sqrt{1 - \beta_t} \mathbf{z}_{t-1} + \sqrt{\beta_t} \boldsymbol{\epsilon}_t$$

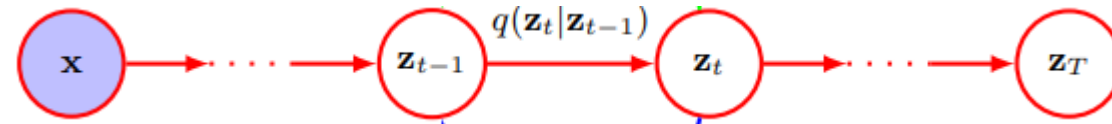
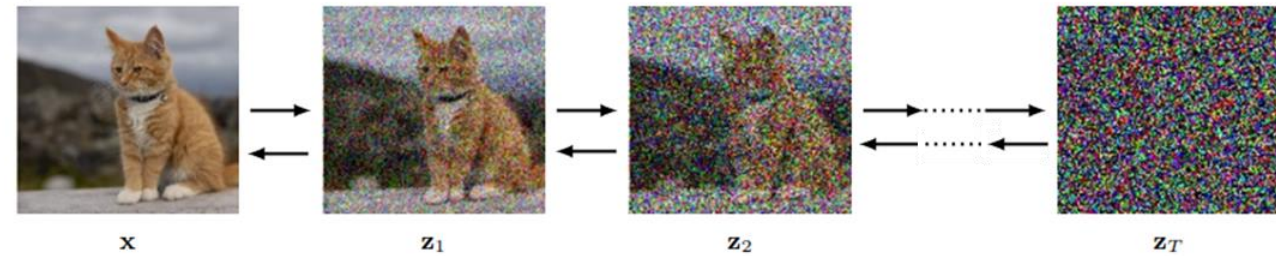
Noise add each step $\boldsymbol{\epsilon}_t \sim N(\boldsymbol{\epsilon}_t | 0, \mathbf{I})$

Noise distribution variance on step t

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Noise distribution variance on step t

$$\beta_1 < \beta_2 < \dots < \beta_T$$

Increase each step

Both coefficients ensures that, for each step, mean and variance of \mathbf{z}_t gets closer 0 and to \mathbf{I}

For $T \rightarrow \infty$, $\beta_T \rightarrow 1$ and \mathbf{z}_T would be a variable within the prior defined noise distribution $N(0, \mathbf{I})$

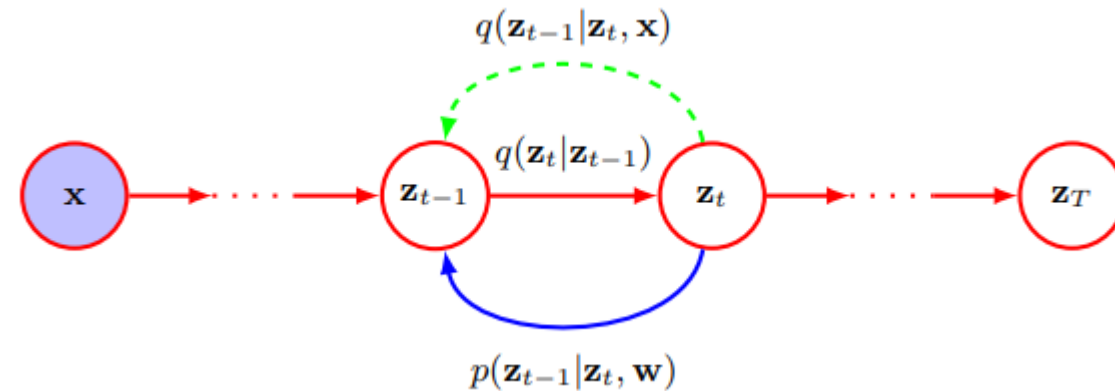
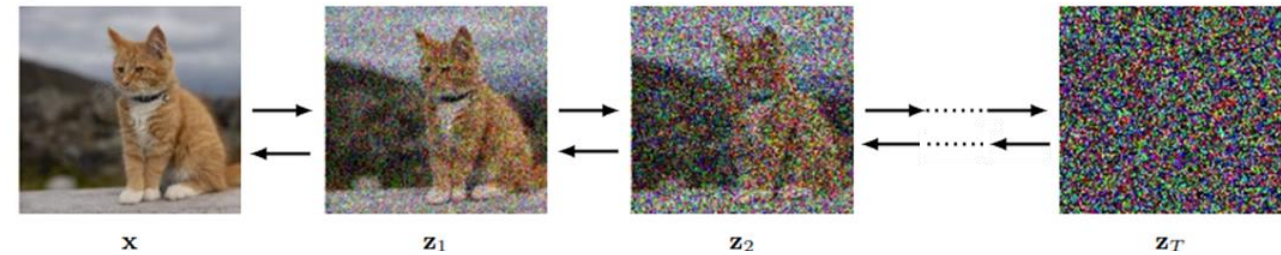
- Diffusion models [46] typically involve two stages:

- Forward encoder**

- Gradually corrupts input data by adding noise to it
- After several steps input data is transformed on a known noise-like distribution

- Reverse decoder**

- Progressively denoise the data to reconstruct
- $p(x)$ is unknown!
- Often parametrized by a Neural Network



$$\mathbf{z}_t = \sqrt{1 - \beta_t} \mathbf{z}_{t-1} + \sqrt{\beta_t} \boldsymbol{\epsilon}_t$$

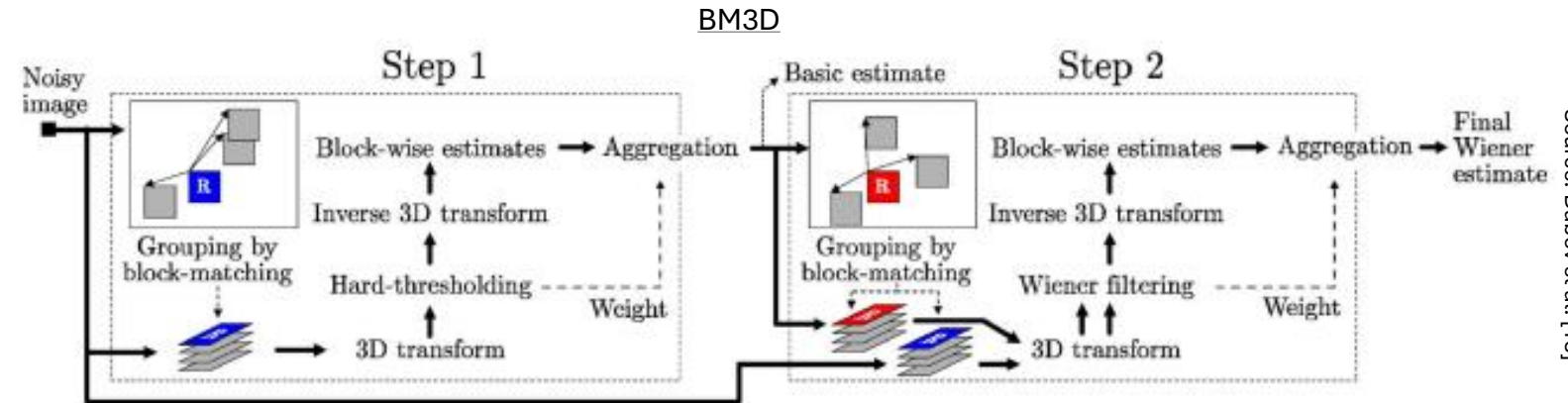
$$\beta_t \ll 1$$

- ↑ Inference time
 - ↓ shift between \hat{x} and x
- ↑ T →
- ↓ Inference time
 - ↑ shift between \hat{x} and x
- ↓ T →

Reverting each step should be easier as the distance between $q(\mathbf{z}_{t-1})$ and $q(\mathbf{z})$ distributions would be narrower

• Non-Learning Classical Methods

- Non-Local Means [8]
- Block-Matching and 3D Filtering [13]



Source: Dabov et al. [13]

Posters / 146

Accelerating Neutron Tomography Ring Artifact Removal Using BM3DORNL

Author: Chen Zhang¹Co-authors: Dmitry Ganyushin¹; Jose Borreguero-Calvo¹; Pete Peterson¹¹ Oak Ridge National Laboratory

