

Graph Denoising for Molecular Imaging

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Outline

1 Molecular Imaging Methods

2 Graphs

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

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1 Molecular Imaging Methods

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Cryo-Electron Microscopy (Cryo-EM)

- Major motivation
- Enables observation of molecules in near atomic resolution

Cryo-Electron Microscopy (Cryo-EM)

- › Major motivation
- › Enables observation of molecules in near atomic resolution
- › Observation through an electron microscope
- › Frozen state of molecules required for observation
 - › Frozen molecules are fragile \mapsto electron microscope low power
 - › During freezing, molecules rotate randomly
- › Observations can be reconstructed to a 3D model
- › Single particle Cryo-EM

Cryo-Electron Microscopy (Cryo-EM) - Illustration

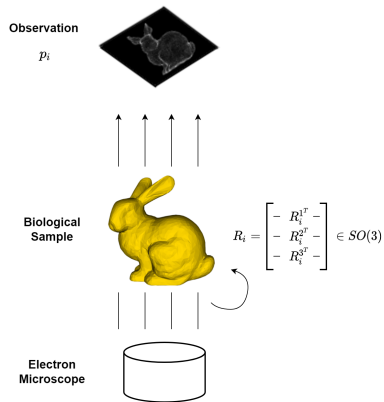


Figure: Cryo-EM overview

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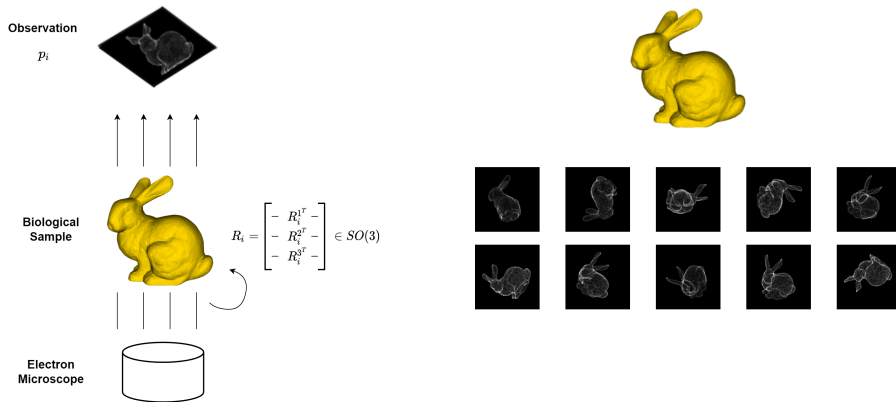
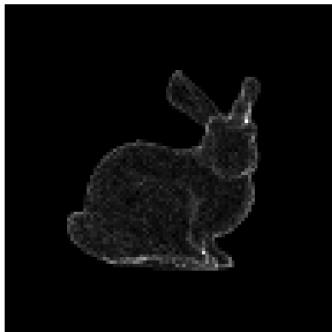
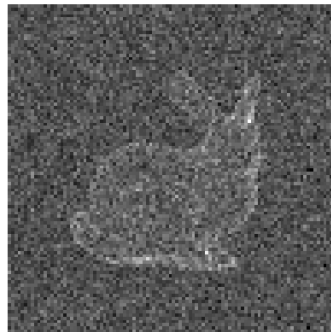


Figure: Cryo-EM overview

Cryo-Electron Microscopy (Cryo-EM) - Illustration



(a) Clean micrograph



(b) Noisy micrograph

Computed Tomography (CT)

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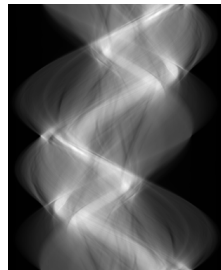
- › Similar to Cryo-EM
- › Can be seen as a simpler version in 2D with known observation angles
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(a) Biological sample



(b) Clean observation (sinogram)

Shared Observation Model

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Observation

$$y_i = p_i + \eta_i \quad \text{with } 1 \leq i \leq N \quad (1)$$

- > y : noisy observation
- > p : noiseless observation
- > η : noise, assumed $\eta_i \sim \mathcal{N}(0, \sigma^2)$
- > $y_i \in \mathbb{R}^M$, M : observation dimension
- > N : number of observations

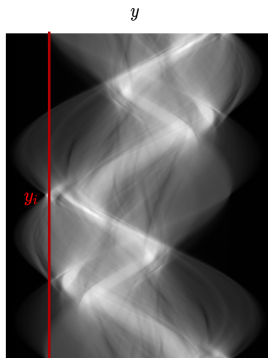
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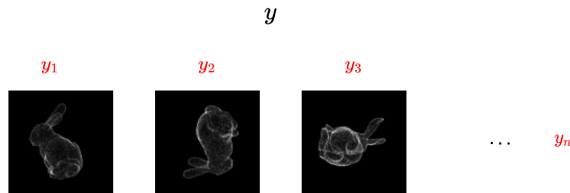
$$\begin{aligned} y_i &= p_i + \eta_i && \text{with } 1 \leq i \leq N \\ y_i &= A(x, \theta_i) + \eta_i && \text{with } 1 \leq i \leq N \end{aligned} \tag{1}$$

- > y : noisy observation
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- > η : noise, assumed $\eta_i \sim \mathcal{N}(0, \sigma^2)$
- > x : biological sample
- > $y_i \in \mathbb{R}^M$, M : observation dimension
- > N : number of observations
- > $A : x \mapsto A(x; \theta_i) \in \mathbb{R}^M$: a non-linear operator
- > θ_i : observation angle

Observation - Illustration



(a) CT Observation -
sinogram



(b) Cryo-EM Observation - micrographs

Reconstruction

Reconstruction

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$$Recon : \mathbb{R}^{M \times N} \rightarrow \mathbb{R}^{M \times M} \quad y \mapsto Recon(y; \theta) \quad (2)$$

Signal-to-noise-ratio (SNR)

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- SNR is a measure, which compares the power of an input signal to the power of the undesired noise
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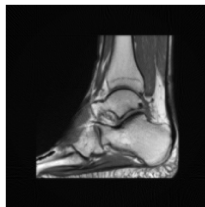
SNR is used as a metric for the quality of reconstructions.

Reconstruction - Computed Tomography

- Filter Backprojection (FBP)
 - Can be considered historical approach
 - Enables reconstruction for moderate noise

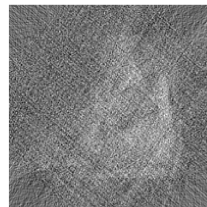
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$$\text{Recon}(p, \theta) \approx x$$



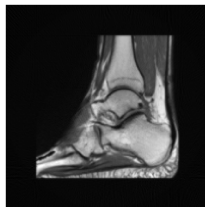
(b) Reconstruction noisy

with SNR_y 5 dB:

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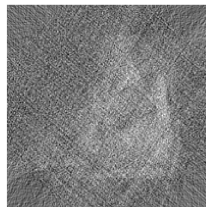
Reconstruction - Computed Tomography

- Filter Backprojection (FBP)
 - Can be considered historical approach
 - Enables reconstruction for moderate noise
- Neural Network Approaches
 - Today state-of-the art
 - Using result of FBP and further denoise
 - U-Net Ronneberger, Fischer, and Brox 2015



(a) Reconstruction clean:

$$\text{Recon}(p, \theta) \approx x$$



(b) Reconstruction noisy

with SNR_y 5 dB:

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Problem and Goal

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Problem

p not observable directly only access to y .

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Goal

$$\text{denoiser} : y_i \mapsto y_i^* \approx p_i$$

$$\text{Recon}(\text{denoiser}(y; \theta)) \approx x$$

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Nodes

$(n_1, n_2, \dots) \in \mathbb{R}^F$, with F as node feature dimensions

Edges

Edges are defined as a set of tuples (i, j) , where i and j determine the index of the nodes.

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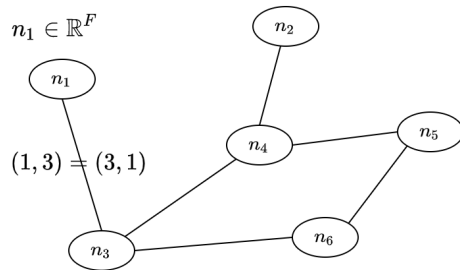


Figure: Sample graph

Graph - Definitions - Adjacency Matrix

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Adjacency Matrix

The binary adjacency matrix of graph $G = \langle V, E \rangle$ is defined as:

$$A_{ij} = \begin{cases} 1 & \text{if } (i, j) \in E \\ 0, & \text{otherwise} \end{cases} \quad (3)$$