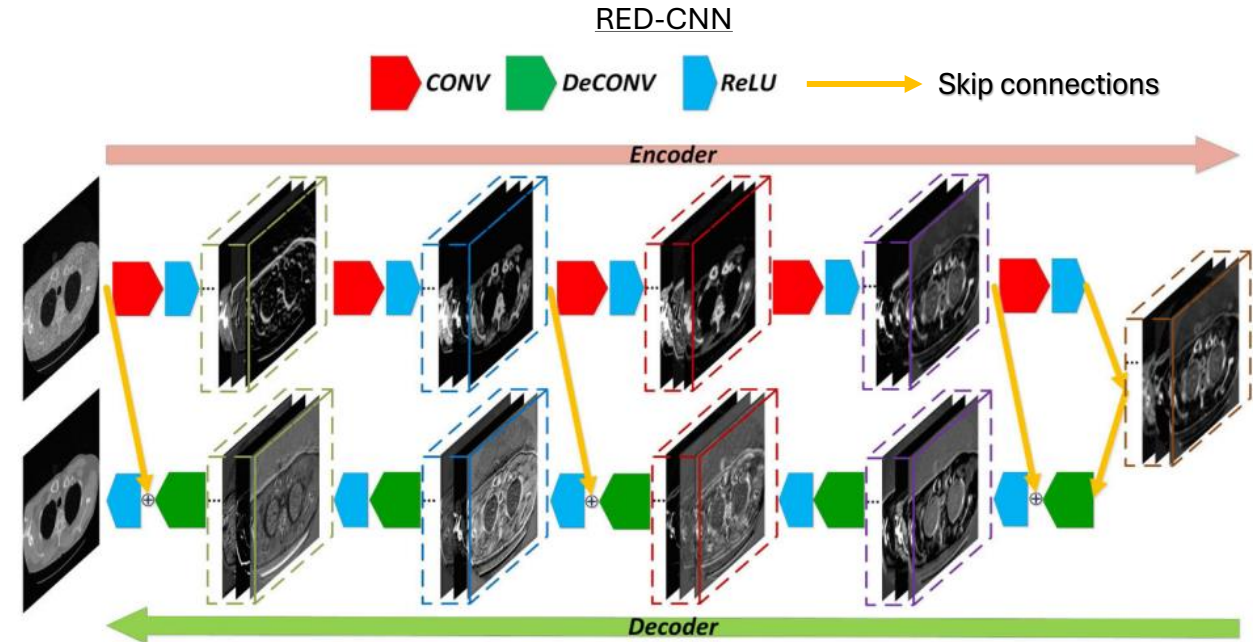
Neutron tomography is a crucial tool for material examination, but ring artifacts can significantly decrease data quality and complicate tasks like segmentation and morphological analysis. The Block-Matching and 3D filtering (BM3D) algorithm, known for mitigating vertical streaks in sinograms and addressing the root cause of ring artifacts, is unfortunately slow and CPU-intensive. We introduce a unique, open-source software solution that eliminates ring artifacts in neutron tomography using the BM3D algorithm. By leveraging both CPU acceleration through Numba and GPU acceleration through CuPy, our approach significantly improves computational efficiency while maintaining data integrity. This dual-acceleration framework drastically speeds up BM3D processing, allowing researchers to quickly obtain refined results and streamline segmentation and morphological analysis.

Abstract publication:

I agree that the abstract will be published on the web site

Source: NOBUGS 2024

- Non-Learning Classical Methods
 - Non-Local Means [8]
 - Block-Matching and 3D Filtering [13]
- **CNN-based Methods**
 - RED-CNN [12]



Source: Chen et al. [12]

- Non-Learning Classical Methods

- Non-Local Means [8]
- Block-Matching and 3D Filtering [13]

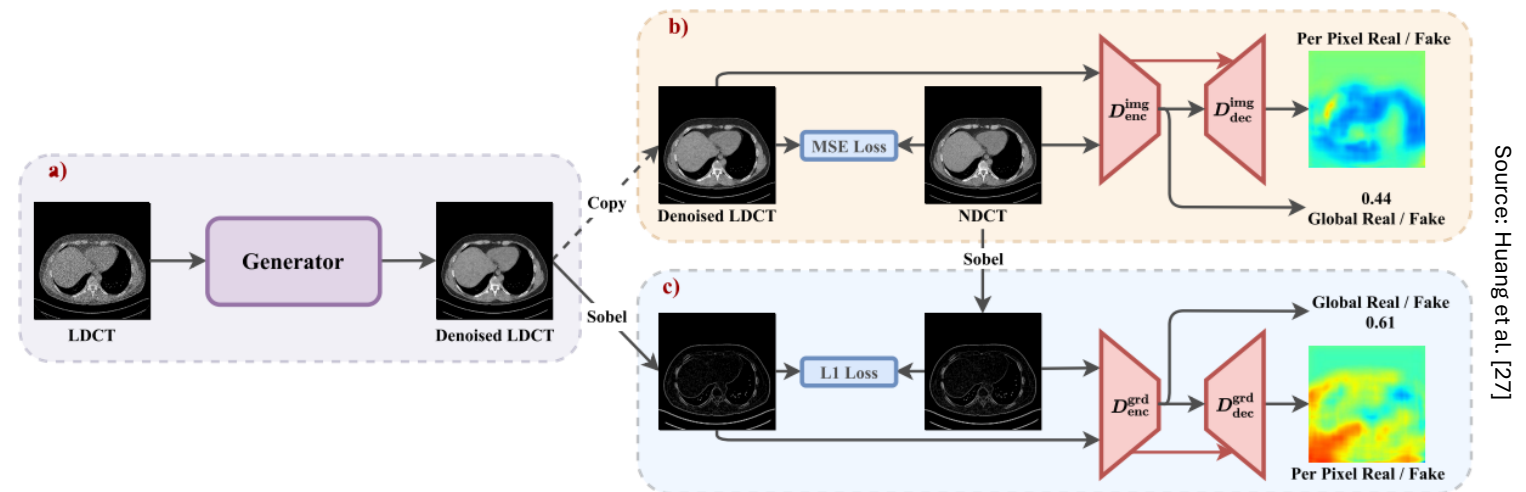
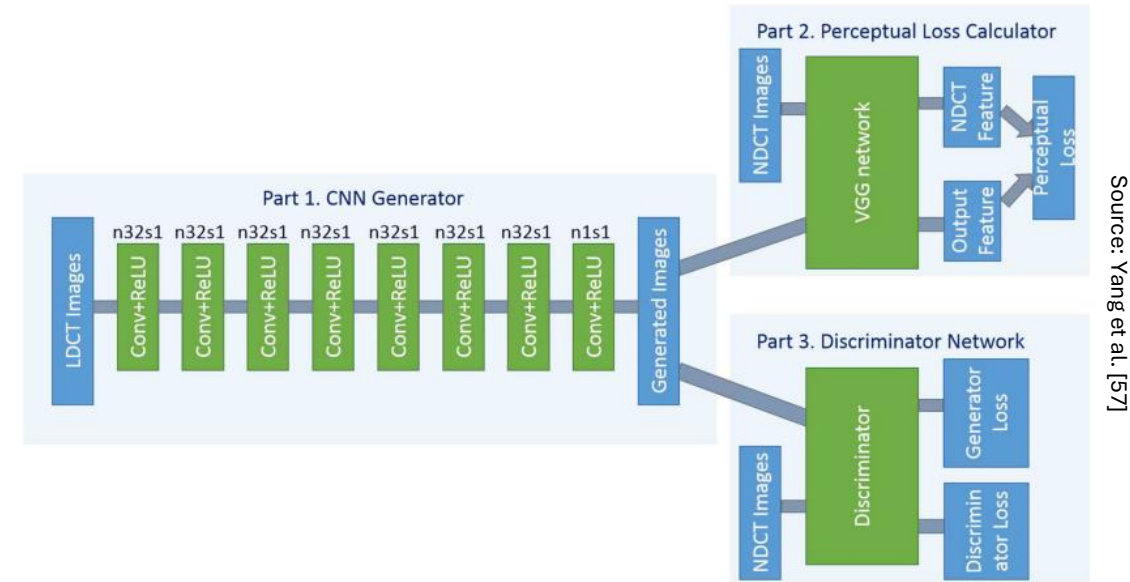
- CNN-based Methods

- RED-CNN [12]

- GAN-based Methods**

- WGAN-VGG [57]
- DU-GAN [27]