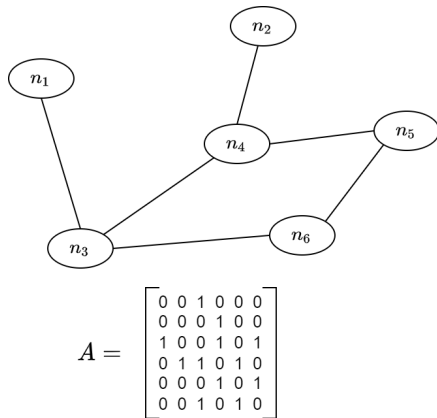


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Graph - Definitions - Degree

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Degree of a node

The *degree* of a node is defined as the number of (incoming) edges.

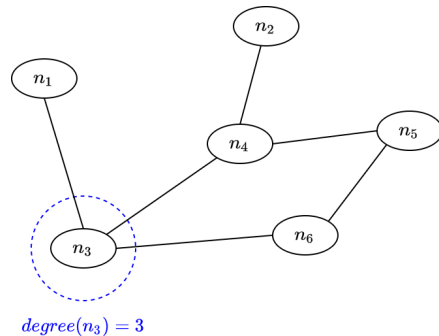


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Degree Matrix of Graph G

Is a diagonal matrix with degree of nodes as entries.

$$D_{ii} = \text{degree}(n_i)$$

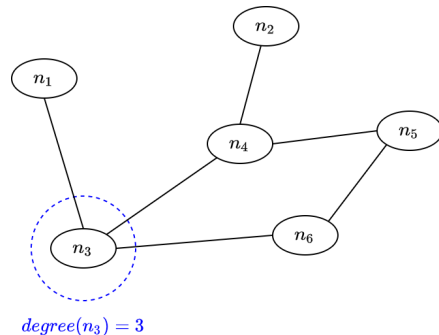


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Graph for Molecular Imaging Observation

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› Nodes: Single observation y_i

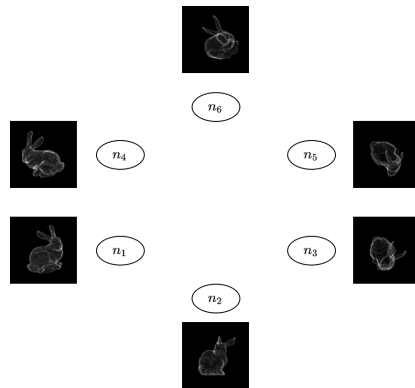


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How to construct a graph for molecular imaging?

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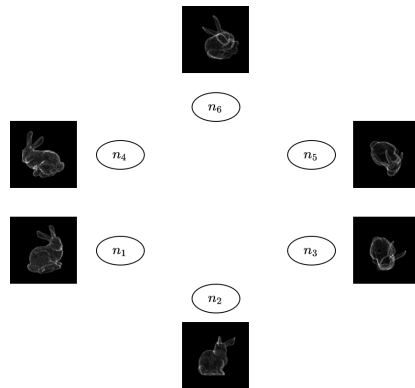


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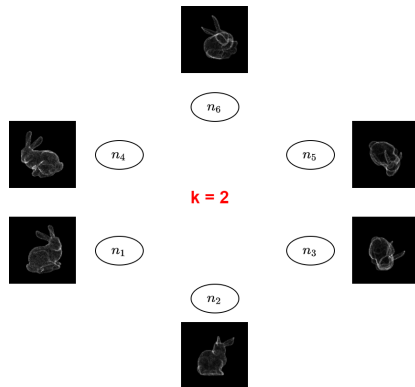


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Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

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 - Define similarity measure for nodes: ℓ_2 -norm

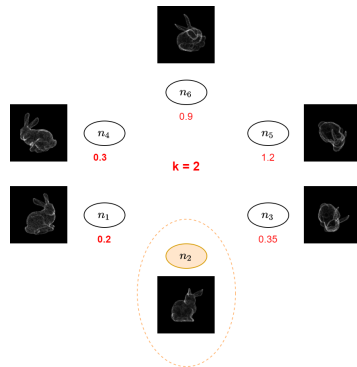


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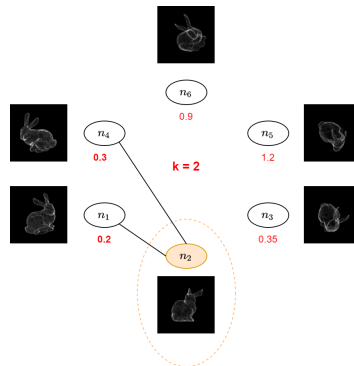


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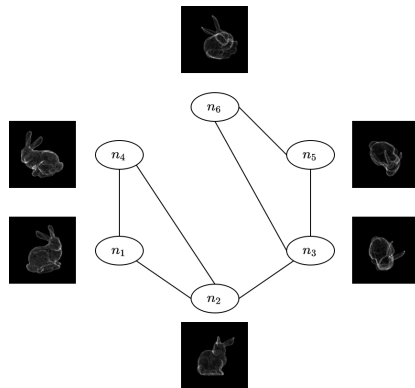


Figure: Sample graph for cryo-EM observation

Graph for Molecular Imaging Observation - Noise

What happens with our noisy observations?

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What happens with our noisy observations?

- > With noise, graph will capture neighborhood inaccurately.

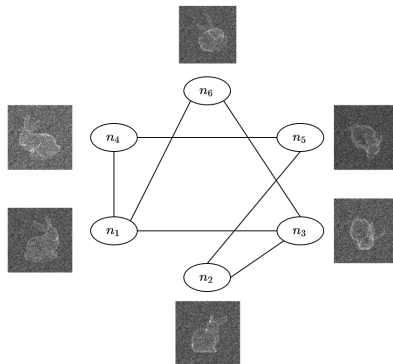


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Graph Laplacian (GL)

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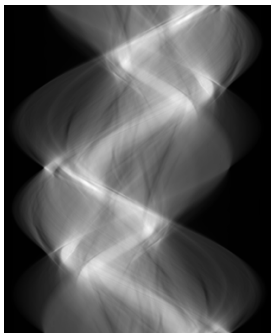
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Low-dimensional Embedding

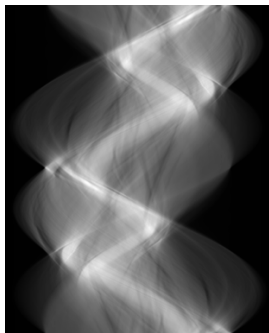
1. Construct a k-NN graph from observations.
2. Calculate $L = D - A$
3. Get 2nd and 3rd smallest eigenvalue with corresponding eigenvectors.

Low-dimensional Embedding for Computed Tomography



(a) Clean CT observation

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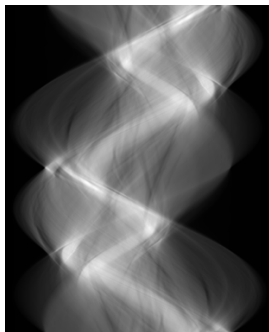


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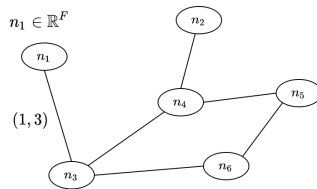


(b) Building k-NN graph with $k = 2$

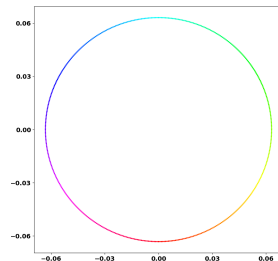
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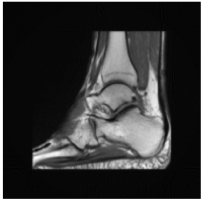
(c) 2_{nd} and 3_{rd} smallest eigenvectors of $L = D - A$

Computed Tomography with unknown angles

- GL-Embedding estimates observation angles

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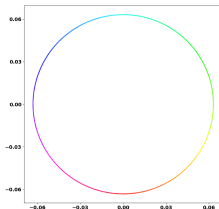
(a) Reconstruction known angles

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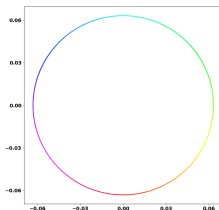
(b) GL-Embedding from $k = 2$

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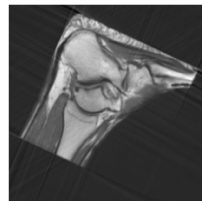
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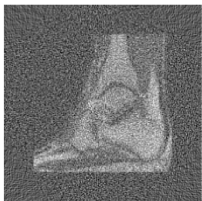
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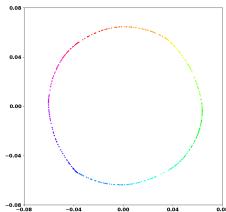
What happens in the noisy case?