WGAN-VGG



- Non-Learning Classical Methods

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- CNN-based Methods

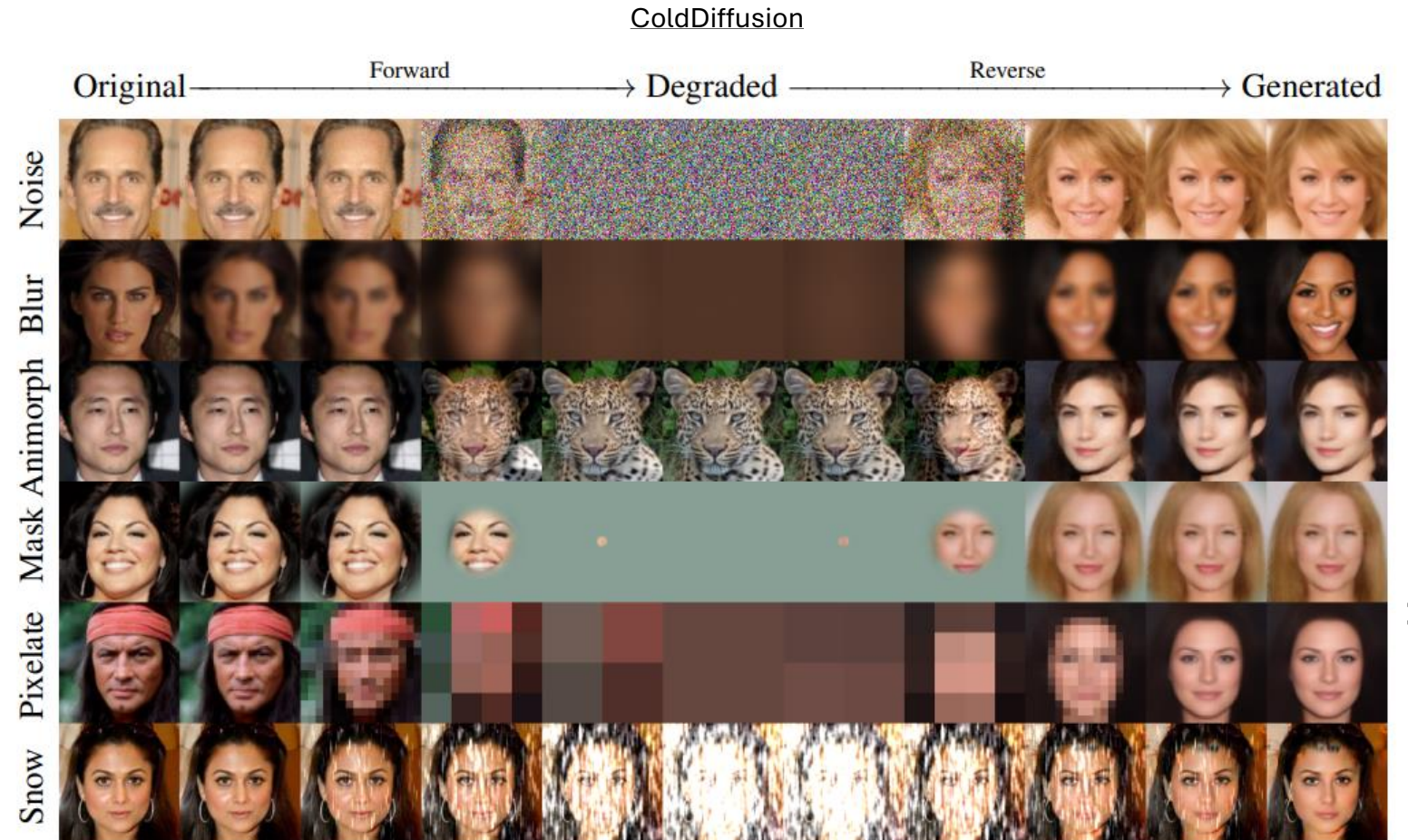
- RED-CNN [12]

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- Diffusion-based Methods**

- Cold Diffusion [2]
- DDPM [56]
- Contextual Conditional Diffusion model (CoCoDiff) [18]
- Denoising with Diffusion Prior (Dn-Dp) [35]
- Contextual Error-modulated Generalized Diffusion Mode (CoreDiff) [17]



- Non-Learning Classical Methods

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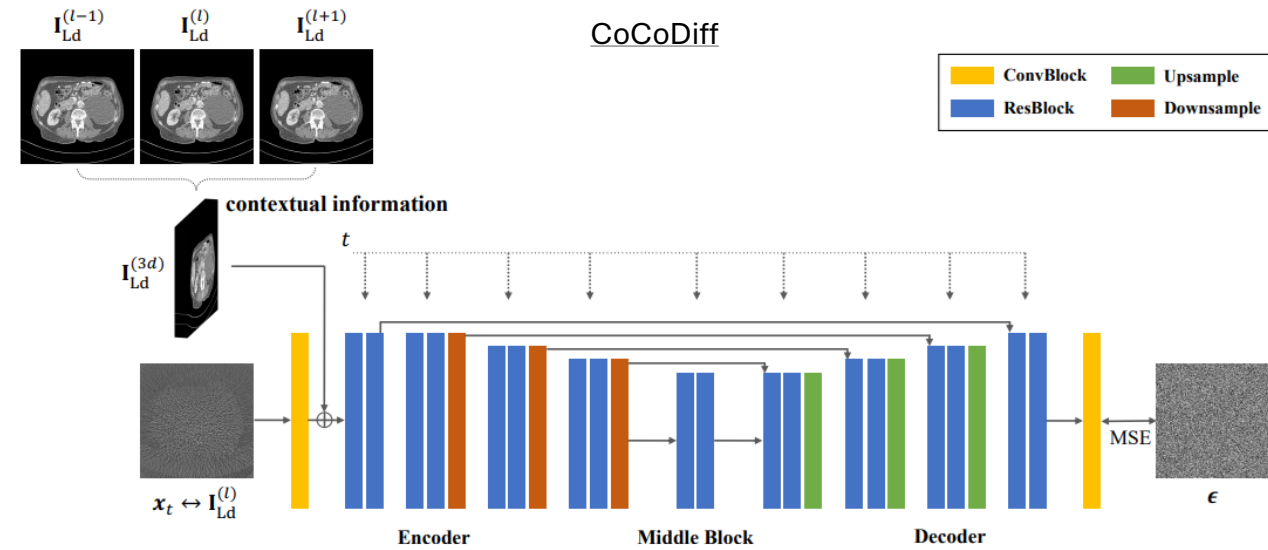
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- GAN-based Methods

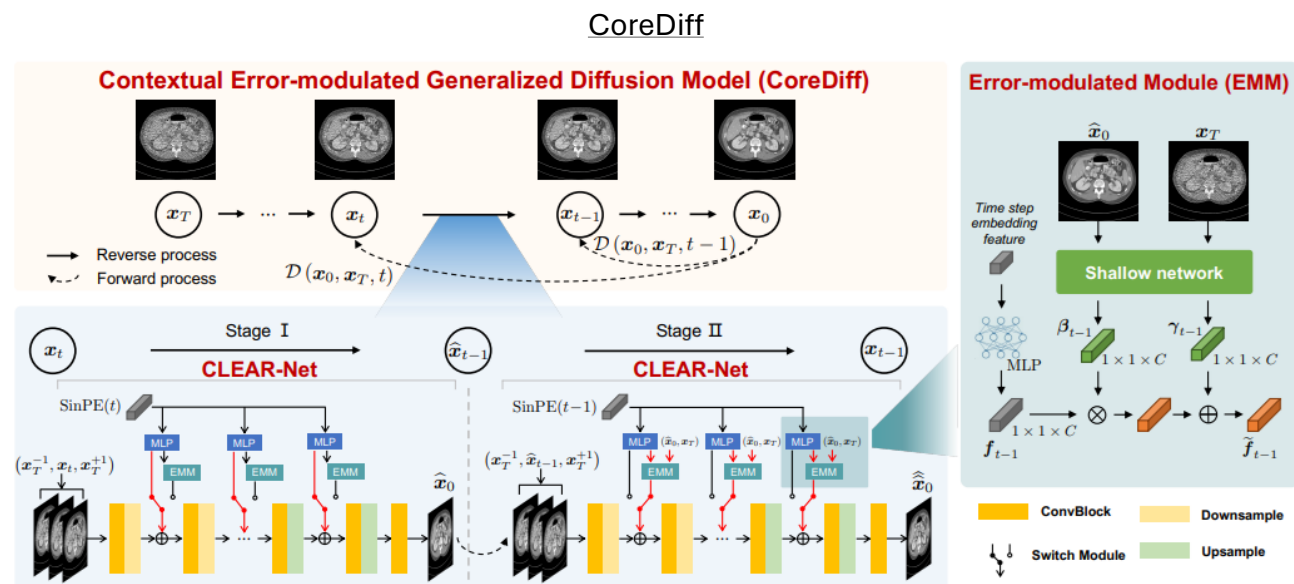
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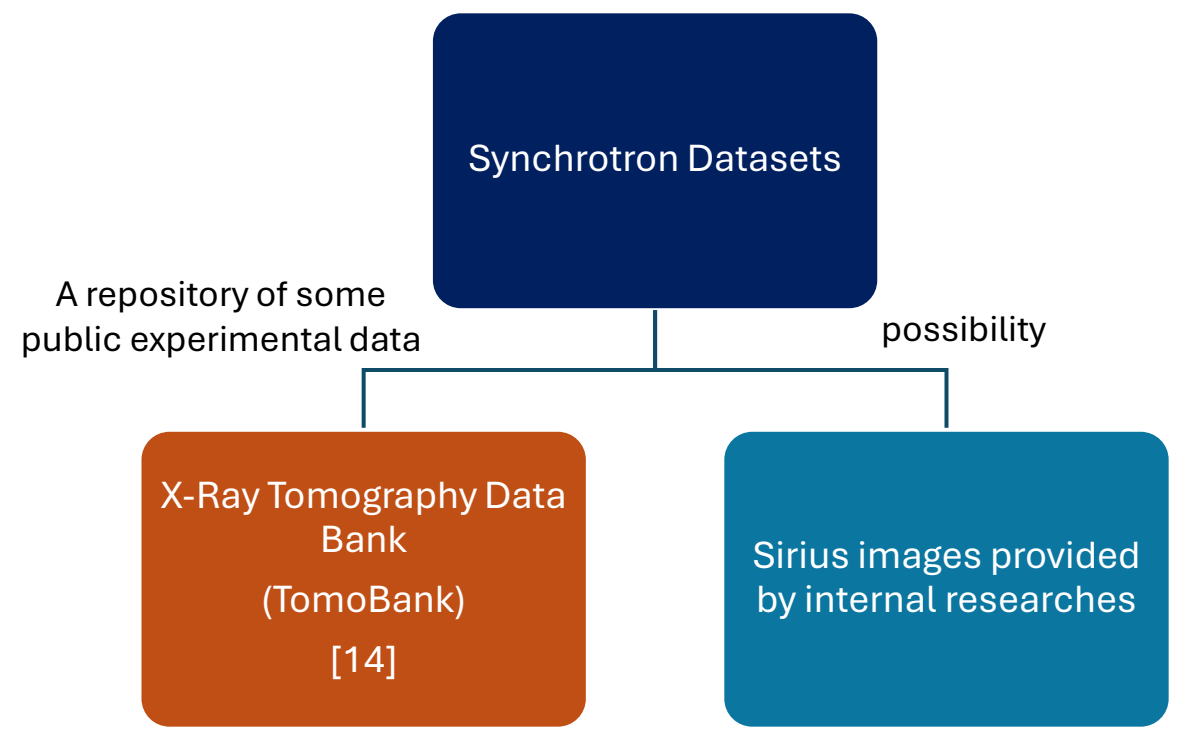
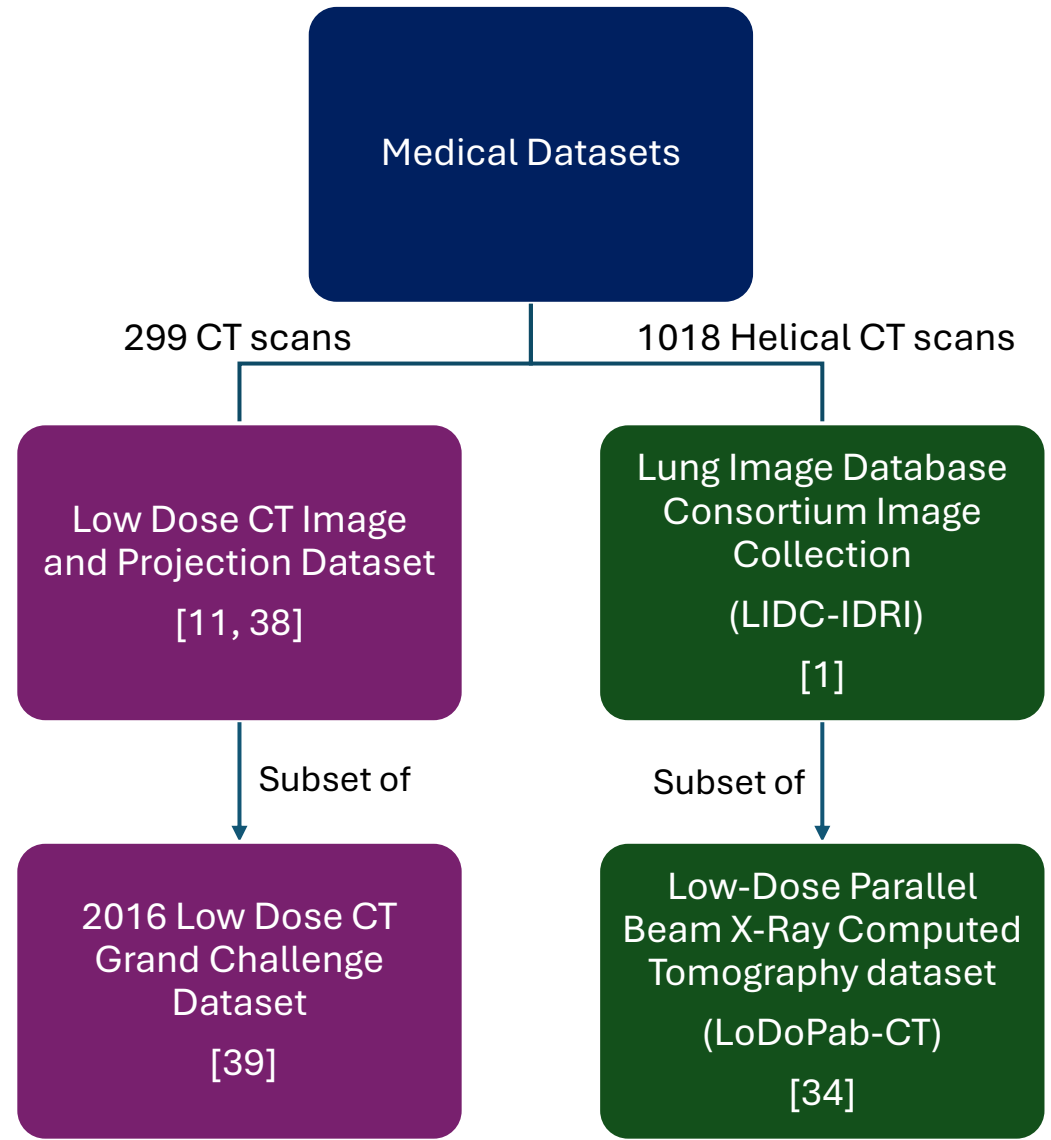
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Source: Gao and Shan [18]



Source: Gao et al. [17]



- **Number of Medical CT scans available is much higher than synchrotron CT scans!**
- **Low-dose data are simulated**

Root Mean Square Error [20]

$$\text{MSE} = \frac{1}{M \times N} \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} \left(I_{ij} - \hat{I}_{ij} \right)^2$$

$$\text{RMSE} = \sqrt{\text{MSE}}$$

Peak Signal-to-Noise Ratio [25]

$$\text{PSNR} = 10 \times \log_{10} \left(\frac{L_I^2}{\text{MSE}} \right)$$

Structural Similarity Index Measure [52]

$$\text{SSIM}(\mathbf{x}, \mathbf{y}) = [l(\mathbf{x}, \mathbf{y})]^\alpha \times [c(\mathbf{x}, \mathbf{y})]^\beta \times [s(\mathbf{x}, \mathbf{y})]^\gamma$$

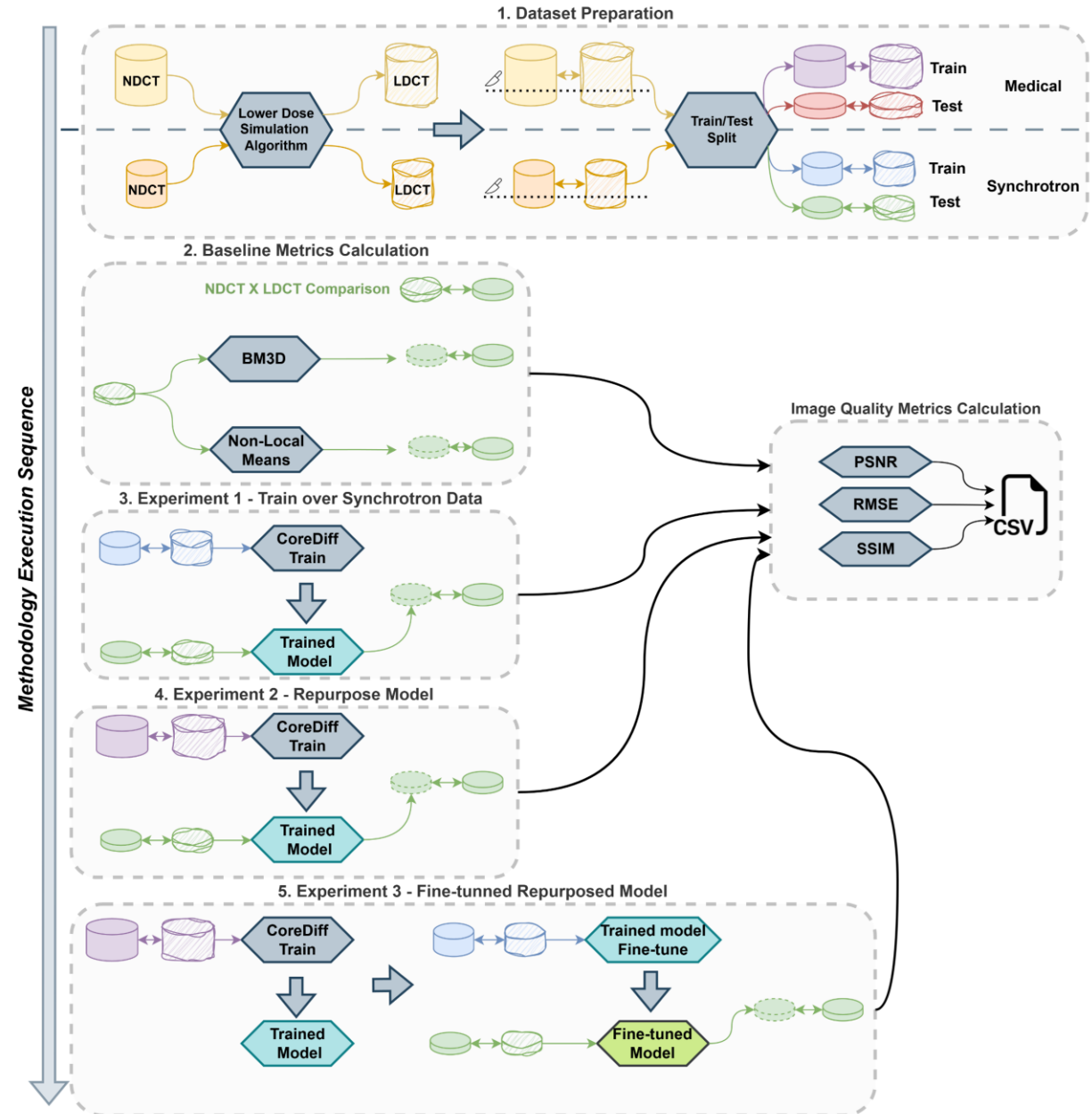
$$l(\mathbf{x}, \mathbf{y}) = \frac{2\mu_x\mu_y + C_1}{\mu_x^2 + \mu_y^2 + C_1} \quad C_1 = (k_1L)^2$$

$$c(\mathbf{x}, \mathbf{y}) = \frac{2\sigma_x\sigma_y + C_2}{\sigma_x^2 + \sigma_y^2 + C_2} \quad C_2 = (k_2L)^2$$

$$s(\mathbf{x}, \mathbf{y}) = \frac{\sigma_{xy} + C_3}{\sigma_x\sigma_y + C_3} \quad C_3 = \frac{C_2}{2}$$