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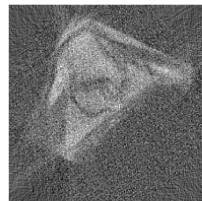
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(a) Reconstruction known angles $SNR_y : 10$ dB



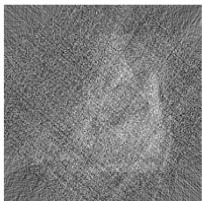
(b) GL-Embedding from $k = 8$ and $SNR_y : 10$ dB



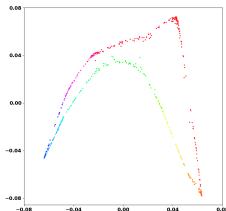
(c) Reconstruction unknown angles $SNR_y : 10$ dB

Computed Tomography with unknown angles

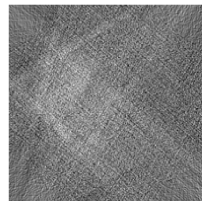
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The fewer noise is available in the observation, the better reconstruction is possible.

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 - No graph as data structure
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Exploit graph as a data structure and the GL-Embedding

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- › GAT-Denoiser is a graph neural network (GNN) to denoise observations
- › Consists of three components:
 - › Convolution
 - › Graph Attention Network (GAT) Veličković et al. 2017
 - › End-to-End Learning

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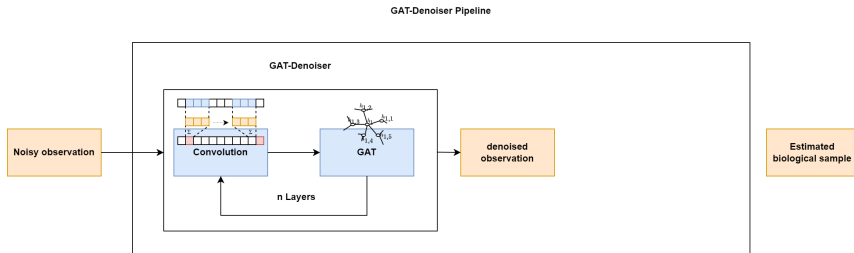
GAT-Denoiser Pipeline

Noisy observation

Estimated
biological sample

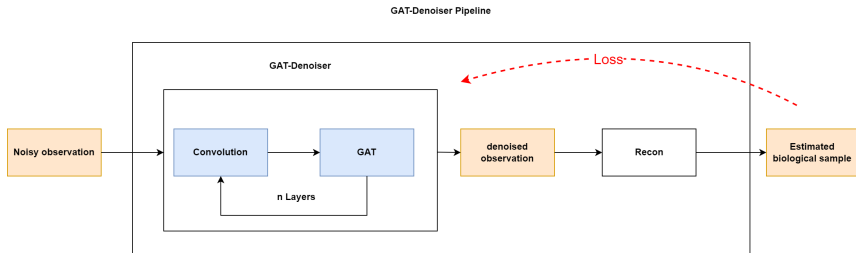
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- > Consists of three components:
 - > **Convolution**
 - > Denoise single observation
 - > **Graph Attention Network (GAT)** Veličković et al. 2017
 - > Denoise neighboring observation
 - > End-to-End Learning



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- > Consists of three components:
 - > Convolution
 - > Graph Attention Network (GAT) Veličković et al. 2017
 - > End-to-End Learning
 - > Optimize for reconstruction quality
 - > Loss: $\mathcal{L}_{reconstruction} = \|x - Recon(GAT-Denoiser(y))\|_2^2$



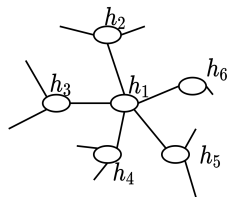
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- › Computes new node features
- › Averages graph over neighborhood
- › Multi-head available, motivated by Vaswani et al. 2017
- › α : normalized attention coefficients
- › W : learnable weight matrix
- › σ : activation function (Exponential Linear Unit)



$$h'_1 = \sigma \left(\sum_{i=1}^6 \alpha_i W h_i \right)$$

Input Graph

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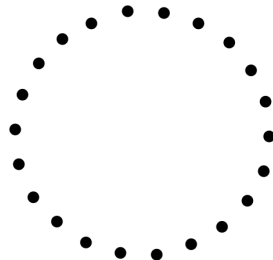
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- › Construct graph from observation angles