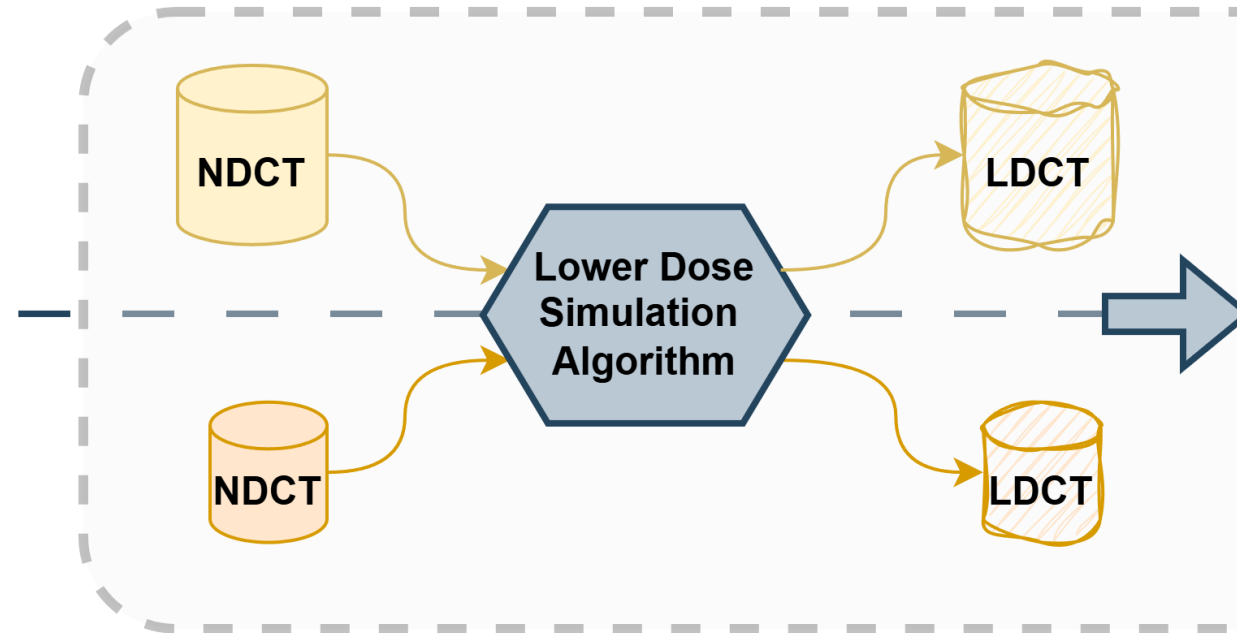
$$k_1 = 0.01 \quad k_2 = 0.03$$

- Methodology was divided into 5 stages:



- Methodology was divided into 5 stages:
- **Stage 1: Dataset Preparation**
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1. Dataset Preparation

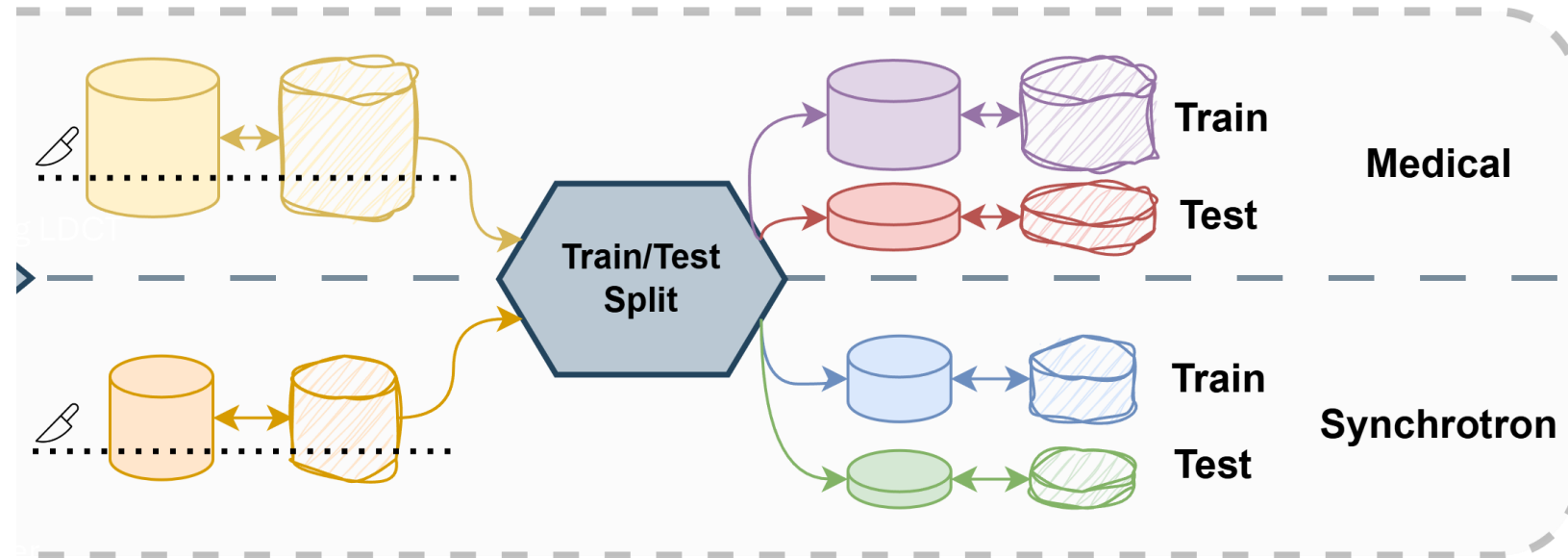


- Methodology was divided into 5 stages:

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- Simulate LDCT data for a range of different dose levels using the algorithm proposed by Yu et al. [58]
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1. Dataset Preparation





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• **Stage 2: Baseline Metrics Calculation**

- Compare simulated LDCT and NDCT data for baseline metrics
- Test BM3D and Non-Local Means for denoising LDCT synchrotron data

2. Baseline Metrics Calculation

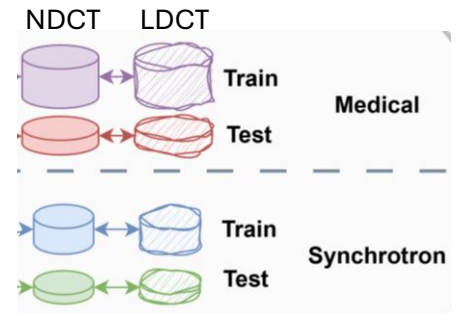
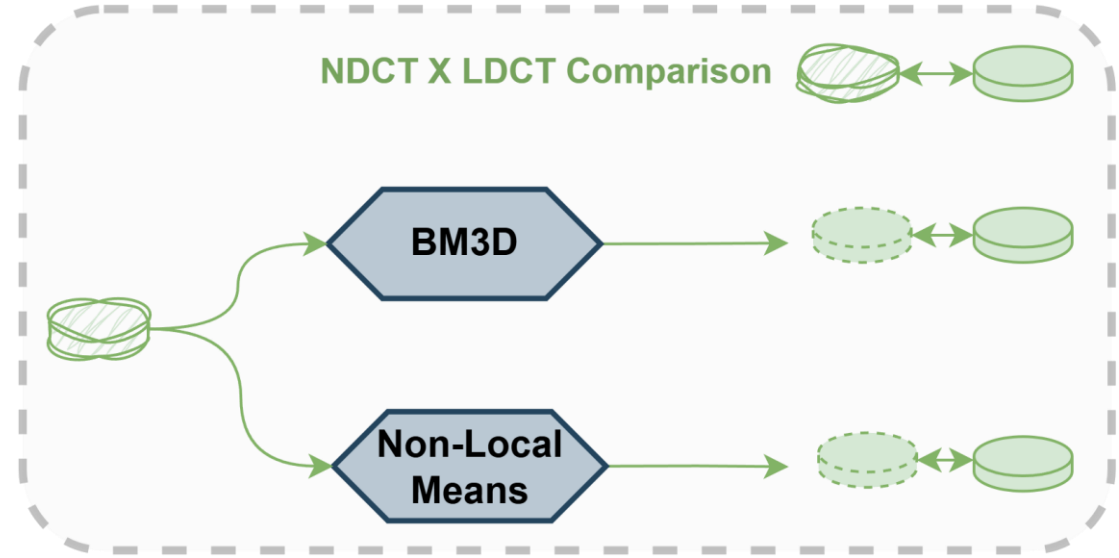
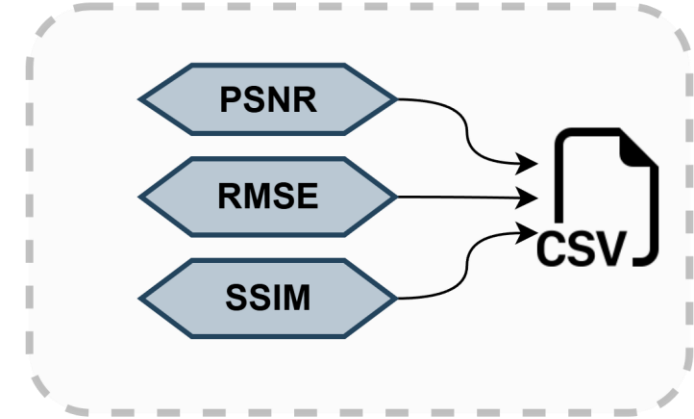
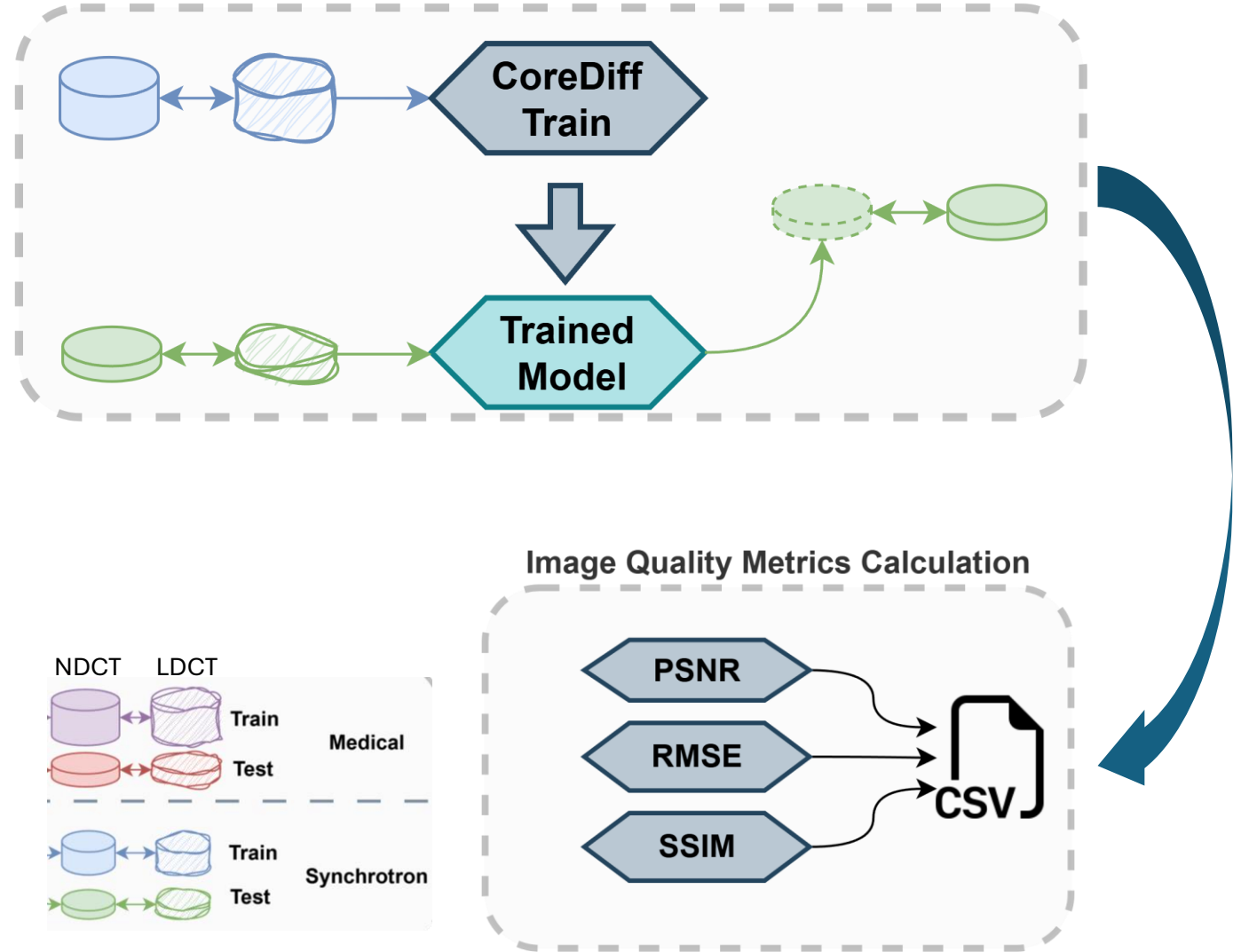


Image Quality Metrics Calculation



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 - Test BM3D and Non-Local Means for denoising LDCT synchrotron data
- Stage 3: Experiment 1**
 - Train and Test CoreDiff over synchrotron data

3. Experiment 1 - Train over Synchrotron Data





• Methodology was divided into 5 stages:

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• Stage 2: Baseline Metrics Calculation

- Compare simulated LDCT and NDCT data for baseline metrics
- Test BM3D and Non-Local Means for denoising LDCT synchrotron data

• Stage 3: Experiment 1

- Train and Test CoreDiff over synchrotron data

• Stage 4: Experiment 2

- Train CoreDiff over Medical data and test it over synchrotron data

4. Experiment 2 - Repurpose Model

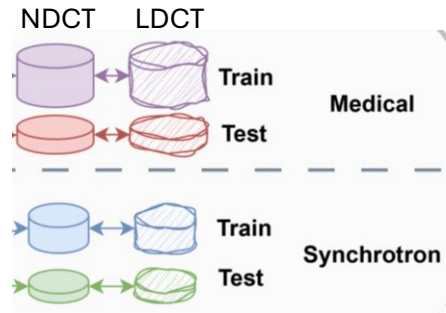
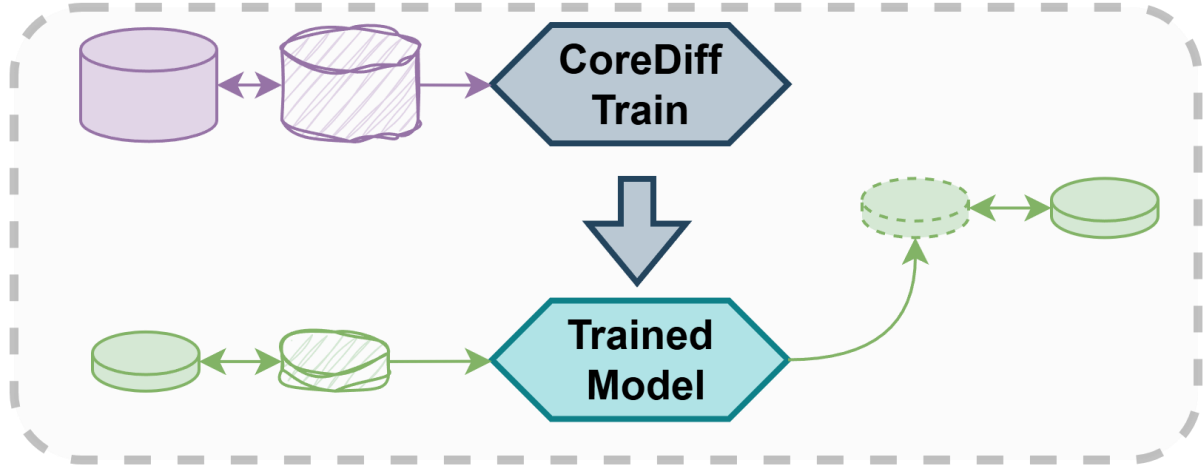
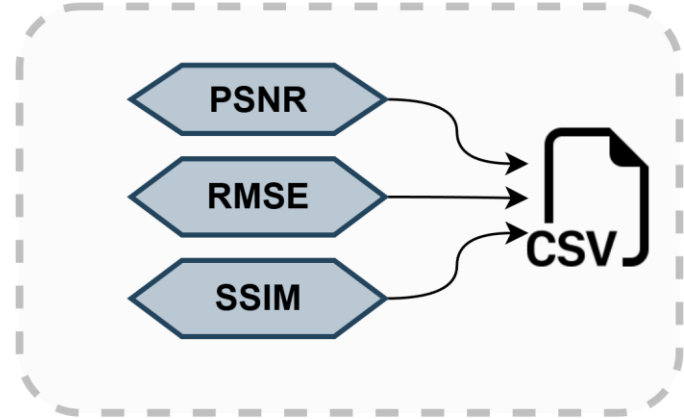


Image Quality Metrics Calculation



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- Stage 3: Experiment 1

- Train and Test CoreDiff over synchrotron data

- Stage 4: Experiment 2

- Train CoreDiff over Medical data and test it over synchrotron data

- Stage 5: Experiment 3**

- Train CoreDiff over Medical data, finetune it using synchrotron data and test it over synchrotron