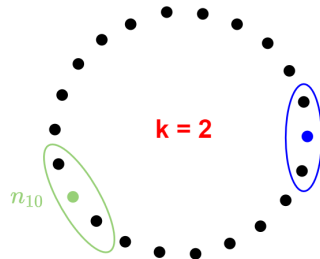
Input Graph

- Exploit information from GL
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 - Map angles to unit-circle / unit-sphere



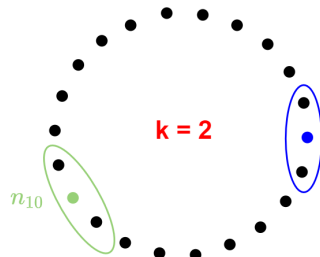
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Observation angles θ are assumed to be equally spaced on the unit-circle / unit-sphere.

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LoDoPaB-CT dataset

- › Dataset for low-dose Computed Tomography
- › 35'820 train samples
- › 3'553 test samples
- › Resolution 64×64
- › BM3D as baseline algorithm

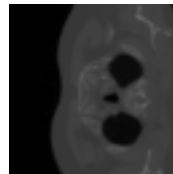
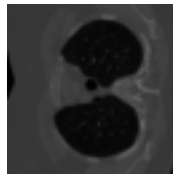
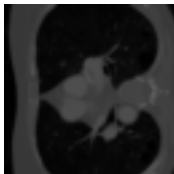


Figure: Some samples from the LoDoPaB-CT dataset.

GAT-Denoiser - Implementation for Computed Tomography

- › PyTorch Geometric
- › U-Net used for reconstruction
 - › Pre-trained with complete dataset and SNR_y in $[-10, 0]$ dB for 200 epochs
 - › Joint U-Net training possible
- › Modular architecture which allows disabling of Convolution and U-Net
- › Mini-batch gradient descent

Evaluation

> Small Scale Experiments

- > 1024 train samples
- > 100 test samples
- > 200 epochs

> Large Scale Experiments

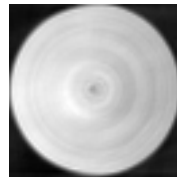
- > Complete LoDoPaB-CT dataset
- > 20 - 40 epochs

Evaluation

- › Small Scale Experiments
 - › 1024 train samples
 - › 100 test samples
 - › 200 epochs
 - › Goal: Find most promising architecture
- › Large Scale Experiments
 - › Complete LoDoPaB-CT dataset
 - › 20 - 40 epochs
 - › Goal: Find best model

Small Scale Experiments - Overall Results

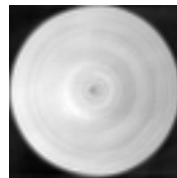
- Learning fails with random graph (Erdős–Rényi)
- Learning succeeds with defined input graph



(a) Reconstruction with random Erdős–Rényi graph with $p = 0.01$

Small Scale Experiments - Overall Results

- › Learning fails with random graph (Erdős–Rényi)
- › Learning succeeds with defined input graph
- › Single components contribute to success of GAT-Denoiser:
 - › $N=1024$
 - › $M=64$
 - › GAT: 2 layers and 4 heads
 - › Convolution: kernel size 3 and padding 1
 - › k-NN with $k=2$
 - › Joint U-Net training

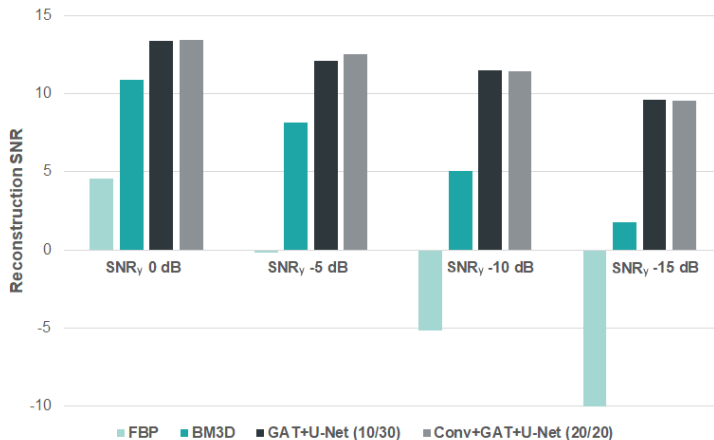


(a) Reconstruction with random Erdős–Rényi graph with $p = 0.01$

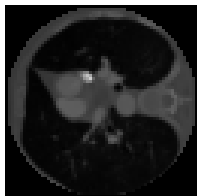
Large Scale Experiment - SNR Results

- Different algorithms / GAT-Denoiser models:
 - FBP
 - BM3D
 - GAT + U-Net (10/30)
 - Conv + GAT + U-Net (20/20)
- 3'553 test samples
- Reconstruction SNR averages over all test samples