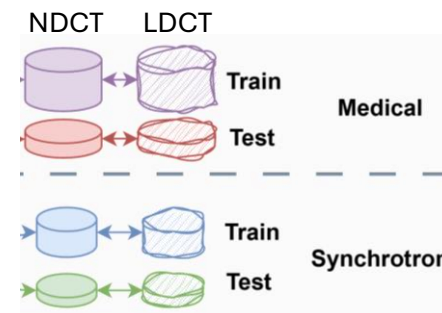
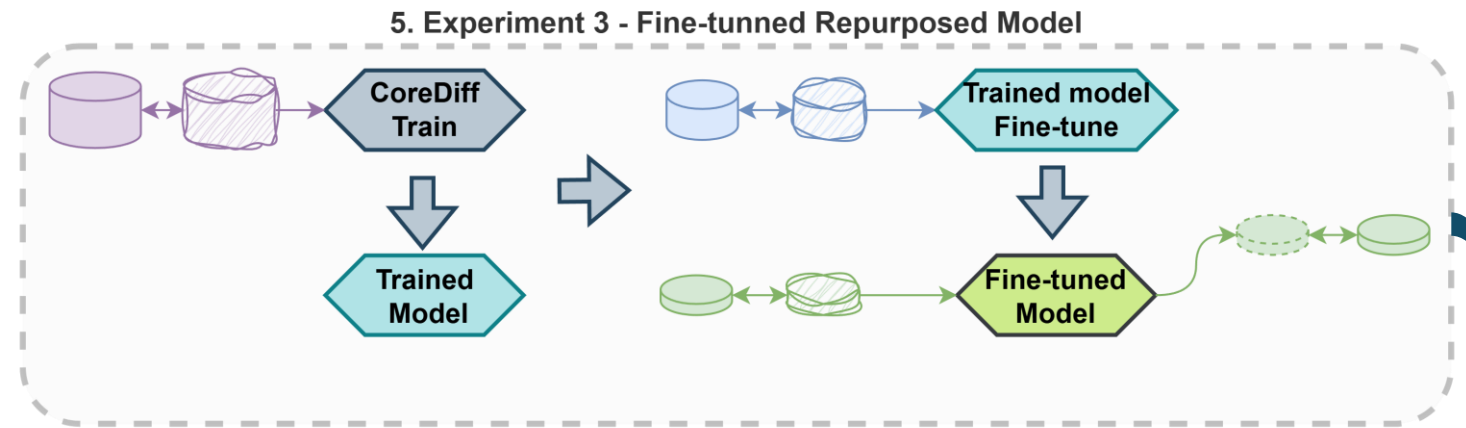
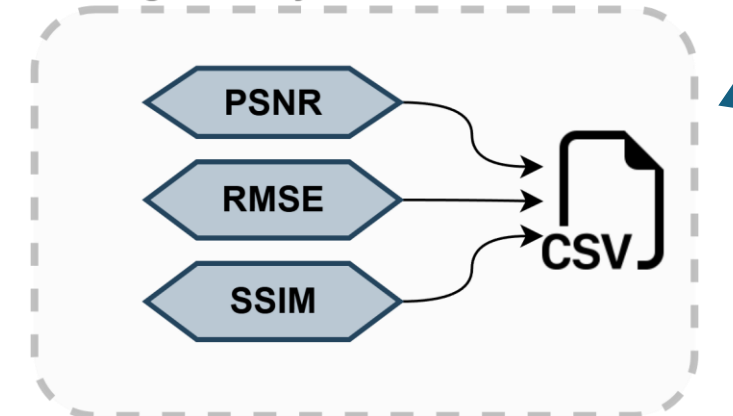
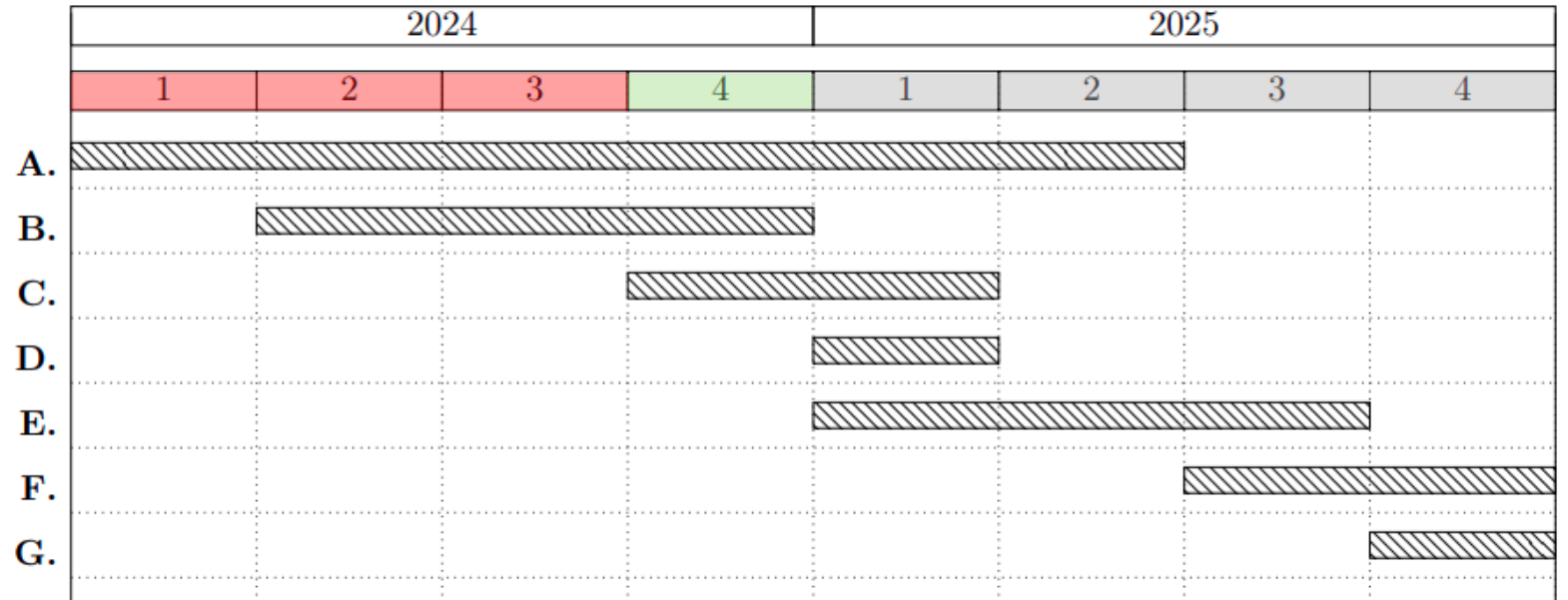


Image Quality Metrics Calculation



- A. Literature review
- B. Proposal writing and realization of the qualification exam
- C. Dataset preparation
- D. Baseline metric calculation
- E. Experiments execution and results comparison
- F. Results documentation and publishing
- G. Dissertation writing and defense



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Enhancing Synchrotron Low-Dose Computed Tomography Image Quality Using Diffusion-Based Generative Models

Candidate: Paulo Baraldi Mausbach

Supervisor: Prof. Dr. Zanoni Dias

Co-supervisor: Prof. Dr. Hélio Pedrini

Master's Degree Proposal

Institute of Computing - University of Campinas

October 17th of 2024

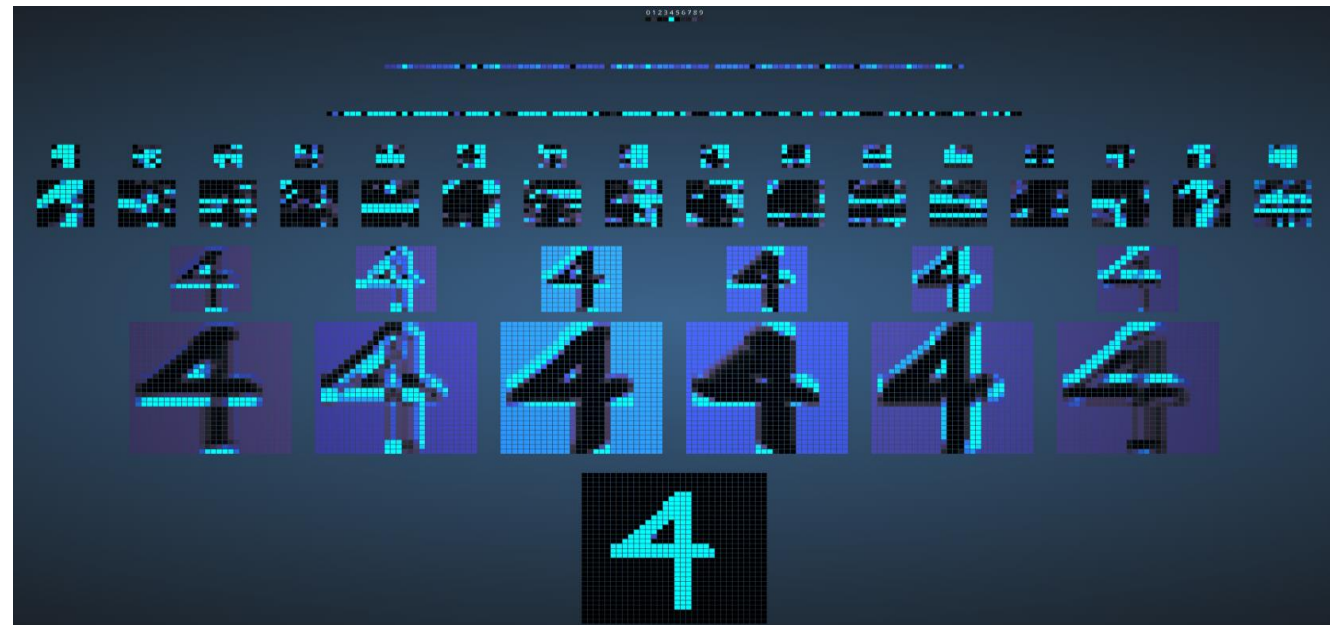
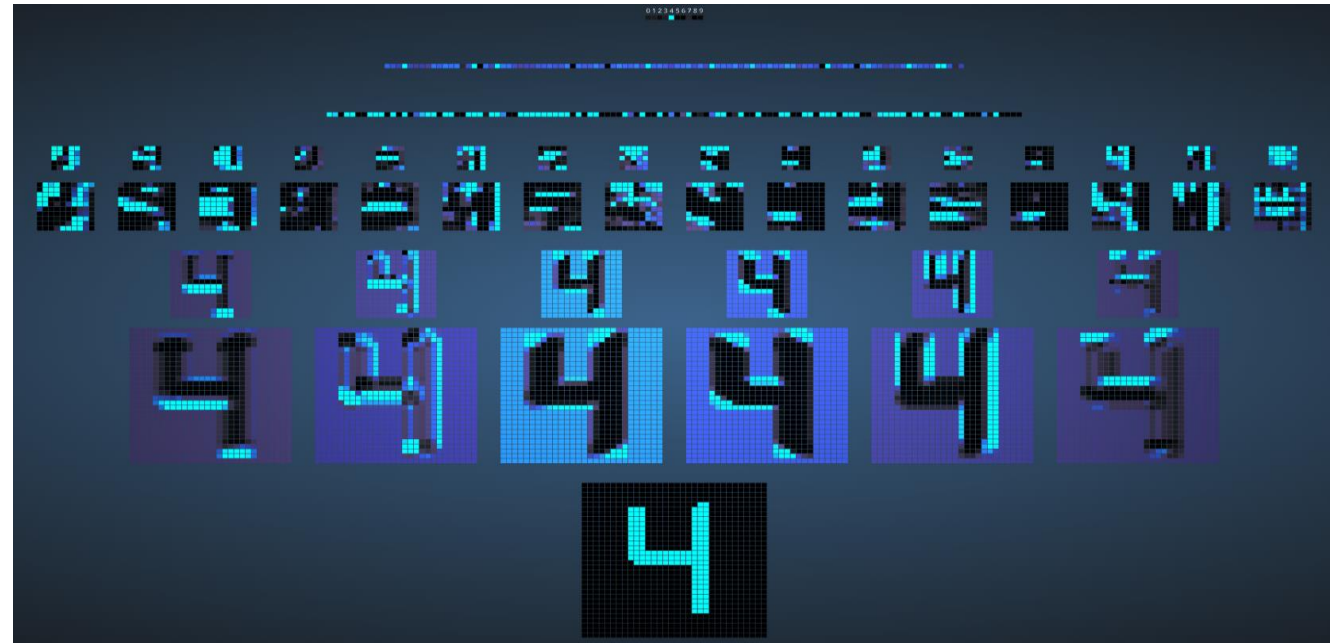
- Classical approaches:
 - Extensive feature extraction work
 - Handcrafted filters for edge, texture, shape
 - Some approaches required pre-processing data to work
- Convolutional Neural Networks (CNNs):
 - Processing done directly on raw data
 - Learn filters based on data
 - Learn simple and complex filters that optimize target task
- CNN revolutionized Computer Vision (CV) area [5]
- It is also extensively used for other data dimension applications such as time series, 3D images and videos [5]



$$\begin{array}{|c|c|c|} \hline a & b & c \\ \hline d & e & f \\ \hline g & h & i \\ \hline \end{array} * \begin{array}{|c|c|} \hline j & k \\ \hline l & m \\ \hline \end{array} = \begin{array}{|c|c|} \hline aj + bk + & bj + ck + \\ dl + em & el + fm \\ \hline dj + ek + & ej + fk + \\ gl + hm & hl + im \\ \hline \end{array}$$

I *K* *C*

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- ↓RMSE (Root Mean Square Error) [20]
- Difference between all pixel values from the “noise-free” image and the denoised estimation
- Low RMSE value indicates high similarity between “noise-free” and denoised estimation image

Image height Image width

$$\text{MSE} = \frac{1}{M \times N} \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} \left(I_{ij} - \hat{I}_{ij} \right)^2$$

Pixel value from position (i,j) in the noise-free image

Pixel value from position (i,j) in the denoised estimation

$$\text{RMSE} = \sqrt{\text{MSE}}$$

- \uparrow Peak Signal-to-Noise Ratio [25]
- Commonly used for assessing quality of image reconstruction, compression and denoising algorithms
- Ratio between the maximum possible signal value ($L_{\hat{I}}$) and the noise corrupting it (MSE between I and \hat{I})
- $MSE \rightarrow 0, PSNR \rightarrow \infty$
- High PSNR indicates low degradation of the signal by the existing noise

Maximum pixel value from noise-free image

$$PSNR = 10 \times \log_{10} \left(\frac{L_I^2}{MSE} \right)$$

$$MSE = \frac{1}{M \times N} \sum_{i=0}^{M-1} \sum_{j=0}^{N-1} (I_{ij} - \hat{I}_{ij})^2$$

Image height

Image width

Pixel value from position (i,j) in the noise-free image

Pixel value from position (i,j) in the denoised estimation

- SSIM (Structural Similarity Index Measure) [52]
- Perception-based metric to measure similarity between two images
- Calculated by comparing the degradation between two same size windows taken from the same position in the “noise-free” and the denoised estimation image.
- Evaluates three image aspects:
 - $l(x, y) \rightarrow$ Luminance
 - $c(x, y) \rightarrow$ Contrast
 - $s(x, y) \rightarrow$ Structure
- Calculated as a weighted combination of all three aspects

$$\text{SSIM}(\mathbf{x}, \mathbf{y}) = [l(\mathbf{x}, \mathbf{y})]^{\alpha} \times [c(\mathbf{x}, \mathbf{y})]^{\beta} \times [s(\mathbf{x}, \mathbf{y})]^{\gamma}$$

Mean pixel intensity

Weight factors

$$l(\mathbf{x}, \mathbf{y}) = \frac{2\mu_x\mu_y + C_1}{\mu_x^2 + \mu_y^2 + C_1}$$

Standard deviation

$$c(\mathbf{x}, \mathbf{y}) = \frac{2\sigma_x\sigma_y + C_2}{\sigma_x^2 + \sigma_y^2 + C_2}$$

Variance

Covariance

$$s(\mathbf{x}, \mathbf{y}) = \frac{\sigma_{xy} + C_3}{\sigma_x\sigma_y + C_3}$$

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Mean pixel intensity \rightarrow Weight factors \rightarrow Stabilization Variables

$$l(\mathbf{x}, \mathbf{y}) = \frac{2\mu_x\mu_y + C_1}{\mu_x^2 + \mu_y^2 + C_1}$$

Standard deviation \rightarrow

$$c(\mathbf{x}, \mathbf{y}) = \frac{2\sigma_x\sigma_y + C_2}{\sigma_x^2 + \sigma_y^2 + C_2}$$

Variance \rightarrow

Covariance \rightarrow

$$s(\mathbf{x}, \mathbf{y}) = \frac{\sigma_{xy} + C_3}{\sigma_x\sigma_y + C_3}$$

$$C_1 = (k_1L)^2$$

$$C_2 = (k_2L)^2$$

$$C_3 = \frac{C_2}{2}$$

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Stabilization Variables

$$C_1 = (k_1 L)^2$$

$$C_2 = (k_2 L)^2$$

$$C_3 = \frac{C_2}{2}$$

Maximum pixel value

$$k_1 = 0.01 \quad k_2 = 0.03$$

Small constants ($\ll 1$) with arbitrary value

SSIM authors reported “fairly insensitive” variations of SSIM caused by variations on values from k_1 and k_2 [52]

- SSIM (Structural Similarity Index Measure) [52]
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$$\text{SSIM}(\mathbf{x}, \mathbf{y}) = [l(\mathbf{x}, \mathbf{y})]^\alpha \times [c(\mathbf{x}, \mathbf{y})]^\beta \times [s(\mathbf{x}, \mathbf{y})]^\gamma$$

SSIM can be simplified as

if $\alpha = \beta = \gamma = 1$

$$\text{SSIM}(\mathbf{x}, \mathbf{y}) = \frac{(2\mu_x\mu_y + C_1)(2\sigma_{xy} + C_2)}{(\mu_x^2 + \mu_y^2 + C_1)(\sigma_x^2 + \sigma_y^2 + C_2)}$$

$$C_1 = (k_1L)^2 \quad C_2 = (k_2L)^2 \quad C_3 = \frac{C_2}{2}$$

$$k_1 = 0.01 \quad k_2 = 0.03$$