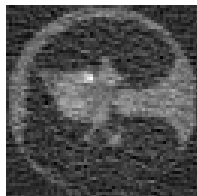
Large Scale Experiment - SNR Results



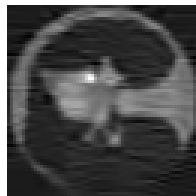
Large Scale Experiment - Visual Results - SNR_y 0 dB



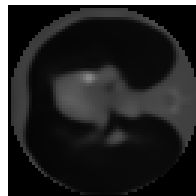
(a) Clean sample



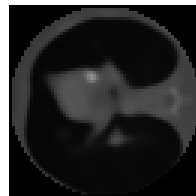
(b) FBP



(c) BM3D



(d) *GAT+U-Net(10/30)*

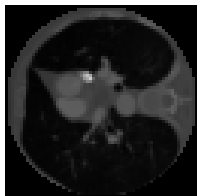


(e) *Conv+GAT+U-Net(20/20)*

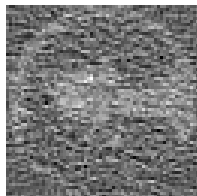
Figure: Large Scale Experiment: Visual results for SNR_y 0 dB.

GAT-Denoiser improves BM3D for SNR_y 0 dB by 27.6%.

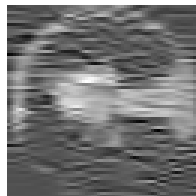
Large Scale Experiment - Visual Results - SNR_y -10 dB



(a) Clean sample



(b) FBP



(c) BM3D



(d) $GAT+U-$
 $Net(10/30)$

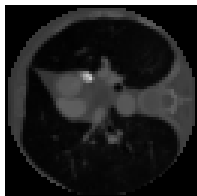


(e)
 $Conv+GAT+U-$
 $Net(20/20)$

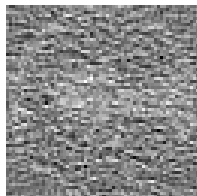
Figure: Large Scale Experiment: Visual results for SNR_y -10 dB.

GAT-Denoiser improves BM3D for SNR_y -10 dB by 126.0%.

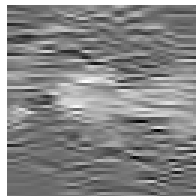
Large Scale Experiment - Visual Results - SNR_y -15 dB



(a) Clean sample



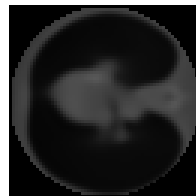
(b) FBP



(c) BM3D



(d) *GAT+U-Net(10/30)*



(e) *Conv+GAT+U-Net(20/20)*

Figure: Large Scale Experiment: Visual results for SNR_y -15 dB.

GAT-Denoiser improves BM3D for SNR_y -15 dB by 379.9%.

Outline

1 Molecular Imaging Methods

2 Graphs

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

Summary

- › GAT-Denoiser enables denoising of observations
 - › Convolution
 - › GAT
 - › End-To-End Learning

Summary

- › GAT-Denoiser enables denoising of observations
 - › Convolution
 - › GAT
 - › End-To-End Learning
- › Evaluated on LoDoPaB-CT dataset
 - › Outperformed baseline BM3D by up to 379.9 %
 - › Joint U-Net training boost performance

Future Work

- Computed Tomography
 - Improve current GAT-Denoiser
 - Derive GAT-Denoiser for 3D
 - Make it work for unknown angles

Future Work

- › Computed Tomography
 - › Improve current GAT-Denoiser
 - › Derive GAT-Denoiser for 3D
 - › Make it work for unknown angles
- › Cryo-EM
 - › Known angles
 - › Unknown angles
 - › Work with structural variety in observations

Outline

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Questions



References (1)

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